## AN ABSTRACT OF THE THESIS OF

Connie J. Hubbard for the degree of Master of Science in Forest Science presented on _ April 15. 1991. Title: A Plant Association Classification for McDonaldDunn Forest

Signature redacted for privacy.
Abstract approved:
William H. Emmingham

The purpose of this study was to develop a plant association classification for the mature coniferous forests of the McDonald and Paul M. Dunn Research Forests. A secondary objective was to quantify diversity within the plant associations. This Forest is approximately 11,000 acres and is located 6 miles to the north of Corvallis Oregon. This forest is owned and managed by the College of Forestry at oregon State University.

This study, and a companion study by Leavell (1991), was funded by the Research Forest to broaden their understanding of the plant communities on the Forest. The companion study developed relationships between the plant associations and the environment.

Classification was based on 108 plots, using percent cover of trees, shrubs, forbs, and grasses. This sampling density of approximately 1 plot per 100 acres is much
greater than most community classifications in the area (Hemstrom \& Logan, 1986; Hemstrom et al., 1987; Topik et al., 1988; Juday, 1976). A total of 117 vascular plant species were encountered; 68 of these were used in classification. Stands sampled for this classification were primarily seral, yet the most mature available. Few climax stands were available for sampling.

Plant associations were developed using TWINSPAN, a two-way indicator species analysis (Hill, 1979b). Six plant associations were developed and described in this study:

Tsuga heterophylla/Acer circinatum-Gaultheria shallon Abies grandis/Acer circinatum-Gaultheria shallon Abies grandis/Disporum hookeri-Thalictrum occidentale Abies grandis/Polystichum munitum

Abies grandis/Rubus ursinus-Rhus diversiloba
Abies grandis/Brachypodium sylvaticum
One plant association, Abies grandis/Acer circinatumGaultheria shallon, parallels a previously described plant association described by Juday in the Valley Margin Zone (Juday, 1976). The other 5 plant associations described in this study are original descriptions.

The TSHE/ACCI-GASH, ABGR/ACCI-GASH, and the ABGR/RUURRHDI plant associations are shrub-dominated. ABGR/DIHOTHOC and ABGR/POMU are forb-dominated associations, and ABGR/BRSY is a grass-dominated plant association.

Plant associations were shown to have significant differences in species richness, heterogeneity, evenness, and in vertical structure. Average species richness per plot is highest in the ABGR/ACCI-GASH and ABGR/DIHO-THOC plant associations, which also has high relative diversity (Shannon's index) and evenness. Species richness is lowest in the TSHE/ACCI-GASH plant association, probably because it was represented only by seral stands.

Structural diversity included an analysis of snags. Snag number, size, and distribution were not related to plant associations. Current snag levels indicate that at most, 30 percent of the maximun populations for six woodpecker species could be supported on this Forest.

A description of each plant association is given. The description for each includes: extent, location, characteristic species, species richness and percent cover within four strata (trees, shrubs, forbs, and grasses), heterogeneity (Shannon's diversity index), and an evenness measure.

# A Plant Association Classification 

for McDonald-Dunn Forest
by
Connie J. Hubbard

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## APPROVED:



Signature redacted for privacy.
Head of Deparyment of Forest Science
Signature redacted for privacy.


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Typed by researcher for: Connie J. Hubbard

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# A PLANT ASSOCIATION CLASSIFICATION FOR MCDONALD-DUNN FOREST 

## INTRODUCTION

This study is a plant association classification for the McDonald and Paul M. Dunn Forests which includes an analysis of diversity measures. The College of Forestry at Oregon State University owns and manages the 11,000-acre Forest. For simplicity, these forests will be referred to as McDonald-Dunn Forest, or the Forest. The Forest is located northwest of Corvallis, Oregon (see Figure 1.).

## BACKGROUND

It is human nature to classify; to categorize and organize information. Classification in general aims at grouping, condensing, or summarizing data in order to reveal an underlying structure, or organization.

This study is a result of the managers' desire to improve their knowledge of the Forests resources. Prior to this study quantitative information on vegetation was limited to that of trees. Plant association classification will improve the managers' knowledge of their floristic resource. Vegetation not only reflects the environment, but also provides wildife habitat and aesthetic qualities to the Forest.

Figure 1. Location of McDonald-Dunn Forest.


A companion project (Leavell, 1991) relates environmental variables to the vegetation structure and composition of the plant associations that were derived in this study for McDonald-Dunn Forest (Figure 2). These environmental variables are both simple and complex, and include measures of productivity. Together the two studies provide additional information for Forest managers.

Forest management agencies such as the USDA Forest Service use vegetation classification to provide managers with productivity, wildlife, and reforestation information (Hemstrom \& Logan, 1986; Halverson et al., 1986, Hemstrom et al., 1987). Plant associations also provide information to evaluate resource condition, and to predict response to management (Hall, 1989).

Available USDA Forest Service plant association guides for this area (Hemstrom and Logan, 1986; Hemstrom et al., 1987) do not cover the floristic composition or environment of McDonald-Dunn Forest. These classifications do not apply to McDonald-Dunn Forest. McDonald-Dunn Forest has a drier climate than the Forest Service lands. Other classifications in the Oregon Coast and Cascade Ranges (Halverson et al., 1986; Dyrness et al., 1974) also do not adequately describe the communities in McDonald-Dunn Forest.

Figure 2. Location of McDonald-Dunn Forest in relation to the Siuslaw and Willamette National Forests.


## OBJECTIVES

There were two objectives to this study.
The first objective was to develop and describe plant associations for the upland forest areas based on vegetation composition and cover in the Forest. This was accomplished through classification procedures.

The second objective was to quantify diversity within the plant associations. Measures of diversity such as; species richness, heterogeneity, evenness, and vertical structure (including snags) were calculated within plant associations.

## LITERATURE REVIEW

There are two broadly conceived research methods dealing with the understanding of the relationships of plant communities to one another and to the environment. These are classification and ordination (also called gradient analysis) (Whittaker, 1973).
"Classification groups communities into classes that may be characterized and treated as discontinuous with each other. Gradient analysis studies vegetation in terms of gradients of environment, species populations, and community characteristics in relation to one another. (Whittaker, 1973)."

The primary objective of this study was to organize vegetation data collected on McDonald-Dunn Forest into
units, plant associations, by means of classification. The companion study (Leavell, 1991) used indirect gradient analysis to determine relationships between this vegetation (plant associations) and the environment.

A literature review on classification was conducted to accomplish three things: 1) to gain a historical perspective on the development of classification; 2) to determine which method of classification to use for this study; 3) to determine if any vegetation classifications have been previously done in McDonald-Dunn Forest.

Krajina (1959) synecologically classified forests at three levels: environmental, biocoenotic, and ecosystematic. According to Krajina, environmental classifications were sometimes limited to only macroclimatic differences. Cowles (1899), Clements (1902), and Tansley (1920) were some of the earliest ecologists to use this approach in the concept of plant formation (Krajina, 1959). Krajina cited Svoboda (1949) in saying this method of classification is artificial and incomplete because of a lack of understanding of the real effects of ecological action (reaction or coaction), though it provides good information on significant characteristics of the environment.

Biocoenotic classification is dominated by a phytocoenotic approach and is usually independent of environmental information (Krajina, 1959). According to

Krajina, this method of classification was initiated by Cajander (1909), Ilvesalo (1920), and Lakari (1920), all ecologists from Finland. The Zurich-Montpelier school, represented by Braun-Blanquet was also part of the phytocoenotic method of classification. Krajina cited Dansereau (1957) in saying the Zurich-Montpelier school:

> "invokes the floristic composition first and places almost exclusive emphasis upon it, trusting, as it seems, that associationcharacteristics embody the very ecologicalpedological-climatic factors which determine them".
> The ecosystematic or biogeocoenotic classification method combines both floristic and environmental classification methods (Krajina, 1959). Odum (1953), Woodbury (1954), and Oosting (1956) believe that the crux of ecological thinking is the holocoenotic (floristic and environmental) point of view (Krajina, 1959).

Shimwell (1971) reviewed the historical development of the units of classification. According to Shimwell, the concept of the plant association is "one of the oldest in plant geography, even pre-dating use of the term 'ecology' ". Shimwell cited Humboldt (1805) as being the first to use the term. Shimwell divided different trends in vegetation classification into three traditions: the Zurich-Montpelier Tradition, the Northern Tradition, and the English Tradition.

The Zurich-Montpelier Tradition used the term association as "the fundamental vegetation unit characterized by physiognomic dominants" (Shimwell, 1971). This definition was adopted from Flahaut (1893). Shimwell cited Brockman-Jerosch (1907) as being the first to advocate classification of vegetation by dominant indicator species, but also by constancy as a basis for characterization. "Constant" species were those species occurring in better than $50 \%$ of samples in an association. The Zurich-Montpelier Traditionalists encouraged the 1910 International Botanical Congress to adopt the following definition of association (Shimwell, 1971):
"An association is a plant community of definite floristic composition, presenting a uniform physiognomy, and growing in uniform habitat conditions."

According to Shimwell, Braun-Blanquet (a leading representative of the Zurich-Montpelier Tradition), put increasing importance on vegetative classification while placing environmental influence into the background. Shimwell said the term "differential" species was initiated by Braun-Blanquet and Jenny (1926). Differential species applied to vegetation units lower than the level of plant association.

The English Tradition, according to Shimwell, combines British and American ecological influence. Pound and Clements (1898) adhered to the formation concept. Shimwell referred to Drude's (1896) definition of formation:
"... (a) any principal association which has found its natural termination in itself, (b) which consists of biologically related life forms, (c) (and) which is confined to similar substrata...This association...has reached its climax development." (Shimwell, 1971).

According to Shimwell, Clement's work became overshadowed by the continuum approach of Whittaker (1962). The Northern Tradition started with the Scandinavian concept of ecological series of communities along environmental gradients (Norrlin, 1870; Cajander, 1903). The Northern Tradition gradually merged in conceptual philosophy with the Zurich-Montpelier Tradition. Whittaker (1973) also gave a history of classification similar to Shimwell's, but with the addition of the Russian Tradition and a separate American Tradition. According to Whittaker, early classifications completed by the Russian Traditionalists were mainly physiognomic. Sukacev (1928) and Alekhin (1936) were early community ecologists who arranged communities in series along environmental gradients. According to Whittaker, the Russian Traditionalists mainly view forest communities as landscape units or biogeocoenoses (Sukacev, 1945).

Whittaker (1973) also wrote of the development of the American Tradition with the debate of the "unit" versus the "continuum" concept. Cowles (1899) and Clements (1905) advocated vegetative organization made up of discrete units of similar vegetation (associations). According to

Whittaker, these units were climax communities adapted to the "climates of geographic regions". The American Tradition developed from the unit concept of vegetation organization to the continuum concept, initially advocated by Gleason (1926). The continuum concept places vegetative species independently along environmental gradients (Gleason, 1926). Whittaker (1967), Curtis (1959), and MacIntosh (1967) are all proponents of the continuum concept of community analysis, and all representatives of the American Tradition.

Community ecologists have applied plant community classification in the Pacific Northwest (Daubenmire, 1952; Hall, 1973; Pfister et al., 1977; Hemstrom et al., 1987; and many more). These classifications place plant communities into units such as habitat types or plant associations. Community ecologists in the Pacific Northwest are using both classification and gradient analysis to discover and describe plant associations (Atzet, 1978; Hall, 1989).

Historical insight was important to understand the role of classification in this study. A further review of more recent literature was done to decide which classification method would be used in this study.
"Classification techniques used in community ecology may be considered in three groups : table arrangement, hierarchical, and nonhierarchical classification (Gauch,
1982). Braun-Blanquet (1932) initiated the table arrangement method. The table arrangement approach orders samples-by-species data by placing samples and species into an order that best illustrates community organization. Similar species listed are placed together, dissimilar species are placed apart. Braun-Blanquet based classification on the differential species in the communities. Whittaker (1973) said the Braun-Blanquet method:
"...is the most widely applied and most effectively standardized of all approaches to classification, and has been adapted to diverse kinds of vegetation."

Although this method is widely used, it has the following limitations:

1. Ecologists need to be trained for the method;
2. It is fairly subjective; and
3. It is not suited for large data sets.

Gauch (1982) stated nonhierarchical classification places similar samples or species into clusters. These clusters demonstrate no inherent relationships between each other. Gauch further suggested nonhierarchical classification should be used as an initial clustering for large data sets to reduce outliers and redundancy.

According to Gauch (1982), hierarchical classification puts similar samples or species into groups (as in the nonhierarchical method), but the groups are also arranged
hierarchically. The hierarchy indicates relationships among the groups. Gauch described three methods of hierarchical classification: monothetic divisive, polythetic agglomerative, and polythetic divisive.

The monothetic divisive approach starts with all plots in a single cluster and then divides them into groups based on presence or absence of a single species (monothetic $=1$ species). Polythetic means information on greater than one species is used. In the polythetic agglomerative method, information on more than one species is used. It starts out with each plot in its own cluster and systematically links the plots together on the basis of similarity or other criterion. The polythetic divisive method also uses information on more than one species. The plots start out in one cluster and are subsequently subdivided into groups.

Orloci (1966) said monothetic classifications have the disadvartage of producing uninformative subdivisions on the basis of unimportant attributes. Madgwick and Desrochers (1971) used a monothetic association-analysis method to classify forest vegetation of the Jefferson National Forest. One conclusion from their classification study was "...all monothetic classifications suffer from problems arising from divisions on the presence and absence of a single (monothetic) characteristic".

Lambert et al., (1973) developed two polytheticdivisive methods for hierarchical classification. They
said the only polythetic methods in use as of their writing had been agglomerative. They went on to say there are "...theoretical advantages in that all the available information is used to make the critical topmost divisions".

Hill et al., (1975) used a divisive polythetic method of classification to classify native pinewoods in Scotland. According to the authors, agglomerative methods of classification can be "...strongly dependent on the way in which stands are clustered at the lower levels." Divisive methods have an advantage in using the overall structure of the data set initially, with higher levels of the hierarchy being insensitive to the lower levels. The authors also stated that the monothetic divisive method of associationanalysis "makes far too many misclassifications". The authors developed a polythetic divisive classification method based on the iterative algorithm technique used in the ordination method of reciprocal averaging (Hill, 1973). The method was called indicator species analysis (Hill et al., 1975).

Hill (1979b) modified the indicator species analysis program to produce a two-way indicator species analysis program (TWINSPAN). This FORTRAN program differed from the indicator species program by the following:

1. The program first constructs a classification of the samples, then uses this classification to obtain a classification of species:
2. The two classifications (species and samples) are used together to produce a two-way table which illustrates the synecological relationships of the species.

The TWINSPAN program creates a "tabular matrix arrangement which approximates the results of the BraunBlanquet tablework" (Gauch, 1982). TWINSPAN incorporates two of the three basic methods of classification. It is hierarchical and includes a tablework arrangement. Gauch also said TWINSPAN is objective as compared to the subjectivity of the Braun-Blanquet tablework method.

From the literature review it was determined that the hierarchial polythetic divisive method and the program TWINSPAN would be used for classification in this study. Ecologists in the Pacific Northwest such as Hemstrom (1990), Atzet (1990), Smith (1990), and Diaz (1990), all personal communication, have used or are using the TWINSPAN program for vegetation classification.

The literature was also consulted to determine whether or not there were any existing classifications for the McDonald-Dunn Forest, or if any of the classifications for the surrounding country could be applied to this forest. Literature was reviewed for Tsuga heterophylla, Abies
grandis, and Pseudotsuga menziesii communities in Western Oregon.

Hall and Alaback (1982) surveyed vegetation on McDonald-Dunn Forest and developed a checklist if vascular plants. West (1964) mapped vegetation on McDonald-Dunn Forest. West's project was a classification, but based only for trees. West devised an elaborate mapping system and map for the Forest illustrating shrub, forb, and grass cover for taxa with greater than $5 \%$ cover. Unfortunately, neither the Forest nor West (personal communication) could locate the maps or any of the original data in 1989.

A successional study of McDonald-Dunn Forest was completed by Sprague and Hansen (1946). This study concentrated on arboreal vegetation rather than all strata of vegetation. Sprague and Hansen provided insight into succession of tree species on the Forest. Their study indicated a "successional trend of white oak to Douglas-fir followed possibly by a climax forest of lowland white fir (sic, Abies grandis) or a Douglas-fir lowland white fir association" (Sprague and Hansen, 1946).

Sabhasri and Ferrell (1960) did a study "to determine the effects of some environmental variables on the species, numbers, degree of cover, age, and growth of shrub species on south slopes in McDonald and Dunn Forests." Sabhasri and Ferrell's work provided some information on succession for a few shrub species, but not on classification.

Bigleaf maple, poison-oak, hazel, snowberry, and wild rose were studied. It was found that bigleaf maple, hazel, and snowberry had greater cover in openings than under the canopy. Poison-oak decreased in percent cover when a stand is opened. "Poison oak could very well be considered a climax community shrub on south slopes in this area" (Sabhasri \& Ferrell, 1960). Wild rose showed no significant reaction to openings in the canopy. There was no existing plant association classification for McDonald-Dunn Forest prior to this study. Vegetation studies in the Coast Range have been done by Juday (1976), Merkle (1948), Anderson (1967), Bailey (1966), Hemstrom and Logan (1986), and Thilenius (1968). I have reviewed these studies and vegetation studies in the Oregon Cascade Range (Hemstrom et al., 1987; Halverson et al., 1986; Topic et al., 1988; Means, 1980; and Dyrness et al., 1974. From review of the above vegetation studies it was determined that McDonald-Dunn was locationally, climatically, and vegetationally different from these existing classifications. There were very few Abies grandis associations in these classifications.

## THE STUDY AREA

The location, climate, soils, and land use history of McDonald-Dunn Forest contribute to its unique complex of plant communities.

## LOCATION

The study area is the McDonald and Paul M. Dunn Research Forest. The Forest is owned and managed by oregon State University's College of Forestry. The Forest consists of approximately 11,000 acres of predominantly forested land on the western edge of the Willamette Valley in Oregon, and on the eastern foothills of the Coast Range (Figure 3).

McDonald-Dunn Forest is in Townships 10 and 11 South, and Range 5 West, Willamette Meridian. It lies west of U.S. Highway 99 just to the north of Corvallis. The Forest occupies a ridge system that projects eastward into the Willamette Valley (Figure 3). In general, the streams and ridges extend northwest and southeast from the main ridge (Sprague and Hansen, 1946). The Forest is somewhat isolated topographically from the rest of the Oregon Coast Range, residing in the rain shadow created by it. McDonald-Dunn Forest is in the "Valley Margin Zone" as defined by Juday (1976) (Figure 4).

Figure 3. McDonald-Dunn Forest shown in relation to the Willamette Valley.


Figure 4. Valley Margin Zone'.


## CLIMATE

The climate of the McDonald-Dunn Forest is different than adjacent forests. The forests to the west have a greater coastal influence, with greater annual precipitation. McDonald-Dunn Forest is in the rain shadow of the Coast Range, receiving 100 to 150 cm of rain annually (Hall and Alaback, 1982), while the heart of the Oregon Coast Range receives 150 to 300 cm annually (Franklin and Dyrness, 1984). The Forest's climate is more influenced by the drier Willamette Valley climate than the typically wet Oregon Coast Range climate.

The macroclimate is summer-dry and winter-wet (Waring and Franklin, 1979). Precipitation occurs only occasionally in the summer. Rain gear was worn only once during the June-October field season in which the data for this study were collected. Most precipitation occurs in the winter.

## SOILS

The soils of this study area have been intensively surveyed (Rowley and Jorgensen, 1983). Thirteen soil series were identified and mapped. These are: McAlpina, Abiqua, Waldo, Jory, Price, Ritner, Witzel, Dixonville, Philomath, Dupee, Hazelair, Panther, and steiwer. Soil
series descriptions in Rowley and Jorgensen follow those of Knezevich (1975) in the soil survey for Benton County. Only nine of these series were sampled in this study. MCAlpina, Waldo, Dupee and Panther were not sampled because they are relatively uncommon on the Forest. Rowley and Jorgensen (1983) provided a description of the parent materials for the Forests soils:
"The parent material for most of McDonald and Dunn Forests soils is from the Siletz River Volcanics, a basalt formation. This rock formation is the foundation for the ridges and underlies most of the valleys. It underlies the Jory, Price, Ritner, Witzel, Dixonville, and Philomath series. The Flourney Formation (Tyee sandstone) is concentrated in the northwest corner of Dunn Forest and is the base for Dupee, Hazelair, Panther and Steiwer series. The wide flat drainage bottoms are recent alluvium which form the basis for MCAlpina, Abiqua and Waldo series."

The soil survey, completed in 1983, produced the soils map used in this study. Soil types were one criteria used in stratifying plot location.

## HISTORY

The McDonald-Dunn Forest has a complex history of settlement, ownership, use, and management. Prior to the migration and settlement of Anglos into the Willamette Valley in about 1845, Indians had burned the countryside repeatedly to facilitate game hunting and to maintain certain plants for food (Sprague and Hansen, 1946). Growth ring studies show that the country was frequently burned
since 1647, and less frequently burned after 1848 (Sprague and Hansen, 1946). Burning had kept the vegetation along the fringe of the Willamette valley in an early successional stage, where open, savanna-like Quercus garryana groves persisted. With Anglo settlements came fire suppression. With fire suppression came the slow conversion of the Quercus garryana savanna into Quercus garryana-Pseudotsuga menziesii forests. Abies grandis was also a component of these forests. Today there are fewer Quercus garryana forests and more Pseudotsuga menziesii/Abies grandis forests, although much of the Forest still contains Quercus garryana remnants.

Human settlement in the area brought cattle, sheep, pigs, horses, mules, and oxen into the foothills of Benton County in the 1850's (Jackson, 1981). Wild goats, originating from abandoned domestic herds, ranged the Forest in the 1930's. These foraging animals (along with deer) had an impact on vegetation, especially to young seedlings and small trees (Jackson 1981). Domestic livestock grazing has ceased on the Forest.

In 1953, the Oregon Game Commission organized a special deer hunt in the McDonald-Dunn Forest. One objective for this special hunt was to provide relief from deer browsing damage on Pseudotsuga menziesii seedlings (DeCalesta, 1985). Browsing pressure from deer still exists. Approximately 260 black-tailed deer were harvested
from the Forest annually between 1959 and 1982 (DeCalesta, 1985).

Although most of the funds for acquisition of the McDonald-Dunn Forest were donated by Mary J. L. McDonald, the tracts of land came from mixed previous ownership, and therefore have a variety of previous impacts. "Much of the land had been logged prior to being acquired by the school" (Jackson, 1981).

In 1989 Marvin Rowley, McDonald-Dunn Forest manager from 1973 to 1987, wrote histories for all stands in the Forest, recollecting most activities occurring prior to and during his management regime. Rowley's histories are the best source of historical logging activities available for the stands sampled in this study.

Forest management has had an impact on the species composition of the Forest. Harvesting activities under different Forest managers varied. Harry Nettleton, forest manager from 1948-1959, was characterized as "custodial", and emphasized protection of forest resources (Jackson, 1981). William Davies (manager from 1959-1973) and Marvin Rowley (manager from 1973-1987) emphasized productivity and implemented thinning programs. Current management under Jeff Garver (1987 to present) is more intensive, with clearcuts becoming more frequent than in the past. Besides harvesting and homesteading, Forest lands were used for
many other activities such as the Civilian Conservation Corps, research, recreation, and military activities.

## METHODS

Data were collected on 115 plots during the field season beginning in June and ending in september of 1989. Plots completed in this study are called ecology plots. Location of ecology plots is found in Figure 5. The Forests tract, compartment, inventory plot number, and Township, Range, Section, and are identified for each ecology plot in Appendix 1.

## STAND SELECTION

Ecology plots were placed within stands representing the overall resource variability in the Forest (excluding riparian areas). An initial ground reconnaissance indicated potential plant community variability. Topographic maps were consulted for physiographic variability. The soils inventory completed on the Forest (Rowley and Jorgensen, 1983) provided possible soil series differences. The most current timber type map for the forest was also studied. Timber typing was done according to the type mapping system devised by the USDA Forest Service (USFS, 1962).

The ecology plots were stratified on the basis of community type, physiography, soils, and timber type. The sampling objective was to obtain representative plots in

Figure 5. Location of ecology plots in McDonald-Dunn Forest.

all combinations of these features. Stands selected met the following criteria:

1. Mature; oldest stands available on the Forest.
2. Relatively undisturbed; stands not disturbed within the last 10 years, or stands where understory vegetation did not reveal any recent disturbance.
The relatively small area encompassed by McDonald-Dunn Forest allowed sampling of all stands that met the above criteria. Approximately $25 \%$ of the Forest area was initially eliminated by stands clearly too young (sapling size or smaller) to be considered in this classification. Many more stands were eliminated because of apparent recent disturbance.

## ECOLOGY PLOTS

In this study, the plots are called ecology plots but are often referred to as just "plots". The Forests' timber inventory plots are referred to as inventory plots.

## LOCATING ECOLOGY PLOTS

Once a qualified stand was located, Forest inventory plots already in place were selected for sampling before the stand was entered. An inventory plot in the interior
of the stand was chosen. If this inventory plot did not fit desired criteria (relatively undisturbed, relatively homogeneous in vegetation, soils, and physiography) the next plot in the transect was visited and evaluated. We continued down the transect in this manner till an appropriate inventory plot was located. If this transect failed to provide an appropriate plot, plots on an adjacent transect were evaluated. Ecology plots were not located on ecotones (obvious changes in vegetation composition within a short distance) or within riparian areas. Riparian areas require a different sampling scheme than the one used for this classification.

Ecology plot centers were placed on existing Forest inventory plot centers. This was done for several reason: 1. Non-bias; Inventory plot centers were systematically "surveyed in" by Forest engineers. Plots occur as frequently as every 200 (1 plot/acre) feet along predetermined transects. 2. Relocation; Plots will be relocated and measured every 10 years by the Forest. Each inventory plot has two bearing trees with aluminum tags giving azimuth and distance to plot center. This assists in locating ecology plots in the future for successional or other vegetation studies. 3. Simplified data collection; Information on the trees (species, diameter, height, site index,
basal area, age, growth, etc) was available from the Forest's inventory data base and did not have to be measured. This facilitated timely collection of a sufficient number of ecology plots.

## ECOLOGY PLOT IDENTIFICATION

Ecology plots were circular, with an area of $5382 \mathrm{ft}^{2}$ ( $500 \mathrm{M}^{2}$ ) (uncorrected for slope), or a radius of 41 feet. Ecology plot centers were marked with 2 " X 2 " X 18" wooden stakes. An aluminum identification tag with ecology plot number, Forest inventory plot number, and date of establishment was nailed to the top of this stake.

Two black and white photographs and two color slides were taken from each plot center for a permanent record of the site. These photographs and slides were taken in the two opposing directions that best captured the floristic composition of the stand. Photographs and slides were taken with 35 mm cameras; photographs with a wide angle lens ( 28 mm ), slides with a 50 mm lens. The tripod for the cameras was placed approximately 4 feet above ground level directly over plot center. Occasionally the tripod could not be placed on plot center because a mature tree was too close, or blocked the best view of the stand. When this happened, the distance and azimuth that the camera was
offset from plot center was noted on ecology plot sheet 1. The photo-point record is on file at the Research Forest headquarters in Peavy Arboretum.

## ECOLOGY PLOT DATA

The four data sheets and a detailed description of data collected appear in Appendices 2 and 3. A short description of data collected on each plot sheet follows.

1) Ecology plot card 1 included information on plot location, physiography (slope, azimuth, microposition), vegetation structure, surface characteristics, average tree height and diameter, site index, stand density index, snags, and vertical complexity of vegetation. Total live basal area, average stand height, average site tree age, quadratic mean diameter, site index, and stand density index were obtained or calculated from the Forests' inventory data bank.
2) Ecology plot card 2 consist of percent canopy coverage of all trees, shrubs, forbs, and grasses (complete species list in Appendix 4). Percent canopy cover was ocularly estimated. Canopy cover is defined as:
"the percentage of ground covered by a polygon drawn around the extremities of the undisturbed canopy of a plant species. Individual canopy coverages are then summed to represent the total canopy coverage for that particular species. (Daubenmire, 1959)"

Tree species had three categories of canopy cover: trees <12 feet, 12-50 feet, and >50 feet in height. Heights were estimated.
3) Soil description were completed for the top 60 inches of the soil profile (less if the profile was shallow). A soils pit was dug to 18 inches, and the remainder of the 60 inches of soil was described from an auger core. Depth, color, texture, and percent coarse fragments of major horizons, were recorded. Soil series were determined from those described in the 1983 soil mapping guide for McDonald-Dunn Forests (Rowley and Jorgensen, 1983).
4) The historical commentary included observations of disturbance history, stand succession and structure, and degree of recovery from disturbance. Stand histories written by Marvin Rowley were also consulted.

## DATA STORAGE

Data from plot card 1 and the soil descriptions were entered in dBASE III (1985). Data from plot card 2, vegetation cover percents, were entered into a Quatro Pro spreadsheet (Quatro Pro Manual, 1987). Original plot sheets, photos, and slides are in the Research Forest office at Peavy Arboretum. The data collected in this study, with the exception of the historical commentary, was
entered into the Forest Science Data Bank (Stafford et al., 1984) in the College of Forestry's Forest Science Department.

## ANALYSIS

Analysis of the data collected for this study was accomplished in two parts: classification and diversity. Classification facilitated the primary objective of this study: developing and describing plant associations. The classification program used was TWINSPAN (Hill, 1979b). Once plant associations were derived, significant differences between them were tested. This verification test was accomplished through a non-parametric procedure called MRPP, multi-response permutation procedure, to test the hypothesis of no significant difference among the associations. Quantification of diversity includes measures of richness, heterogeneity (Shannon's diversity), evenness, and vertical structure (which includes snags).

## CLASSIFICATION

Species percent cover, as ocularly estimated on the $5382 \mathrm{ft}^{2}\left(500 \mathrm{~m}^{2}\right)$ field plots, was used for classification into plant associations. Since there were many ubiquitous species throughout the sampling area, percent cover was
more meaningful than presence/absence data. Plant association, as used here, primarily follows the 1910 International Botanical Congress definition:
"An association is a plant community of definite floristic composition, presenting a uniform physiognomy, and growing in uniform habitat conditions."

But the term association for this study also refers to potential climax. The aim is to be able to identify the same plant associations in the future when communities are closer to climax. Most plant communities in McDonald-Dunn Forest have not attained a climax state and are seral due to past disturbance and age. Even though most of the Forest communities are seral, it was important to highlight potential climax tree species in order to place the associations described in this study within the context of western Oregon classifications. The potential climax tree species for the purposes of this study is the most shade tolerant tree species present and reproducing successfully on the site.

Plant association classifications developed in the Pacific Northwest use climax tree species to identify series, and shrubs, forbs, or grasses to identify associations (Hemstrom and Logan, 1986; Halverson et al., 1986; Topik et al., 1988; Hemstrom et al., 1987). This study follows the same naming pattern. Hemstrom et al. (1987) defined plant associations as follows:
"After a relatively long disturbance-free period, only those plants which can grow and reproduce in competition with their neighbors remain. This long-term stable collection of plants is the plant association."

Communities sampled in this study are as close to the climax state as available, taking into account the full range of vegetative diversity found in the Forest.

A total of 117 species were identified on the 115 ecology plots. Hitchcock and cronquist (1973) were the taxonomic authority for nomenclature. Common names not listed in Hitchcock and Cronquist were found in either Gilkey and Dennis (1980) or Garrison et al. (1976). A complete species list with numerical codes, acronym, scientific and common names is in Appendix 4. Acronym names are from Garrison et al. (1976). Acronym names are usually represented by the first two letters of both the genus and species of a taxa. Species not identified in Garrison et al. were generally named by the first two letters of both the genus and species. A complete list of species and cover percent on each plot is in Appendix 5.

## Twinspan

TWINSPAN (Hill, 1979b) was the program used for classification analysis. Species cover data was formatted to be compatible with this classification program by a program called CONDENSE (Singer and Gauch, 1979). TWINSPAN
is a program in the Cornell Ecology Program series (Gauch, 1981). This program was a development of a previously published classification method called "indicator species analysis" (Hill et al., 1975). TWINSPAN is a hierarchial polythetic divisive method of classification. Being hierarchial facilitates the construction of a dichotomous key for the associations, which can be used to identify associations in the field.

The name TWINSPAN stands for Two-Way INdicator SPecies ANalysis. It is called two-way because it classifies both plots and species. It is called indicator species analysis because it identifies one or more species, called indicator species, that are diagnostic of each division created in the classification.

The basic method of TWINSPAN involves the division of three ordinations. The first ordination is called the "primary ordination". It involves reciprocal averaging and orders plots. The second ordination is called the "refined ordination". The refined ordination is produced by making a dichotomy of the plots in the primary ordination. The dichotomy of the plots is made through identification of differential species, or species that show preference to one side or the other of the dichotomy (because of habitat or environmental preference). The refined ordination is divided in such a manner until the desired number of levels in the dichotomy obtained.

The third ordination is called the "indicator ordination". This ordination is based on the most strongly differential species and is used for a succinct characterization of the dichotomy (Hill, 1979b). It is used to identify indicator species. The indicator species are used in the dichotomous key which can be used to identify associations in the field.

Twinspan constructs two-way tables by identifying differential species ${ }^{1}$. Differential species drive the classification. Species are ordered according to their ecological preferences within this table. "The table created by TWINSPAN is ordered to exhibit the relationship between species and samples as clearly as possible" (Hill, 1979b).

The species percent cover data file was systematically modified to derive a classification (through TWINSPAN) that revealed as much structure as reasonable within the data set. The first runs through TWINSPAN included all plots and species, and used default input parameters. One very important input parameter is the pseudospecies cut levels. Pseudospecies are defined as the quantitative equivalent of differential species (Hill et al., 1975). Pseudospecies cut levels determine the particular scale of cover used during the classification process. The TWINSPAN program

[^0]makes divisions (in the ordinations) by implementing assigned pseudospecies cut levels. The default pseudospecies cut levels are: 0, 2, 5, 10, 20. These pseudospecies cut levels relate directly to a five point percent cover scale:
\[

$$
\begin{array}{ll}
1=>0-1 \% & 4=10-19 \% \\
2=2-4 \% & 5=20-100 \% \\
3=5-9 \% &
\end{array}
$$
\]

These default species levels did not illuminate satisfactory pattern or organization in the data set, and other pseudospecies cut levels were used. To simulate the Braun-Blanquet (Westhoff and Maarel, 1973) classification method, cut levels were changed to 0, 5, 26, 51, and 76. These cut levels also did not illuminate satisfactory organization in the data. Differentiation in percent cover greater than 20 into a number of abundance levels was necessary to illuminate sufficient patterns in this data set. Many pseudospecies cut levels were tried.

The pseudospecies cut levels providing the most effective and useful TWINSPAN output were: $0,6,11,21$, 31, 41, 51, 61, and 76. These cut levels relate directly to species percent cover on this nine point scale:

$$
\begin{array}{lll}
1=>0-5 \% & 4=21-30 \% & 7=51-60 \% \\
2=6-10 \% & 5=31-40 \% & 8=61-75 \% \\
3=11-20 \% & 6=41-50 & 9=76-100 \%
\end{array}
$$

The nine point scale also directly relates to the values in
the body of the two-way table produced by TWINSPAN (Table 2, page 51). The values 1-9 represent the cover of each species within each plot in this table. The output produced from TWINSPAN based on these cut levels, derived vegetative groupings that felt intuitively accurate, yet were reached primarily objectively.

TWINSPAN analysis was also made giving rare species greater weight. This was done by changing pseudospecies cut levels to $0, .2, .6,2,6,11,21$, and 31 , or $0, .6,2$, $6,11,21,31,51$, and 76 . These cut levels did not reveal additional floristic distinction in the data. Giving rare species more weight as indicators could also make it more difficult for field crews to identify the plant associations.

After a few initial TWINSPAN runs it became apparent that all species were not necessary to derive a classification. A total of 68 species were used for the majority of the classification analysis. A list of species used in the classification analysis are listed in Table 1. Species not used in the classification were trees, undesirable plants, and species that occurred in fewer than three plots.

Tree species were not used in the final TWINSPAN analysis. With or without tree species, preliminary analyses produced two-way TWINSPAN tables with similar results. Removing trees from the analysis also allowed

Table 1. Species used in TWINSPAN classification ${ }^{1}$.

## shrubs

ACCI
AMAL
BEAQ
BENE
COCO
GASH
HODI
LOCI
LOHI RHPU
RHDI
RIDI
ROSA
RULA
RULE
RUPA
RUPR
RUUR
SAGL
SYAL
SYMO
VAPA

POMU
PTAQ
ACTR
ACRU
ADBI
ANDE
ARMA
ASCA
CASC
COLA
DIFO
DIHO
FRVE
GAAP
GATR
GOOB
HIAL
HYOC
IRTE

Acer circinatum
Amelanchier alnifolia
Berberis aquifolium
Berberis nervosa
Corylus cornuta Gaultheria shallon Holodiscus discolor Lonicera ciliosa

Lonicera hispidula
Rhamnus purshiana Rhus diversiloba Ribes divaricatum Rosa spp.
Rubus laciniatus
Rubus leucodermis
Rubus parviflorus
Rubus discolor
Rubus ursinus
Sambucus glauca
Symphoricarpos albus
Symphoricarpos mollis
Vaccinium parviflorum
vine maple
western serviceberry
Oregon hollygrape
Cascade hollygrape
hazel
salal
ocean-spray
western trumpet
honeysuckle
hairy honeysuckle cascara buckthorn
poison oak
straggly gooseberry
rose
cutleaf blackberry
black cap raspberry
western thimbleberry
Himalaya blackberry
trailing blackberry
blue elderberry
common snowberry mountain snowberry
red whortleberry

## FORBS

Polystichum munitum
Pteridium aquilinum
Achlys triphylla
Actaea rubra
Adenocaulon bicolor
Anemone deltoidea
Arenaria macrophyllum
Asarum caudatum
Campanula scouleri
Coptis laciniata
Dicentra formosa
Disporum hookeri
Fragaria vesca
Galium aparine Galium triflorum Goodyera oblongifolia Hieracium albiflorum
Hydrophyllum occidentale Iris tenax
western swordfern bracken fern
deerfood vanillaleaf
baneberry
pathfinder
wind-flower
bigleaf sandwort
wild ginger
Scouler's hairbell cutleaf golden thread Pacific bleedingheart
Hooker's fairybells
common strawberry
cleavers
sweetscented bedstraw
rattlesnake plantain
hairy hawkweed
western waterleaf
Oregon iris

Table 1. (cont.)
FORBS (cont.)

| L_AMU | Lactuca muralis | lettuce |
| :---: | :---: | :---: |
| LATHY | Lathyrus spp. |  |
| LIAP | Ligusticum apiifolium | lovage |
| MOSI | Montia siberica | Siberian montia |
| NEPA | Nemophilia parviflorus | small flowered nemophilia |
| OSCH | Osmorhiza chilensis | sweet mountain cicely |
| PRVU | Prunella vulgaris | common selfheal |
| SACR | Sanicula crassicaulis | snakeroot |
| SADO | Satureja douglasii | yerba buena |
| SEJA | Senecio jacobaea | tansy ragwort |
| SMRA | Smilacina racemosa | false solomon's seal |
| SMST | Smilacina stellata | stary false solomon's seal |
| STCR | Stellaria crispa | chickweed |
| SYRE | Synthyrus reniformis | spring queen |
| TEGR | Tellima grandiflora | fringecup |
| THOC | Thalictrum occidentale | western meadowrue |
| TRIA | Trientalis latifolia | western starflower |
| TROV | Trillium ovatum | Pacific trillium |
| VAHE | Vancouveria hexandra | inside-out flower |
| VECAC | Veratrum californicum caudatum | false helibore |
| VICIA | Vicia spp. |  |
| VIGL | Viola glabella | pioneer violet |
| VISE | Viola sempervierns | redwoods violet |

GRASSES (including grasslike plants)
BRSY Brachypodium sylvaticum false brome
BRVU Bromus vulgaris Columbia brome
CAREX Carex spp.
FEOC Festuca occidentalis western fescue

1 Hitchcock and Cronquist (1973) was the taxomonic authority for nomenclature.
more emphasis to be placed on understory vegetation, which more accurately characterizes sites, since tree species cover and presence have been affected by management operations. Only 16 out of 115 ecology plots were in stands that had no known timber management disturbance in the past. sixty-nine plots were in stands that were thinned once, 26 plots were in stands that were thinned twice, and 4 plots were in stands that were thinned three times (Rowley, 1989). In the stands sampled, thinnings removed an average of 5 to 8 MBF (thousand board feet) per acre (Rowley, 1989).

Weedy species were removed as they are transitory in nature and are not a natural part of the community. Species occurring in fewer than three plots were removed from the analysis because they created noise in the results; they did not add any additional interpretational value to the classification.

At one point in the analysis, ubiquitous species and species with less apparent indicator value were removed. This reduced the species list to 28 of the 68 species used in most of the TWINSPAN runs. This further reduced species list did not improve the classification, as classification results were essentially the same as with 68 species.

One hundred and eight of the 115 ecology plots were used in the final classification analysis. Seven plots ( $10,47,49,69,72,87$, and 114 ) were removed because they
were excessively disturbed. Final analysis through TWINSPAN was made with 108 plots, 68 species, and default values for all program options except pseudospecies cut levels. It may be helpful for future users of TWINSPAN to know that best the results were made with default options (except pseudospecies cut levels), and that the results are likely more subjective and repeatable than if more parameters had been varied from defaults.

## Associations

The TWINSPAN analysis was not the end of the classification process. TWINSPAN analysis produced plant groupings (plant associations) that retained borderline and misclassified plots. In other words, there were likely some plots that TWINSPAN classified into the wrong plant association. This was apparent by the fact that in successive TWINSPAN runs with slight modification in pseudospecies cut levels, there were some plots that jumped back and forth between associations. Percent similarity between plots was used to determine within which association these difficult plots would be located. These plots were placed in the plant association that had plots with the highest percent similarity to the plot in question.

After each plot had its proper place in an association, each association was studied in more detail, and comparisons were made among associations. This process was facilitated by a constancy table of the plant associations. Constancy tables illustrate species composition relationships within and between plant associations.

## Verification of plant associations

Once the plant associations have been identified through TWINSPAN, a multi-response permutation procedure (MRPP) was used to test the hypothesis of no difference among the groups of plots within the plant associations. MRPP is a "non-parametric procedure for testing the hypothesis of no significant difference among two or more groups of entities" (McCune, 1987). The MRPP procedure used is part of a package of programs called PC-ORD (McCune, 1987). MRPP measures the distances between all pairs of plots within each association, and calculates a within-group average distance. Average distance was obtained using the Euclidean distance measure. Comparisons are made between these within-group averages and all other partitions (possible groupings) of the same number of plots in the same number of groups (Mielke et al., 1981). MRPP does not require data to be normally distributed, or to
have equal variances and covariances (McCune, 1987; Mielke et al., 1981). Normality and equal variances are required by many other statistical analyses, but rarely exists in community data.

Procedurally, MRPP requires each plot to be identified to its group, or association. The program then reports if the average between point distance (Euclidian distance) within each group (plant association), and tests the hypothesis of no significant difference among groups (Figure 7).

## DIVERSITY

After the plant associations were identified, it was possible to analyze their composition and structure. Some measures of diversity were used to do this analysis. Diversity is defined as: "The relative degree of abundance of (wildlife) species, communities, habitats, or habitat features per unit area" (Thomas, 1979). Management for diversity preserves viable populations of as many species as possible throughout a landscape. Vegetative and habitat diversity relates directly to the availability of ecological niches. Niches are habitats which supply factors necessary for the existence of an organism or species (Hanson, 1962).

Richness and equitability are the two components of heterogeneity, a measure of diversity. Richness equals numbers per unit area. Equitability is evenness in relative abundance of items per unit area (Westman, 1990). Three types of diversity are defined by Whittaker (1972). These are:

1) Alpha diversity - The number of species and the evenness of distribution of those species within a single habitat or community. This is micro-scale diversity, generally to be found in the area of a stand, a plot, or community, etc.
2) Beta diversity - The extent of differentiation of communities (or associations) along habitat (or environmental) gradients. This is between-community diversity.
3) Gamma diversity - A product of the alpha diversity of communities and the degree of beta differentiation among them. This is landscape or macro-diversity.

The terms alpha, beta, and gamma diversity have a broad acceptance in the field of ecology. These terms are used by Whittaker (1972), Moral and Flemming (1979), Noss (1983), and Schroeder (1987). There are two other types of diversity defined by Thomas (1979). These are:

1) Vertical diversity - The diversity in an area resulting from the complexity of aboveground vegetation stratification. This could be either on a micro-scale and
associated with alpha diversity or on a macro-scale and associated with beta diversity. Micro-scale vertical diversity can be found, for example, in a forest and takes into account all the various grass, forb, shrub, and tree strata. Macro-scale vertical diversity, for example, can be made up of different stands of different age and size classes spread throughout the landscape (Thomas, 1979). 2) Horizontal diversity - This is dispersion (juxtaposition) of vegetation over an area. An example is the various age and size classes of trees over the landscape. Horizontal diversity includes edge. The greater amount of edge, the higher the degree of horizontal diversity (Thomas, 1979).

This study concentrates on alpha and vertical diversity in mature coniferous forests. The Forest as a whole is represented by a wider range of alpha, beta, and gamma (landscape) diversity, which includes many different size classes of coniferous forest stands, hardwood stands, riparian areas, and meadows. Quantifying beta, gamma, and horizontal diversity of McDonald Forest was beyond the scope of this study.

Although there are many measures of diversity, the measures of diversity calculated in this study are species richness, heterogeneity (Shannon's diversity measure), evenness, and vertical structure. Species richness and vertical structure have a number of values affiliated with
them for each plant association, whereas heterogeneity (Shannon's diversity measure) and evenness are both single values. All of these measures are calculated as the average of individual plot measures within each plant association.

Species richness was calculated as the number of species that were encountered within each association, and as an average number of species per plot within each association. Average species richness was also calculated for each strata within plant associations. strata were divided into trees, shrubs, forbs, and grasses. These species richness values are measuring alpha diversity.

Shannon's diversity index is a measure of heterogeneity, involving species richness and equitability. The equation for Shannon's diversity index is:

$$
H^{\prime}=-\Sigma p_{i} \log p_{i}
$$

where,

$$
p_{i}=n_{i} / N_{i} ;
$$

that is, $p_{i}$ is the proportion of the total abundance occurring in species i. Logarithmic base 10 was used. Shannon's index was chosen because it best satisfies important criteria according to Elliott (1990). Shannon's diversity index is widely published (Elliott, 1990; Smith, 1980; Schroeder, 1987; and Brower et al., 1990). Evenness is calculated using Shannon's diversity index. It is
calculated as the ratio of the observed diversity index value to its maximum value. The evenness equation is:

$$
J^{\prime}=\text { Evenness }=H^{\prime} / H_{\max }^{\prime}
$$

where,

$$
\mathrm{H}_{\max }^{\prime}=\log \mathrm{s}
$$

where,

$$
s=\text { total number of species }
$$

Both of these values, Shannon's diversity index and evenness were calculated using the AID1 program (Overton et al., 1987). The three strata of tree species were added to include only measure for each tree species. Both the heterogeneity and evenness measures are measure of alpha diversity.

Analysis for vertical diversity was limited to the vertical component of vegetation structure. The average percent coverage of each vegetation strata (grass, forb, shrub, and tree strata) was calculated within each association. The tree strata was divided into three categories: >50 feet tall, 12-50 feet tall, and <12 feet tall. The shrub strata was also divided into in three categories: shrubs <2 feet tall, 2-6 feet tall, and >6 feet tall. Average forb and grass percent coverage was also calculated. Vertical profile diagrams were constructed to illustrate vertical structure for five height classes; $0-2,2-6,6-12,12-50$, and $50+$ feet. Evenness was also calculated using the percent cover of
vegetation in each of these five height classes for each association.

Snags were also considered part of the vertical structure of the Forest. Snag composition for the associations and for the plots overall were analyzed. Number, type, and size of snags were recorded on a 150-foot radius plot. Distance to the snag plot boundary was estimated. Three size classes of snags were defined. These were: 4-12" DBH, 12-21" DBH, and 21+" DBH. These size classes were also separated into two types, hardwood and conifer snags. A snag was defined as any dead tree >4" in diameter and $10^{\prime}$ tall. The average number of snags/acre in each size class and type was calculated for each association. "

An unbalanced analysis of variance and multiple means comparison (Fisher's Protected LSD) was used to identify significant differences in these averages among and between associations for all the measures discussed above. The statistical package SAS, General Linear Models procedure, accomplished this analysis (SAS Institute Inc., 1987).

Snag data were also analyzed independent of associations. Average number of snags per 100 acres in the size and type classes was calculated for the plots overall. Snag availability for some wildlife species on McDonald Forest was also analyzed.

A number of wildife species that are found in McDonald Forest use snags for nesting, breeding, feeding, and other purposes. Wildlife species on McDonald Forest that use snags were determined from a recent survey in a project titled "Comparisons of Terrestrial Vertebrate Communities and Tree Regeneration Among 3 Silvicultural Systems in the East-Central Coast Range, Oregon" (McComb and Chambers, 1989) (Appendix 6). It was determined that 15 bird and 3 mammal species from this list use snags for breeding, feeding, or resting (Neitro et al., 1985). Appendix 7 lists these species and the suggested size class of snag suitable for nesting for those species that use cavities (Neitro et al., 1985).

Specific snag requirements for six species of woodpeckers on the Forest were calculated by a program called the Snag Recruitment Simulator, or SRS (Marcot, 1989). These species are: downy woodpecker (Picoides pubescens), red-breasted sapsucker (Sphyrapicus ruber), hairy woodpecker (Picoides villosus), northern flicker (Colaptes auratus), red-breasted nuthatch (Sitta canadensis), and the pileated woodpecker (Dryocopus pileatus). SRS calculates the density and size classes of snags required to support populations of these woodpeckers (Marcot, 1989). This program was run to compare snag requirements for these species against the snags that were recorded in this study.

## RESULTS

Determination of plant associations on McDonald-Dunn Forest was done through TWINSPAN classification analysis and percentage similarity between plots. These plant association groupings were verified to be significant through a multi-response permutation procedure. Diversity measures; species richness, heterogeneity (Shannon's diversity), evenness, vertical structure, and snag composition results are presented. The associations are described by their location, species composition, and diversity measures. A picture representing a typical plot within each association is also included with plant association descriptions.

## CLASSIFICATION

Six plant associations were derived with the aid of TWINSPAN analysis and a percentage similarity matrix for the plots. The complete two-way table for the final TWINSPAN results is in Appendix 8. A simplification of the two-way table for this TWINSPAN run is shown in Table 2. The species listed in this table (Table 2) are mostly differential species, of which some are indicator species. Acronyms for species names are listed along the left-hand

Table 2. Simplified TWINSPAN classification using

| Scale for cover percent values in body of table |  |  |
| :--- | :--- | :--- | :--- |
| $1=$ |  |  |
| $2=60-5 \%$ | $4=21-30 \%$ | $7=51-60 \%$ |
| $3=11-20 \%$ | $6=41-50 \%$ | $9=76-100 \%$ |

Table 2 .

PLOT NUMBER


PLOT CLASSIFICATION
margin, and plot numbers along the top of Table 2. The acronym names for the associations are listed at the bottom (Table 2). Full association names and acronym name are listed in Table 3. The values within the chart are categories for species cover values. These categories are defined with the legend (Table 2).

The divisions for the associations were determined through the hierarchy shown along the bottom margins of the two-way table. The hierarchy is depicted in binary notation. The first division in the hierarchy is between plots 70 and 17 . The plots on the right-hand side of this division (all $1^{\prime}$ s) are then divided between plots 22 and 113. The plots on the left-hand side of the first division (all $0^{\prime} s$ ) were at the same time divided between plots 100 and 82. Divisions continued in this fashion. Determinations of plant associations were formed from these hierarchial divisions.

Five plant associations came from the third level of hierarchial divisions, one from the second level. ABGR/BRSY was formed from the second level. Further division of this plant associations was not practical for the intent of this study. Of the five plots that were in the one side of the division that broke off from the main ABGR/BRSY association, one was a misclassified plot, one had $5 \%$ cover of BRSY and another $3 \%$. It was felt that the difference of these plots from the main ABGR/BRSY plant association would not be
distinguishable on the ground. That the groupings be easily identifiable on the ground was an objective of this study to make it useful to management. The 5 plots that were in this division also did not fit into the environmental gradient by Leavell (1991).

The division between the ABGR/ACCI-GASH and TSHE/ACCIGASH plant associations does not follow a strict hierarchial division. At the third level of divisions plot numbers 113 and 115 are separated out into their own grouping. A grouping of two plots was not desirable or practical. A comparison of percentage similarity (discussed further below) between these two plots with the plots in both TSHE/ACCI-GASH and ABGR/ACCI-GASH showed that they had higher percent similarity to the plots in the ABGR/ACCI-GASH plant association.

The plots within each plant association are listed in Table 3. The plots listed for each plant association in Table 3 do not match the number of plots for each association in Table 2. Some borderline or misclassified plots were put into a different plant association than shown in Table 2 because they showed higher percentage similarity to the plots in the groupings listed in Table 3 than the plots in the groupings listed in Table 2. For example, plot 46 is in the ABGR/BRSY plant association in Table 2, but is in the ABGR/RUUR-RHDI plant association in Table 3, the final groupings used for this classification. Plot 46 was
considered a misclassified plot within the ABGR/BRSY association in the TWINSPAN analysis. Plot 46 had higher percent similarities to the plots in the ABGR/RUUR-RHDI plant association than in the ABGR/BRSY plant association. Plot 46 also looks out of place in the ABGR/BRSY plant association because it has less than five percent cover of Brachypodium sylvaticum, whereas the other plots in that association have substantially higher percent cover of this species.

Table 3. Plant associations described on McDonald-Dunn Forest.

| Plant association | NUMBER OF PLOTS IN ASSOCIATION | PLOTS IN ASSOCIATION |
| :---: | :---: | :---: |
| Isuga heterophylla/Acer circinatum-Gaultheria shallon TSHE/ACCI-GASH | 6 | 107, 108, 109, 110, 111, 112 |
| Abies arandis/Acer circinatum-Gaultheria shallon ABGR/ACCI-GASH | 12 | $\begin{aligned} & 21,22,28,35,62,63,73 \\ & 74,81,103,113,115 \end{aligned}$ |
| Abies grandis/Disporum hookeri-Thalictrum occidentale ABGR/DIHO-THOC | 21 | $\begin{aligned} & 4,17,20,23,24,26,29,30,34, \\ & 36,37,53,61,64,67,68,78,84, \\ & 97,98,102 \end{aligned}$ |
| Abies qrandis $/ \frac{\text { Polystichum munitum }}{\text { ABGR/POMU }}$ A | 16 | $\begin{aligned} & 7,11,12,14,18,19,27,39,41, \\ & 50,54,65,70,71,75,77 \end{aligned}$ |
| Abies grandis/Rubus ursinus-Rhus diversiloba ABGR/RUUR-RHDI | 27 | $\begin{aligned} & 13,15,16,33,38,40,44,45,46, \\ & 48,51,52,55,56,57,58,59,60, \\ & 66,76,79,80,82,83,85,86,101 \end{aligned}$ |
| $\frac{\text { Abies grandis } / \text { Brachypodium sylvaticum }}{\text { ABGR/BRSY }}$ | 26 | $\begin{aligned} & 1,2,3,5,6,8,9,25,31,32,42, \\ & 43,88,89,90,91,92,93,94,95, \\ & 96,99,100,104,105,106 \end{aligned}$ |

There are one or two indicator species for each of the divisions (plant associations) in the TWINSPAN classification. Indicator species are species indicative of a particular association. Species do not have $100 \%$ fidelity to be indicators. Indicators can be species that occur in all plots, but occur at a higher abundance levels (pseudospecies cut levels) in a specific division of the hierarchy. These species have been highlighted in the simplified TWINSPAN run (Table 2). The main indicator species for the ABGR/BRSY plant association is Brachypodium sylvaticum (see purple highlight). The indicator species for the ABGR/RUUR-RHDI plant association are Rubus ursinus and Rhus diversiloba (see orange highlight). These indicator species are the same species by which the plant associations are named. The other four associations are named similarly, by indicator species.

The two-way table also gives a species classification, located along the right margin (Table 2). The species classification is formed through a similar process as the plot classification. Species classification was not used in this study.

A dichotomous key for field identification of associations was made using the indicator species in the TWINSPAN analysis (Figure 6). Instructions on using the key are included at the bottom of the key. This key includes tree species. Tree species are used to identify

Figure 6. Key to plant associations on McDonaldDunn Forest.
1A. Western hemlock present and reproducing successfully ${ }^{1}$
TSHE/ACCI-GASH
1B. Western hemlock not present ................................ 2


4A. False brome $>10 \%$ cover ........................... ABGR/BRSY
4B. False brome not as above .................................... 5
5A. Sword fern $>40 \%$ cover .............................. ABGR/POMU
5B. Sword fern not as above ..................ABGR/RUUR-RHDI

1 Tsuga heterophylla is not listed in Table 6, but is used in the key. Even though tree species were not used in the classification, Tsuga heterophylla is the best indicator species for this plant association.

## Instructions for key

1. This key is not the classification. Plant association must always be confirmed with written description.
2. User must follow the key from 1A and comply with all conditions as successive steps in the key are followed.
3. It is possible that not every location will key precisely to one plant association. When the key does not work as written, take the route that most closely fits the vegetation, and read descriptions of associations.
4. If area appears to not accurately fit an association, list the association(s) that fits best, or note the reasons why one does not fit. This will aid in future refinement of the classification and/or key.
5. If user gets to the ABGR/RUUR-RHDI plant association by default and the description does not fit, go back through the key and be more flexible using cover standards. If problem persists, read descriptions and find the association that fits best and note as in 4 above.
"series", and shrubs, forbs, or grasses added to define the associations. Tsuga heterophylla and Abies grandis were found to be the climax tree species on McDonald-Dunn Forest, and name the series of plant associations developed in this study.

Tsuga heterophylla series is identified by the presence of Tsuga heterophylla successfully reproducing in an isolated area in the Forest. Most ecology plots had Abies grandis present and successfully reproducing, indicating most of the forest (that portion without presence of Tsuga heterophylla) belonged in the Abies grandis series. A Pseudotsuga menziesii plant association was not identified.

The only plant association in the Tsuga heterophylla series is Tsuga heterophylla/Acer circinatum-Gaultheriashallon. The other 5 plant associations described here are in the Abies grandis series.

Plant associations have been determined to lie on an environmental gradient (Leavell, 1991). The Tsuga heterophylla/Acer circinatum-Gaultheria shallon plant association is at the moist end of the environmental gradient. The Abies grandis/Brachypodium sylvaticum plant association is on the dry end of the environmental gradient. The other four plant associations are placed between these two associations along the environmental gradient. The list of plant associations (Table 3) is
arranged according to this environmental gradient as determined by Leavell (1991).

A Constancy table for associations is found in Table 4. Two numbers are listed for each species and association. The first number is constancy. Constancy is the percentage of plots in the association in which the species occurs. The second number is average percent cover for the plots within the association in which the species occurred. Species are listed alphabetically within four strata: trees, shrubs, forbs, and grasses. Constancy tables were used in describing plant associations, as well as making comparisons of the associations described in this study to other classification studies in western Oregon.

Table 4. Constancy ${ }^{1}$ and average percent cover ${ }^{2}$ of plants in plant associations on McDonald-Dunn Forest.


Shrub species

| ACCI | 100 / 23.8 |  | 75 / 39.8 | 14 | 1.7 |  |  | 41 | 6.0 | $4 /$ | 3.0 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| AMAL |  |  | 17 / . 3 | 9 | 1.6 | 37 | . 2 | 301 | . 1 | $38 /$ | . 8 |
| BEAQ |  |  | $17 / 1.2$ | 24 | 9.0 | 19 | . 2 | $41 /$ | . 6 | 15 / | . 3 |
| BENE | $100 /$ | 8.5 | $75 / 22.1$ | 43 | 4.3 | 19 | . 5 | $11 /$ | . 8 | $8 /$ | . 5 |
| COCO | 501 | 3.0 | 100 / 15.3 | 100 | 12.0 | $94 /$ | 11.0 | $96 /$ | 13.7 | $100 /$ | 7.5 |
| GASH | 100 / | 7.8 | $75 / 15.6$ |  |  |  |  |  |  |  |  |
| HODI | 501 | 2.4 | 92 / 3.9 | 71 | 2.4 | 44 | 2.1 | 67 / | 3.3 | $31 /$ | 1.3 |
| LOCI |  |  | $25 / .2$ | 43 | . 3 | 31 | . 2 | $11 /$ | 1.4 | 19 / | . 1 |
| LOHI | $17 /$ | . 1 | $25 / 1.3$ | 24 | 1.2 | 56 | 1.1 | 67 / | 2.3 | 69 / | 1.0 |

[^1]Table 4. (cont.)

|  | $\begin{gathered} \text { TSHE/ } \\ \text { ACCI-GASH } \end{gathered}$ | $\begin{gathered} \text { ABGR/ } \\ \text { ACCI-GASH } \end{gathered}$ | ABGR/DIHO | ABGR/POMU | $\begin{gathered} \text { ABGR/ } \\ \text { RUUR-RHDI } \end{gathered}$ | ABGR/BRSY |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| Number of Plots | $\mathrm{N}=6$ | $N=12$ | $N=21$ | $N=16$ | $\mathrm{N}=27$ | $\mathrm{N}=26$ |

Shrub species (cont.)

| RHPU |  |  | 331 | . 7 | $33 /$ | . 4 | 69 / | . 4 | 67 / | . 5 | $50 /$ | . 2 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| RHDI |  |  | $33 /$ | 1.4 | $57 \%$ | 5.8 | $100 \%$ | 6.4 |  | 15.0 | 100 / | 14.6 |
| RIDI |  |  |  |  |  |  | 121 | . 3 | 71 |  |  |  |
| ROSA | $100 /$ | 1.4 | $100 /$ | 3.00 | $100 /$ | 2.8 | 75 / | . 9 | 89 | 1.6 | 921 | 1.1 |
| RULA |  |  |  |  | $5 /$ | . 1 | 19 / | . 4 | $14 /$ | . 4 | 27 / | . 3 |
| RULE |  |  |  |  | $14 /$ | . 2 | 251 | 1.3 | $48 /$ | 2.0 | 81 | 3.5 |
| RUPA | $67 /$ | . 2 | 83 / | 3.3 | $71 /$ | 2.1 | 50 / | 1.5 | $25 /$ | 2.8 | $31 /$ | . 5 |
| RUPR |  |  |  |  |  |  | 19 / | 1.3 | 191 | 3.4 | 81 | 2.8 |
| RUUR | $100 /$ | . 5 | 1001 | 3.2 | $90 /$ | 5.3 | 100 / | 7.9 | $92 /$ | 21.0 | 88 / | 7.0 |
| SAGL | 17 / | . 1 | 81 | 4.0 |  |  |  |  | 4 | 3.0 |  |  |
| SYAL | $50 /$ | . 1 | $83 /$ | 5.0 | $90 /$ | 5.7 | 87 / | 5.9 | $96 /$ | 5.9 | $88 /$ | 3.3 |
| SYMO |  |  |  |  | $5 /$ |  | 25 / | 1.1 |  |  | 15/ | . 8 |
| VAPA | $50 /$ | 1.0 | 25 / | . 4 |  |  |  |  |  |  |  |  |


| Forb species |  |  |  |  |  |  |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| POMU | 100/18.3 |  | 100 / 10.5 |  | 100/14.2 |  | $100 / 57.2$ |  | 100 / 12.8 |  | $\begin{aligned} & 92 / \\ & 58 / \end{aligned}$ | 6.14.5 |
| PTAQ |  |  | 50 / | 2.7 | 71 | 3.0 | 44 | 3.1 | 74 | 3.2 |  |  |
| ACTR | 33 | . 3 | $83 /$ | 4.2 | 100 | 9.1 | 44 | 2.4 | 26 | 1.5 | 15 | 1.2 |
| ACRU |  |  | $33 /$ | 1.8 | 43 | 5.7 | 6 | . 1 |  |  | $11 /$ | 2.7 |
| ADBI |  |  | 83 / | 1.3 | 100 | 3.6 | 100 | 1.8 | $100 /$ | 2.0 | $81 /$ | 2.8 |
| ANDE | 50 | . 1 | $58 /$ | . 8 | 76 | . 9 | 25 | . 1 | 30 / | . 1 | $11 /$ | . 4 |
| ARMA |  |  | 58 / | . 6 | 81 | 2.1 | 37 |  | $52 /$ | 2.9 | 19 / | . 3 |
| ASCA | 17 | . 1 | 81 | . 1 | 5 | 5.0 |  |  |  |  |  |  |
| CASC | 17 |  | $100 /$ | . 8 | 76 | . 8 | $31 /$ |  | $33 /$ | . 7 | $27 /$ | . 3 |
| COLA | 67 |  | $17 /$ |  |  |  |  |  |  |  |  |  |
| DIHO |  |  | $100 \%$ | 2.2 | 95 | 2.7 | $25 /$ 19 | 2.3 .9 | 111 | 4.2 1.5 | 41 | . 1 |
| FRVE | 17 | . 1 | 25 / | . 1 | 48 | . 6 | 56 | . 6 | 67 / | 1.5 | 46 / | . 4 |
| GAAP |  |  | $33 /$ | . 8 | 24 | . 2 | $31 /$ | . 3 | 301 |  |  |  |
| GATR | 67 | . 2 | 92 / | 1.1 | $90 \%$ | 1.6 | $100 /$ | 1.9 | 100 / | 3.6 | $92 /$ | 1.1 |
| GOOB |  |  | $50 /$ |  | 62 | . 2 | 501 | . 3 | 63 / | . 3 | $58 /$ | . 3 |
| HIAL |  |  | 42 / |  | $43 \%$ | . 3 | 50 / | . 3 | $37 /$ | . 2 | $35 \%$ | . 1 |
| HYOC |  |  |  |  | 241 | 1.6 | 19 / | . 1 | $11 /$ | . 7 |  |  |
| IRTE |  |  |  |  |  |  | 61 | . 1 | $4 /$ | 2.0 |  |  |
| LAMU |  |  |  |  | 19 / | . 9 | $37 /$ | 4.3 | $33 /$ | 4.6 | $19 /$ | 1.6 |
| LATHY |  |  | 17 \% | . 3 | $57 /$ | . 6 | $31 /$ | . 1 | 26 / | . 2 | 191 | . 3 |
| LIAP |  |  | 17 / | 1.5 | $38 /$ | 2.4 | 25 / | 2.0 | 37 / | 1.7 | $35 /$ | 1.4 |
| MOSI | $17 /$ | . 1 | 25 / | . 2 | $57 /$ | 2.5 | 75 / | 2.2 | 55 / | 1.4 | $15 /$ | . 3 |
| NEPA |  |  |  |  | $24 /$ | . 1 | $31 /$ | . 1 | $7 /$ | . 1 |  |  |
| OSCH PRVU | 17 / | . 1 | 83 / |  | 100 / | 2.1 | $100 \%$ | 2.4 | $100 \%$ | 3.7 | $92 /$ | 1.6 |
| SACR |  |  |  |  |  |  | $6 /$ 19 | . 1 | 41 | . 1 | 15 / | .2 .1 |
| SADO |  |  | 251 | 1.0 | 331 | 2.1 | 69 / | 1.1 | 781 | 2.9 | $61 /$ | 2.0 |
| SEJA |  |  | $25 /$ | . 1 | 51 | . 5 | 19 / | . 1 | $33 /$ | . 2 | $11 /$ | . 1 |
| SMRA SMST | 501 | . 1 | $42 /$ | . 3 | $19 /$ | . 3 | $12 /$ | . 3 | 71 | . 3 | 41 | . 1 |
| SMST | 17 / | . 1 | 33 / | 2.5 | $43 /$ | 3.6 | 19 / | 1.7 | $22 /$ | 2.0 | 81 | . 5 |
| SYRE |  |  | 25 / | . 7 | 52 / | . 9 | 19 / | . 4 | 18/ | 1.2 | 41 | . 1 |
| TEGR | $17 /$ | . 1 | 17 / | . 6 | $43 /$ | 1.2 | 56 / | . 8 | 26 / | . 2 . | 41 | . 1 |
| THOC |  |  | $67 /$ | 2.1 | 90 / | 3.3 | 6 / | . 5 | 41 | . 5 | 41 | . 1 |
| TRLA | $83 /$ | . 4 | $100 /$ |  | $90 /$ |  | 69 / |  | 78 / | 2.7 | $31 /$ | 1.2 |

Table 4. (cont.)


## VERIFICATION OF ASSOCIATIONS

Multi-response permutation procedure (MRPP) results are illustrated in Figure 7. Each plant association is reported as a group with an average distance. Each distance represents a within-group average of pairwise distance measures (Zimmerman et al., 1985). MRPP results demonstrated significant differences (alpha < .05) among the groupings of plots (associations).

The average distance for the TSHE/ACCI-GASH association (group 6) is 25.4. This is the smallest average distance among the six associations. This shows that the TSHE/ACCI-GASH plots are more similar to each other than the plots within the other associations.

Figure 7. MRPP output for six plant associations.

> MULTI-RESPONSE PERMUTATION PROCEDURES (MRPP)

INPUT HAS 108 plots
THERE WERE 68 species
WEIGHTING OPTION: $\quad C(I)=n(I) / \operatorname{sum}(n(I))$
DISTANCE MEASURE = Euclidean

GROUP NUMBER 1 OF SIZE 26 HAS AN AVERAGE DISTANCE $=39.437039$ (ABGR/BRSY)

GROUP NUMBER 2 OF SIZE 21 HAS AN AVERAGE DISTANCE $=48.375329$ (ABGR/DIHO-THOC)

GROUP NUMBER 3 OF SIZE 16 HAS AN AVERAGE DISTANCE $=34.103703$ (ABGR/POMU)

GROUP NUMBER 4 OF SIZE 27 HAS AN AVERAGE DISTANCE $=42.853919$ (ABGR/RUUR-RHDI)

GROUP NUMBER 5 OF SIZE 12 HAS AN AVERAGE DISTANCE $=62.215145$ (ABGR/ACCI-GASH)

GROUP NUMBER 6 OF SIZE 6 HAS AN AVERAGE DISTANCE $=25.433365$ (TSHE/ACCI-GASH)

THE TEST STATISTIC IS $=\quad-39.184033$
THE OBSERVED DELTA IS $=\quad 42.992055$
THE EXPECTED DELTA IS $=\quad 60.220312$
THE VARIANCE OF DELTA $=\quad .19331449$
THE SKEWNESS OF DELTA $=\quad-.72922439$
PROBABILITY OF A SMALLER OR EQUAL DELTA $=\quad .00000000$

## DIVERSITY

## SPECIES RICHNESS

Total species richness values for associations as well as average per plot species richness by strata (tree, shrub, forb, grass) within each association are presented (Table 5). The six plant associations defined in this study differ significantly ${ }^{2}$ in species richness values.

Total species richness is lowest for the TSHE/ACCIGASH plant association which has a total of 38 species. The other plant associations have higher, similar total species richness values that range from 63 to 70 species.

The two associations on either end of the environmental gradient (Leavell, 1991), TSHE/ACCI-GASH and ABGR/BRSY, have the lowest average per plot species richness values (21.7 and 24.4). These two association values are significantly different from the other four associations which have higher and similar species richness values of 33.2 (ABGR/ACCI-GASH), 34.0 (ABGR/DIHO-THOC), and 30.2 (for both ABGR/POMU and ABGR/RHDI-RUUR).

[^2]Table 5. Species richness by plant association and strata.

| Plant Association | of plots in association | Total Species Richness in Association | Average Species Richness/plot' in association | Ave. Tree Richness/ plot | Ave. Shrub Richness/ plot | Ave. Forb Richness/ plot | Ave. Grass Richness/ plot |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| TSHE/ACCI-GASH | 6 | 38 | $\begin{aligned} & 21.7 \\ & (2.28)^{2} \end{aligned} C^{3}$ | $\begin{aligned} & 5.5 \\ & (0.76) \end{aligned}$ | $\begin{array}{ll} 8.0 & B \\ (1.15) \end{array}$ | ${ }_{(8.60)}^{7.8} \mathrm{C}$ | $\begin{aligned} & .3 \quad{ }^{\circ}{ }^{(0.236)} \end{aligned}$ |
| ABGR/ACCI-GASH | 12 | 63 | $\begin{array}{ll} 33.2 & A B \\ (5.32) & \end{array}$ | ${ }_{(1.61)}^{3.9}$ | $\begin{array}{ll} 9.7 \\ (2.39) \end{array} \quad \mathrm{A}$ | ${\underset{(2.63)}{17.7} \quad 8}^{8}$ | $\begin{aligned} & 2.1 \\ & (0.759)^{A} \end{aligned}$ |
| ABGR/DIHO-THOC | 21 | 65 | $\begin{array}{ll} 34.0 & A \\ (3.75) \end{array}$ | $\begin{aligned} & 3.7 \\ & (0.891)^{8} \end{aligned}$ | $\begin{array}{lr} 8.0 & B \\ 1.57) \end{array}$ | $\begin{aligned} & 20.2 \\ & (2.52) \end{aligned}$ | $\begin{aligned} & 1.7 \\ & (0.69) \end{aligned} \mathrm{AB}$ |
| ABGR/POMU | 16 | 70 | $\begin{array}{ll} 30.2 & B \\ (5.61) \end{array}$ | $\begin{aligned} & 4.0 \\ & (0.94)^{8} \end{aligned}$ | ${ }_{(1.98)}^{9.1} A B$ | $\begin{aligned} & 15.5 \\ & (4.02) \end{aligned}$ | $\begin{aligned} & 1.6 \\ & (0.61) \end{aligned} \quad A B$ |
| ABGR/RHDI-RUUR | 27 | 69 | $\begin{aligned} & 30.2 \\ & (4.13) \end{aligned} \quad \text { B }$ | $\begin{aligned} & 4.0 \\ & (1.17)^{8} \end{aligned}$ | ${\underset{(1.57)}{9.4} \quad A B}^{\text {AB }}$ | $\underset{(3.84)}{15.1}{ }^{8}$ | $\begin{aligned} & 1.7 \\ & (0.67) \end{aligned} \text { AB }$ |
| ABGR/BRSY | 26 | 64 | $\begin{array}{ll} 24.4 & C \\ (5.28) & \end{array}$ | ${ }_{(1.18)}{ }^{8}$ | $\begin{aligned} & 8.3 \\ & (1.99) \end{aligned} \quad A B$ | $\underset{(3.93)^{10.2} \mathrm{C}}{ }$ | $\begin{aligned} & 1.4 \\ & (0.68) \end{aligned}$ |

' All average species richness values are means calculated from the plots within the association.
${ }^{2}$ All values in parentheses are standard deviations.
3 Within each column, means with the same letter are not significantly different ( $P=0.05$, GLM, multiple means comparison, unbalanced ANOVA, Fisher's Protected (SD).

Species richness within tree, shrub, and grass strata are fairly similar between associations. One plant association, TSHE/ACCI-GASH, had 5.5 tree species/plot. This is significantly higher than the other plant associations.

The shrub richness was higher in the ABGR/ACCI-GASH plant association is than in the TSHE/ACCI-GASH and ABGR/DIHO-THOC plant associations.

The TSHE/ACCI-GASH association had the lowest species richness in the grass strata, with 0.3 species/plot for an association average. This is significantly lower than the other five plant associations. The other associations had grass species richness values of 1.4 to 2.1 species/plot.

The most significant difference in species richness among associations was within the forbaceous strata (F = 70.8, $\mathrm{p}=.0001$ ). The ABGR/DIHO-THOC plant association had the highest forbaceous species richness, 20.2 species/plot. This was significantly higher from all other associations. ABGR/ACCI-GASH, ABGR/POMU, and ABGR/RUUR-RHDI plant associations had the next highest average forbaceous species richness per plot values (17.7, 15.5, 15.1). The TSHE/ACCI-GASH and ABGR/BRSY plant associations had the lowest forbaceous species richness with 7.8 and 10.2 forbaceous species/plot.

## DIVERSITY AND EVENNESS

Shannon's diversity index and an evenness measure were calculated for each plant association (Table 6). Shannon's diversity is significantly higher in the ABGR/ACCI-GASH, ABGR/DIHO-THOC, and ABGR/RUUR-RHDI plant associations. Evenness was shown to be significantly lower for the ABGR/POMU and the ABGR/BRSY plant associations.

## VERTICAL STRUCTURE

Vertical structure, giving average percent cover by strata for each association, is shown in Table 7. There were no significant differences in percent cover in any tree strata among associations. There were significant differences among associations in cover percent of the shrub, forb, and grass strata. The dominance of percent cover in certain strata support characterizing associations as shrub, forb, or grass types.

Table 6. Shannon's diversity ${ }^{1}$ and an evenness ${ }^{2}$ measure.

| Association | $\mathrm{N}=$ | Diversity ${ }^{1}$ mean (SD) | Evenness ${ }^{2}$ mean (SD) |  |
| :---: | :---: | :---: | :---: | :---: |
| TSHE/ACCI-GASH | 6 | $\underset{(.868}{(.090)^{3}} \quad \mathrm{~B}^{4}$ | $\begin{aligned} & .650 \\ & (.067) \end{aligned}$ | A |
| ABGR/ACCI-GASH | 12 | $\begin{array}{ll} .995 \\ (.110) \end{array} \quad \text { A }$ | $\begin{aligned} & .656 \\ & (.067) \end{aligned}$ | A |
| ABGR/DIHO-THOC | 21 | $\begin{array}{ll} 1.020 \\ (.078) \end{array} \quad \mathrm{A}$ | $\begin{aligned} & .670 \\ & (.055) \end{aligned}$ | A |
| ABGR/POMU | 16 | $\begin{array}{ll} .869 \\ (.117) \end{array} \quad B$ | $\begin{aligned} & .592 \\ & (.066) \end{aligned}$ | B |
| ABGR/RUUR-RHDI | 27 | $\begin{gathered} .966 \\ (.092) \end{gathered}$ | $\begin{gathered} .651 \\ (.054) \end{gathered}$ | A |
| ABGR/BRSY | 26 | $\begin{array}{ll} .827 \\ (.103) \end{array} \quad \text { B }$ | $\begin{aligned} & .601 \\ & (.065) \end{aligned}$ | B |

1 Shannon's diversity index - measure of heterogeneity. Calculated as average for plots within association. ${ }_{2}$ Evenness using Shannon's diversity index - averages for plots within association.
Standard deviation for averages (means).
4 Within each column, means with the same letter are not significantly different ( $P=0.05$, GLM, multiple means comparison, unbalanced ANOVA, LSD).

Table 7. Vertical structure: percent cover by strata.


All values are average percent cover calculated using values from each plot within the association.
2 All values in parentheses are standard deviations.
3 There were no significant differences between associations in the Tree strata.
4 Within each column, means with the same letter are not significantly different ( $P=0.05$, GLM, multiple means comparison, unbalanced ANOVA, Fisher's Protected LSD).

The ABGR/ACCI-GASH plant associations had the highest shrub cover percent 73.3. TSHE/ACCI-GASH and ABGR/RUURRHDI plant associations were the next highest in shrub percent cover ( 55.6 and 43.3 percent). Three associations; ABGR/ACCI-GASH, ABGR/RUUR-RHDI, and TSHE/ACCI-GASH are characterized as shrub types, as they have greater percent cover of shrubs than any other understory strata.

The ABGR/POMU plant association had the highest percent cover of forbs (71.1 \%). The next highest was ABGR/DIHO-THOC with 55.5 percent cover forbs. The forb strata was the most dominant understory strata for these two associations. These associations were characterized as forbaceous types. The next highest percent cover of forbs was in the ABGR/ACCI-GASH and ABGR/RUUR-RHDI plant associations with 33.3 and 27.1 percent cover. The TSHE/ACCI-GASH and ABGR/BRSY plant associations had the lowest cover of forbs (19.3 and 16.1).

The ABGR/BRSY association had the highest percent cover in the grass strata, with 54.7 percent. This was significantly higher than all other associations. The ABGR/BRSY association was characterized as a grass type as the grass strata had the highest percent cover of the understory strata within this association.

A vertical profile and its affiliated evenness value for each association is illustrated in Table 8. Vertical profiles were made on the basis of 5 vertical strata
categories: 0-2', 2-6', 6-12', 12-50', and 50+'. It is evident from these figures that ABGR/ACCI-GASH and ABGR/RUUR-RHDI have the highest percent cover in the tall shrub category (6-12 feet). ABGR/POMU has the highest percent coverage in the forb strata (0-2 feet), followed by ABGR/BRSY and ABGR/DIHO-THOC. The TSHE/ACCI-GASH association has the least amount of cover in the 0-6 foot strata. ABGR/ACCI-GASH has the highest total percent cover of all strata combined. TSHE/ACCI-GASH has the lowest total persent cover of all strata. The ABGR/ACCI-GASH plant association has the highest evenness of .914. ABGR/POMU and ABGR/BRSY have the lowest evenness values, .771 and . 734 respectively.

Figure 8. Vertical profiles for plant associations.

Vertical Protile tor TSHE/ACCI-GASH


Verfical Protile tor ABGR/DIHO-THOC


Verlical Protile for ABGR/ACCH-GASH


Vertical Protile tor ABGR/POMU


## Figure 8. (cont.)

## Vertical Profila tor ABGR/RUUR-RHDI

Vertical Prolile for ABGR/BRSY


## Snags

Snag information for each plant association is summarized in Table 9. Average number of snags per plot within each association are listed. There were no significant differences in average number of snags between associations at the level tested (alpha=.05). In spite of no significant differences between plant associations, snag inventory results revealed the size and type of snags found within McDonald-Dunn Forest. Eighty-seven percent (8.1/acre) of the snags were in the 4 to 12 -inch class, 11 percent (1.1/acre) in the 12 to 21 -inch class, and 1.5 percent ( 0.2 /acre) in the $21+$-inch class. Most snags were in the smallest diameter class.

The proportion of hardwood to conifer snags was fairly equal in the 4 to 12 -inch class; $45 \%$ and $42 \%$ of total snags respectively. The 12 to 21 -inch class had a higher percentage of hardwood snags; $9 \%$ versus $2 \%$ of total snags respectively. The $21+-i n c h$ class contains the least snags; . $5 \%$ hardwood and $1 \%$ conifer of the total snags sampled. Snags are also listed by snags per 100 acres in each size class (Table 9). There were 810 snags per 100 acres in the 4 to 12 -inch size class, 100 in the 12 to 21 -inch size class, and 14 in the $21+-i n c h$ size class. These values are used for comparison to snags required for populations levels calculated by the Snag Recruitment simulator
(Marcot, 1989). The results from the Snag Recruitment Simulator (Marcot, 1989) are in Table 10. This table lists six species of primary cavity-nesters found on the Forest, and the number and size of snags that are required for different species populations levels. Analysis through SRS (Snag Recruitment Simulator, Marcot, 1989) indicated that a total of 383 snags would be needed to support $100 \%$ populations of theses six species of cavity-nesters (Table 10). The size of snags necessary to support 100 percent populations were: 16 snags 11+ inches, 237 snags 15+ inches, 124 snags $17+$ inches, and 6 snags $25+$ inches in diameter per 100 acres. Snags necessary to support 90 to $30 \%$ population levels are also listed. Comparisons of these levels to the current snag composition on McDonald Forest can be found in the discussion section of this study.

Table 9. Snag summary for ecology plots on McDonald-Dunn Forest.


| \# of snags per <br> 100 acres in <br> size class |  | 810 | 100 | 14 |
| :--- | :--- | :--- | :--- | :--- |

2 values in parentheses are standard deviations - $95 \frac{\%}{c}$ conf.
${ }^{2}$ calculations based on a total of 1739 sampled snags
'lable 10. Number and size of snags required for selected cavity-nesters.


1 Snag diameter class as defined by Neitro(1985).
2 Percent of population size values from Marcot (1989). Percent of population is the population level at which you want to manage the species (Marcot, 1989). $\infty_{0}$

## PLANT ASSOCIATION DESCRIPTIONS

Coniferous tree species composition is fairly similar throughout the plant associations except for the Tsuga heterophylla/Acer circinatum-Gaultheria shallon plant association. For this reason, tree species composition will be described briefly here and individual plant association descriptions will focus on understory shrub, forb, and grass species.

The most consistent component of the coniferous overstory is Pseudotsuga menziesii; it occurs with 100 percent constancy (Table 4). Abies grandis is the next most frequently encountered conifer species in the overstory. These two species dominate the overstory in all plant associations described in this study. The Tsuga heterophylla/Acer circinatum-Gaultheria shallon plant association is unique in the addition of Tsuga heterophylla as a component in the overstory. Taxus brevifolia can be found occasionally throughout much of the forest, but occurs most frequently in the Tsuga heterophylla/Acer circinatum-Gaultheria shallon plant association. Thuja plicata was encountered in the Tsuga heterophylla/Acer circinatum-Gaultheria shallon and the Abies grandis/Polystichum munitum plant associations.

Acer macrophyllum dominates the deciduous tree species component within the plant associations, and is found
throughout the Forest. Acer macrophyllum often dominates the midstory strata in the Forest (12 to 50 foot class). The next most consistent deciduous tree species found in the Forest are cornus nuttallii and Quercus garryana. Quercus qarryana was encountered most often in and is a fairly consistent component of the Abies
grandis/Brachypodium sylvaticum plant association. Other hardwoods that can be found occasionally in the plant associations are Prunus spp, Fraxinus latifolia, Arbutus menziesii. Alnus rubra was encountered on only one plot within this study. Understory tree species composition generally follows the trends of the overstory tree species, but is more variable.

Descriptions of the six plant associations classified in this study for the McDonald-Dunn Forest follow. The associations are described in the order in which they are placed along the environmental gradient (Leavell, 1991), moist to dry. The geographic range where an association was encountered are given by Tract. Tracts are maps for geographic areas within the Forest, each Tract is on a separate Forest map. There are 8 Tracts within McDonaldDunn Forest. Tract locations are outlined in Appendix 9.

Tsuga heterophylla/Acer circinatum-Gaultheria shallon Western hemlock/vine maple-salal

TSHE/ACCI-GASH


Figure 8. Tsuga heterophylla/Acer circinatum-Gaultheria shallon plant association.

This plant association is very limited geographically. It occurs in only one drainage located in the Soap Creek Tract, and is scattered over an area of less than one square mile. The plant association is located within the west half of Sec. 6, TllS.R5W., Willamette Meridian.

The TSHE/ACCI-GASH association is characterized by the presence of Tsuga heterophylla as well as shrub species Acer circinatum, Gaultheria shallon, and Berberis nervosa. Tsuga heterophylla was not encountered in any other area on the McDonald-Dunn Forest, and has not been found anywhere
else by Forest workers. The TSHE/ACCI-GASH association is characterized as a shrub-dominated association, with an average of $44.3 \%$ cover in the shrub strata (Table 9). The association has low cover in the forb and grass strata (19.3 and 0.2\% respectively).

The TSHE/ACCI-GASH association is low in species richness as compared to the other associations on McDonaldDunn Forest. There were a total of 38 different species encountered in the plots making up this association. The highest number of species in an association on this Forest is 70 (ABGR/POMU) (Table 5). This plant association and ABGR/BRSY have the lowest total richness among associations, 21.7 and 24.4 species per plot respectively. The TSHE/ACCI-GASH association has the highest species richness in the tree strata. This plant association is in the upper half of the range for evenness and in the lower half of the range for Shannon's diversity for associations (Table 6).

Species found in this plant association but not in other associations described for McDonald-Dunn are Tsuga heterophylla and Coptis laciniata. Taxus brevifolia, Acer circinatum, Gaultheria shallon, Vaccinium parviflorum, and Viola sempervirens, are generally limited to the TSHE/ACCIGASH and the ABGR/ACCI-GASH plant associations. These species are all indicative of moist sites in this area (Leavell, 1991).

This association is also characterized by the absence of the following species: Brachypodium sylvaticum, Rhus diversiloba, Rhamnus purshiana, Adenocaulon bicolor, Goodyera oblongifolia, Disporum hookeri, and Thalictrum occidentale. Even though these species were not located within this association, they are nearly ubiquitous throughout the rest of the Forest.
stands that are within this association have the most recent intensive disturbance history of all stands sampled. Stands within this association were $35-40$ years old, the youngest of identified plant associations. This area was clearcut in 1946 (approximately 50 MBF/acre were removed), and was slash burned in 1949 (Rowley, 1989). Part of the area was planted with 2-0 Douglas-fir in 1959 with $8^{\prime}$ X $8^{\prime}$ spacing. Other parts of this area were allowed to naturally regenerate. Even though this area has been very disturbed, it has an identifiably unique association in which Tsuga heterophylla is an indicator species.

The TSHE/ACCI-GASH association is on the moist end of the environmental gradient (Leavell, 1991). A combination of environmental factors including elevation, solar radiation, soils, and aspect make this geographical location the most moist part of the McDonald-Dunn Forest (Leavell, 1991).

Abies grandis/Acer circinatum-Gaultheria shallon
Grand fir/vine maple-salal
ABGR/ACCI-GASH


Figure 9. Abies grandis/Acer circinatum-Gaultheria shallon plant association.

The ABGR/ACCI-GASH plant association is also limited in extent. This plant association geographically surrounds the TSHE/ACCI-GASH association. It is just out of the range where Tsuga heterophylla can exist, or has migrated. This association can also be found south of the Lewisberg saddle area, and in the northern part of the Forest in the bottom of the South Fork of Berry Creek tract.
ABGR/ACCI-GASH is characterized by the presence of

Acer circinatum, Gaultheria shallon, and Berberis nervosa. The main floristic difference between the ABGR/ACCI-GASH
association and TSHE/ACCI-GASH is the lack of Tsuga heterophylla. The ABGR/ACCI-GASH association is shrubdominated, with an average of $73.3 \%$ cover in the shrub strata. This is the highest shrub cover among associations. Forb and grass cover is comparatively low (21.7 and 2.3\% respectively).

This is a species rich association. There is an average of 33.2 species per plot within the association, which is similar to all but the TSHE/ACCI-GASH and the ABGR/BRSY plant associations which have 21.7 and 24.4 species per plot respectively. The ABGR/ACCI-GASH association is in the upper half of the range for both evenness and Shannon's diversity among associations.

This association has a presence of the following species that are absent in the TSHE/ACCI-GASH association: Brachypodium sylvaticum, Rubus ursinus, Adenocaulon bicolor, Disporum hookeri, and Thalictrum occidentale. Rhus diversiloba and Rubus ursinus were also found in low abundance. ABGR/ACCI-GASH has the highest coverage of Corylus cornuta of any of the associations. Achlys triphylla and Vancouveria hexandra are generally found in this association.

The ABGR/ACCI-GASH association is on the moist end of the environmental gradient, having moist site indicators such as Acer circinatum and Gaultheria shallon (Leavell, 1991). Stands in the ABGR/ACCI-GASH association are some
of the oldest stands on the Forest, with an average stand age of 129 years. Many of the old-growth stands on the Forest are within this plant association.

Abies grandis/Disporum hookeri-Thalictrum occidentale
Grand fir/hooker's fairybells-western meadowrue
ABGR/DIHO-THOC


Figure 10. Abies grandis/Disporum hookeri-Thalictrum occidentale plant association.

The ABGR/DIHO-THOC plant association is more widespread geographically than the TSHE/ACCI-GASH and ABGR/ACCI-GASH associations. It occurs throughout the Forest where preferred environmental conditions exist. This association occurs primarily in the Oak Creek and Jackson Creek tracts, but is also found to a limited extent in Peavy and Forest Peak Ridge Road tracts.

> The ABGR/DIHO-THOC type is characterized by the
presence of both Disporum hookeri and Thalictrum occidentale. Berberis nervosa is present and dominates the
understory within approximately half of the ecology plots in this association. Most forbs occurring in this type also occur in the ABGR/ACCI-GASH association. Viola glabella, Vancouveria hexandra, Anemone deltoidea, Campanula scouleri, and Achlys triphylla are also commonly present in this association. The ABGR/DIHO-THOC association is forb-dominated. It has an average of $55.5 \%$ cover in the forb strata. The only association with higher forb percent is ABGR/POMU, with $77.1 \%$ cover. The ABGR/DIHO-THOC association is relatively low in both shrub and grass cover ( 34.5 and $4.9 \%$ cover respectively).

This is a species rich association. There were 65 species encountered within this plant association. ABGR/DIHO-THOC and ABGR/ACCI-GASH have the highest average species richness per plot among associations (34.0 and 33.2 species per plot). The ABGR/DIHO-THOC association also has the highest forb species richness associations, with 20.2 forbs per plot. This plant association is in the high half of the range for both evenness and Shannon's diversity between associations.

Species on the dry end of the gradient such as Ribes divaricatum, Rubus discolor, Stellaria crispa, and Sanicula crassicaulis (Leavell, 1991) are absent within this association. Species on the moist end of the gradient such as Viola sempervirens, Coptis laciniata, Vaccinium parviflorum, Gaultheria shallon, and Tsuga heterophylla
are also absent. Acer circinatum occurs only occasionally in this association. The ABGR/DIHO-THOC association is located towards the moist end of the environmental gradient, but is drier than the TSHE/ACCI-GASH and ABGR/ACCI-GASH associations (Leavell, 1991).

Abies grandis/Polystichum munitum
Grand fir/sword fern
ABGR/POMU


Figure 11. Abies grandis/Polystichum munitum plant association.

This association occurs throughout the Forest, and was found in all tracts except Soap Creek.

The ABGR/POMU association is characterized by a high percentage cover (generally $>40 \%$ ) of Polystichum munitum. Rhus diversiloba, Rubus ursinus, and corylus cornuta are a consistent component of this association, but generally have less than $10 \%$ cover each. ABGR/POMU is a forbdominated association. It has an average of $71.1 \%$ cover in the forb strata, the majority of which is Polystichum munitum. This is a significantly higher percent forb cover than all other associations on McDonald-Dunn Forest. This
association has low cover of shrubs and grasses (31.2 and 2.7\% respectively).

This association is also species rich. A total of 70 species were encountered within the association, which is the highest among associations. But it has an average of 30.2 species per plot within the association which is significantly higher than the TSHE/ACCI-GASH and ABGR/BRSY plant associations. This plant association in similar in species richness per plot to ABGR/ACCI-GASH and ABGR/RUURRHDI. This association is in the lower half of the range for both evenness and Shannon's diversity between associations.

Besides the dominance of Polystichum munitum, a number of forb species are occasionally present in low abundance. These are: Thalictrum occidentale, Disporum hookeri, Vancouveria hexandra, Actaea rubra, and Smilacina stellata. Species on the moist end of the gradient such as Viola sempervirens, Coptis laciniata, Vaccinium parviflorum, Gaultheria shallon, and Tsuga heterophylla are absent. The species on the dry end of the gradient, such as Ribes divaricatum, Rubus discolor, Stellaria crispa, and sanicula crassicaulis (Leavell, 1991) are occasionally present in low abundance in this association. The ABGR/POMU association is mesic on the environmental gradient for the Forest (Leavell, 1991).

Abies Grandis/Rubus ursinus-Rhus diversiloba
Grand fir/trailing blackberry-poison oak
ABGR/RUUR-RHDI


Figure 12. Abies grandis/Rubus ursinus-Rhus diversiloba plant association.

The ABGR/RUUR-RHDI association was found throughout the Forest except in the Soap Creek tract. This and the ABGR/BRSY associations are the most common plant associations encountered in this study.

ABGR/RUUR-RHDI is characterized by a high coverage of Rhus diversiloba and Rubus ursinus and generally $<5 \%$ coverage of Brachypodium sylvaticum. This association
lacks the high cover of Polystichum munitum, which occurs consistently in the ABGR/POMU association. ABGR/RUUR-RHDI is a shrub-dominated association. It has an average of
$55.6 \%$ cover of shrubs. The only other association with a higher cover in the shrub strata is ABGR/ACCI-GASH, with $73.3 \%$ cover. The shrub cover of this associations is similar to the TSHE/ACCI-GASH plant association, both are higher in shrub cover than the ABGR/DIHO-THOC, ABGR/POMU, and ABGR/BRSY plant associations.

This association is also species rich. A total of 69 species were encountered within the plots. There is an average of 30.2 species per plot, which is in the middle of the range of species per plot among plant associations. This plant association is in the upper half of the range for both evenness and Shannon's diversity between associations.

Dry-site species such as Rubus discolor, Stellaria crispa, and Sanicula crassicaulis, Rubus leucodermis, Lactuca muralis, and Senecio jacobaea can occur in this association. Many species on the moist end of the environmental gradient such as Viola sempervirens, Coptis laciniata, Vaccinium parviflorum, Gaultheria shallon, Thalictrum occidentale, and Tsuga heterophylla are absent. Acer circinatum and Disporum hookeri are other moist-site indicators that occur occasionally in this association.

The ABGR/RHDI-RUUR association occurs in the mesic to dry environments in the Forest (Leavell, 1991). Both Rhus diversiloba and Rubus ursinus, which are indicator species,
are located towards the dry end of the species environmental gradient (Leavell, 1991).

Abies grandis/Brachypodium sylvaticum
Grand fir/false brome
ABGR/BRSY


Figure 13. Abies grandis/Brachypodium sylvaticum plant association.

The ABGR/BRSY association was found mostly in the southern third of McDonald-Dunn Forest, in the Oak Creek and Jackson Creek tracts. Brachypodium sylvaticum occurs throughout the Forest, except in the TSHE/ACCI-GASH association. Eighty-nine out of 115 ecology plots have Brachypodium sylvaticum. But, coverages are mostly low outside of the ABGR/BRSY plant association.

The ABGR/BRSY association and is characterized by a high coverage ( $>40 \%$ ) of Brachypodium sylvaticum. Brachypodium sylvaticum is a well-established understory
dominant within portions of McDonald-Dunn Forest. However, it is not indigenous to this area (Hubbard, 1954). This species is not a transitory part of the community. It appears to be spreading and increasing in dominance in parts of the Forest (Leavell and Hubbard, 1989).

ABGR/BRSY is a grass-dominated association. It has an average of $54.7 \%$ cover of grass ( $53.3 \%$ Brachypodium sylvaticum), which is the highest among plant associations. Cover of shrubs and forbs is relatively low (34.0 and 16.1\% respectively).

A total of 64 species were encountered within the association, which makes it moderate in total species richness. But, the ABGR/BRSY association has an average of 24.4 species per plot, sharing the spot of lowest species richness per plot with the TSHE/ACCI-GASH plant association. Forb species richness in this association is relatively low (10.2 species per plot). This plant associations is in the lower half of the range for both evenness and Shannon's diversity between associations.

ABGR/BRSY association often has a component of mature Quercus garryana, but is not limited to stands with evidence of Oak remnants. The most common shrubs in this type are Rhus diversiloba, Rubus ursinus, and Corylus cornuta. Coverage of Polystichum munitum is considerably less (generally <10\%) compared to the ABGR/RUUR-RHDI or ABGR/POMU associations.

The ABGR/BRSY association is on the dry end of the environmental gradient for the plant associations in McDonald Forest (Leavell, 1991).

## DISCUSSION

## CLASSIFICATION

## SERIES DETERMINATION

Two series were identified in McDonald-Dunn Forest, Tsuga heterophylla, and Abies grandis. A Pseudotsuga menziesii series was considered, but not supported by the data in this study.

The Tsuga heterophylla series is identified by the presence of Tsuga heterophylla being present and successfully reproducing in an isolated area on the Forest. Tsuga heterophylla is the most shade tolerant tree species on this Forest, and therefore the climax tree species where it is present and reproducing successfully.

In the areas where Tsuga heterophylla was not present, Abies grandis was determined to be the climax tree species (less shade tolerant than Tsuqa heterophylla, but more shade tolerant than Pseudotsuga menziesii). Most ecology plots had Abies grandis present and successfully reproducing, indicating most of the forest (that portion without the presence of Tsuga heterophylla) belonged in the Abies grandis series.

Twenty-three plots did not contain presence of Tsuga heterophylla or Abies grandis, and could have potentially
been in a Pseudotsuga menziesii series. Even though Pseudotsuga menziesii is the dominant tree species in the stands sampled, a Pseudotsuqa menziesii series was not identified. Series for the plant associations are identified by the potential climax coniferous tree species (conifers are the dominant life form in the stands sampled). Pseudotsuga menziesss is not considered a potential climax species in the portion of the McDonaldDunn Forest sampled in this study.

The plots without Tsuga heterophylla or Abies grandis (plot numbers $8,11,12,13,22,28,35,44,61,63,65$, $70,71,77,85,86,91,92,98,107,110,114)$ would have been the plots conducive to a Pseudotsuga menziesii series. These plots should have clustered at the dry end of the environmental gradient for the classification if Pseudotsuga menziesii was to be a climax species on the Forest, just as the plots in the Tsuga heterophylla series were clustered at the moist end of the environmental gradient. TWINSPAN analysis failed to cluster these plots into an association(s) independent of plots with presence of Tsuga heterophylla or Abies grandis. These plots were scattered throughout the classification, failing to cluster at any point. This trend occurred whether or not tree species were included in the analysis. The following hypotheses were posed:

1) These plots are part of the Abies grandis series, or
2) The classification is not based on an environmental gradient, or
3) All plots, except the Tsuga heterophylla plots, are in a Pseudotsuga menziesii series, or
4) The classification procedure does not work.

The first hypothesis is preferred. The plots without Tsuga heterophylla or Abies grandis are part of the Abies grandis series. Tree species have been managed in this Forest for a substantial period of time as indicated by an earlier discussion of the thinning that has occurred in the stands sampled in this study. The presence of Abies grandis could have been affected by management. Although Abies grandis was not found in every plot, it is hypothesized that it could grow on the entire Forest. Therefore the plots mentioned above are in the Abies grandis series.

The other hypotheses appear to be less feasible. The second hypothesis was eliminated by the work of Leavell. Leavell (1991) has determined an environmental gradient does exist for the associations in this classification.

It is doubtful that the third hypothesis is viable since Abies grandis is more shade tolerant than Pseudotsuga menziesii which makes Abies grandis the more likely climax species. Abies grandis and it is abundant and reproducing successfully (where allowed) in the majority of the Forest. Also, if a Pseudotsuga menziesii series did exist, it
should be in the driest areas within the Forest. But the most southern, lowest elevation sites in the forest have abundant Abies grandis regeneration as well as Pseudotsuqa menziesii. Abies grandis appears to be climax in these areas.

That the classification procedure does not work does not appear to be a valid hypothesis. The procedures used for this classification, mainly TWINSPAN, are being used and promoted by many ecologists in the Pacific Northwest (Hemstrom, 1990; Smith, 1990; Halpern, 1990; Atzet, 1990). Also, the results of this classification, the plant associations, appear to match what is visible and interpretable on the ground. Time will be needed to prove or disprove this hypothesis.

## COMPARISONS

Existing classifications and plant association/community descriptions for western oregon were studied in detail to find parallels to the associations described for McDonald-Dunn Forest. of the six plant associations described on McDonald-Dunn Forest, only one, the ABGR/ACCI-GASH, adequately matched a previous description. The association that matched ABGR/ACCI-GASH came from Juday (1976); his old-growth Pseudotsuga menziesii-Acer circinatum/Corylus cornuta-Adenocaulon
bicolor association in the Valley-Margin zone of the oregon Coast Range. There was no precedent for the descriptions of the other five plant associations. The TSHE/ACCI-GASH, ABGR/DIHO-THOC, ABGR/POMU, ABGR/RUUR-RHDI, and ABGR/BRSY plant associations described in this study appear to not have been previously described. This section will compare the plant associations on McDonald-Dunn Forest to existing classifications and community descriptions.

Juday (1976) described a Pseudotsuga menziesii-Acer circinatum/Corylus cornuta-Adenocaulon bicolor association (PSME-ACCI/COCO-ADBI) in the Valley-Margin zone of the Coast Range that floristically matches the ABGR/ACCI-GASH plant association on the McDonald-Dunn Forest. According to Juday (1976) this association is "the most widespread and abundant community in the Valley-Margin Zone." Minor differences exist between these two associations: Acer circinatum and corylus cornuta have more coverage, and POMU has less coverage in ABGR/ACCI-GASH than in Judays' PSME-ACCI/COCO-ADBI association. There are also some minor forbaceous species that occur in trace amounts in Judays' association that are missing in ABGR/ACCI-GASH. This could be because Juday sampled over a larger area, the entire Valley-Margin Zone (Figure 4), and was doing a classification based on old-growth stands. To define and describe the associations on McDonald-Dunn as succinctly as possible, the plant association name, ABGR/ACCI-GASH will
be retained. It is desirable mainly because Abies grandis, not Pseudotsuga menziesii, is considered the climax tree species.

There were no parallels found for the floristic composition of the TSHE/ACCI-GASH plant association described on McDonald-Dunn. However, there were other classifications that described a TSHE/ACCI-GASH association, but their floristic compositions were different from the TSHE/ACCI-GASH plant association described for McDonald-Dunn Forest. Juday (1976) described a TSHE/ACCI-GASH plant association in his old-growth study in the Coast Range, and Hemstrom and Logan (1986) described TSHE/ACCI-GASH plant association in the Siuslaw National Forest.

The TSHE/ACCI-GASH plant association described by Judy (1976) did not contain Abies grandis which is common in the MCDonald-Dunn TSHE/ACCI-GASH association. Alnus rubra, Oxalis oregana are species in Juday's description that are not found in the TSHE/ACCI-GASH association on McDonaldDunn. The total species richness for the TSHE/ACCI-GASH association on McDonald Forest was 38 species, whereas there were a total of 56 species found in Juday's association. There are other plant associations described by Juday that show some similarities to the TSHE/ACCI-GASH association. They are:

Pseudotsuga menziesii-(Tsuga heterophylla)/Corylus cornuta, Tsuga heterophylla-Pseudotsuga menziesii/Acer
circinatum/Polystichum munitum, and Tsuga
heterophylla/Polystichum munitum. These associations are missing species such as Abies grandis or Acer macrophyllum, and have other species not found in TSHE/ACCI-GASH in McDonald Forest such as oxalis oregana, Oploplanax horridum, Rubus ursinus, Bromus vulgaris.

The TSHE/ACCI-GASH plant association described by Hemstrom and Logan (1986) for the Siuslaw National Forest also has a different floristic composition from the TSHE/ACCI-GASH association on MCDonald Forest. Hemstrom and Logan's TSHE/ACCI-GASH association contain Alnus rubra, Picea sitchensis, Sambucus racemosa, oxalis oregana, Vaccinium ovatum, Athyrium felix-femina and Blechnum spicant which are not in the TSHE/ACCI-GASH association in McDonald Forest. Abies grandis is also missing from Hemstrom and Logan's description. Abies grandis is a consistent component of the TSHE/ACCI-GASH plant association in McDonald Forest. The TSHE/ACCI-GASH plant association on McDonald-Dunn Forest is not like the other TSHE/ACCI-GASH plant associations described elsewhere (Hemstrom and Logan, 1986; Juday, 1976).

It must be kept in mind that there are some problems in the plots from McDonald Forest for the TSHE/ACCI-GASH association. First the stands averaged only 35-40 years
old. Stands this young are generally found to have less species diversity because the tight canopy inhibits the full expression of understory vegetation. Many classification schemes would have avoided stands at this stage of development. These plots were retained within this classification as they identify a unique plant association for which more mature stands were unavailable. Second, there were only 6 plots from which to describe this association. A sample size this small is likely to underestimate the total number of species found in the association. More plots within the Tsuga heterophylla area in McDonald Forest were not available.

There were no parallel descriptions for the other four plant associations described on McDonald Forest. The following is a discussion of other classifications in the western Oregon, Coast and Cascade Ranges, which were studied for communities with floristic similarity to those found on McDonald Forest. Comparisons are made to Pseudotsuga menziesii associations throughout, even though none were identified on McDonald-Dunn Forest because some Pseudotsuga menziesii associations in these other studies may contain Abies grandis. Associations were chosen for comparison if:

1) it was possible to key plots on McDonald-Dunn Forest to them, or
2) if their plant association or community name contained species occurring on McDonald-Dunn Forest.

Other associations described by Juday (1976) in the Valley-Margin Zone were compared to Abies grandis associations on McDonald Forest (Table 11). These associations do not fit the Abies grandis associations on the Forest as they do not contain Rhus diversiloba, Rubus parviflorus, Rubus ursinus, or ouercus garryana.

Table 11. Association from Juday (1976) that were reviewed for similarity to McDonald-Dunn plant associations.

```
                    Valley-Margin zone
Pseudotsuga menziesii-Acer macrophyllum/Corylus cornuta v.
    californica/Bromus vulqaris
Pseudotsuqa menziesii/Holodiscus discolor
Pseudotsuga menziesii-Thuja plicata/Gaultheria shallon/Linnaea borealis
```

Merkle (1948) described four communities on Marys Peak in the Oregon Coast Range. Marys Peak is located approximately 12-15 miles to the southwest of McDonald-Dunn Forest. Merkle's communities (1948) were defined as north slope, east slope, south slope, and meadow communities. These communities were found to be different from the associations described on McDonald-Dunn Forest as all four communities contained oxalis oregana, and three contained Abies procera. The Marys Peak area is substantially higher in elevation from McDonald-Dunn Forest as all four
communities lie above 2,500 feet, and Marys Peak itself is at 4097 feet. Ecology plots taken on McDonald-Dunn Forest range from 340 to 1,520 feet.

Anderson (1967) described six plant communities in the Marys Peak watershed that were reviewed for similarity to McDonald-Dunn Forest plant associations (Table 12).

Table 12. Plant communities from Anderson (1967) that were reviewed for similarity to McDonald-Dunn plant associations.

```
Corylus californica/Bromus vulgaris
Acer circinatum/Gaultheria shallon (Corylus californica-Holodiscus discolor subtype
Holodiscus discolor/Gaultheria shallon
Acer circinatum/Polystichum munitum
```

The Holodiscus discolor/Gaultheria shallon, and Acer circinatum/Polystichum munitum communities are dissimilar to all McDonald-Dunn Forest associations because these two communities do not contain Abies grandis. Although the other two associations, Corylus californica/Bromus vulgaris and Acer circinatum/Gaultheria shallon (Corylus californica-Holodiscus discolor subtype) have species composition somewhat similar to the McDonald-Dunn Forest TSHE/ACCI-GASH and ABGR/ACCI-GASH associations, they contain a substantial amount of Bromus vulgaris which is not found in TSHE/ACCI-GASH, and is rare in ABGR//ACCIGASH. Tsuga heterophylla is also lacking from these associations.

Bailey (1966) described five associations in the Southern Oregon Coast Range on the Millicom Tree Farm. There were only two of these that were compared for similarity to McDonald-Dunn Forest associations, an Acer circinatum/Berberis nervosa and a Holodiscus discolor/Gaultheria shallon association. These two associations are unlike those on McDonald-Dunn as they do not contain Abies grandis, and contain either oxalis oregana which is absent in McDonald-Dunn Forest associations, or Bromus vulgaris, which is scarce.

The Siuslaw National Forest plant association and management guide (Hemstrom and Logan, 1986) contains six plant association descriptions for Tsuga heterophylla that were reviewed for to the McDonald-Dunn TSHE/ACCI-GASH plant association (Table 13). The TSHE/ACCI-GASH association was already compared above. There were no Abies grandis plant associations described on the Siuslaw National Forest (Hemstrom and Logan, 1986). Abies grandis was not mentioned within these associations. Therefore comparisons were made with only the TSHE/ACCI-GASH plant association on McDonald-Dunn Forest. All of the comparable associations described by Hemstrom and Logan (Table 13) have the following dissimilarities to the TSHE/ACCI-GASH association. They all contain Alnus rubra, Picea sitchensis, Sambucus racemosa, oxalis oregana, Vaccinium ovatum, Athyrium felix-femina and Blechnum spicant which

Table 13. Plant associations in the siuslaw National Forest (Hemstrom and Logan, 1986) reviewed for similarity to McDonald-Dunn Forest plant associations.

Tsuga heterophylla/Berberis nervosa
Tsuga heterophylla/Berberis nervosa-Gaultheria shallon
Tsuga heterophylla/Gaultheria shallon
Tsuga heterophylla/Polystichum munitum
Tsuga heterophylla/Acer circinatum-Gaultheria shallon
Tsuga heterophylla/Acer circinatum-Polystichum munitum
are not in the TSHE/ACCI-GASH association in McDonald Forest. Abies grandis is also missing from Hemstrom and Logan's descriptions.

Thilenius (1968) described four communities in a study of the Quercus garryana forests of the Willamette Valley (Table 14). These communities are different from the associations on McDonald-Dunn Forest. They contain a number of species that were not found in this study such as Philadelphus lewisii, Torilis arvensis, Cynosurus echinatus, Holcus lanatus and Poa pratensis. Proportions of species were also quite different, as many stands had Quercus garryana as the dominant in the tree layer, and species such as Galium spp, Dactylis glomerata, and Torilis arvensis as dominants in the forbaceous layer. But these communities contain Rhus diversiloba in significant quantities, and also many other species that are found in McDonald-Dunn Forest. No other classifications reviewed had communities or associations with a high cover of Rhus

Table 14. The Quercus garryana communities of the Willamette Valley (Thilenius, 1968).

```
Quercus garryana/Corylus cornuta/Polystichum munitum
Quercus garryana/Prunus avium/Symphoricarpos albus
Quercus garryana/Amelanchier alnifolia/Symphoricarpos albus
Quercus garryana/Rhus diversiloba
```

diversiloba. Precipitation in the area Thilenius (1968) studied is approximately 40 inches. This is likely the lower limit of precipitation for McDonald-Dunn Forest. These quercus forests are therefore drier than those on McDonald-Dunn, whereas most of the classifications studies reviewed are in areas of higher precipitation. The striking similarity of Thilenius' descriptions is in the high abundance of Rhus diversiloba which is not found in many other studies.

The plant association and management guide for the Willamette National Forest in the Cascade Range (Hemstrom et al., 1987) is one of two classifications reviewed that describes Abies grandis associations. Abies grandis associations are also found in Topik et al. (1988). The Willamette National Forest guide has Pseudotsuga menziesii, Abies grandis, and Tsuga heterophylla associations that were reviewed for similarity to McDonaldDunn Forest plant associations (Table 15).

All of the Pseudotsuga menziesii associations, except for Pseudotsuga menziesii/Symphoricarpos mollis, contain pines (Pinus lambertiana, Pinus ponderosa, or Pinus
monticola), which the associations on McDonald-Dunn Forest do not contain. Castinopsis chrysophylla, and Whipplea modesta are in all five of the Pseudotsuga menziesii associations, which are also absent in McDonald-Dunn associations. The Abies grandis/Berberis nervosa

Table 15. Plant associations on the Willamette National Forest (Hemstrom et al., 1987) reviewed for similarity to associations on McDonaldDunn Forest.

[^3]The five Tsuga heterophylla associations by Hemstrom et al. (1987) have combinations of Pinus lambertiana, Pinus ponderosa, Pinus monticola, calocedrus decurrens, Rhododendron macrophyllum, Chimaphila umbellata, Vaccinium alaskaense, Whipplea modesta, oxalis oregana, and Xerophyllum tenax, all of which do not occur on the TSHE/ACCI-GASH association on the McDonald-Dunn Forest.

The plant association and management guide for the Western Hemlock Zone on the Mt. Hood National Forest (Halverson, et al., 1986) contains seven plant associations that were reviewed for similarity to the TSHE/ACCI-GASH plant association on McDonald-Dunn Forest (Table 16). As all these associations contain Tsuga heterophylla, they are compared only to the TSHE/ACCI-GASH association on McDonald-Dunn Forest. These associations contain combinations of Abies amabilis, Abies procera, Alnus rubra, Castanopsis chrysophylla, Rhus diversiloba, Rhododendron macrophyllum, Vaccinium alaskaense, oxalis oregana, Xerophyllum tenax, and oploplanax horridum. All of these species are absent in the TSHE/ACCI-GASH association on McDonald-Dunn Forest.

Dyrness et al. (1974) did a preliminary classification of forest communities in the central portion of the western Cascades. They described six plant associations that I compared for similarity to associations on McDonald-Dunn Forest (Table 17). These six plant associations described

Table 16. Plant associations in the Western Hemlock Zone on the Mt. Hood National Forest (Halverson et al., 1986) reviewed for similarity to McDonaldDunn plant associations.

Tsuga heterophylla/Acer circinatum/achlys triphlla
Tsuqa heterophylla/achlys triphlla
Tsuga heterophylla/Berberis nervosa
Tsuqa heterophylla/Berberis nervosa-Gaultheria shallon-MTH* Tsuga heterophylla/Berberis nervosa/Polystichum munitum
Tsuga heterophylla-Pseudotsuqa menziesii/Holodiscus discolor
Tsuga heterophylla/Polystichum munitum-MTH*

* MTH stands for associations that use the same name elsewhere in the Pacific Northwest, but are not identical to those described in Mt. Hood National Forest (Halverson et al., 1986).
by Dyrness et al. (1974), contain Tsuga heterophylla, and were therefore comparable to the TSHE/ACCI-GASH plant association on McDonald-Dunn Forest. These associations were unlike TSHE/ACCI-GASH as most contain Castanopsis chrysophylla, Rhododendron macrophyllum, and Whipplea modesta, which do not occur in the TSHE/ACCI-GASH association on McDonald-Dunn Forest. These associations also lack Abies grandis, except for the Pseudotsuga menziesii/Acer circinatum-Berberis nervosa association that has only a trace of percent cover. All of McDonald-Dunn plant associations contain Abies grandis.

A study was done on the dry coniferous forests in the western Cascades (Means, 1980). Means (1980) described two communities that I reviewed for similarity to plant associations on McDonald-Dunn Forest. These are:

Table 17. Plant associations in the central portion of the western Cascades (Dyrness et al., 1974) reviewed for similarity to plant associations on McDonald-Dunn Forest.

Pseudotsuga menziesii/holodiscus discolor
Pseudotsuga menziesii-Tsuga heterophylla/Corylus cornuta
Pseudotsuga menziesii/Acer circinatum/Gaultheria shallon
Pseudotsuga menziesii/Acer circinatum/Berberis nervosa
Tsuga heterophylla/Acer circinatum/Polystichum munitum
Tsuga heterophylla/polystichum munitum
Pseudotsuqa menziesii/Holodiscus discolor/Acer circinatum, and Pseudotsuga menziesii/Berberis aquifolium/Disporum. The PSME/HODI/ACCI community does not contain Abies grandis, which all McDonald-Dunn plant associations contain. Also, most plots from which the community type was described contain either Castinopsis chrysophylla, Libocedrus decurrens, or Pinus lambertiana, which do not occur in McDonald-Dunn plant associations. Since Tsuga heterophylla is absent from the PSME/BEAQ/Disporum community type, it was comparable only to the Abies grandis plant associations on McDonald-Dunn Forest. Only two out of nine plots used to describe this community have Abies grandis, and with only 0.1 and $1 \%$ cover. This cover percent was too low to match a McDonald-Dunn Forest association. There was also insufficient cover of Rhus diversiloba, Polystichum munitum, Rubus ursinus, and no Brachypodium sylvaticum, which was necessary to match most McDonald-Dunn Forest plant associations.

The plant association and management guide for the ponderosa pine, Douglas-fir, and grand fir zones in the Mt. Hood National Forest (Topik et al., 1988) describes plant associations that are on the east side of the Cascade Range. These associations contain pinus ponderosa, Larix occidentalis, Pinus contorta, Picea engelmannii, or Pinus monticola, all of which are absent in McDonald-Dunn Forest (except some species that are planted). This classification is not comparable to the classification developed for McDonald-Dunn Forest.

I conclude that five of the plant associations described for McDonald-Dunn do not fit associations in other classifications. These five plant associations are different because the vegetation composition and structure is distinct in some way from these other classifications. Namely, the presence of Abies grandis and the lack of species indicating a moister environment than McDonaldDunn. Adjacent forests on the eastern flanks of the Oregon Coast Range could reasonably be expected to contain plant communities similar to the ones that have distinguished the plant associations on McDonald-Dunn.

## DIVERSITY

Diversity is important to managers because it provides a baseline of alpha and vertical diversity from which they
can make decisions. Managers can use the baseline diversity levels to measure the effects of their management on diversity. Some activities may increase diversity, while others may decrease it. Managers can choose which aspects of diversity they are interested in and have measures for comparing and evaluating the effects of management upon diversity. Although the stands sampled for this study are representative of the most mature communities available on the Forest at this time, they are still seral stands. This baseline data necessarily represents the diversity of plant associations derived from seral stands and not climax stands. Therefore, the data provides a baseline of information for the mature forests of McDonald-Dunn at this point in time.

## SPECIES RICHNESS, SHANNON'S DIVERSITY, AND EVENNESS

Species richness, Shannon's diversity, and evenness values can provide Forest managers with some insight into diversity of the mature stands in the Forest. For instance, of the six plant associations identified, two are lower in average species richness per plot than the rest (Table 5, page 65), the TSHE/ACCI-GASH and ABGR/BRSY plant associations. There is an explanation why these associations are low in species richness.

The TSHE/ACCI-GASH plant association had the lowest per plot species richness (21.7 ave. species/plot) likely because of the stage of development of the stands sampled within this plant association. Stands sampled for the TSHE/ACCI-GASH association had an average age of 34 years, which indicates these stands are in an early successional stage. This and the fact that there are only 6 plots (small sample size) representing this plant association means that species richness could be underestimated. This plant association also has only been located in a geographically limited area, which also may also related to low species richness. Species diversity could increase as stands in this association develop towards climax. It should be interesting to measure species richness in these stands 50 years from now.

The explanation is somewhat different for the low per plot species richness of the ABGR/BRSY plant association. The stands sampled in this plant association are more developed successionally than TSHE/ACCI-GASH, although they are still seral. Low per plot species richness in ABGR/BRSY is more than likely related to the tough competition afforded by Brachypodium sylvaticum, the dominant understory vegetation. Brachypodium sylvaticum averaged $55.5 \%$ cover on plots within the association (Table 7, page 69). Note that the total species richness for this association, 64 species, is very similar to total species
richness of the other associations (except TSHE/ACCI-GASH) which range from 63 to 70. Sixty-four is not a low total species richness value. Yet average species richness per plot is significantly lower than these other associations. This indicates that a variety of species are still found within this plant association, but that they are fewer and farther in between, possibly being slowly pushed out of the community by competition with Brachypodium sylvaticum.

The ABGR/BRSY association has a low value for Shannon's diversity and evenness between associations (Table 6, page 69). The species composition is skewed by the predominance of Brachypodium sylvaticum. This would not be considered a problem if Brachypodium sylvaticum was a desirable species. But, it is not a desirable species. It also is not indigenous to the Forest. It is an invader that has persisted and spread. It is not palatable as forage (Hubbard, 1954), and provides competition to indigenous species, likely including conifer regeneration. This species also appears to be spreading in it's range on the Forest (Leavell and Hubbard, 1989).

Evenness is low, on a relative scale, for the ABGR/POMU plant association, which has it's understory dominated by Polystichum munitum. This association is high in species richness, with 70 species total species within the association. Yet the low evenness signifies that there are fewer of many species and more dominance shown by a few
species. Polystichum munitum is likely providing competition for other species within this association. The lower species diversity found in the ABGR/BRSY or the ABGR/POMU plant associations is not inherently less desirable than that found in other plant associations. But it would not be desirable to have the entire Forest have their characteristics, as is true with any of the plant associations. At this point in time the mature stands of McDonald-Dunn Forest do have a range of diversity. It would be desirable to continue to have that range of diversity on the Forest. The range of diversity that these plant associations represent (in species richness, Shannon's diversity, evenness, and structural diversity) create part of the between community, or beta, diversity of the Forest.

It is also important to recognize that the stands sampled in this study represent only a part of the Forest, and therefore only a part of the diversity. There were a total of 117 species identified within this study, but 339 species were identified by Hall and Alaback (1982) in their survey of the Forest's species, and 227 were identified by West (1964). The Forest as a whole has higher species richness. Where are those species? Likely in riparian areas, meadows, microsites, hardwood stands, and roadsides that were not sampled in this study. These species richness comparisons highlight the fact that a significant
amount of the species richness lies in areas other than the mature stands that were sampled. Therefore, areas such as riparian areas, meadows, microsites, and hardwood stands are important in maintaining the species diversity of McDonald-Dunn Forest.

Species richness values were characterized by data taken at one point in time. If stands sampled were in an undisturbed climax state, then the species richness would be representative of climax communities. The species richness values do not represent that of undisturbed climax communities. It is the intention of this study to be able to identify these same plant associations in the future. But changes in composition and structure may change over time, which may affect the associations.

The species richness values mentioned represent a part of alpha diversity, or within community diversity. Species richness of most plant associations of McDonald Forest is comparable to species richness found in climax communities in other classification studies. TSHE/ACCI-GASH is the one plant association on McDonald Forest that is substantially lower than the other associations on McDonald Forest as well as lower than associations in other classifications. It's has a total of 38 species in the association, whereas the other associations on McDonald Forest are represented by 63 to 70 species. Associations reviewed by Juday have total species richness values of 41 to 66 species. The
associations studied on the Siuslaw National Forest (Hemstrom and Logan, 1986) are represented by 54 to 61 species. The communities described by Anderson (1967) had approximately 36 to 49 total species within the communities. Plant associations studied on the willamette National Forest (Hemstrom et al., 1987) showed total species richness values of 59 to 90 for the associations reviewed. This high number of species in the Willamette National Forest is reasonable as it has a larger beta diversity representing a wider range of environments. From these ranges we can see that although total species richness is the lowest in the TSHE/ACCI-GASH plant association on McDonald Forest, at least one other association by Anderson (1967) has been shown to have even fewer species. These figures also show that the species richness of these associations on McDonald Forest is as high and in some cases higher than that of associations in surrounding areas.

In most associations (this study and others reviewed), species richness is highest in the forb strata. Associations that are lower in species richness usually appear to have similar numbers of species in the tree and shrub strata. The variation in forb species richness more strongly determining the total species richness of the associations.

The plant associations described on McDonald-Dunn Forest also add to the gamma, or landscape, diversity of the Coast Range. These associations are on the dry end of the environmental gradient for the Coast Range, being robbed of Coastal moisture by the mountains to the west. Drier site species such as Abies grandis, Quercus garryana, Amelanchier alnifolia, Rubus laciniatus, Rubus leucodermis, Arenaria macrophyllum, and Galium aparine were not even mentioned in the Siuslaw National Forest plant association guide (Hemstrom, 1986). These and other species not present in the surrounding area increase the species richness on the landscape level.

## VERTICAL STRUCTURE

Different plant associations have concentrations of cover in different vertical strata of vegetation (aside from the tree strata which is high in cover in all associations). The TSHE/ACCI-GASH, ABGR/ACCI-GASH, and ABGR/RUUR-RHDI have their understory dominated by shrub cover (Table 7, page 69). ABGR/POMU and ABGR/DIHO-THOC are dominated by forbs, and ABGR/BRSY is dominated by grass. Having concentrations vegetation in different strata of vegetation in the plant associations increases the structural diversity of the Forest.

The percent cover in the tree strata is fairly consistent between plant associations. Percent cover in the $>50$ ' tree class ranges from 68.5 to 76.3 for the associations (no significant difference). The lack of variation in percent cover within the mature tree strata ( $>50$ feet) was due, in part, to past thinning practices. Tree crowns have been opened up fairly uniformly by thinning practices (Rowley, 1990) in many of the stands sampled. The stands are also fairly similar aged (except TSHE/ACCI-GASH) Pseudotusga menziesii and Abies grandis forests.

Only a part of the range of structural diversity of this Forest is represented by the plant associations in this study. This study concentrates on within (alpha) structural diversity, whereas the Forest as a whole is represented by a much wider range of landscape (gamma) structural diversity. There are many different size and age classes of coniferous forest stands, as well as hardwood stands, riparian areas, and meadows that are integral to structural diversity on the landscape level. Quantifying landscape diversity was beyond the scope of this study.

## Snags

Snag quality and distribution are more related to disturbance and management activities than to plant associations in this study. McDonald-Dunn Forest has been thinned to capture mortality. Firewood gathering has been another activity affecting snags on the Forest. Natural snag development has not taken place on much of this Forest. Analysis showed that there were no significant differences in the number of snags among the plant associations (Table 9). Snag quality and distribution have been affected by activities in the Forest.

Current snag conditions indicate that at most, 30 percent of the maximum populations for the six woodpecker species analyzed could be supported on this Forest in the stands sampled (Table 10). This appraisal is likely an overestimate. These calculations have not included the size of stands or range necessary for individual species. Species also have requirements for specific decay classes of snags that were not evaluated in this study. These figures also do not take into account the other species that use snags on the Forest that are not included in the SRS model. Most of the woodpeckers studied require snags with diameters of 15 inches or more. The tally of snags available for these species of woodpeckers is from two diameter classes, 12-21 and $21+$ inches. If very many snags
were in the 12-15" diameter range, the maximum population of woodpeckers the Forest can support would go down even further. These snag numbers are also likely overestimated because they were from the least-disturbed stands and in many cases the furthest away from roads, where wood gathering and thinning were at a minimum as compared to the Forest as a whole.

## RECOMMENDATIONS

Six plant associations have been identified on the mature coniferous forests of the McDonald-Dunn Research Forest. Five of these six plant associations had not been previously described elsewhere. These plant associations, identified by their vegetation composition and structure, provide a baseline of information for this point in time to which future vegetation composition and structure of these forests can be compared. The McDonald-Dunn Forest management team can use this information to measure the effects of their management. Maintaining the diversity of the plant associations that now exists in these forests will allow for a healthy and diverse ecosystem of plants and animals.

Emphasis must be placed upon the fact that these mature coniferous forests studied represent only part of the diversity of plant communities that exists in McDonaldDunn Forest. Riparian areas, meadows, and successional stands of conifers and hardwoods contribute to the overall alpha, beta, and gamma diversity of these lands. These other plant communities are also an essential part of a healthy, diverse, functioning ecosystem. Many species not encountered in this study will be found in riparian, meadow, and hardwood communities. It is recommended that classification work continue on this Forest for riparian
areas, meadows, and successional processes. A successional study of the coniferous forests is recommended. A successional study will provide valuable insight into the management implications of harvesting practices. A successional study would be difficult without the baseline information provided by this study.

It is very important that the data from this study be preserved to facilitate future vegetation studies in this area. This study would have been enhanced if the data from Neil West's (1964) vegetation study of this Forest could have been located.

The process of understanding these plant associations has only begun. The plant associations identified here need to be followed through time and studied successionally. The plant associations should be followed in their response to management activities and their development thereafter. The series identified in this study, Tsuga heterophylla, and Abies grandis, represent the potential climax tree species. These potential climax species are hypothesized. Time will be necessary to determine the actual climax species on these forests.

It is also recommended that photographs be taken every ten years on the permanent photo points recorded in this study. This would allow the Forest to pictorially follow the development of these plots through time, whether left undisturbed or if harvested. This photograph record would
enhance vegetation studies in the future and help understand successional processes.

The Forest has had the foresight to set aside some old-growth stands (McDonald-Dunn Forest definition of oldgrowth). It is advised that parcels of mature forests representative of all six plant associations described here be included as reserves. These reserves would have a number of functions. Scientifically and educationally they represent the current diversity of the mature forests in this area, a legacy so to speak. These parcels could be left to reach near climax (late successional) conditions. The Forest management team could learn from these reserves. We could learn the true climax species. We could learn what species are lost in the development towards climax, as well as those species gained. Species richness and structural diversity could be followed. It would be interesting to follow the development of the most seral, productive (Leavell, 1991), and unique plant association in this Forest, TSHE/ACCI-GASH.

The Forest management team is interested in maintaining their old-growth. But merely setting aside these stands will not ensure the maintenance of old-growth forests in McDonald-Dunn. Providing for replacement oldgrowth stands would be necessary. Having set asides that can be allowed to proceed through natural succession will provide for future old-growth stands. Another way to
provide for old for old-growth replacement stands is to first evaluate the composition and structure of their existing old-growth stands in more detail as this study had a limited sample of true old-growth. Then this composition and structure needs to be compared to the structure of the rest of their mature stands (stands sampled in this study). Then they can determine what needs to be done to develop old-growth qualities from these mature, yet younger, stands that now exist. The development through time of the reserves suggested earlier would facilitate the understanding of compositional and structural development towards old-growth forests.

The key for field identification of plant associations on McDonald-Dunn Forest will be tested by field crews in the summer of 1991. Identification of plant associations by the key will be attempted as inventory plots are remeasured (all inventory plots are remeasured every 5 or 10 years). Plot size for occular estimate of percent cover of trees, shrubs, forbs, and grasses will be smaller than the plots used in this study (exact size to be determined by the Forest). All trees, shrubs, forbs, and grasses and cover percent will be recorded. From this information the Forest can construct a plant association map and more accurately know the extent of each plant association. In the case that there are problems with the field key, the vegetation information collected on each inventory plot
could be incorporated into new classification analysis. The data collected in this study could be combined with the data collected during plant association identification of the inventory plots to make a revised classification and key.

The Forest has recently been leaving snags and snag replacements in their clearcuts and partial cuts. The Forest may also want to consider some active snag management in their mature forests. This study has shown the current snag levels in these forest to be poor for woodpecker species as a whole on McDonald-Dunn Forest. Most of these woodpeckers require snags of 15 inches in diameter or more. The Forest management team needs to identify the populations of snag-dependent species they want to provide habitat for, and actively work to provide this habitat.

In summary, the plant associations described in this study provide only the first step in understanding vegetation composition, structure, and dynamics on this Forest. Not only does the Forest now have another tool for the purposes described above, but the College of Forestry has a teaching tool for forestry classes. Other disciplines on the University (Botany, Ecology, Soils, etc.) may also use this classification for education and study. Continued effort to study these and other
communities on this Forest will be necessary to formulate specific management implications.

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APPENDICES

Appendix 1. Ecology plot location descriptions.

| Ecology plot \# | Forest Inventory plot \# | Tract-Compartment-Stand |  |  | Township, Section, Range |
| :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | 1015 | 7 | 9 | 7 | T.11S.R.5W.SW1/4SEC. 18 |
| 2 | 1008 | 7 | 9 | 7 | T.11S.R.5W.SE1/4SEC. 18 |
| 3 | 1502 | 7 | 10 | 4 | T.11S.R.5W.SE1/4SEC. 18 |
| 4 | 306 | 7 | 4 | 3 | T.11S.R.5W.SW1/2SEC. 8 |
| 5 | 303 | 7 | 9 | 9 | T.11S.R.5W.NE1/4SEC. 19 |
| 6 | 307 | 7 | 9 | 9 | T.11S.R.5W.NE1/4SEC. 19 |
| 7 | 502 | 7 | 11 | 11 | T.11S.R.5W.NW1/4SEC. 20 |
| 8 | 806 | 7 | 7 | 3 | T.11S.R.5W.NW1/4SEC. 17 |
| 9 | 610 | 7 | 7 | 3 | T.11S.R.5W.NE1/4SEC. 18 |
| 10 | 301 | 4 | 19 | 1 | T.10S.R.5W.NE1/4SEC. 36 |
| 11 | 1101 | 4 | 17 | 5 | T.10S.R.5W.SE1/4SEC. 36 |
| 12 | 801 | 4 | 17 | 5 | T.10S.R.5W. W1/2SEC. 36 |
| 13 | 501 | 4 | 23 | 1 | T.10S.R.5W.SE1/4SEC. 36 |
| 14 | 303 | 4 | 22 | 1 | T.10S.R.5W.SW1/4SEC. 36 |
| 15 | 400 | 4 | 21 | 7 | T.10S.R.5W.SE1/4SEC. 35 |
| 16 | 700 | 4 | 6 | 5 | T.10S.R.5W.SW1/4SEC. 25 |
| 17 | 201 | 4 | 10 | 6 | T.10S.R.5W.NE1/4SEC. 36 |
| 18 | 700 | 4 | 7 | 3 | T.10S.R.5W.SE1/4SEC. 25 |
| 19 | 1301 | 4 | 6 | 5 | T.10S.R.5W.SW1/4SEC. 25 |
| 20 | 607 | 4 | 17 | 1 | T.10S.R.5W.NW1/4SEC. 36 |
| 21 | 407 | 7 | 6 | 5 | T.11S.R.5W.NW1/4SEC. 18 |
| 22 | 411 | 7 | 6 | 3 | T.11S.R.5W.NW1/4SEC. 18 |
| 23 | 405 | 7 | 6 | 5 | T.11S.R.5W.NW1/4SEC. 18 |
| 24 | 300 | 7 | 6 | 6 | T.11S.R.5W.NE1/4SEC. 18 |
| 25 | 1403 | 7 | 11 | 1 | T.11S.R.5W.SW1/4SEC. 17 |
| 26 | 500 | 7 | 5 | 1 | T.11S.R.5W.SE1/4SEC. 8 |
| 27 | 403 | 6 | 5 | 3 | T.11S.R.5W.SE1/4SEC. 8 |
| 28 | 406 | 6 | 5 | 1 | T.11S.R.5W.SE1/4SEC. 8 |
| 29 | 504 | 6 | 5 | 3 | T.11S.R.5W.SE1/4SEC. 8 |
| 30 | 302 | 7 | 6 | 5 | T.11S.R.5W.SW1/4SEC. 9 |
| 31 | 902 | 6 | 8 | 10 | T.11S.R.5W.NW1/4SEC. 16 |
| 32 | 402 | 6 | 8 | 5 | T.11S.R.5W.NW1/4SEC. 16 |
| 33 | 802 | 6 | 8 | 9 | T.11S.R.5W.NW1/4SEC. 16 |
| 34 | 601 | 6 | 3 | 8 | T.11S.R.5W.NE1/4SEC. 9 |
| 35 | 201 | 6 | 2 | 1 | T.11S.R.5W.NE1/4SEC. 9 |
| 36 | 501 | 6 | 2 | 4 | T.11S.R.5W.NE1/4SEC. 9 |
| 37 | 602 | 6 | 2 | 3 | T.11S.R.5W.NW1/4SEC. 9 |
| 38 | 605 | 1 | 7 | 3 | T.10S.R.5W.SW1/4SEC. 9 |
| 39 | 600 | 1 | 7 | 1 | T.10S.R.5W.SE1/4SEC. 8 |
| 40 | 600 | 1 | 5 | 11 | T.10S.R.5W.SW1/4SEC. 9 |
| 41 | 607 | 1 | 1 | 7 | T.10S.R.5W.SW1/4SEC. 5 |
| 42 | 104 | 7 | 4 | 6 | T.11S.R.5W.SW1/4SEC. 8 |
| 43 | 212 | 7 | 7 | 3 | T.11S.R.5W.NW1/4SEC. 19 |
| 44 | 303 | 1 | 9 | 3 | T.10S.R.5W.NE1/4SEC. 9 |
| 45 | 502 | 1 | 9 | 3 | T.10S.R.5W.NW1/4SEC. 9 |
| 46 | 704 | 1 | 9 | 5 | T.10S.R.5W.NW1/4SEC. 16 |
| 47 | 901 | 2 | 2 | 8 | T.10S.R.5W.NW1/4SEC. 16 |
| 48 | 1007 | 2 | 2 | 5 | T.10S.R.5W.NE1/4SEC. 16 |
| 49 | 808 | 2 | 2 | 5 | T.10S.R.5W.NE1/4SEC. 16 |
| 50 | 302 | 2 | 4 | 1 | T.10S.R.5W.NE1/4SEC. 21 |
| 51 | 1105 | 2 | 3 | 2 | T.10S.R.5W.NW1/4SEC. 15 |
| 52 | 805 | 3 | 8 | 12 | T.10S.R.5W.NW1/4SEC. 15 |
| 53 | 1104 | 3 | 7 | 1 | T.10S.R.5W.SE1/4SEC. 22 |
| 54 | 705 | 1 | 5 | 1 | T.10S.R.5W.NE1/4SEC. 8 |
| 55 | 1401 | 1 | 4 | 1 | T.10S.R.5W.NW1/4SEC.8 |
| 56 | 900 | 1 | 6 | 1 | T.10S.R.5W.NW1/4SEC.8 |
| 57 | 1101 | 1 | 3 | 3 | T.10S.R.5W.NW1/4SEC.8 |
| 58 | 307 | 1 | 2 | 5 | T.10S.R.5W.NW1/4SEC. 15 |




COMMENTS:

[^4]Appendix 2. (cont.)

BCOLOGY PLOT CARD - 2
PLOT"
TREES

|  | $\begin{aligned} & \hline<12^{\prime} \\ & \mathrm{CC} / \% \\ & \hline \end{aligned}$ | $\begin{gathered} 12-50 \\ \mathrm{cc} / \% \\ \hline \end{gathered}$ | $\begin{aligned} & 50+ \\ & \mathrm{CC} / \% \end{aligned}$ | Name Code |
| :---: | :---: | :---: | :---: | :---: |
|  |  |  |  | PSME |
|  |  |  |  | ABGR |
|  |  |  |  | ACMA |
|  |  |  |  |  |
| - |  |  |  |  |
| I |  |  |  |  |



DATE:


SHRUBS


Appendix 2. (cont.)

SOILS DESCRIPTION
PLOT"
INV.PLOT" $\qquad$ DATE $\qquad$

| DEPTH | COLOR | TEXTURE | \% COARSE FRAGMENTS <br> (RSIZE) | COMMENTS |
| :--- | :--- | :--- | :--- | :--- | :--- |
|  |  |  |  |  |

SOIL SERIES
SOIL DEPTH REACHED
\% COARSE FRAGMENTS IN TOP 18" OF SOIL
COLOR IN TOP 18"
TEXTURE OF TOP 18"
ADDITIONAL COMMENTS ON SOILS: $\qquad$
$\longrightarrow$

$\qquad$
Appendix 2. (cont.)
HISTORICAL COMMENTAFY
FLOT* INV PLOT \# ..... DATE:
_-_-_-_-_-_-
LOGGING DISTUREANCEOTHEF DISTUFEANCE
$\qquad$
TIME SINCE DISTUREANCE
$\qquad$
DEGREE OF RECOVERY
$\qquad$
SILVICULTURAL HISTOFY
NATURAL HISTORF

STAND SUCCESSION

STAND STRUCTUFE0
OTHEF COMMENTS:

    -  .....       ..... 

Appendix 3. Description of ecology plot data.
**** IMPORTANT NOTE: Field numbers on plot sheets will not correspond to the field numbers listed below, but these numbers will correspond to the field numbers in dBASE III.

Field Name Description and Ranges

| 1. ECOPLOT | Ecology Plot number ( 1-115) Limited to 3 numbers. |
| :---: | :---: |
| 2. INVPLOT | Forest Inventory plot number ( ex. 0701 ) <br> Limited to 4 numbers, the first two numbers indicate transect \#, the 2nd two numbers indicate plot \# within transect. |
| 3.TT | Timber Type. (ex. D4=/gf-) <br> Standard type codes used for Forest timber typing. |
| 4.DATE | Date plot data recorded. (ex. 09/26/89) |
| 5.TRACT | Tract in which plot is located. (ex. 07 ) |
| 6. COMPARTM | Compartment within tract where plot is located. (ex.05) |
| 7. STAND | Stand \# within tract and compartment. (ex. 11) |
| 8. PLANTCOM | Plant Community as defined by dominant tree species, dominant shrub species, dominant herb species, and dominant grass species. (ex. PSME-ACMA/COCO/POMU) |
| 9.TRSEC | Township, Section, and Range; legal description of plot location. Correct way to document: (ex. <br> T.11S.R.5W.SW1/4Sec.18) |
| 10.ELEV | Elevation. Limited to 4 numbers. (ex. 0925) |
| 11. ASPECT | Azimuth. 1-360 degrees. (ex. 285) |
| 12. SLOPE | Slope percent. (ex. 36) |
| 13. MICPOS | ```Microposition: Where on slope plot is located. l=Top 2=Top of slope 3=Mid-slope 4=Bottom of slope 5=Bench 6=Toe of slope 7=Bottom of slope 8=Basin 9=Draw``` |
| 14.MICROV | Vertical Microrelief: Relief on vertical axis, up and down the slope. ```1=Convex 2=Flat 3=Concave 4=Undulating``` |


| Appendix | t.) |
| :---: | :---: |
| 15. MICROH | Horizontal Microrelief: Relief on horizontal axis across the slope. <br> Same 4 descriptions as in MICROV (\#14). |
| 16. TREEM | Percent Thermal Tree Cover: percent of tree crown cover from trees 50' tall or taller. |
| 17. THERM | Percent Hiding Tree Cover: percent of tree crown cover from trees between 12 and 50 feet tall. |
| 18.TREER | Percent Regenerating Tree Cover: percent of tree crown cover from trees less than 12 feet tall. |
| 19. SHRUB | Percent Shrub Cover. |
| 20.FORB | Percent Herb Cover. |
| 21. GRASS | Percent Grass Cover. |
| 22. MOSS | Percent Moss cover. |
| 23.BRGRND | Percent Bare Ground \&/or ground covered by surface coarse fragments that are less than $1 / 16$ inch in diameter. |
| 24.GRAVEL | Percent Gravel. Percent of ground covered by surface coarse fragments that are between $1 / 16$ inch and 3 inches in diameter. |
| 25. ROCK | Percent Rock. Percent of ground covered by coarse fragments that are greater than 3 inches in diameter. |
| 26. BDROCK | Percent Bedrock. Percent of ground covered by exposed bedrock. |
| 27. BROCK | Percent surface bedrock that is covered with moss or litter. |
| 28.TLBA | Total Live Basal Area, expressed in board feet, for conifers only. |
| 29. AGE | Age of stand. |
| 30.HEIGHT | Average height of trees in variable plot. |
| 31. DBH | Average Diameter, in inches, at Breast Height (4.5') of trees in variable plot. |
| 32.SI | Site Index. |
| 33. SDI | Stand Density Index. |
| 34.SNAGS | Number of snags within a 150 foot radius from plot center. |




Appendix 4. Complete species list.
Species used in the analysis are marked with an (*).
List includes all weeds and species that only occurred once or twice in plots.

TREES

| SPP \# | SPP. CODE |
| :--- | :--- |
| $1,2,3$ | ABGR |
| $4,5,6$ | ACMA |
| 7,8, | ALRU |
| 13,14, | ARME |
| 16,17, | CONU |
| $19,20,21$ | FRLA |
| 24, | PIPO |
| $31,32,33$ | PRUNU |
| $34,35,36$ | PREM |
| $40,41,42$ | PSME |
| $46,47,48$ | QUGA |
| $49,50,51$ | TABR |
| $52,53,54$ | THPL |
| $176,177,178$ TSHE |  |

SCIENTIFIC NAME
Abies grandis
Acer macrophyllum
Alnus rubra
Arbutus menziesii
Cornus nuttallii
Fraxinus latifolia
Pinus ponderosa
Prunus spp.
Prunus emarginata
Pseudotsuga menziesii
Quercus garryana
Taxus brevifolia
Thuja plicata
Tsuga heterophylla

COMMON NAME
--------------
grand fir
bigleaf maple
red alder
Pacific madrone
Pacific dogwood
Oregon ash
ponderosa pine
cherry
bitter cherry
Douglas-fir
Oregon white oak
Pacific yew
western redcedar
western hemlock

## SHRUBS

| 55 | ACCI | * |
| :---: | :---: | :---: |
| 56 | AMAL | * |
| 57 | BEAQ | $\star$ |
| 58 | BENE | * |
| 61 | COCO | * |
| 62 | COST |  |
| 63 | CRATA |  |
| 64 | CRMO |  |
| 65 | GASH | * |
| 66 | HODI | * |
| 174 | HOLLY |  |
| 68 | LOCI | * |
| 69 | LOHI | * |
| 70 | OECE |  |
| 71 | RHPU | * |
| 72 | RHDI | * |
| 73 | RIDI | * |
| 74 | ROSA | * |
| 76 | RULA | * |
| 77 | RULE | * |
| 78 | RUPA | * |
| 79 | RUDI | * |
| 81 | RUUR | * |
| 82 | SAGL | * |
| 83 | SALI |  |
| 84 | SAMBU |  |
| 86 | SYAL | * |
| 87 | SYMO | * |
| 88 | VAPA | * |

Acer circinatum
Amelanchier alnifolia
Berberis aquifolium
Berberis nervosa
Corylus cornuta
Cornus stolonifera
Crataegus spp.
Crataegus monogyna
Gaultheria shallon
Holodiscus discolor
Holly spp.
Lonicera ciliosa

Lonicera hispidula
Oemleria cerastiformis
Rhamnus purshiana
Rhus diversiloba
Ribes divaricatum
Rosa spp.
Rubus laciniatus
Rubus leucodermis
Rubus parviflorus
Rubus discolor
Rubus ursinus
Sambucus glauca
Salix spp.
Sambucus spp.
Symphoricarpos albus
Symphoricarpos mollis
Vaccinium parviflorum
vine maple
western serviceberry
Oregon hollygrape
Cascade hollygrape
hazel
red-osier dogwood
hawthorn
one-seed hawthorn
salal
ocean-spray
holly
western trumpet
honeysuckle
hairy honeysuckle
indian plum
cascara buckthorn
poison oak
straggly gooseberry
rose
cutleaf blackberry
black cap raspberry
western thimbleberry
Himalayan blackberry
trailing blackberry
blue elderberry
willow
elderberry spp
common snowberry
red whortleberry

Appendix 4. (Cont.)
FORBS (ferns)

| 91 | BLSP |  |
| :--- | :--- | :--- |
| 92 | POGL |  |
| 93 | POMU | * |
| 94 | PTAQ | * |

Blechnum spicant
Polypodium glycyrrhiza
Polystichum munitum
Pteridium aquilinum
deer fern
licoricefern
western swordfern
bracken fern
FORBS

| 96 | ACTR | * | Achlys triphylla | deerfoot vanillaleaf |
| :---: | :---: | :---: | :---: | :---: |
| 97 | ACRU | * | Actaea rubra | baneberry |
| 98 | ADBI | * | Adenocaulon bicolor | pathfinder |
| 99 | ANDE | * | Anemone deltoidea |  |
| 100 | ANMA |  | Anaphalis margaritaceae | pearly everlasting |
| 101 | AQFO |  | Aquilegia formosa | sitka columbine |
| 102 | ARMI |  | Arctimum minus | common burdock |
| 103 | ARMA | * | Arenaria macrophyllum | bigleaf sandwort |
| 104 | ASCA | * | Asarum caudatum | wild ginger |
| 107 | CASC | * | Campanula scouleri | Scouler's hairbell |
| 108 | CHLE |  | Chrysanthemum |  |
|  |  |  | leucanthemum | oxeye daisy |
| 109 | CIAL |  | Circaea alpina | alpinae circaea |
| 110 | CIEL |  | Cimicifuga elata |  |
| 111 | CIVU |  | Cirsium vulgare |  |
| 113 | COLA | * | Coptis laciniata | cutleaf golden thread |
| 114 | DACA |  | Daucus carrota |  |
| 115 | DIFO | * | Dicentra formosa | Pacific bleedingheart |
| 116 | DIHO | * | Disporum hookeri | Hooker's fairybells |
| 118 | EPILO |  | Epilobium spp. | willow weed |
| 119 | EPWA |  | Epilobium watsonii |  |
| 120 | FRVE | * | Fragaria vesca | common strawberry |
| 121 | FRVI |  | Fragaria virginiana | Virginia strawberry |
| 122 | GAAP | * | Galium aparine | cleavers |
| 123 | GABO |  | Galium boreale | northern bedstraw |
| 124 | GATR | * | Galium triflorum | sweetscented bedstraw |
| 125 | GEMA |  | Geum macrophyllum | largeleaf avens |
| 126 | GERO |  | Geranium rogertianum |  |
| 127 | GOOB | * | Goodyera oblongifolia | rattlesnake plantain |
| 128 | HIAL | * | Hieracium albiflorum | hairy hawkweed |
| 129 | HYOC | * | Hydrophyllum occidentale | western waterleaf |
| 130 | HYPE |  | Hypercum perforatum | common St. Johnswort |
| 131 | HYRA |  | Hypochaeris radicata | spotted catsear |
| 133 | IRTE | * | Iris tenax | Oregon iris |
| 134 | LAMU | * | Lactuca muralis | lettuce |
| 136 | LATHY | * | Lathyrus spp. | vetch |
| 137 | LIAP | * | Ligusticum apifoliium | lovage |
| 138 | LIBO |  | Linnaea borealis | twinflower |
| 175 | MAGR |  | Madia gracilis | spreading tar-weed |
| 139 | MAOR |  | Marah organus | wild cucumber |
| 140 | MOSI | * | Montia siberica | Siberian montia |
| 141 | MOUN |  | Monotropa uniflora | indianpipe |
| 142 | NEPA | * | Nemophilia parviflora | small flowered nemophilia |
| 143 | OSCH | * | Osmorhiza chilensis | sweet mountain cicely |
| 144 | PHACE |  | Phacelia spp. | phacelia |
| 145 | PRVU | * | Prunella vulgaris | common selfheal |
| 146 | PYPI |  | Pyrola picta | whitevein pyrola |
| 147 | RUAC |  | Rumex acetosella | sheep sorrell |
| 148 | SACR | * | Sanicula crassicaulis | snakeroot |

Appendix 4. (Cont.)

| 149 | SADO * | Satureja douglasii |
| ---: | :--- | :--- |
| 150 | SEJA * | Senecio jacobaea |
| 151 | SMRA * | Smilacina racemosa |
| 152 | SMST * | Smilacina stellata |
| 153 | STCR * | Stellaria crispa |
| 154 | SYRE * | Synthyrus reniformis |
| 155 | TEGR * | Tellima grandiflora |
| 156 | THOC * | Thalictrum occidentale |
| 158 | TRLA * | Trientalis latifolia |
| 159 | TROV * | Trillium ovatum |
| 80 | URTIC | Urtica spp. |
| 160 | VAHE * | Vancouveria hexandra |
| 161 | VECAC * | Veratrum californicum |
| 162 | VICIA * | Vaudatum |
| 163 | VIGL * | Vicia spp. |
| 164 | VISE * | Viola sempervierns |

yerba buena
tansy ragwort
false solomon's seal
stary false " "
spring queen
fringecup
western meadowrue
western starflower
Pacific trillium
nettles
inside-out flower
false helibore
vetch
pioneer violet
redwoods violet

GRASSES
166 BRSY *
Brachypodium sylvaticum
168 BRVU *
169 CAREX *
170 DAGL
171 FEOC *
172 HOLA
Bromus vulgaris
Carex spp.
Dactylis glomerata
Festuca occidentalis Holcus lanatus
false brome Columbia brome sedge
orchard grass
western fescue
velvetgrass

Appendix 5. Complete species percent cover by plot matrix ${ }^{3}$.

|  | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | $\begin{aligned} & 1 \\ & 0 \end{aligned}$ | $\begin{aligned} & 1 \\ & 1 \end{aligned}$ | $\begin{aligned} & 1 \\ & 2 \end{aligned}$ | $\begin{aligned} & 1 \\ & 3 \end{aligned}$ | $\begin{aligned} & 1 \\ & 4 \end{aligned}$ | 5 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| ABGRA ${ }^{4}$ | 1.0 | 0.5 | 0.5 | 0.5 | 1.0 | 3.0 | 5.0 | 0.0 | 2.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 1.0 |
| ABGRB | 4.0 | 0.0 | 0.0 | 10.0 | 0.0 | 0.0 | 3.0 | 0.0 | 15.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 |
| ABGRC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 30.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 |
| ACMAA | 0.1 | 1.0 | 2.0 | 0.0 | 0.1 | 0.1 | 3.0 | 1.0 | 0.0 | 1.0 | 1.0 | 2.0 | 2.0 | 2.0 | 1.0 |
| ACMAB | 4.0 | 10.0 | 0.5 | 0.0 | 0.0 | 0.0 | 3.0 | 5.0 | 0.0 | 0.0 | 20.0 | 80.0 | 12.0 | 20.0 | 20.0 |
| ACMAC | 15.0 | 40.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 8.0 | 15.0 | 5.0 | 5.0 | 5.0 | 0.0 | 25.0 | 5.0 |
| ALRUA | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ALRUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ARMEA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 4.0 | 0.0 |
| ARMEB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CONUA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 |
| CONUB | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 |
| FRLAA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 9.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PIPOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 |
| PRUNUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 |
| PRUNUC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMA | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 |
| PREMB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 |
| PREMC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PSMEA | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.5 | 0.0 | 0.0 | 2.0 |
| PSMEB | 1.0 | 0.0 | 20.0 | 0.0 | 0.0 | 4.0 | 0.0 | 2.0 | 4.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PSMEC | 30.0 | 40.0 | 50.0 | 70.0 | 70.0 | 70.0 | 75.0 | 50.0 | 35.0 | 80.0 | 60.0 | 30.0 | 65.0 | 40.0 | 35.0 |
| QUGAA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 |
| QUGAB | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| QUGAC | 5.0 | 1.0 | 25.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TABRA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TABRB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TABRC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ACCI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| AMAL | 0.0 | 0.0 | 0.1 | 3.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| BEAQ | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 3.0 |
| BENE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| COCO | 2.0 | 2.0 | 8.0 | 20.0 | 2.0 | 12.0 | 10.0 | 1.0 | 2.0 | 10.0 | 35.0 | 5.0 | 12.0 | 12.0 | 8.0 |
| GASH | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| HODI | 1.0 | 0.0 | 1.0 | 1.0 | 0.0 | 0.5 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 | 15.0 |
| LOCI | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| LOHI | 0.1 | 0.0 | 0.5 | 0.0 | 0.0 | 2.0 | 0.1 | 0.1 | 0.1 | 0.0 | 0.5 | 0.0 | 2.0 | 1.0 | 0.5 |
| RHPU | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.5 | 0.0 | 0.0 | 0.1 | 0.0 |
| RHDI | 12.0 | 10.0 | 4.0 | 3.0 | 25.0 | 15.0 | 10.0 | 30.0 | 2.0 | 50.0 | 5.0 | 0.5 | 12.0 | 5.0 | 2.0 |

3
Species listed by acronym names along left hand margin. plot numbers listed along the top of page.
4 The $A, B$, or $C$ after the four leter acronym refers to the different strata of trees: $A=$ regeneration (<12' tall)
$B=$ midstory (12-50' tall)
C = overstory (>50'tall)

Appendix 5. (cont.)

|  | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | $\begin{aligned} & 1 \\ & 0 \end{aligned}$ | $\begin{aligned} & 1 \\ & 1 \end{aligned}$ | $\begin{aligned} & 1 \\ & 2 \end{aligned}$ | $\begin{aligned} & 1 \\ & 3 \end{aligned}$ | $\begin{aligned} & 1 \\ & 4 \end{aligned}$ | 5 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| RIDI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ROSA | 0.0 | 1.0 | 0.5 | 5.0 | 1.0 | 2.0 | 1.0 | 0.5 | 0.1 | 0.5 | 0.0 | 0.5 | 3.0 | 0.5 | 0.5 |
| RULA | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| RULE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| RUPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 3.0 | 0.5 | 0.0 | 0.0 | 2.0 |
| RUDI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 3.0 | 0.5 | 0.0 |
| RUUR | 2.0 | 2.0 | 0.5 | 10.0 | 10.0 | 0.0 | 8.0 | 1.0 | 2.0 | 10.0 | 10.0 | 4.0 | 10.0 | 8.0 | 8.0 |
| SAGL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SYAL | 0.1 | 1.0 | 0.0 | 25.0 | 1.0 | 5.0 | 2.0 | 2.0 | 8.0 | 8.0 | 8.0 | 7.0 | 20.0 | 5.0 | 3.0 |
| SYMO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VAPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| POMU | 2.0 | 2.0 | 2.0 | 2.0 | 5.0 | 1.0 | 70.0 | 1.0 | 2.0 | 5.0 | 50.0 | 35.0 | 4.0 | 45.0 | 7.0 |
| PTAQ | 0.0 | 1.0 | 1.0 | 8.0 | 0.0 | 4.0 | 0.0 | 8.0 | 0.0 | 0.0 | 0.5 | 1.0 | 2.0 | 0.0 | 4.0 |
| ACTR | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 2.0 | 0.0 | 0.0 | 0.0 |
| ACRU | 0.1 | 3.0 | 0.0 | 20.0 | 5.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ADBI | 0.1 | 0.1 | 1.0 | 3.0 | 0.0 | 0.1 | 3.0 | 0.0 | 1.0 | 0.5 | 1.0 | 2.0 | 1.0 | 1.0 | 3.0 |
| ANDE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ARMA | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 |
| ASCA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CASC | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| COLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIFO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.1 | 0.1 |
| DIHO | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRVE | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 |
| GAAP | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.1 |
| GATR | 0.1 | 0.1 | 0.5 | 1.0 | 2.0 | 2.0 | 1.0 | 0.5 | 1.0 | 2.0 | 1.0 | 0.1 | 2.0 | 1.0 | 1.0 |
| G00B | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| HIAL | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| HYOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| IRTE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LAMU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LATHY | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| LIAP | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| MOSI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.5 | 1.0 | 0.0 | 0.5 | 0.5 |
| NEPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 |
| OSCH | 0.1 | 0.1 | 1.0 | 1.0 | 1.0 | 2.0 | 0.1 | 0.0 | 1.0 | 5.0 | 2.0 | 1.0 | 2.0 | 2.0 | 1.0 |
| PRVU | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| SACR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.1 | 1.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| SADO | 0.1 | 0.0 | 0.1 | 0.0 | 3.0 | 2.0 | 0.0 | 0.5 | 0.0 | 2.0 | 0.5 | 0.0 | 1.0 | 1.0 | 0.0 |
| SEJA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SMRA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.1 |
| SMST | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 |
| STCR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SYRE | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 |
| TEGR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.5 | 0.1 |
| THOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 |
| TRLA | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 2.0 | 2.0 | 1.0 |
| TROV | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 |
| VAHE | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VECAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VICIA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VIGL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 5.0 | 0.0 | 7.0 |
| VISE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| BRSY | 50.0 | 15.0 | 55.0 | 2.0 | 60.0 | 60.0 | 15.0 | 80.0 | 35.0 | 5.0 | 1.0 | 1.0 | 2.0 | 1.0 | 1.0 |
| BRVU | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 |
| CAREX | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.5 | 0.0 | 0.0 | 0.0 | 2.0 |
| FEOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |

Appendix 5. (cont.)


Appendix 5. (cont.)

|  | 1 | 1 | 1 | 1 | 2 | 2 | 2 | 2 | 2 | 2 | 2 | 2 | 2 | 2 | 3 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 0 |
| POMU | 20.0 | 45.0 | 55.0 | 35.0 | 35.0 | 20.0 | 15.0 | 10.0 | 3.0 | 3.0 | 2.0 | 65.0 | 20.0 | 3.0 | 2.0 |
| PTAQ | 0.1 | 2.0 | 0.5 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.1 | 0.0 | 1.0 | 1.0 | 0.0 |
| ACTR | 0.1 | 2.0 | 2.0 | 3.0 | 8.0 | 5.0 | 4.0 | 3.0 | 4.0 | 0.0 | 7.0 | 4.0 | 0.1 | 1.0 | 3.0 |
| ACRU | 0.0 | 2.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 20.0 | 0.0 | 2.0 | 0.0 | 0.0 |
| ADBI | 2.0 | 0.5 | 5.0 | 3.0 | 4.0 | 2.0 | 1.0 | 3.0 | 3.0 | 2.0 | 5.0 | 3.0 | 0.5 | 2.0 | 6.0 |
| ANDE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.5 | 0.0 | 0.5 | 0.0 | 0.0 | 1.0 | 1.0 |
| ARMA | 2.0 | 0.5 | 0.0 | 3.0 | 0.0 | 0.5 | 1.0 | 0.0 | 1.0 | 0.0 | 1.0 | 0.0 | 0.5 | 1.0 | 2.0 |
| ASCA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CASC | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 1.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.5 | 1.0 |
| COLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIFO | 0.0 | 0.0 | 0.5 | 0.5 | 0.5 | 0.1 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| OIHO | 0.0 | 0.1 | 0.0 | 0.0 | 2.0 | 2.0 | 1.0 | 0.5 | 3.0 | 0.0 | 3.0 | 0.0 | 2.0 | 3.0 | 2.0 |
| FRVE | 1.0 | 0.1 | 0.5 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.1 | 1.0 | 0.5 | 0.1 | 0.1 | 0.0 |
| GAAP | 0.1 | 0.5 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| GATR | 2.0 | 2.0 | 3.0 | 3.0 | 1.0 | 1.0 | 0.0 | 0.5 | 2.0 | 0.5 | 0.0 | 2.0 | 0.5 | 1.0 | 1.0 |
| GOOB | 0.1 | 0.1 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.1 |
| HIAL | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.1 | 1.0 | 0.5 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.1 |
| HYOC | 1.0 | 1.0 | 0.1 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| IRTE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LAMU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 4.0 | 0.0 | 0.0 | 0.0 |
| LATHY | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.5 | 0.0 | 0.0 | 0.5 | 0.0 | 0.5 | 0.0 | 0.0 | 1.0 | 0.1 |
| LIAP | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 10.0 | 4.0 |
| MOSI | 0.1 | 2.0 | 3.0 | 3.0 | 2.0 | 0.5 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| NEPA | 0.1 | 0.1 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 |
| OSCH | 3.0 | 2.0 | 3.0 | 3.0 | 2.0 | 2.0 | 1.0 | 3.0 | 2.0 | 1.0 | 0.5 | 1.0 | 0.1 | 1.0 | 0.5 |
| PRVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SACR | 0.2 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SADO | 3.0 | 0.5 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 2.0 | 0.5 | 0.5 | 0.5 | 0.0 | 0.0 | 0.0 |
| SEJA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 |
| SMRA | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.1 | 0.0 |
| SMST | 0.0 | 25.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| STCR | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SYRE | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.5 | 0.0 | 0.0 | 1.0 | 0.0 |
| TEGR | 0.0 | 0.0 | 0.1 | 0.0 | 2.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 |
| THOC | 0.0 | 3.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 2.0 | 0.0 | 1.0 | 5.0 | 2.0 |
| TRLA | 0.0 | 0.0 | 3.0 | 0.0 | 3.0 | 2.0 | 1.0 | 1.0 | 1.0 | 0.0 | 2.0 | 2.0 | 0.1 | 2.0 | 2.0 |
| TROV | 0.1 | 0.0 | 0.1 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 |
| VAHE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 1.0 | 5.0 | 10.0 | 0.0 | 3.0 | 8.0 | 2.0 | 7.0 | 3.0 |
| VECAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.5 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 |
| VICIA | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VIGL | 0.1 | 15.0 | 3.0 | 1.0 | 25.0 | 3.0 | 0.0 | 0.5 | 1.0 | 0.0 | 2.0 | 0.0 | 0.0 | 2.0 | 2.0 |
| VISE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| BRSY | 1.0 | 1.0 | 2.0 | 2.0 | 1.0 | 5.0 | 0.5 | 3.0 | 30.0 | 20.0 | 25.0 | 2.0 | 0.5 | 2.0 | 2.0 |
| BRVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CAREX | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FEOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |

Appendix 5. (cont.)

|  | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 4 | 4 | 4 | 4 | 4 | 4 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 |
| ABGRA | 0.5 | 1.0 | 1.0 | 0.0 | 0.0 | 0.5 | 0.5 | 0.5 | 1.0 | 5.0 | 0.0 | 0.0 | 0.5 | 0.0 | 3.0 |
| ABGRB | 2.0 | 3.0 | 1.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 3.0 | 0.0 | 0.0 |
| ABGRC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 10.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ACMAA | 1.0 | 0.1 | 2.0 | 1.0 | 0.0 | 3.0 | 0.5 | 3.0 | 4.0 | 5.0 | 1.0 | 0.5 | 1.0 | 2.0 | 5.0 |
| ACMAB | 3.0 | 0.0 | 10.0 | 17.0 | 0.0 | 6.0 | 1.0 | 35.0 | 3.0 | 2.0 | 4.0 | 30.0 | 0.0 | 0.0 | 40.0 |
| ACMAC | 0.0 | 20.0 | 12.0 | 25.0 | 0.0 | 0.0 | 0.0 | 1.0 | 40.0 | 0.0 | 0.0 | 30.0 | 0.0 | 0.0 | 8.0 |
| ALRUA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ALRUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ARMEA | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ARMEB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CONUA | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CONUB | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PIPOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 |
| PRUNUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PSMEA | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 3.0 | 0.5 | 0.5 | 2.0 | 0.0 | 0.0 |
| PSMEB | 12.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 6.0 | 2.0 | 4.0 |
| PSMEC | 80.0 | 55.0 | 75.0 | 65.0 | 95.0 | 80.0 | 75.0 | 75.0 | 55.0 | 80.0 | 65.0 | 55.0 | 60.0 | 65.0 | 60.0 |
| QUGAA | 0.5 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 |
| QUGAB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 |
| QUGAC | 1.0 | 10.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 20.0 | 0.0 | 0.0 |
| TABRA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TABRB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TABRC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ACCI | 0.0 | 0.0 | 0.0 | 0.0 | 40.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| AMAL | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| BEAQ | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 |
| BENE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| COCO | 3.0 | 4.0 | 7.0 | 6.0 | 45.0 | 2.0 | 70.0 | 3.0 | 4.0 | 25.0 | 8.0 | 20.0 | 15.0 | 30.0 | 3.0 |
| GASH | 0.0 | 0.0 | 0.0 | 0.0 | 8.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| HODI | 0.0 | 0.0 | 0.5 | 1.0 | 3.0 | 0.5 | 2.0 | 2.0 | 0.0 | 4.0 | 0.5 | 0.0 | 1.0 | 0.0 | 0.5 |
| LOCI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.5 | 2.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| LOHI | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.5 | 0.0 | 1.0 | 1.0 | 1.0 | 0.0 | 0.0 | 4.0 | 0.0 | 0.0 |
| RHPU | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 | 0.5 | 0.1 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 |
| RHDI | 1.0 | 5.0 | 25.0 | 1.0 | 0.0 | 0.5 | 0.1 | 7.0 | 10.0 | 12.0 | 10.0 | 5.0 | 35.0 | 3.0 | 2.0 |
| RIDI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ROSA | 0.1 | 0.1 | 2.0 | 3.0 | 3.0 | 2.0 | 8.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.5 | 1.0 | 0.0 | 2.0 |
| RULA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 |
| RULE | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 | 2.0 |
| RUPA | 0.0 | 0.0 | 0.0 | 0.1 | 13.0 | 2.0 | 15.0 | 0.0 | 7.0 | 15.0 | 0.0 | 3.0 | 0.0 | 1.0 | 0.0 |
| RUDI | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 10.0 | 2.0 |
| RUUR | 3.0 | 2.0 | 35.0 | 17.0 | 3.0 | 6.0 | 5.0 | 35.0 | 20.0 | 35.0 | 12.0 | 7.0 | 0.0 | 65.0 | 9.0 |
| SAGL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 |
| SYAL | 0.1 | 0.0 | 0.1 | 15.0 | 15.0 | 3.0 | 10.0 | 12.0 | 4.0 | 5.0 | 4.0 | 25.0 | 1.0 | 0.0 | 10.0 |
| SYMO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 |
| VAPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |

Appendix 5. (cont.)

|  | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 3 | 4 | 4 | 4 | 4 | 4 | 4 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 |
| POMU | 2.0 | 8.0 | 3.0 | 20.0 | 12.0 | 10.0 | 2.0 | 10.0 | 75.0 | 8.0 | 85.0 | 12.0 | 1.0 | 30.0 | 7.0 |
| PTAQ | 0.0 | 0.0 | 2.0 | 0.5 | 3.0 | 3.0 | 0.5 | 0.0 | 1.0 | 10.0 | 0.0 | 7.0 | 1.0 | 3.0 | 0.5 |
| ACTR | 0.5 | 0.0 | 1.0 | 0.5 | 5.0 | 1.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ACRU | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ADBI | 5.0 | 25.0 | 3.0 | 4.0 | 3.0 | 3.0 | 2.0 | 1.0 | 0.5 | 0.5 | 2.0 | 3.0 | 2.0 | 0.1 | 1.0 |
| ANDE | 1.0 | 0.1 | 0.1 | 0.5 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ARMA | 0.1 | 0.1 | 2.0 | 0.5 | 0.5 | 1.0 | 2.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 |
| ASCA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CASC | 0.0 | 0.0 | 0.0 | 0.1 | 1.0 | 2.0 | 1.0 | 0.5 | 0.0 | 2.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 |
| COLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIFO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIHO | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 2.0 | 0.5 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRVE | 0.0 | 0.0 | 2.0 | 2.0 | 0.0 | 0.5 | 0.5 | 0.5 | 0.1 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| GAAP | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 |
| GATR | 3.0 | 3.0 | 3.0 | 2.0 | 2.0 | 2.0 | 1.0 | 1.0 | 2.0 | 2.0 | 3.0 | 2.0 | 3.0 | 6.0 | 2.0 |
| GOOB | 0.5 | 0.5 | 0.0 | 0.5 | 0.0 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.1 |
| HIAL | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.5 | 0.1 | 0.0 |
| HYOC | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| IRTE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LAMU | 5.0 | 2.0 | 1.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.5 | 0.0 | 1.0 | 0.0 | 0.0 | 4.0 | 2.0 |
| LATHY | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 1.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LIAP | 0.0 | 0.0 | 0.1 | 0.1 | 1.0 | 0.5 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.1 | 0.0 | 0.0 |
| MOS I | 0.1 | 0.1 | 0.1 | 10.0 | 0.0 | 0.5 | 0.5 | 0.0 | 0.5 | 0.0 | 1.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| NEPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| OSCH | 2.0 | 1.0 | 1.0 | 3.0 | 0.1 | 3.0 | 3.0 | 2.0 | 2.0 | 1.0 | 2.0 | 2.0 | 2.0 | 3.0 | 3.0 |
| PRVU | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SACR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SADO | 0.1 | 0.0 | 0.0 | 3.0 | 0.1 | 0.5 | 0.0 | 1.0 | 1.0 | 1.0 | 3.0 | 0.0 | 2.0 | 0.0 | 2.0 |
| SEJA | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| SMRA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SMST | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.5 | 3.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 |
| STCR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| SYRE | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 2.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TEGR | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| THOC | 0.0 | 0.0 | 0.0 | 0.5 | 0.5 | 1.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TRLA | 0.0 | 0.0 | 0.1 | 2.0 | 3.0 | 2.0 | 1.0 | 2.0 | 1.0 | 3.0 | 0.0 | 2.0 | 0.0 | 0.0 | 2.0 |
| TROV | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.0 | 0.1 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| VAHE | 0.1 | 0.1 | 7.0 | 1.0 | 1.0 | 4.0 | 22.0 | 0.0 | 0.0 | 0.0 | 0.0 | 7.0 | 0.0 | 0.0 | 0.0 |
| VECAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VICIA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VIGL | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 2.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VISE | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| BRSY | 50.0 | 55.0 | 20.0 | 2.0 | 3.0 | 2.0 | 4.0 | 2.0 | 0.1 | 2.0 | 0.5 | 45.0 | 70.0 | 0.5 | 0.5 |
| BRVU | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| CAREX | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FEOC | 0.0 | 0.1 | 0.0 | 0.0 | 0.5 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |

Appendix 5. (cont.)

|  | 4 | 4 | 4 | 4 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 6 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 0 |
| ABGRA | 4.0 | 0.1 | 3.0 | 0.5 | 3.0 | 3.0 | 2.0 | 3.0 | 2.0 | 0.5 | 0.5 | 2.0 | 1.0 | 4.0 | 0.0 |
| ABGRB | 0.0 | 1.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 | 0.0 | 0.0 | 3.0 | 2.0 | 2.0 | 3.0 |
| ABGRC | 0.0 | 10.0 | 15.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ACMAA | 5.0 | 2.0 | 5.0 | 3.0 | 1.0 | 3.0 | 8.0 | 2.0 | 3.0 | 10.0 | 3.0 | 2.0 | 4.0 | 2.0 | . 0 |
| ACMAB | 12.0 | 3.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 30.0 | 30.0 | 5.0 | 7.0 | 0.5 | 7.0 | 4.0 | 0 |
| ACMAC | 0.0 | 15.0 | 0.0 | 0.0 | 0.0 | 0.0 | 6.0 | 55.0 | 0.0 | 0.0 | 0.0 | 7.0 | 5.0 | 4.0 | 0 |
| ALRUA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0 |
| ALRUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ARMEA | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ARMEB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CONUA | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CONUB | 0.0 | 5.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAA | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PIPOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUA | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUC | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | . 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 4.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PSMEA | 11.0 | 0.1 | 0.1 | 0.5 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| PSMEB | 8.0 | 0.0 | 1.0 | 0.0 | 3.0 | 0.0 | 3.0 | 0.0 | 0.0 | 4.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 |
| PSMEC | 65.0 | 60.0 | 55.0 | 90.0 | 85.0 | 65.0 | 80.0 | 35.0 | 65.0 | 75.0 | 80.0 | 90.0 | 50.0 | 80.0 | 50.0 |
| QUGAA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 2.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| QUGAB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0 |
| QUGAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 35.0 | 0.0 | 0.0 |
| TABRA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TABRB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TABRC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ACCI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| AMAL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 |
| BEAQ | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.1 | 1.0 |
| BENE | 0.0 | 2.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| COCO | 2.0 | 2.0 | 10.0 | 2.0 | 7.0 | 4.0 | 7.0 | 8.0 | 35.0 | 3.0 | 17.0 | 2.0 | 6.0 | 2.0 | 6.0 |
| GASH | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| HODI | 0.0 | 0.0 | 0.5 | 0.0 | 4.0 | 1.0 | 0.0 | 0.0 | 1.0 | 0.1 | 0.5 | 5.0 | 9.0 | 2.0 | 10.0 |
| LOCI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LOHI | 2.0 | 0.0 | 10.0 | 7.0 | 0.0 | 1.0 | 1.0 | 0.1 | 4.0 | 0.0 | 2.0 | 2.0 | 0.0 | 8.0 | 0.0 |
| RHPU | 0.5 | 0.1 | 0.1 | 0.1 | 0.1 | 0.0 | 2.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.5 | 0.1 | 0.0 |
| RHOI | 70.0 | 0.0 | 2.0 | 20.0 | 0.5 | 7.0 | 1.0 | 0.5 | 5.0 | 5.0 | 9.0 | 4.0 | 35.0 | 14.0 | 1.0 |
| RIDI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ROSA | 3.0 | 0.0 | 2.0 | 1.0 | 1.0 | 0.5 | 0.1 | 3.0 | 0.1 | 1.0 | 1.0 | 1.0 | 0.1 | 4.0 | 2.0 |
| RULA | 0.0 | 0.0 | 0.5 | 1.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| RULE | 2.0 | 0.0 | 2.0 | 1.0 | 0.5 | 4.0 | 0.5 | 0.0 | 0.0 | 0.1 | 2.0 | 2.0 | 0.0 | 0.0 | 0.0 |
| RUPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| RUDI | 0.0 | 0.0 | 1.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| RUUR | 10.0 | 0.1 | 25.0 | 3.0 | 7.0 | 80.0 | 15.0 | 4.0 | 7.0 | 12.0 | 25.0 | 10.0 | 11.0 | 0.0 | 3.0 |
| SAGL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 8.0 |
| SYAL | 4.0 | 2.0 | 8.0 | 3.0 | 3.0 | 5.0 | 2.0 | 3.0 | 15.0 | 3.0 | 2.0 | 3.0 | 10.0 | 12.0 0.0 | 0.0 |
| SYMO | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VAPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| POMU | 3.0 | 5.0 | 20.0 | 20.0 | 75.0 | 25.0 | 5.0 | 15.0 | 45.0 | 4.0 | 35.0 | 20.0 | 22.0 | 8.0 | 20.0 |

Appendix 5. (cont.)

|  | 4 | 4 | 4 | 4 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 5 | 6 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 0 |
| PTAQ | 0.0 | 3.0 | 8.0 | 10.0 | 4.0 | 2.0 | 6.0 | 0.5 | 10.0 | 9.0 | 3.0 | 0.0 | 4.0 | 0.0 | 3.0 |
| ACTR | 0.0 | 0.1 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| ACRU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| AD8I | 0.5 | 0.0 | 4.0 | 2.0 | 2.0 | 1.0 | 2.0 | 6.0 | 0.5 | 1.0 | 1.0 | 2.0 | 10.0 | 6.0 | 2.0 |
| ANDE | 0.0 | 0.1 | 0.5 | 0.1 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.1 |
| ARMA | 0.0 | 0.5 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 8.0 | 0.5 | 1.0 | 1.0 | 1.0 | 0.0 | 10.0 | 5.0 |
| ASCA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CASC | 0.0 | 0.0 | 0.1 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.5 | 2.0 | 0.0 | 0.5 |
| COLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIFO | 0.0 | 0.0 | 0.0 | 0.0 | 8.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 12.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIHO | 0.0 | 2.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 6.0 | 0.0 | 0.0 | 0.0 | 0.0 | 4.0 | 0.1 | 0.0 |
| FRVE | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 3.0 | 0.5 | 2.0 | 6.0 | 3.0 | 0.5 |
| GAAP | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 |
| GATR | 2.0 | 1.0 | 4.0 | 8.0 | 3.0 | 3.0 | 7.0 | 4.0 | 2.0 | 3.0 | 2.0 | 4.0 | 10.0 | 15.0 | 5.0 |
| G008 | 0.0 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.0 | 0.5 | 0.0 | 1.0 | 0.1 | 0.1 | 0.1 |
| HIAL | 0.0 | 0.0 | 0.1 | 0.1 | 0.5 | 0.1 | 0.0 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.0 | 0.0 | 1.0 |
| HYOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| IRTE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LAMU | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 20.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LATHY | 0.0 | 0.1 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 |
| LIAP | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.1 | 1.0 |
| MOS I | 0.0 | 0.0 | 0.0 | 0.5 | 0.5 | 0.5 | 0.1 | 4.0 | 0.0 | 0.1 | 3.0 | 3.0 | 3.0 | 3.0 | 4.0 |
| NEPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| OSCH | 3.0 | 0.0 | 4.0 | 5.0 | 0.5 | 1.0 | 2.0 | 3.0 | 3.0 | 3.0 | 2.0 | 1.0 | 10.0 | 8.0 | 2.0 |
| PRVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| SACR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SADO | 3.0 | 0.0 | 5.0 | 1.0 | 3.0 | 6.0 | 0.0 | 0.0 | 1.0 | 1.0 | 3.0 | 2.0 | 8.0 | 7.0 | 3.0 |
| SEJA | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.5 | 0.5 | 0.0 | 0.0 | 0.0 |
| SMRA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 |
| SMST | 0.0 | 0.0 | 0.0 | 0.0 | 4.0 | 7.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 3.0 |
| STCR | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| SYRE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.1 |
| TEGR | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.5 | 0.1 | 0.1 |
| THOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 6.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TRLA | 3.0 | 10.0 | 5.0 | 6.0 | 3.0 | 2.0 | 4.0 | 2.0 | 3.0 | 4.0 | 0.5 | 0.0 | 0.0 | 5.0 | 1.0 |
| TROV | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VAHE | 0.0 | 1.0 | 0.0 | 0.5 | 1.0 | 0.0 | 2.0 | 25.0 | 0.0 | 0.1 | 0.0 | 0.0 | 2.0 | 1.0 | 0.0 |
| VECAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VICIA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VIGL | 0.0 | 0.5 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 5.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 | 0.0 |
| VISE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| BRSY | 2.0 | 1.0 | 1.0 | 4.0 | 0.1 | 1.0 | 5.0 | 2.0 | 1.0 | 8.0 | 2.0 | 3.0 | 2.0 | 2.0 | 1.0 |
| 8RVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CAREX | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.5 | 0.0 | 0.1 | 0.0 | 0.5 | 1.0 | 0.0 | 0.0 | 0.0 | 1.0 |
| FEOC | 0.0 | 1.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 4.0 | 3.0 | 0.1 |

Appendix 5. (cont.)

|  | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 7 | 7 | 7 | 7 | 7 | 7 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 |
| ABGRA | 0.0 | 2.0 | 0.0 | 9.0 | 0.0 | 0.1 | 0.5 | 0.0 | 0.5 | 0.0 | 0.0 | 0.1 | 0.1 | 2.0 | 0.0 |
| ABGRB | 0.0 | 3.0 | 0.0 | 20.0 | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 7.0 | 2.0 |
| ABGRC | 0.0 | 0.0 | 0.0 | 7.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 |
| ACMAA | 0.5 | 0.0 | 0.0 | 0.0 | 2.0 | 4.0 | 1.0 | 2.0 | 1.0 | 3.0 | 1.0 | 2.0 | 0.0 | 0.1 | 4.0 |
| ACMAB | 5.0 | 0.0 | 0.0 | 0.0 | 10.0 | 15.0 | 1.0 | 2.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 15.0 |
| ACMAC | 20.0 | 0.0 | 0.0 | 70.0 | 30.0 | 0.0 | 50.0 | 0.0 | 0.0 | 0.0 | 27.0 | 20.0 | 3.0 | 25.0 | 13.0 |
| ALRUA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ALRUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ARMEA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ARMEB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CONUA | 2.0 | 3.0 | 0.0 | 0.0 | 4.0 | 3.0 | 0.0 | 6.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 |
| CONUB | 0.0 | 12.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 25.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 12.0 | 0.0 |
| FRLAA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0 | 1.0 |
| FRLAB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |  | 0.0 |
| FRLAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PIPOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNU8 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 2.0 | 0.0 | 0.0 | 0.0 |
| PREMA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PSMEA | 0.0 | 0.5 | 0.0 | 0.0 | 5.0 | 1.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.5 | 0.0 |
| PSMEB | 3.0 | 0.0 | 3.0 | 0.0 | 10.0 | 5.0 | 0.0 | 0.0 | 2.0 | 1.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PSMEC | 45.0 | 60.0 | 80.0 | 10.0 | 55.0 | 55.0 | 40.0 | 75.0 | 60.0 | 45.0 | 60.0 | 65.0 | 70.0 | 40.0 | 68.0 |
| QUGAA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| QUGAB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | . 0 |
| QUGAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0 | . 0 |
| TABRA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | . 0 |
| TABRB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 12.0 | 0.0 | 0 |
| TABRC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | . 0 |
| THPLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 8.0 | 0.0 | 0.0 | 0.0 | 0 | 0 |
| THPLB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 10.0 | 0.0 | 0.0 | 0.0 | 0.0 | . 0 |
| THPLC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0 | 0 |
| TSHEA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0 |
| TSHEB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0 | 0 |
| TSHEC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | . 0 |
| ACCI | 0.0 | 4.0 | 75.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 75.0 | 30.0 | 0 |
| AMAL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| BEAQ | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.5 | 3.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 |
| BENE | 0.0 | 35.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 0.0 | 8.0 | 0.1 | 0.0 |
| COCO | 25.0 | 11.0 | 25.0 | 4.0 | 20.0 | 15.0 | 6.0 | 15.0 | 5.0 | 2.0 | 7.0 | 2.0 | 3.0 | 15.0 | 0.0 |
| GASH | 0.0 | 6.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 30.0 | 10.0 | 0.0 |
| HODI | 0.0 | 7.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 1.0 | 0.1 | 0.5 | 7.0 | 0.0 | 5.0 | 2.0 | 0.0 |
| LOCI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 |
| LOHI | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 3.0 | 0.0 | 0.0 | 2.0 | 1.0 | 2.0 | 2.0 | 1.0 | 1.0 | 0.0 |
| RHPU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.5 | 2.0 | 0.1 | 0.1 |
| RHDI | 0.0 | 0.0 | 0.0 | 0.0 | 7.0 | 16.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.5 | 1.0 | 0.0 | 0.1 | 0.1 |
| RIDI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ROSA | 6.0 | 8.0 | 4.0 | 0.1 | 2.0 | 2.0 | 1.0 | 2.0 | 1.0 | 0.1 | 1.0 | 0.5 | 2.0 | 3.0 | 0.1 |
| RULA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| RULE | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.5 | 0.0 | 0.5 | 0.0 | 4.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| RUPA | 5.0 | 9.0 | 2.0 | 0.5 | 0.1 | 0.5 | 0.5 | 0.5 | 2.0 | 1.0 | 0.5 | 0.1 | 7.0 | 0.1 | 0.0 |
| RUDI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| RUUR | 2.0 | 6.0 | 6.0 | 4.0 | 10.0 | 12.0 | 12.0 | 0.0 | 15.0 | 4.0 | 6.0 | 7.0 | 5.0 | 4.0 | 3.0 |
| SAGL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5 | 0.0 | 0.0 |
| SYAL | 2.0 | 15.0 | 1.0 | 2.0 | 2.0 | 3.0 | 10.0 | 5.0 | 3.0 | 0.0 | 7.0 | 0.0 | 0.5 | 0.0 | 0.0 |
| SYMO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 |
| VAPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 |

Appendix 5. (cont.)

|  | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 6 | 7 | 7 | 7 | 7 | 7 | 7 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 |
| POMU | 1.0 | 6.0 | 3.0 | 40.0 | 40.0 | 30.0 | 10.0 | 15.0 | 55.0 | 83.0 | 32.0 | 5.0 | 4.0 | 6.0 | 80.0 |
| PTAQ | 15.0 | 6.0 | 0.0 | 2.0 | 0.0 | 0.0 | 5.0 | 2.0 | 9.0 | 0.0 | 5.0 | 2.0 | 1.0 | 1.0 | 0.0 |
| ACTR | 5.0 | 5.0 | 10.0 | 10.0 | 4.0 | 6.0 | 4.0 | 55.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 |
| ACRU | 3.0 | 2.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 3.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| ADBI | 7.0 | 2.0 | 0.1 | 3.0 | 0.1 | 1.0 | 6.0 | 2.0 | 2.0 | 1.0 | 3.0 | 12.0 | 0.1 | 0.5 | 0.5 |
| ANDE | 1.0 | 0.1 | 1.0 | 3.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.1 | 0.1 |
| ARMA | 6.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.5 | 0.5 | 0.5 | 0.0 | 6.0 | 15.0 | 0.1 | 1.0 | 0.0 |
| ASCA | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CASC | 0.5 | 1.0 | 1.0 | 0.0 | 2.0 | 0.0 | 1.0 | 0.1 | 1.0 | 2.0 | 2.0 | 0.5 | 1.0 | 2.0 | 0.0 |
| COLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIFO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| DIH0 | 10.0 | 3.0 | 12.0 | 8.0 | 0.1 | 0.0 | 3.0 | 4.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 |
| FRVE | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 1.0 | 0.5 | 0.0 | 0.0 | 0.1 | 0.5 | 0.0 | 0.1 | 0.1 | 0.0 |
| GAAP | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 3.0 | 0.0 | 2.0 | 1.0 |
| GATR | 3.0 | 1.0 | 1.0 | 3.0 | 2.0 | 5.0 | 2.0 | 2.0 | 2.0 | 2.0 | 1.0 | 0.0 | 1.0 | 1.0 | 1.0 |
| GOOB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.5 | 0.1 | 0.1 | 0.1 | 0.0 |
| HIAL | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.1 | 0.0 | 0.1 | 0.1 | 0.0 |
| HYOC | 0.0 | 0.0 | 0.0 | 4.0 | 0.1 | 1.0 | 0.5 | 0.0 | 0.5 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| IRTE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 |
| LAMU | 0.0 | 0.0 | 0.0 | 0.0 | 20.0 | 3.0 | 1.0 | 1.0 | 0.0 | 0.0 | 0.0 | 38.0 | 0.0 | 0.0 | 0.1 |
| LATHY | 0.5 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 1.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| LIAP | 0.0 | 2.0 | 0.0 | 0.0 | 2.0 | 0.0 | 1.0 | 2.0 | 0.1 | 0.0 | 5.0 | 0.5 | 0.0 | 0.0 | 0.1 |
| MOSI | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.5 | 1.0 | 0.5 | 1.0 | 0.0 | 1.0 | 3.0 | 0.0 | 0.0 | 2.0 |
| NEPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| OSCH | 2.0 | 0.0 | 2.0 | 3.0 | 2.0 | 2.0 | 2.0 | 0.5 | 2.0 | 1.0 | 5.0 | 2.0 | 0.1 | 0.0 | 2.0 |
| PRVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SACR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SADO | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.1 | 1.0 | 0.0 | 0.0 |
| SEJA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 |
| SMRA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| SMST | 0.0 | 6.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 1.0 | 0.1 | 3.0 | 0.0 | 0.0 |
| STCR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 |
| SYRE | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| TEGR | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.1 | 2.0 | 1.0 | 0.0 | 0.5 | 0.1 | 0.0 | 0.0 | 0.0 | 4.0 |
| THOC | 12.0 | 4.0 | 8.0 | 6.0 | 0.0 | 0.0 | 4.0 | 3.0 | 8.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 |
| TRLA | 0.1 | 0.1 | 1.0 | 0.0 | 2.0 | 0.0 | 0.5 | 2.0 | 1.0 | 0.0 | 2.0 | 0.0 | 1.0 | 2.0 | 0.5 |
| TROV | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.5 | 0.0 | 0.1 | 0.0 | 0.1 | 0.1 | 0.1 |
| VAHE | 30.0 | 15.0 | 15.0 | 20.0 | 3.0 | 15.0 | 20.0 | 5.0 | 16.0 | 0.0 | 0.1 | 0.0 | 0.5 | 0.1 | 0.0 |
| VECAC | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| VICIA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VIGL | 8.0 | 5.0 | 7.0 | 0.0 | 0.0 | 1.0 | 3.0 | 3.0 | 2.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.1 | 0.0 |
| VISE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 |
| BRSY | 6.0 | 2.0 | 3.0 | 1.0 | 4.0 | 3.0 | 3.0 | 3.0 | 6.0 | 0.5 | 3.0 | 25.0 | 1.0 | 2.0 | 0.5 |
| BRVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CAREX | 0.0 | 1.0 | 0.0 | 2.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 2.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| FEOC | 1.0 | 0.0 | 2.0 | 2.0 | 2.0 | 0.1 | 0.0 | 0.1 | 0.1 | 0.1 | 1.0 | 0.1 | 0.1 | 0.0 | 0.1 |

Appendix 5. (cont.)

|  | $6$ | $7$ |  | $\begin{aligned} & 7 \\ & 9 \end{aligned}$ | $\begin{aligned} & 8 \\ & 0 \end{aligned}$ | $\begin{aligned} & 8 \\ & 1 \end{aligned}$ | $\begin{aligned} & 8 \\ & 2 \end{aligned}$ | $\begin{aligned} & 8 \\ & 3 \end{aligned}$ | $\begin{aligned} & 8 \\ & 4 \end{aligned}$ | $5$ | $6$ | $\begin{aligned} & 8 \\ & 7 \end{aligned}$ | $\begin{aligned} & 8 \\ & 8 \end{aligned}$ | $\begin{aligned} & 8 \\ & 9 \end{aligned}$ | $\begin{aligned} & 9 \\ & 0 \end{aligned}$ |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| ABGRA | 4.0 | 0.0 | 5.0 | 4.0 | 5.0 | 3.0 | 4.0 | 0.5 | 2.0 | 0.0 | 0.0 | 0.1 | . 0 | 3.0 | 2.0 |
| Abgrb | 0.0 | 0.0 | 35.0 | 0.0 | 2.0 | 5.0 | 0.0 | 0.0 | 6.0 | 0.0 | 0.0 | 0.0 | 0.0 | 7.0 | 0.0 |
| ABGRC | 20.0 | 0.0 | 0.0 | 7.0 | 0.0 | 0.0 | 0.0 | 0.0 | 30.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ACMAA | 6.0 | 0.1 | 3.0 | 1.0 | 3.0 | 0.0 | 5.0 | 30.0 | 0.1 | 2.0 | 5.0 | 0.1 | 2.0 | 0.5 | 0.5 |
| mab | 15.0 | 1.0 | 35.0 | 20.0 | 1.0 | 0.0 | 5.0 | 3.0 | 8.0 | 15.0 | 55.0 | 15.0 | 0.0 | 8.0 | 2.0 |
| ACMAC | 2.0 | 35.0 | 0.0 | 15.0 | 8.0 | 35.0 | 0.0 | 25.0 | 50.0 | 15.0 | 0.0 | 20.0 | 38.0 | 0.1 | 75.0 |
| alrua | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ALRUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| armea | 0.0 | 0.0 | 0.0 | 0.1 | 0.5 | 0.0 | 0.5 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.5 |
| ARMEB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | . 0 |
| conua | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 1.0 | 0.0 | 2.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 |
| CONUB | 0.0 | 3.0 | 2.0 | 2.0 | 0.0 | 4.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 |
| FRLAB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FrLac | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PIPOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| prunua | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| Prunue | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| prunuc | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PSMEA | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 7.0 | 4.0 | 0.0 |
| PSMEB | 3.0 | 0.0 | 0.0 | 10.0 | 5.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 | 12.0 | 1.0 | 0.0 |
| PSMEC | 50.0 | 65.0 | 38.0 | 60.0 | 80.0 | 70.0 | 80.0 | 20.0 | 20.0 | 80.0 | 55.0 | 65.0 | 55.0 | 85.0 | 45.0 |
| Qugat | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.5 | 0.1 | 0.1 |
| qugab | 0.0 | 0.0 | 0.0 | 4.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 8.0 | 0.5 | 0.0 |
| qugac | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 |
| tabra | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| tabrb | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 4.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| tabrc | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ACCI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 25.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| AMAL | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 | 0.0 | 0.1 |
| beap | 0.5 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.1 | 0.0 | 0.5 |
| bene | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 60.0 | 1.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| coco | 12.0 | 1.0 | 7.0 | 25.0 | 12.0 | 8.0 | 20.0 | 0.0 | 20.0 | 22.0 | 50.0 | 3.0 | 3.0 | 8.0 | 10.0 |
| GASH | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| HOOI | 0.5 | 0.0 | 4.0 | 0.0 | 2.0 | 1.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 |
| LOCI | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 2.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| LOHI | 0.0 | 0.0 | 0.0 | 4.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.5 | 0.1 | 0.1 | 1.0 | 0.0 |
| RHPU | 2.0 | 1.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.5 | 0.5 | 0.1 | 0.5 | 0.1 | 0.1 | 0.1 | 0.1 |
| RHDI | 45.0 | 1.0 | 0.5 | 10.0 | 30.0 | 0.0 | 27.0 | 2.0 | 2.0 | 15.0 | 5.0 | 6.0 | 10.0 | 15.0 | 20.0 |
| RIDI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ROSA | 0.0 | 0.0 | 3.0 | 3.0 | 1.0 | 3.0 | 2.0 | 1.0 | 1.0 | 1.0 | 2.0 | 0.5 | 0.5 | 0.5 | 0.5 |
| RULA | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| RULE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| RUPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.5 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| RUDI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 |
| RUUR | 12.0 | 10.0 | 6.0 | 0.0 | 14.0 | 3.0 | 15.0 | 5.0 | 4.0 | 40.0 | 8.0 | 0.0 | 1.0 | 12.0 | 18.0 |
| SAGL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SYAL | 3.0 | 3.0 | 4.0 | 1.0 | 10.0 | 8.0 | 8.0 | 0.1 | 0.0 | 3.0 | 3.0 | 0.1 | 0.5 | 0.1 | 2.0 |
| SYM0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VAPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |

Appendix 5. (cont.)

|  | 7 | 7 | 7 | 7 | 8 | 8 | 8 | 8 | 8 | 8 | 8 | 8 | 8 | 8 | 9 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 0 |
| POMU | 3.0 | 45.0 | 4.0 | 20.0 | 15.0 | 10.0 | 1.0 | 6.0 | 40.0 | 5.0 | 10.0 | 1.0 | 0.0 | 2.0 | 8.0 |
| PTAQ | 0.1 | 0.0 | 4.0 | 0.1 | 0.0 | 0.0 | 5.0 | 0.5 | 0.0 | 0.1 | 0.0 | 0.0 | 0.5 | 0.0 | 20.0 |
| ACTR | 0.0 | 0.0 | 10.0 | 1.0 | 0.0 | 10.0 | 0.0 | 2.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ACRU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ADBI | 3.0 | 0.5 | 7.0 | 1.0 | 0.5 | 2.0 | 6.0 | 2.0 | 0.5 | 0.1 | 0.1 | 0.1 | 1.0 | 1.0 | 2.0 |
| ANDE | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 4.0 | 0.0 | 0.1 | 1.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 |
| ARMA | 0.0 | 0.0 | 1.0 | 0.0 | 8.0 | 0.0 | 5.0 | 0.0 | 0.0 | 5.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| ASCA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CASC | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.5 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| COLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIF0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIH0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.1 | 0.5 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRVE | 0.1 | 0.0 | 0.1 | 0.0 | 0.5 | 0.0 | 3.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 |
| GAAP | 1.0 | 0.1 | 0.1 | 0.0 | 2.0 | 1.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| GATR | 2.0 | 3.0 | 1.0 | 2.0 | 2.0 | 2.0 | 0.5 | 6.0 | 0.0 | 2.0 | 3.0 | 0.5 | 0.5 | 0.1 | 2.0 |
| G00B | 0.1 | 1.0 | 0.1 | 1.0 | 0.1 | 0.1 | 0.0 | 0.5 | 0.1 | 0.1 | 0.0 | 0.1 | 0.1 | 0.1 | 0.0 |
| HIAL | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 | 0.0 |
| HYOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| IRTE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LAMU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 10.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LATHY | 0.0 | 0.0 | 2.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LIAP | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 8.0 | 0.0 | 0.0 | 3.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.5 |
| MOSI | 0.5 | 12.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 3.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| NEPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| OSCH | 8.0 | 8.0 | 0.5 | 2.0 | 15.0 | 3.0 | 1.0 | 6.0 | 2.0 | 3.0 | 2.0 | 4.0 | 2.0 | 2.0 | 2.0 |
| PRVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| SACR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SADO | 3.0 | 1.0 | 0.0 | 2.0 | 7.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.5 | 8.0 |
| SEJA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SMRA | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SMST | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| STCR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SYRE | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 5.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TEGR | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THOC | 0.0 | 0.0 | 5.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TRLA | 1.0 | 1.0 | 1.0 | 2.0 | 2.0 | 4.0 | 6.0 | 0.0 | 4.0 | 2.0 | 2.0 | 3.0 | 0.0 | 0.0 | 0.0 |
| TROV | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VAHE | 0.0 | 0.0 | 10.0 | 0.5 | 0.0 | 3.0 | 2.0 | 10.0 | 10.0 | 2.0 | 0.0 | 0.1 | 0.1 | 0.0 | 4.0 |
| VECAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VICIA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VIGL | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VISE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| BRSY | 2.0 | 2.0 | 1.0 | 2.0 | 3.0 | 0.5 | 3.0 | 4.0 | 2.0 | 5.0 | 2.0 | 2.0 | 35.0 | 35.0 | 35.0 |
| BRVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CAREX | 0.5 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 |
| FEOC | 1.0 | 0.0 | 0.1 | 0.1 | 0.5 | 2.0 | 1.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 |

Appendix 5. (cont.)

|  |  |  |  |  |  |  |  |  |  | 1 | 1 | 1 | 1 | 1 | 1 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 9 | 9 | 9 | 9 | 9 | 9 | 9 | 9 | 9 | 0 | 0 | 0 | 0 | 0 | 0 |
|  | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 |
| ABGRA | 0.0 | 0.0 | 7.0 | 1.0 | 10.0 | 2.0 | 2.0 | 0.0 | 8.0 | 5.0 | 7.0 | 3.0 | 10.0 | 6.0 | 1.0 |
| ABGRB | 0.0 | 0.0 | 4.0 | 0.0 | 0.0 | 5.0 | 10.0 | 0.0 | 6.0 | 15.0 | 15.0 | 3.0 | 20.0 | 0.0 | 0.0 |
| ABGRC | 0.0 | 0.0 | 45.0 | 0.0 | 0.0 | 0.0 | 8.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 | 25.0 | 0.0 | 0.0 |
| ACMAA | 1.0 | 0.0 | 0.1 | 0.0 | 0.0 | 1.0 | 0.1 | 1.0 | 0.1 | 2.0 | 0.1 | 1.0 | 0.1 | 0.1 | 2.0 |
| ACMAB | 8.0 | 5.0 | 3.0 | 5.0 | 0.0 | 0.0 | 20.0 | 30.0 | 5.0 | 1.0 | 10.0 | 4.0 | 0.0 | 40.0 | 25.0 |
| ACMAC | 0.0 | 25.0 | 30.0 | 20.0 | 0.0 | 0.0 | 25.0 | 30.0 | 25.0 | 0.0 | 70.0 | 18.0 | 0.0 | 20.0 | 30.0 |
| ALRUA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ALRUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ARMEA | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| ARMEB | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 4.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 |
| CONUA | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.5 | 0.5 | 4.0 | 0.0 | 0.0 | 0.1 |
| CONUB | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 12.0 | 0.0 | 0.0 | 0.0 | 15.0 | 4.0 | 0.0 | 0.0 |
| FRLAA | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 |
| FRLAB | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PIPOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PSMEA | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 3.0 | 0.0 | 1.0 | 1.0 | 0.5 | 2.0 | 0.0 | 0.5 |
| PSMEB | 0.0 | 3.0 | 0.0 | 10.0 | 2.0 | 0.0 | 1.0 | 1.0 | 6.0 | 5.0 | 0.0 | 0.0 | 1.0 | 1.0 | 0.0 |
| PSMEC | 65.0 | 45.0 | 40.0 | 40.0 | 50.0 | 70.0 | 60.0 | 35.0 | 55.0 | 60.0 | 40.0 | 60.0 | 40.0 | 60.0 | 65.0 |
| QUGAA | 1.0 | 0.1 | 2.0 | 0.5 | 0.1 | 0.1 | 0.1 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 |
| QUGAB | 0.0 | 0.0 | 0.0 | 0.0 | 15.0 | 1.0 | 0.5 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 |
| QUGAC | 0.0 | 3.0 | 0.0 | 20.0 | 25.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TABRA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TABRB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TABRC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ACCI | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 6.0 | 0.0 | 70.0 | 0.0 | 0.0 |
| AMAL | 0.0 | 0.0 | 0.1 | 3.0 | 3.0 | 0.5 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| BEAQ | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 40.0 | 2.0 | 0.0 | 0.0 |
| BENE | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.5 | 15.0 | 0.0 | 0.0 | 0.0 | 0.0 | 8.0 | 0.0 | 0.0 |
| COCO | 10.0 | 1.0 | 15.0 | 0.5 | 1.0 | 8.0 | 10.0 | 3.0 | 7.0 | 18.0 | 12.0 | 15.0 | 9.0 | 15.0 | 6.0 |
| GASH | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 3.0 | 0.0 | 0.0 |
| HOOI | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 4.0 | 0.5 | 0.0 | 0.5 | 0.5 | 7.0 | 9.0 | 0.0 | 0.1 |
| LOCI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.5 | 0.1 | 0.1 | 0.0 | 0.1 | 0.5 | 0.0 | 0.0 |
| LOHI | 0.1 | 0.1 | 1.0 | 0.1 | 1.0 | 0.0 | 5.0 | 0.0 | 0.0 | 6.0 | 1.0 | 0.0 | 2.0 | 0.1 | 0.1 |
| RHPU | 0.0 | 0.1 | 0.1 | 0.1 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 0.5 |
| RHOI | 10.0 | 8.0 | 15.0 | 20.0 | 25.0 | 12.0 | 4.0 | 0.0 | 0.1 | 30.0 | 25.0 | 0.0 | 5.0 | 3.0 | 3.0 |
| RIOI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ROSA | 2.0 | 0.5 | 1.0 | 4.0 | 4.0 | 0.5 | 4.0 | 0.5 | 1.0 | 2.0 | 2.0 | 8.0 | 2.0 | 1.0 | 2.0 |
| RULA | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| RULE | 4.0 | 3.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 |
| RUPA | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.1 | 0.5 | 1.0 | 0.1 | 0.1 | 0.1 |
| RUDI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 |
| RUUR | 15.0 | 0.0 | 8.0 | 10.0 | 3.0 | 18.0 | 7.0 | 0.0 | 3.0 | 12.0 | 9.0 | 5.0 | 6.0 | 6.0 | 12.0 |
| SAGL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SYAL | 0.1 | 0.1 | 0.5 | 3.0 | 8.0 | 7.0 | 0.0 | 0.5 | 0.0 | 3.0 | 5.0 | 5.0 | 0.0 | 3.0 | 0.5 |
| SYMO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 25.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| VAPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |

Appendix 5. (cont.)

|  |  |  |  |  |  |  |  |  |  | 1 | 1 | 1 | 1 | 1 | 1 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 9 | 9 | 9 | 9 | 9 | 9 | 9 | 9 | 9 | 0 | 0 | 0 | 0 | 0 | 0 |
|  | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 |
| POMU | 30.0 | 4.0 | 3.0 | 4.0 | 0.0 | 6.0 | 5.0 | 14.0 | 14.0 | 4.0 | 5.0 | 20.0 | 3.0 | 10.0 | 13.0 |
| PTAQ | 0.0 | 4.0 | 0.0 | 0.0 | 0.0 | 8.0 | 0.0 | 0.0 | 0.5 | 1.0 | 1.0 | 0.0 | 0.0 | 4.0 | 4.0 |
| ACTR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 65.0 | 2.0 | 0.1 | 0.1 | 9.0 | 0.0 | 2.0 | 0.0 |
| ACRU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ADBI | 0.0 | 0.1 | 4.0 | 0.0 | 0.1 | 0.0 | 2.0 | 4.0 | 0.1 | 2.0 | 2.0 | 2.0 | 2.0 | 5.0 | 3.0 |
| ANDE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.5 | 0.0 | 0.0 |
| ARMA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 10.0 | 0.5 | 0.1 | 0.0 | 0.5 | 0.5 | 0.5 | 1.0 | 0.0 |
| ASCA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CASC | 0.0 | 0.0 | 0.1 . | 0.0 | 0.0 | 0.0 | 2.0 | 3.0 | 0.0 | 0.5 | 0.5 | 0.5 | 1.0 | 0.5 | 0.0 |
| COLA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIFO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIHO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 1.0 | 0.1 | 0.0 | 0.0 | 2.0 | 2.0 | 0.0 | 0.0 |
| FRVE | 0.0 | 0.0 | 0.1 | 0.1 | 1.0 | 0.0 | 0.5 | 0.0 | 0.1 | 2.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| GAAP | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| GATR | 0.1 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 1.0 | 0.5 | 0.1 | 1.0 | 2.0 | 1.0 | 2.0 | 0.5 | 2.0 |
| GOOB | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 | 0.0 | 0.5 | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 |
| HIAL | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.1 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 |
| HYOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| IRTE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LAMU | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 |
| LATHY | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LIAP | 0.0 | 0.0 | 3.0 | 0.1 | 7.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.5 | 2.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| MOSI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 5.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 |
| NEPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| OSCH | 0.1 | 2.0 | 2.0 | 2.0 | 3.0 | 0.0 | 4.0 | 3.0 | 1.0 | 2.0 | 8.0 | 4.0 | 4.0 | 2.0 | 2.0 |
| PRVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SACR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SADO | 0.1 | 0.5 | 0.0 | 3.0 | 8.0 | 0.0 | 5.0 | 0.0 | 0.0 | 3.0 | 2.0 | 0.0 | 2.0 | 0.0 | 0.0 |
| SEJA | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| SMRA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SMST | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 1.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 1.0 | 0.0 | 0.0 |
| STCR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| SYRE | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 3.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 |
| TEGR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| THOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 7.0 | 0.0 | 0.0 | 0.0 | 4.0 | 0.0 | 0.1 | 0.0 |
| TRLA | 0.0 | 0.1 | 3.0 | 2.0 | 0.0 | 0.0 | 4.0 | 6.0 | 0.5 | 1.0 | 8.0 | 2.0 | 6.0 | 1.0 | 0.1 |
| TROV | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 | 0.0 | 0.0 |
| VAHE | 0.0 | 0.0 | 3.0 | 0.0 | 2.0 | 0.0 | 4.0 | 3.0 | 4.0 | 3.0 | 2.0 | 7.0 | 2.0 | 4.0 | 2.0 |
| VECAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.5 | 0.0 | 0.0 |
| VICIA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VIGL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 |
| VISE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| BRSY | 50.0 | 90.0 | 25.0 | 75.0 | 85.0 | 85.0 | 2.0 | 1.0 | 75.0 | 12.0 | 10.0 | 1.0 | 1.0 | 90.0 | 68.0 |
| BRVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CAREX | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FEOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 1.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 |

Appendix 5. (cont.)

|  | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 0 | 0 | 0 | 0 | 1 | 1 | 1 | 1 | 1 | 1 |
|  | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 |
| ABGRA | 1.0 | 0.0 | 1.0 | 0.0 | 0.0 | 1.0 | 0.0 | 3.0 | 0.0 | 2.0 |
| ABGRB | 11.0 | 0.0 | 0.0 | 3.0 | 0.0 | 5.0 | 2.0 | 2.0 | 0.0 | 2.0 |
| ABGRC | 4.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 5.0 | 6.0 | 0.0 | 4.0 |
| ACMAA | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 4.0 | 0.1 |
| ACMAB | 2.0 | 5.0 | 0.5 | 0.5 | 15.0 | 0.0 | 8.0 | 3.0 | 2.0 | 8.0 |
| ACMAC | 55.0 | 10.0 | 35.0 | 8.0 | 10.0 | 0.0 | 0.0 | 25.0 | 0.0 | 25.0 |
| ALRUA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ALRUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ARMEA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| ARMEB | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CONUA | 0.0 | 0.0 | 5.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CONUB | 0.0 | 0.0 | 0.0 | 23.0 | 5.0 | 5.0 | 18.0 | 7.0 | 0.0 | 0.0 |
| FRLAA | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FRLAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PIPOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PRUNUA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| PRUNUB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 2.0 |
| PRUNUC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PREMC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| PSMEA | 0.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.5 |
| PSMEB | 0.0 | 1.0 | 10.0 | 17.0 | 5.0 | 0.0 | 0.0 | 4.0 | 0.0 | 2.0 |
| PSMEC | 55.0 | 40.0 | 20.0 | 38.0 | 60.0 | 15.0 | 70.0 | 55.0 | 75.0 | 41.0 |
| QUGAA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| QUGAB | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| QUGAC | 4.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TABRA | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.1 | 0.0 |
| TABRB | 0.0 | 5.0 | 5.0 | 12.0 | 2.0 | 5.0 | 0.0 | 5.0 | 0.0 | 2.0 |
| TABRC | 0.0 | 0.0 | 0.0 | 6.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLA | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 |
| THPLB | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| THPLC | 0.0 | 0.0 | 0.0 | 0.0 | 2.0 | 10.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEA | 0.0 | 4.0 | 0.1 | 2.0 | 0.0 | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEB | 0.0 | 15.0 | 0.0 | 1.0 | 5.0 | 6.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| TSHEC | 0.0 | 40.0 | 25.0 | 4.0 | 0.0 | 60.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ACCI | 0.0 | 9.0 | 25.0 | 20.0 | 20.0 | 55.0 | 14.0 | 4.0 | 1.0 | 35.0 |
| AMAL | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 |
| BEAQ | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| BENE | 0.0 | 6.0 | 7.0 | 9.0 | 12.0 | 10.0 | 7.0 | 0.0 | 1.0 | 2.0 |
| COCO | 18.0 | 0.0 | 3.0 | 3.0 | 3.0 | 0.0 | 0.0 | 10.0 | 0.0 | 6.0 |
| GASH | 0.0 | 4.0 | 17.0 | 6.0 | 10.0 | 4.0 | 6.0 | 15.0 | 2.0 | 8.0 |
| HODI | 0.0 | 0.1 | 5.0 | 0.0 | 2.0 | 0.0 | 0.0 | 1.0 | 0.0 | 4.0 |
| LOCI | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LOHI | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 |
| RHPU | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.0 | 0.0 |
| RHDI | 35.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| RIDI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ROSA | 0.0 | 1.0 | 1.0 | 0.5 | 3.0 | 2.0 | 1.0 | 3.0 | 0.1 | 1.0 |
| RULA | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| RULE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| RUPA | 0.0 | 0.1 | 0.5 | 0.0 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 |
| RUDI | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| RUUR | 12.0 | 1.0 | 1.0 | 0.5 | 0.1 | 0.5 | 0.1 | 1.0 | 1.0 | 1.0 |
| SAGL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 4.0 |
| SYAL | 3.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 3.0 | 0.0 | 2.0 |
| SYMO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VAPA | 0.0 | 0.1 | 1.0 | 0.0 | 2.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 |

Appendix 5. (cont.)

|  | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | 0 | 0 | 0 | 0 | 1 | 1 | 1 | 1 | 1 | 1 |
|  | 6 | 7 | 8 | 9 | 0 | 1 | 2 | 3 | 4 | 5 |
| POMU | 7.0 | 12.0 | 30.0 | 10.0 | 30.0 | 8.0 | 20.0 | 12.0 | 5.0 | 15.0 |
| PTAQ | 3.0 | - 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 4.0 | 1.0 | 0.0 |
| ACTR | 0.0 | 0.0 | 0.5 | 0.1 | 0.0 | 0.0 | 0.0 | 0.1 | 0.5 | 2.0 |
| ACRU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ADBI | 2.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| ANDE | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 | 0.1 | 0.5 | 0.1 |
| ARMA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| ASCA | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CASC | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 2.0 | 0.5 |
| COLA | 0.0 | 0.0 | 0.5 | 0.0 | 1.0 | 0.5 | 0.1 | 0.0 | 0.0 | 0.0 |
| DIFO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| DIHO | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | 0.5 | 0.1 |
| FRVE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 |
| GAAP | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| GATR | 1.0 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 0.5 | 0.1 | 0.1 | 0.5 |
| GOOB | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| HIAL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 |
| HYOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| IRTE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| LAMU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| LATHY | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| LIAP | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| MOS I | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 |
| NEPA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| OSCH | 3.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 |
| PRVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SACR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SADO | 1.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SEJA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SMRA | 0.0 | 0.1 | 0.1 | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.1 | 0.5 |
| SMST | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| STCR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| SYRE | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| TEGR | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.0 | 0.1 | 0.0 |
| THOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 1.0 |
| TRLA | 0.0 | 0.5 | 0.0 | 0.1 | 0.1 | 0.1 | 1.0 | 2.0 | 0.1 | 6.0 |
| TROV | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.0 | 0.0 | 1.0 | 0.0 |
| VAHE | 2.0 | 0.0 | 0.5 | 0.0 | 1.0 | 0.0 | 0.0 | 0.5 | 1.0 | 2.0 |
| VECAC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 |
| VICIA | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VIGL | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| VISE | 0.0 | 0.5 | 0.1 | 0.0 | 0.1 | 0.1 | 0.1 | 0.1 | 0.0 | 0.1 |
| BRSY | 30.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.5 |
| BRVU | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CAREX | 0.0 | 0.1 | 0.0 | 0.0 | 0.1 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| FEOC | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.1 | 0.1 | 0.1 |

Appendix 6. Wildlife species list.*

* Compiled from species observed on McDonald-Dunn from a study titled
"Comparisons of terrestrial vertebrate communities and tree
regeneration among 3 silvicultural systems in the east-central Coast
Range, Oregon." by McComb and Chambers (1989).
Bird species
Golden-crowned kinglet
Chestnut-backed chickadee
Hermit warbler
Winter wren
Wilson's warbler
Brown creeper
Dark-eyed junco
Swainson's thrush
American Robin
Black-headed grosbeak
Read-breasted nuthatch
Black-throated gray warbler
Orange-crowned warbler
Western flycatcher
Stellers jay
Bushtit
Evening grosbeak
Rufous hummingbird
Common flicker
Pileated woodpecker
Red-breasted sapsucker
Hairy woodpecker
Gray jay
Mountain quail
Rufous-sided towhee
Downy woodpecker
Hammond's flycatcher
Blue grouse
Red crossbill
Olive-sided flycatcher
Western tanager
Band-tailed pigeon
Pine siskin
Hermit thrush
Hutton's vireo
Townsend's warbler
Common raven
Purple finch
Pygmy owl
Western wood pewee
Willow flycatcher
White-crowned sparrow
Red-tailed hawk
American goldfinch

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Appendix 6. (cont.)
Bird species
California quail
Warbling vireo
Chipping sparrow
MacGillivray's warbler
Mourning dove
Scrub jay
Violet-green swallow
Song sparrow
Sharp-shinned hawk
House wren
Turkey vulture
Ruffed grouse
Varied thrush
Mammals
Douglas squirrel
Townsend's chipmunk
Western gray squirrel
Black-tailed deer
Trowbridge's shrew
Deer mouse
Pacific shrew
Coast mole
Creeping vole
California red-backed vole
Vagrant shrew
Townsend's chipmunk
Red tree vole
Long-tailed weasel
Coyote
Brush rabbit
Garter snake
Rough-skinned newt
Elk*
Bobcat*
*observed by Leavell and Hubbard during data collection in the summer of 1989.
```

Appendix 7. Wildlife species on McDonald-Dunn Forest that use snag cavities.*

| Species that use snags | Cavity users | Diameter ** |
| :---: | :---: | :---: |
| BIRDS |  |  |
| turkey vulture | NO |  |
| northern pygmy owl | YES | 17 |
| red-breasted sapsucker | YES | 15 |
| downy woodpecker | YES | 11 |
| hairy woodpecker | YES | 15 |
| pileated woodpecker | YES | 25 |
| olive-sided flycatcher | NO |  |
| Hammond's flycatcher | NO |  |
| western flycatcher | NO |  |
| chestnut-backed chickadee | YES | 9 |
| northern flicker | YES | 17 |
| red-breasted nuthatch | YES | 17 |
| brown creeper | YES | 15 |
| house wren | YES | 15 |
| winter wren | NO |  |
| MAMMALS |  |  |
| bobcat | YES | 29 |
| western grey squirrel | YES | 17 |
| deer mouse | YES | 15 |

* from Neitro et al. (1985)
** suggested diameter suitable for nesting

Appendix 8. Complete TWINSPAN two-way table (with 68 species, and 108 plots).

| Scale for cover percent values in body of table |  |  |
| :--- | :--- | :--- |
| $1=>0-5 \%$ | $4=21-30 \%$ | $7=51-60 \%$ |
| $2=6-10 \%$ | $5=31-40 \%$ | $8=61-75 \%$ |
| $3=11-20 \%$ | $6=41-50 \%$ | $9=76-100 \%$ |

Appendix 8. Complete TWINSPAN two-way table (with 68 species and 108 plots).

PLOT Numbers

| 1 | 1 | 11 | 1 | 1 | 1111111 |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 339404 | 899990 | 24 | 989090855678677011834455554435868 | 1351117124757126269 | 222339123675680336687702211001011 | 129246391246512568539580630289060519136685815670432563719424879715007048884346067594783142752313431235780912



000000000000000000000000000000000000000000000000000000000000000000001111111111111111111111111111111111111111 00000000000000000000000000011111111111111111111111111111111111110000000000000000000000000000000011111111 0000011111111111111111111100000000000000000000000000011111111111110000000000000000000000011111111100111111 0001100000000000000000111110000000000000111111111111110000000001111100000011111111111111111000000011000001

Appendix 9. Location of Tracts in McDonald-Dunn Forest.



[^0]:    1 Differential species are those with clear ecological preference (Hill, 1979b). Ecological preference is preference for habitat or environment.

[^1]:    , Constancy: the percent of plots in which the species occurred.
    2 Coverage: average cover for those plots in which the species occurred.
    ${ }^{3}$ The $A, B$, and $C$ following the acronym names stand for the different tree strata: $A=$ Regeneration ( $>12^{\prime}$ tall), $B=$ midstory ( $12-50$ ' tall), and C = overstory (50+' tall).

[^2]:    2 Throughout this section "significant" differences refers to alpha $=.05$.

[^3]:    Pseudotsuqa menziesii/Holodiscus discolor-Berberis nervosa
    Pseudotsuga menziesii/Holodiscus discolor/grass
    Pseudotsuqa menziesii/Symphoricarpos mollis
    Pseudotsuqa menziesii-Tsuga heterophylla/Berberis nervosa Pseudotsuqa menziesii-Tsuga heterophylla/Gaultheria shallon Abies grandis/Berberis nervosa

    Tsuga heterophylla/Berberis nervosa
    Tsuga heterophylla/Berberis nervosa/Gaultheria shallon
    Tsuga heterophylla/Berberis nervosa/achlys triphlla
    Tsuga heterophylla/Gaultheria shallon
    Tsuga heterophylla/Polystichum munitum
    Tsuga heterophylla/achlys triphlla
    association described by Hemstrom et al. (1987) also
    contains pines (Pinus lambertiana and Pinus ponderosa)
    which are absent in McDonald-Dunn Forest associations. The
    Abies grandis association also contains Tsuga heterophylla, which excludes it from an Abies grandis association on McDonald-Dunn Forest.

[^4]:    - Indicates field observations

