

Supplemental File 3

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FA19 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
MS11 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
F62 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
1291 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
PID332 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
SK-92-679 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
PID1 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
SK-93-1035 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
35/02 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
DGI18 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
FA6140 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
PID24-1 MKFIDEAKIEVAAGKGGNGATSFREKFFVPRGGPDGGDGGKGGSVWAEADENTNTLVEYR
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FA19 FVKRYQAKNGEKHGSDRYGAGADDIVLKMPVGTILRDLDTDEIVADLTYHGQRVCLAKG
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FA19 GKGGGLGNIHFKSSVNRAPKQSTPGEETRSLOLELKVLDVGLLGMNPAGKSTLITAVS
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FA19 AARPKIANYPF^TTLHPNLGVVRIDENHSFVMADIPGLIEGAAEGAGLGHRFLKHL^SRTGL
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PID1 AARP KIANYPFTTLHPNLGVVRIDENHSFVMADIPGLIEGAAEGAGLGHRFLKHL SRTGL
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FA6140 LLHVVDLAPFDETVNPAAEALAI INELRKYDEELYGKPRWLVLNKLDMLDEEEARARTAA
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DGI2 LLHVVDLAPFDETVNPAAEALAI INELRKYDEELYGKPRWLVLNKLDMLDEEEARARTAA
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SK-92-679 FLEAVGWDYPEPDRFQFDMETPRLFQISAL THQGTQELVHQINQYLAEKKR IEAEKAEA
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FA6140 EKAAANVEIIEQQPKTDTGVFKPE
PID24-1 EKAAANVEIIEQQPKTDTGVFKPE
DGI2 EKAAANVEIIEQQPKTDTGVFKPE
PID18 EKAAANVEIIEQQPKTDTGVFKPE
