

ALS87661.1 1 10 20 30 40 50 60
CDO72823.1:13-345 MGFRSLALSLGLVCNGLANVISKRVCSPRLCCISLLCFLKHAALLTDQAHAAQKIYD
.....

ALS87661.1 70 80 90 100 110 120
CDO72823.1:13-345 FIVIGGGTAGSVIANRLSENPGVRVLLIEAGVNNAQGPHVNELQVPYFVSSIDPSFDWNY
.....

ALS87661.1 130 140 150 160 170 180
CDO72823.1:13-345 TTVPQSALQNQTIPYPRGHVLGGSSSTNFMFYTRGSSDEFDRLARVTGDEGWSWRNIFSY
.....MFYTRGSSDEFDRLARVTGDEGWSWRNIFSY

ALS87661.1 190 200 210 220 230 240
CDO72823.1:13-345 ILKTEMLSPPADGHNTTGQVNPSVHGHAGPLLTSLPGNATVLDSTRVIETTSQISEFPFNL
ILKTEMLSPPADGHNTTGQVNPSVHGHAGPLLTSLPGNATVLDSTRVIETTSQISEFPFNL

ALS87661.1 250 260 270 280 290 300
CDO72823.1:13-345 DMNSGNPLGVGWLQSTIGHGMRSSAATAFLTDVVLGRENLDVLIGTRVTRLIRTEQFLST
DMNSGNPLGVGWLQSTIGHGMRSSAATAFLTDVVLGRENLDVLIGTRVTRLIRTEQFLST

ALS87661.1 310 320 330 340 350 360
CDO72823.1:13-345 PTFGLVELAQSSASGPRISLTARKEVILSAGSIGTAQILMLSGIGDRQELSAHKINTLVLDL
PTFGLVELAQSSASGPRISLTARKEVILSAGSIGTAQILMLSGIGDRQELSAHKINTLVLDL

ALS87661.1 370 380 390 400 410 420
CDO72823.1:13-345 PDVGKNMQDHPWVPLQWMVNSNDTLD SINRNP ELMSQFVALYNATKQGPIANNP GGNQIG
PDVGKNMQDHPWVPLQWMVNSNDTLD SINRNP ELMSQFVALYNATKQGPIANNP GGNQIG

ALS87661.1 430 440 450 460 470 480
CDO72823.1:13-345 WFRLPDDSPILKQFGDPTAGPLSPHFELTFGNSFLSFNQPEPDTGNFMSMCVALVAPTSR
WFRLPDDSPILKQFGDPTAGPLSPHFELTFGNSFLSFNQPEPDTGNFMSMCVALVAPTSR

ALS87661.1 490 500 510 520 530 540
CDO72823.1:13-345 GS VTLASASAFDAPLIDPAFMQTQSDLA ILTEAVKAAHRFAAAPAWKDFIIAPFKDAAQT
GS

ALS87661.1 550 560 570 580 590 600
CDO72823.1:13-345 TSDDDIHAYIRRMVSTFRHPMGTAMMSGPLDSKGVVNPDLRVKKVNGLRVVDASIFPHIF
.....

ALS87661.1 610 620 630
CDO72823.1:13-345 GAHLQAPVYAIAERAADLIKEAHGIPLSHHHHHHH
.....