

Figures

Figure 1

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MGDYKDDDDK MEPRDGSPEA RSSDSESASA SSSGSDERDAG PEPDKAPRRL NKRRFPGLRL FGHRKAITKS GLQHLAPPPP TPGAPCSESE
RQIRSTVDWS ESATYGEHIW FETNVSGDFC YVGEQYCVAR MLKSVSRKRC AACKIVVHTP CIEQLEKINF RCKPSFRESG SRNVREPTFV
RHHWVHRRRQ DGKCRHCGKG FQKFTFHSK EIVAISCSWC KQAYHSKVSC FMLQIEEPC SLGVHAAYVI PPTWILRARR PQNTLKASKK
KKRASFKRKS SKKGPEEGRW RPFIRPTPS PLMKPLLVFV NPKSGGNQGA KIIQSFLWYL NPRQVFDLSQ GGPKEALEMY RKVHNLRLA
CGGDGTVGWI LSTLDQLRLK PPPPVAILPL GTGNDLARTL NWGGGYTDEP VSKILSHVEE GNVVQLDRWD LHAEPNPEAG PEDRDEGATD
RLPLDVFNNY FSLGFDHVT LEFHESREAN PEKFNSRFRN KMFYAGTAFS DFLMGSSKDL AKHIRVVCDG MDLTPKIQDL KPQCVMFLNI
PRYCAGTMPW GHPGEHDFE PQRHDDGYLE VIGFTMTSLA ALQVGGHGER LTQCREVLT TSKAIPVQVD GEPCKLAASR IRIALRNQAT
MVQKAKRRSA APLHSDQQPV PEQLRIQVSR VSMHDYEALH YDEQLKEAS VPLGTVVVPG DSDLELCRAH IERLQQEPDG AGAKSPTCQK
LSPKWCFLDA TTASRFYRID RAQEHLNYVT EIAQDEIYIL DPELLGASAR PDLPTPTSPL PTSPCSPTPR SLQGDAAPPQ GEELIEAAKR
NDFCKLQELH RAGGDLMHRD EQSRTLHHA VSTGSKDVVR YLLDHAPPEI LDAVEENGET CLHQAAALGQ RTICHYIVEA GASLMKTDQQ
GDTPRQRAEK AQDTELAAYL ENRQHYQMIQ REDQETAV

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Figure 2

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Ser Ile Ile Asn Phe Glu Lys Leu
1 5

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Figure 3

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Ser Met Leu Val Leu Leu Pro Asp Glu Val Ser Gly Leu Glu Gln Leu
1 5 10 15
Glu Ser Ile Ile Asn Phe Glu Lys Leu Thr Glu Trp Thr Ser
20 25 30

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Figure 4

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Asp Tyr Lys Asp Asp Asp Asp Lys
1 5

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Figure 5

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Gly Cys Cys Ala Cys Cys
1 5

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Figure 6







