

logo

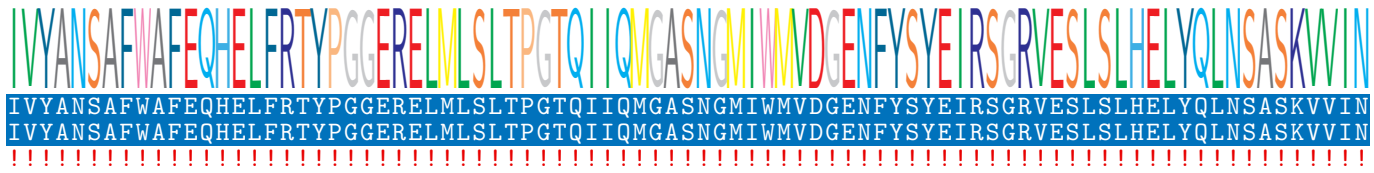
NTDB id 97262 ASZ79 RS10630 WP 001911421.1
NTDB id 1151 GTF74 RS04205 WP 001911421.1
consensus



85
85

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
NTDB id 1151 GTF74 RS04205 WP 001911421.1
consensus



170
170

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
NTDB id 1151 GTF74 RS04205 WP 001911421.1
consensus



255
255

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
NTDB id 1151 GTF74 RS04205 WP 001911421.1
consensus



340
340

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
NTDB id 1151 GTF74 RS04205 WP 001911421.1
consensus



425
425

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
NTDB id 1151 GTF74 RS04205 WP 001911421.1
consensus



510
510

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
NTDB id 1151 GTF74 RS04205 WP 001911421.1
consensus



595
595

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
NTDB id 1151 GTF74 RS04205 WP 001911421.1
consensus

QDNSSLIIIGSQFGVKTNTEELAVSNLPNIQVTFSSQVSVNHTTKLLGYRMSEPLKIAYGDSISFQFGALPSSRSQSLEYKLNDEQ
QDNSSLIIIGSQFGVKTNTEELAVSNLPNIQVTFSSQVSVNHTTKLLGYRMSEPLKIAYGDSISFQFGALPSSRSQSLEYKLNDEQ
QDNSSLIIIGSQFGVKTNTEELAVSNLPNIQVTFSSQVSVNHTTKLLGYRMSEPLKIAYGDSISFQFGALPSSRSQSLEYKLNDEQ
!!

680
680

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
NTDB id 1151 GTF74 RS04205 WP 001911421.1
consensus

QMQRFDMALLNFDQLLPGKYRLKVRSANPTQRYRTEALDFEVLQPWYMTSIALVGYVLSVLGLLALALWWRTRMILAANRKLKA
QWQRFDMALLNFDQLLPGKYRLKVRSANPTQRYRTEALDFEVLQPWYMTSIALVGYVLSVLGLLALALWWRTRMILAANRKLKA
QWQRFDMALLNFDQLLPGKYRLKVRSANPTQRYRTEALDFEVLQPWYMTSIALVGYVLSVLGLLALALWWRTRMILAANRKLKA
!!

765
765

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
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consensus

QVELKTSQLRHQSKIVLSNNHQLRKQLQLHHTILGNVLDNIDASLRHLSSYAKVRAWSEFEAPFNKVKQELACLHFMHRGDEDES
QVELKTSQLRHQSKIVLSNNHQLRKQLQLHHTILGNVLDNIDASLRHLSSYAKVRAWSEFEAPFNKVKQELACLHFMHRGDEDES
QVELKTSQLRHQSKIVLSNNHQLRKQLQLHHTILGNVLDNIDASLRHLSSYAKVRAWSEFEAPFNKVKQELACLHFMHRGDEDES
!!

850
850

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
NTDB id 1151 GTF74 RS04205 WP 001911421.1
consensus

KYHDLHLLIESALKSWQEEYSKALIKLTYSGEPRFVEVRQFNLDVLFNAVLTDIAIKRKYKHQELTIQLEVRNAQAVIVFSDFGAP
KYHDLHLLIESALKSWQEEYSKALIKLTYSGEPRFVEVRQFNLDVLFNAVLTDIAIKRKYKHQELTIQLEVRNAQAVIVFSDFGAP
KYHDLHLLIESALKSWQEEYSKALIKLTYSGEPRFVEVRQFNLDVLFNAVLTDIAIKRKYKHQELTIQLEVRNAQAVIVFSDFGAP
!!

935
935

logo

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consensus

IQNI DKTMTVTLGSSSYPLEELVIRSGGVLHVLALKERNVIELAWPLATLPELETIHKPEEGEQAAKVGIDEVEKEWLSKVEKLIIE
IQNIDKTMVTLGSSSYPLEELVIRSGGVLHVLALKERNVIELAWPLATLPELETIHKPEEGEQAAKVGIDEVEKEWLSKVEKLIIE
IQNIDKTMVTLGSSSYPLEELVIRSGGVLHVLALKERNVIELAWPLATLPELETIHKPEEGEQAAKVGIDEVEKEWLSKVEKLIIE
!!

1020
1020

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
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consensus

LHYSDPNFTTSTAAQALVVSERSLQRRFKAATGKTFKESLNEVRLEKACQSLLSGAKIAQVAFDCGFNDPSYFSQRFKHHFGLSP
LHYSDPNFTTSTAAQALVVSERSLQRRFKAATGKTFKESLNEVRLEKACQSLLSGAKIAQVAFDCGFNDPSYFSQRFKHHFGLSP
LHYSDPNFTTSTAAQALVVSERSLQRRFKAATGKTFKESLNEVRLEKACQSLLSGAKIAQVAFDCGFNDPSYFSQRFKHHFGLSP
!!

1105
1105

logo

NTDB id 97262 ASZ79 RS10630 WP 001911421.1
NTDB id 1151 GTF74 RS04205 WP 001911421.1
consensus

SQFI EDQEN
SQFI EDQEN 1114
SQFI EDQEN 1114
!!!!!!!!!!!!

non conserved
 similar
 $\geq 50\%$ conserved