

logo

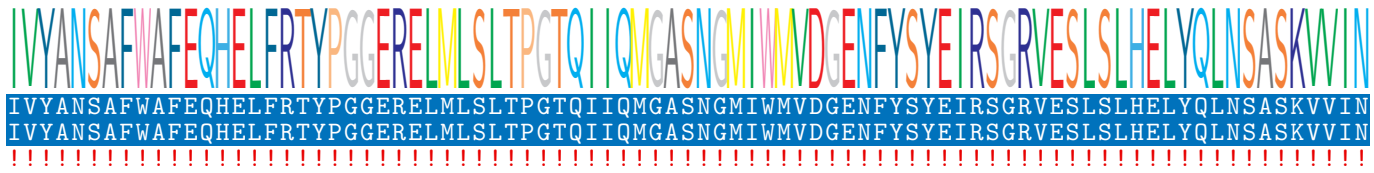
NTDB id 40889 KW3 RS0108645 WP 001911421.1  
NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus



85  
85

logo

NTDB id 40889 KW3 RS0108645 WP 001911421.1  
NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus



170  
170

logo

NTDB id 40889 KW3 RS0108645 WP 001911421.1  
NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus



255  
255

logo

NTDB id 40889 KW3 RS0108645 WP 001911421.1  
NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus



340  
340

logo

NTDB id 40889 KW3 RS0108645 WP 001911421.1  
NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus



425  
425

logo

NTDB id 40889 KW3 RS0108645 WP 001911421.1  
NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus



510  
510

logo

NTDB id 40889 KW3 RS0108645 WP 001911421.1  
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consensus



595  
595

logo

NTDB id 40889 KW3 RS0108645 WP 001911421.1  
NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus

QDNSSLIIIGSQFGVKTNTEELAVSNLPNIQVTFSSQVSVNHHTTKLLGYRMSEPLKIAYGDSISFQFGALPSSRSQSLEYKLNDEQ  
QDNSSLIIIGSQFGVKTNTEELAVSNLPNIQVTFSSQVSVNHHTTKLLGYRMSEPLKIAYGDSISFQFGALPSSRSQSLEYKLNDEQ  
QDNSSLIIIGSQFGVKTNTEELAVSNLPNIQVTFSSQVSVNHHTTKLLGYRMSEPLKIAYGDSISFQFGALPSSRSQSLEYKLNDEQ  
!!

680  
680

logo

NTDB id 40889 KW3 RS0108645 WP 001911421.1  
NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus

QWQRFDWALLNFDQLLPGKYRLKVRSANPTQRYRTEALDFEVLQPWYMTSIALVGYVLSVLGLLALALWWRTRMILAANRKLKA  
QWQRFDWALLNFDQLLPGKYRLKVRSANPTQRYRTEALDFEVLQPWYMTSIALVGYVLSVLGLLALALWWRTRMILAANRKLKA  
QWQRFDWALLNFDQLLPGKYRLKVRSANPTQRYRTEALDFEVLQPWYMTSIALVGYVLSVLGLLALALWWRTRMILAANRKLKA  
!!

765  
765

logo

NTDB id 40889 KW3 RS0108645 WP 001911421.1  
NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus

QVELKTSQLRHQSKIVLSNNHQLRKQLQLHHTILGNVLDNIDASLRHLSSYAKVRAWSEFEAPFNKVKQELAQLFHMRGDEDES  
QVELKTSQLRHQSKIVLSNNHQLRKQLQLHHTILGNVLDNIDASLRHLSSYAKVRAWSEFEAPFNKVKQELAQLFHMRGDEDES  
QVELKTSQLRHQSKIVLSNNHQLRKQLQLHHTILGNVLDNIDASLRHLSSYAKVRAWSEFEAPFNKVKQELAQLFHMRGDEDES  
!!

850  
850

logo

NTDB id 40889 KW3 RS0108645 WP 001911421.1  
NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus

KYHDLHLLIESALKSWQEEYSKALIKLTYSGEPRFVEVRQFNLDVLFNAVLTDIAIKRKYKHQELTIQLEVRNAQAVIVFSDFGAP  
KYHDLHLLIESALKSWQEEYSKALIKLTYSGEPRFVEVRQFNLDVLFNAVLTDIAIKRKYKHQELTIQLEVRNAQAVIVFSDFGAP  
KYHDLHLLIESALKSWQEEYSKALIKLTYSGEPRFVEVRQFNLDVLFNAVLTDIAIKRKYKHQELTIQLEVRNAQAVIVFSDFGAP  
!!

935  
935

logo

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NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus

IQNIDKTMVTLGSSSYPLEELVIRSGGVLHVLALKERNVIELAWPLATLPELETIHKPEEGEQAAKVGIDEVEKEWLSKVEKLIIE  
IQNIDKTMVTLGSSSYPLEELVIRSGGVLHVLALKERNVIELAWPLATLPELETIHKPEEGEQAAKVGIDEVEKEWLSKVEKLIIE  
IQNIDKTMVTLGSSSYPLEELVIRSGGVLHVLALKERNVIELAWPLATLPELETIHKPEEGEQAAKVGIDEVEKEWLSKVEKLIIE  
!!

1020  
1020

logo

NTDB id 40889 KW3 RS0108645 WP 001911421.1  
NTDB id 1151 GTF74 RS04205 WP 001911421.1  
consensus

LHYSDPNFTTSTAAQALVVSERSLQRRFKAATGKTFKESLNEVRLEKACQSLLSGAKIAQVAFDCGFNDPSYFSQRFKHHFGLSP  
LHYSDPNFTTSTAAQALVVSERSLQRRFKAATGKTFKESLNEVRLEKACQSLLSGAKIAQVAFDCGFNDPSYFSQRFKHHFGLSP  
LHYSDPNFTTSTAAQALVVSERSLQRRFKAATGKTFKESLNEVRLEKACQSLLSGAKIAQVAFDCGFNDPSYFSQRFKHHFGLSP  
!!

1105  
1105

logo

NTDB id 40889 KW3 RS0108645 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