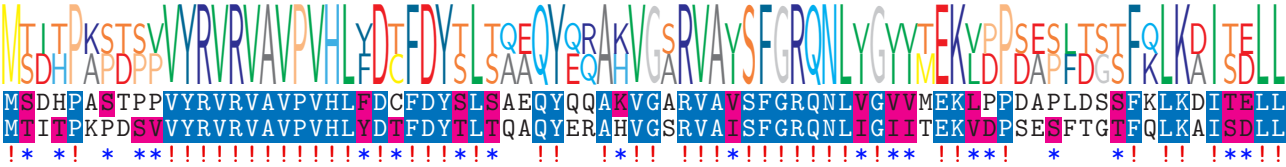


logo

NTDB id 251126 BS636 RS03825 WP 099337588.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



80
80

logo

NTDB id 251126 BS636 RS03825 WP 099337588.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



159
160

logo

NTDB id 251126 BS636 RS03825 WP 099337588.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



239
240

logo

NTDB id 251126 BS636 RS03825 WP 099337588.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



319
320

logo

NTDB id 251126 BS636 RS03825 WP 099337588.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



399
400

logo

NTDB id 251126 BS636 RS03825 WP 099337588.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



479
480

logo

NTDB id 251126 BS636 RS03825 WP 099337588.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus



559
560

logo

NTDB id 251126 BS636 RS03825 WP 099337588.1

NTDB id 1072 ABD1 RS01805 WP 000156662.1

consensus

YVTLVAILDIDSGLLSVD_{RA}PERTAQLIVQVAGRAGRGERKGDVYLQTLRPDHALNTLYTKNYRAFAKQTL_{AE}RQYAR

YVTLVAILDIDSGLLSVD_{FR}TERTAQLIVQVAGRAGRGERKGDVYLQTLRPDHALNTLYTDNYRAFAKQTLEQRQ_{IA}AR

!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!*!!!!!!!!!!!!!!*!!*!!*!!*!!*!!*!!*!!

639

640

logo

NTDB id 251126 BS636 RS03825 WP 099337588.1

NTDB id 1072 ABD1 RS01805 WP 000156662.1

consensus

MPPYRYAVMIRCESRDRTYTHFLQKHA_{EV}LRQYQGLQVEIWGPIPA_{ME}RKAGRY_{RS}HMV_{LS}ADRARLHYYRQ_{AA}AA

MPPYRYAVMIRCESRDQTLNTEFLQKHAAVLRQYQDLQLEIWGPIPA_{ME}RKAGRYQSHMV_{LS}ADRARLHYYR_{AA}WQA

!!!!!!!!!*!*!*!*!!!!!!!!*!*!*!!*!!*!!*!!*!!*!!*!!*!!*!!*!!*!!*!!*!!*!!*!!

719

720

logo

NTDB id 251126 BS636 RS03825 WP 099337588.1

NTDB id 1072 ABD1 RS01805 WP 000156662.1

consensus

MLQD_{LP}SSMKLSLDVDPQE_{LS}

MLQD_{LP}SSMKLSLDVDPQE_{LS}

*** **** **!!!!!!!!!!!!!!

740

742

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved