

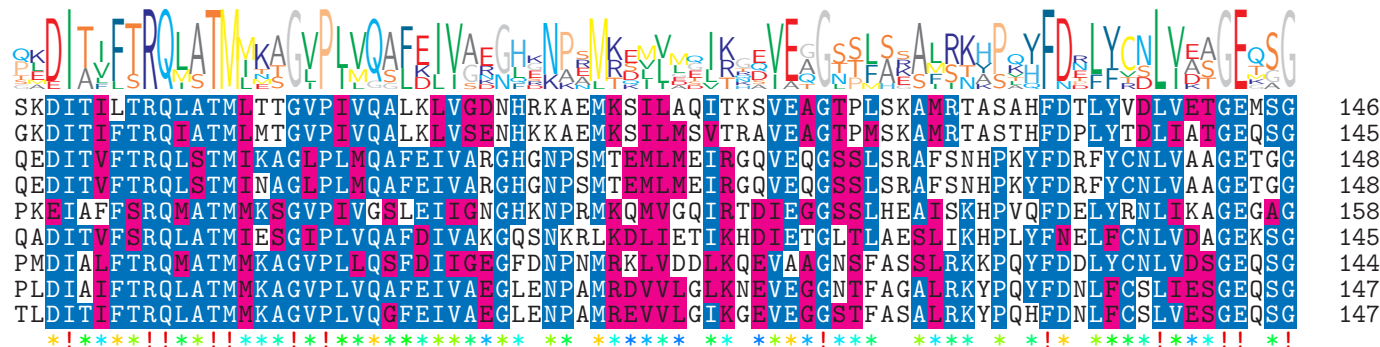
logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451()
NTDB id 1042306 ABQF06 RS05160 WP 228442817.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus



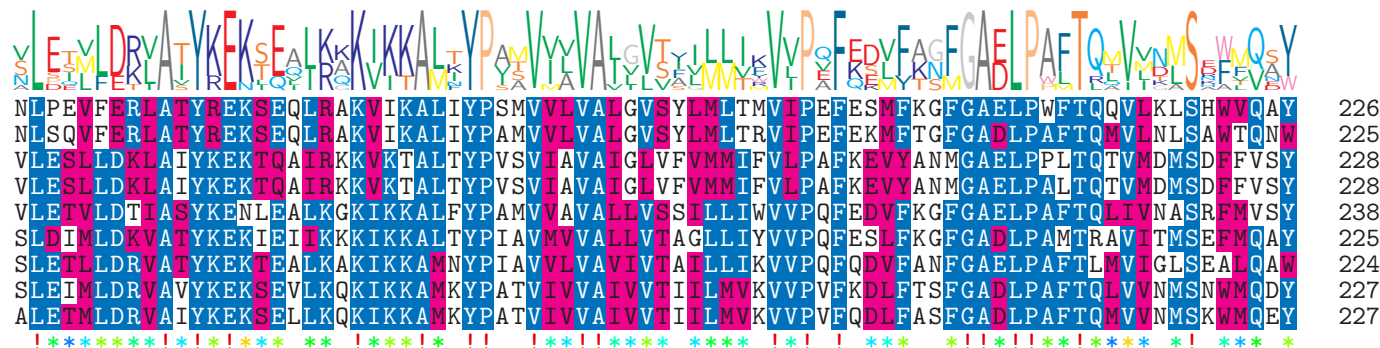
logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451()
NTDB id 1042306 ABQF06 RS05160 WP 228442817.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus



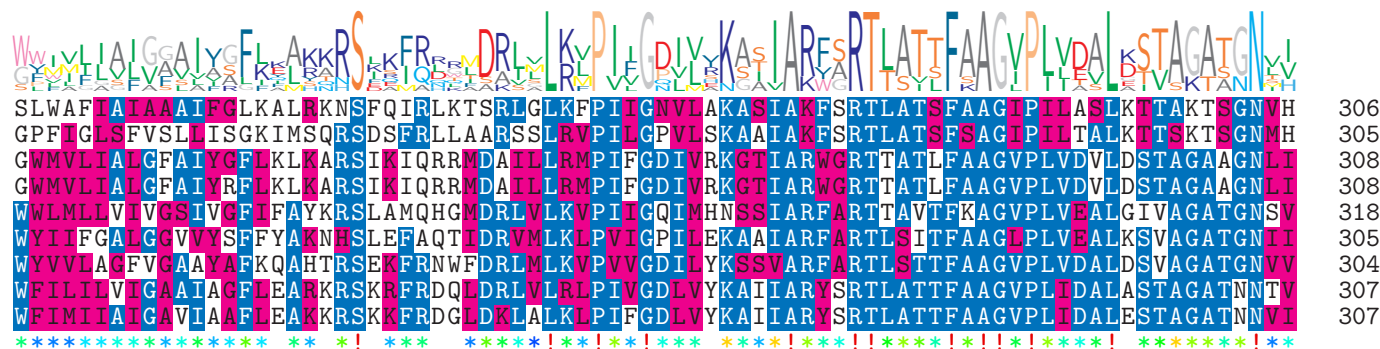
logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451()
NTDB id 1042306 ABQF06 RS05160 WP 228442817.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus



logo

NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451()
NTDB id 1042306 ABQF06 RS05160 WP 228442817.1
NTDB id 1252 GC085 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus



logo

NTDB id 1169	A1552VC	RS11075	WP 000648511.1	YEATKIREDA	386
NTDB id 1402	DSB67	RS12670	WP 010643257.1	YQLAIEEVYRDTAAGMPMYIAMRNTEAFPEMVLQMVMIGEESGQLDDMLNKVATIIYEF	385
NTDB id 1112	NGFG	RS09215	WP 003689811.1	YEEATREIRTRVIQGLSMTSGMRATELFPNMMLQMSSIGEESGSLDDMLNKAAEFY	388
NTDB id 1113	AAA85695.1	219..1451()		YEEATREIRTRVIQGLSMTSGMRATELFPNMMLQMSSIGEESGSLDDMLNKAAEFY	388
NTDB id 1042306	ABQF06	RS05160	WP 228442817.1	YEKAVLRMRREDVSVGYPVNVSMKQVNLFPHMVIQMTAIGEEAGALDAMLFKVAEY	398
NTDB id 1252	GC085	RS07730	WP 011213805.1	YAKATDKIREEVATGQQMFIAMENTHLFPNMVIQMVAIGEESGALEKMLSKVADF	385
NTDB id 1198	PSJM300	03950	AFN76868.1	FRSATNKKISDVSSGMLNFSMRTTGTFFPSMAVQMTAIGEESGALDEMLS	384
NTDB id 1016	ACIAD	RS01680	WP 004920476.1	YEQAVLKIREDVSTGQQLNFAVRVSNRFPTMAIQMVAIGEESGALDEMLDKVAN	387
NTDB id 1059	ABD1	RS01610	WP 000279215.1	YEKAVMKIREDVATGQQLQFAMRISNRFPSTMAIQMVAIGEESGALDSMLDKVAT	387
consensus					

<0

logo

NTDB id 1169	A1552VC	RS11075	WP 000648511.1	TVVGGGLV	408
NTDB id 1402	DSB67	RS12670	WP 010643257.1	VVVGGLVT	407
NTDB id 1112	NGFG	RS09215	WP 003689811.1	LVIGTLLV	410
NTDB id 1113	AAA85695.1	219..1451()		LVIGTLLV	410
NTDB id 1042306	ABQF06	RS05160	WP 228442817.1	TVVGGMVIGMYLP	420
NTDB id 1252	GC085	RS07730	WP 011213805.1	ILVGGLV	406
NTDB id 1198	PSJM300	03950	AFN76868.1	VLVGGLII	405
NTDB id 1016	ACIAD	RS01680	WP 004920476.1	ILVGGLVI	408
NTDB id 1059	ABD1	RS01610	WP 000279215.1	VLVGGLVI	408
consensus					

- non conserved
- similar
- ≥ 50% conserved