

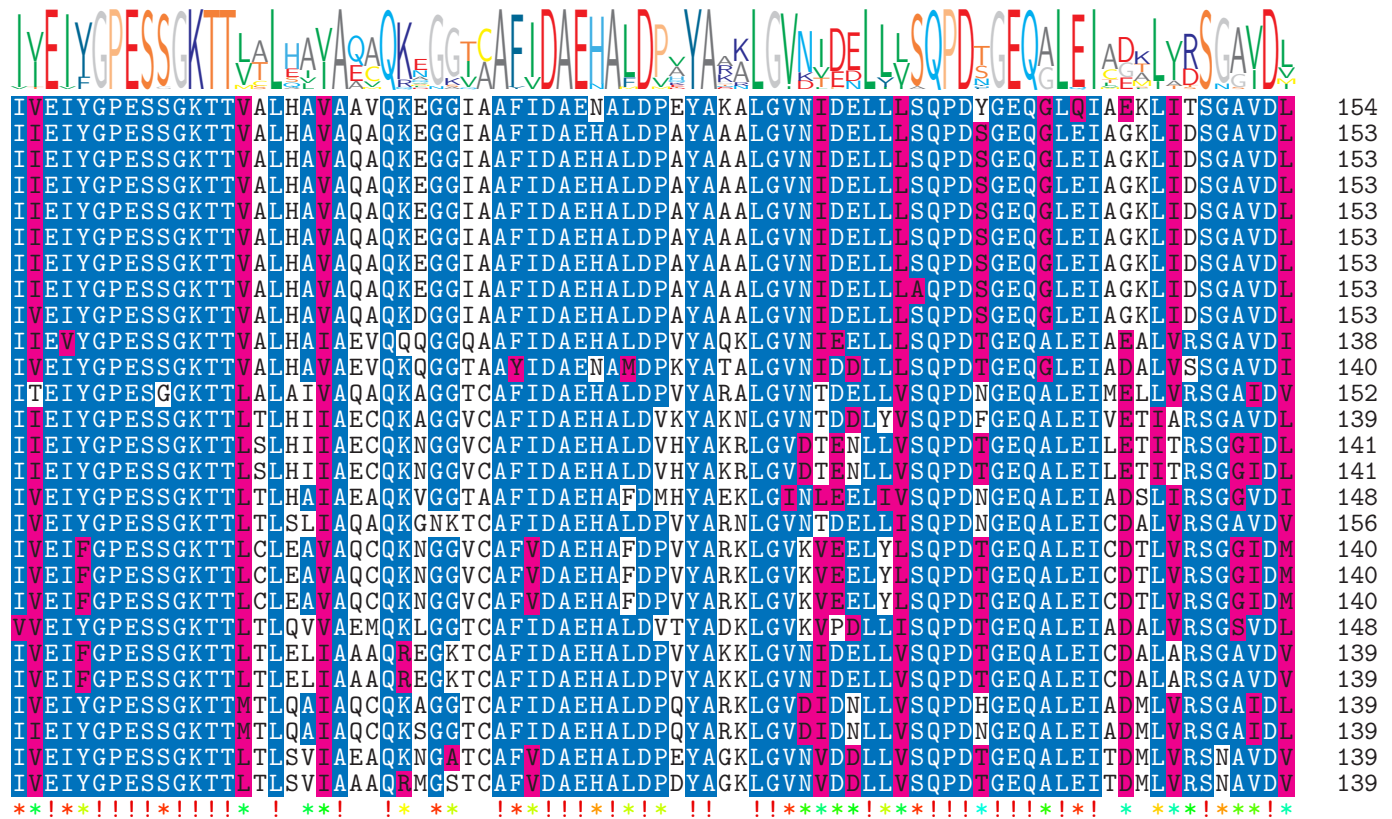
logo

NTDB id 600 KW2 RS01765 WP 011675413.1	MATKKK....TNFDDITKKYGA	ERDKALADALALIEKD	FGKGSIMRLGEA	.ANQKVSVVSSGSLALD	I	ALGAGGYPKGR	74
NTDB id 269 KZH43 RS08665 WP 001085462.1	.MAKKP....KKLEEISKKFGA	EREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALD	I	ALGSGGYPKGR	73
NTDB id 228 SPD RS09265 WP 001085462.1	.MAKKP....KKLEEISKKFGA	EREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALD	I	ALGSGGYPKGR	73
NTDB id 194 SPR RS08825 WP 001085462.1	.MAKKP....KKLEEISKKFGA	EREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALD	I	ALGSGGYPKGR	73
NTDB id 159 SP RS09750 WP 001085462.1	.MAKKP....KKLEEISKKFGA	EREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALD	I	ALGSGGYPKGR	73
NTDB id 499 SM12261 RS08315 WP 078228268.1	.MAKKP....KKLDEISKKFGA	EREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALD	I	ALGSGGYPKGR	73
NTDB id 527 SMSK321 RS11065 WP 001085454.1	.MAKKP....KKLDEISKKFGA	EREKALNDALKLIEKD	FGKGSIMRLGER	.AEQKVQVMSSGSLALD	I	ALGSGGYPKGR	73
NTDB id 453 Spy49 1753c ACI62003.1	.MAKKL....KKNEEITKKFG	DERRKALDDALKNIEKD	FGKGAVMRLGER	.AEQKVQVMSSGSLALD	I	ALGAGGYPKGR	73
NTDB id 408 SMU RS09530 WP 002262392.1	.MAKRI....KKTEEITKKFG	DERKKALDDALKNIEKD	FGKGAVMRLGER	.AEQKVQVMSSGSLALD	I	ALGAGGYPKGR	73
NTDB id 115 BSU 16940 NP 389576.2	MSDRQAALDMALKQ	IEKQFGKGSIMKLGEK	.TDTRISTVPSGSLALD	T	ALGIGGYPRGR	58	
NTDB id 625 LCA RS02525 WP 011374196.1	MAKDERQAALDAALKK	IEKNFGKGSIMRMGEK	.VDTQVSTVSSGSLALD	E	ALGVGGYPRGR	60	
NTDB id 1313 DR RS12030 WP 010888966.1	MSKDATKEISAPTDAK	ERSKATETAM	QIEKAFGKGSIMKLGAE	.SKLDVQVVSTGSLSLD	L	ALGVGGIPRGR	72
NTDB id 1245 Cj1673c YP 002345041.1	MDDNKRKSLDAALKSL	DKTFGKGTILRLGDK	.EVEQIDSIGTGSVGLD	L	ALGIGGVPKGR	59	
NTDB id 1214 NCTC11637 00359 SQJ03206.1	MAIDEDKQKATISLAIK	QIDKVF GKGA LVRLGDK	.QVEKIDISTGSLGLD	L	ALGIGGVPKGR	61	
NTDB id 1211 C694 RS00795 WP 000963128.1	MAIDEDKQKATISLAIK	QIDKVF GKGA LVRLGDK	.QVEKIDISTGSLGLD	L	ALGIGGVPKGR	61	
NTDB id 1334 RAOC RS04870 WP 004918236.1	MAKTE..TTSPND	DKKKALAAVLEKLDKT	YGKGTVMKLG	DGTVDHNI	IEVIPSGSLGLD	ALGVGGYPKGR	68
NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENKKSSKNNTVV..KQVEP	EQKKQALEAALAQIEKQ	FGKGSIMKLGQT	.EQLDIESVSTGSLGLD	L	ALGIGGLPMGR	76
NTDB id 1130 NGFG RS03960 WP 003688695.1	MSDDKSKALAAALAQIEK	SFGKGAIMKMDGSQ	QEEENLEVISTGSLGLD	L	ALGVGGLPRGR	60	
NTDB id 1124 NGFG RS03960 WP 003688695.1	MSDDKSKALAAALAQIEK	SFGKGAIMKMDGSQ	QEEENLEVISTGSLGLD	L	ALGVGGLPRGR	60	
NTDB id 1123 OK783 RS03845 WP 003688695.1	MSDDKSKALAAALAQIEK	SFGKGAIMKMDGSQ	QEEENLEVISTGSLGLD	L	ALGVGGLPRGR	60	
NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....KK..AASMSA	EKQKALAAALAQIEKQ	FGKGSIMKMGDA	.EVEPVQVVSTGSLGLD	V	ALGVGGLPRGR	68
NTDB id 1165 A1552VC RS01565 WP 000344154.1	MDENKQKALAAALGQIEKQ	FGKGSIMRLGDN	.RAMDVETISTGSLSLD	I	ALGAGGLPMGR	59	
NTDB id 1142 GTF74 RS11550 WP 000344154.1	MDENKQKALAAALGQIEKQ	FGKGSIMRLGDN	.RAMDVETISTGSLSLD	I	ALGAGGLPMGR	59	
NTDB id 1022 ACIAD RS06390 WP 004925639.1	MDDNKSKALQAALSQIEKQ	FGKNTVMRLGDN	.TVQAVEAVSTGSLTLD	I	ALGIGGLPKGR	59	
NTDB id 1074 ABD1 RS10195 WP 000344167.1	MDDNKSKALQAALSQIEKQ	FGKNTVMRLGDN	.TVQAVEAVSTGSLTLD	I	ALGIGGLPKGR	59	
NTDB id 40566 X969 RS04225 WP 013971339.1	MDDNKKRALAAALGQIERQ	FGKGAVMRMGDH	.ERQAIPAISTGSLGLD	I	ALGIGGLPKGR	59	
NTDB id 1205 PSJM300 05545 AFN77183.1	MDENKKRALAAALGQIEKQ	FGKGAVMRMGDH	.DRQAIP	SISTGSLGLD	I	ALGIGGLPKGR	59
consensus	*** ** *	*****	*****	*****	*	*****	*****

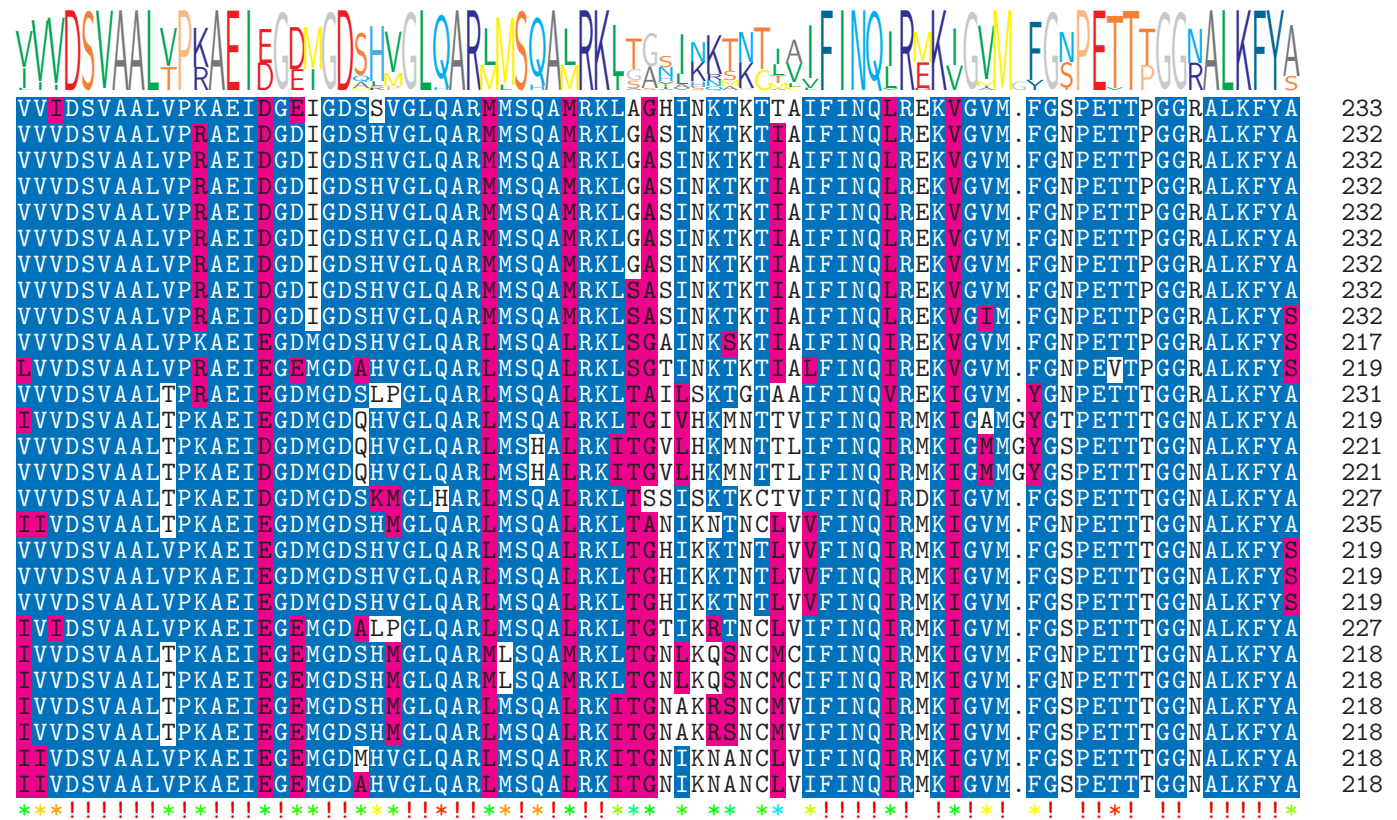
```

NTDB id 600 KW2 RS01765 WP 011675413.1
NTDB id 269 KZH43 RS08665 WP 001085462.1
NTDB id 228 SPD RS09265 WP 001085462.1
NTDB id 194 SPR RS08825 WP 001085462.1
NTDB id 159 SP RS09750 WP 001085462.1
NTDB id 499 SM12261 RS08315 WP 078228268.1
NTDB id 527 SMSK321 RS11065 WP 001085454.1
NTDB id 453 Spy49 1753c ACI62003.1
NTDB id 408 SMU RS09530 WP 002262392.1
NTDB id 115 BSU 16940 NP 389576.2
NTDB id 625 LCA RS02525 WP 011374196.1
NTDB id 1313 DR RS12030 WP 010888966.1
NTDB id 1245 Cj1673c YP 002345041.1
NTDB id 1214 NCTC11637 00359 SQJ03206.1
NTDB id 1211 C694 RS00795 WP 000963128.1
NTDB id 1334 RAOC RS04870 WP 004918236.1
NTDB id 1389 A4U84 RS07560 WP 021112712.1
NTDB id 1130 NGFG RS03960 WP 003688695.1
NTDB id 1124 NGFG RS03960 WP 003688695.1
NTDB id 1123 OK783 RS03845 WP 003688695.1
NTDB id 1427 RS RS02760 WP 01100051.1
NTDB id 1165 A1552VC RS01565 WP 000344154.1
NTDB id 1142 GTF74 RS11550 WP 000344154.1
NTDB id 1022 ACIAD RS06390 WP 004925639.1
NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 40566 X969 RS04225 WP 013971339.1
NTDB id 1205 PSJM300 05545 AFN717183.1
consensus

```



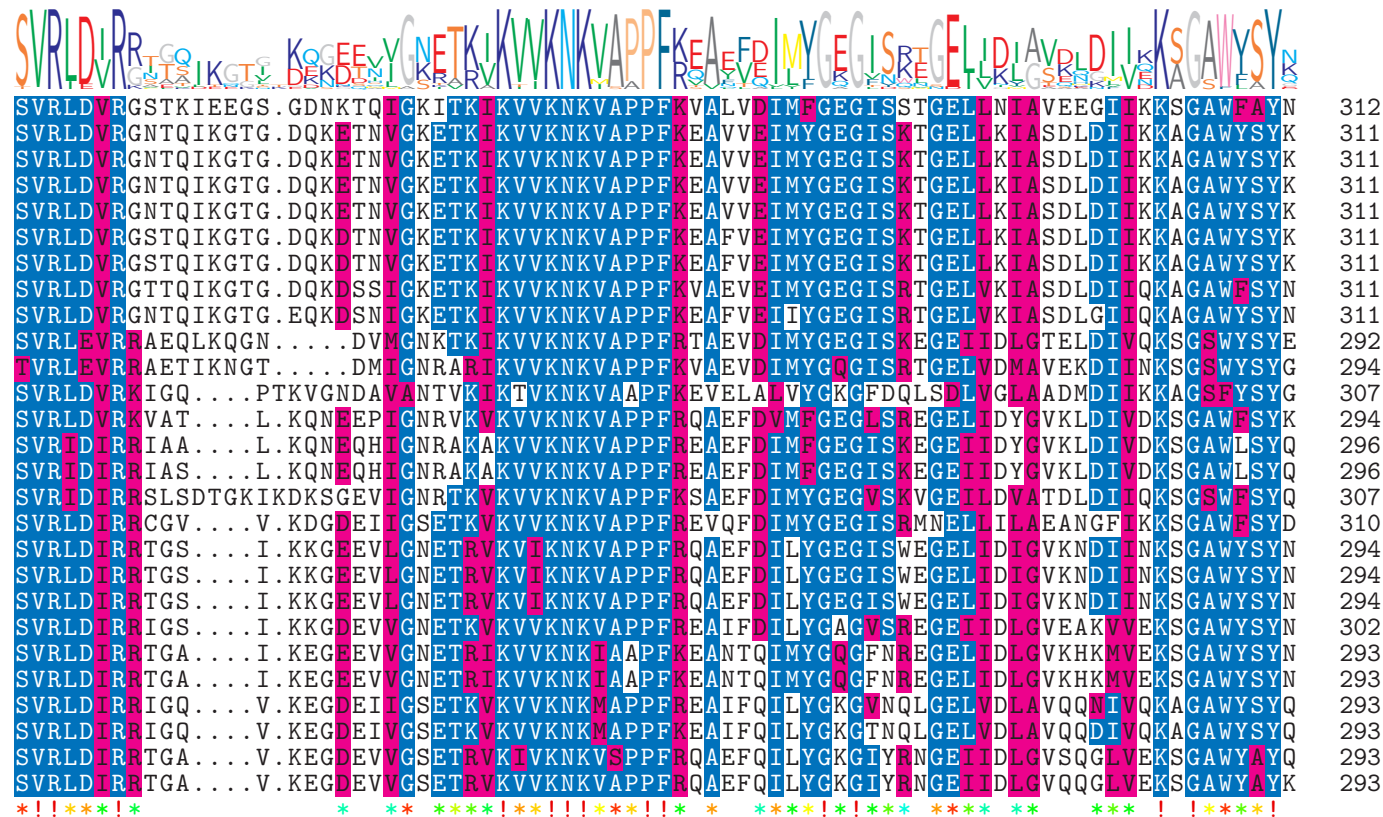
NTDB id 600 KW2 RS01765 WP 011675413.1
NTDB id 269 KZH43 RS08665 WP 001085462.1
NTDB id 228 SPD RS09265 WP 001085462.1
NTDB id 194 SPR RS08825 WP 001085462.1
NTDB id 159 SP RS09750 WP 001085462.1
NTDB id 499 SM12261 RS08315 WP 078228268.1
NTDB id 527 SMSK321 RS11065 WP 001085454.1
NTDB id 453 Spy49 1753c ACI62003.1
NTDB id 408 SMU RS09530 WP 002262392.1
NTDB id 115 BSU 16940 NP 389576.2
NTDB id 625 LCA RS02525 WP 011374196.1
NTDB id 1313 DR RS12030 WP 010888966.1
NTDB id 1245 Cj1673c YP 002345041.1
NTDB id 1214 NCTC11637 00359 SQJ03206.1
NTDB id 1211 C694 RS00795 WP 000963128.1
NTDB id 1334 RAOC RS04870 WP 004918236.1
NTDB id 1389 A4U84 RS07560 WP 021112712.1
NTDB id 1130 NGFG RS03960 WP 003688695.1
NTDB id 1124 NGFG RS03960 WP 003688695.1
NTDB id 1123 OK783 RS03845 WP 003688695.1
NTDB id 1427 RS RS02760 WP 011000510.1
NTDB id 1165 A1552VC RS01565 WP 000344154.1
NTDB id 1142 GTF74 RS11550 WP 000344154.1
NTDB id 1022 ACIAD RS06390 WP 004925639.1
NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 40566 X969 RS04225 WP 013971339.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus



```

NTDB id 600 KW2 RS01765 WP 011675413.1
NTDB id 269 KZH43 RS08665 WP 001085462.1
NTDB id 228 SPD RS09265 WP 001085462.1
NTDB id 194 SPR RS08825 WP 001085462.1
NTDB id 159 SP RS09750 WP 001085462.1
NTDB id 499 SM12261 RS08315 WP 078228268.1
NTDB id 527 SMSK321 RS11065 WP 001085454.1
NTDB id 453 Spy49 1753c ACI62003.1
NTDB id 408 SMU RS09530 WP 002262392.1
NTDB id 115 BSU 16940 NP 389576.2
NTDB id 625 LCA RS02525 WP 011374196.1
NTDB id 1313 DR RS12030 WP 010888966.1
NTDB id 1245 Cj1673c YP 002345041.1
NTDB id 1214 NCTC11637 00359 SQJ03206.1
NTDB id 1211 C694 RS00795 WP 000963128.1
NTDB id 1334 RAOC RS04870 WP 004918236.1
NTDB id 1389 A4U84 RS07560 WP 021112712.1
NTDB id 1130 NGFG RS03960 WP 003688695.1
NTDB id 1124 NGFG RS03960 WP 003688695.1
NTDB id 1123 OK783 RS03845 WP 003688695.1
NTDB id 1427 RS RS02760 WP 011000510.1
NTDB id 1165 A1552VC RS01565 WP 000344154.1
NTDB id 1142 GTF74 RS11550 WP 000344154.1
NTDB id 1022 ACIAD RS06390 WP 004925639.1
NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 40566 X969 RS04225 WP 013971339.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

```




```

NTDB id 600 KW2 RS01765 WP 011675413.1
NTDB id 269 KZH43 RS08665 WP 001085462.1
NTDB id 228 SPD RS09265 WP 001085462.1
NTDB id 194 SPR RS08825 WP 001085462.1
NTDB id 159 SP RS09750 WP 001085462.1
NTDB id 499 SM12261 RS08315 WP 078228268.1
NTDB id 527 SMSK321 RS11065 WP 001085454.1
NTDB id 453 Spy49 1753c ACI62003.1
NTDB id 408 SMU RS09530 WP 002262392.1
NTDB id 115 BSU 16940 NP 389576.2
NTDB id 625 LCA RS02525 WP 011374196.1
NTDB id 1313 DR RS12030 WP 010888966.1
NTDB id 1245 Cj1673c YP 002345041.1
NTDB id 1214 NCTC11637 00359 SQJ03206.1
NTDB id 1211 C694 RS00795 WP 000963128.1
NTDB id 1334 RAOC RS04870 WP 004918236.1
NTDB id 1389 A4U84 RS07560 WP 021112712.1
NTDB id 1130 NGFG RS03960 WP 003688695.1
NTDB id 1124 NGFG RS03960 WP 003688695.1
NTDB id 1123 OK783 RS03845 WP 003688695.1
NTDB id 1427 RS RS02760 WP 011000510.1
NTDB id 1165 A1552VC RS01565 WP 000344154.1
NTDB id 1142 GTF74 RS11550 WP 000344154.1
NTDB id 1022 ACIAD RS06390 WP 004925639.1
NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 40566 X969 RS04225 WP 013971339.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

```

Sequence	Score
DEKIGQGA ⁺ EA ⁺ KA ⁺ KN ⁺ YL ⁺ KE ⁺ HQ ⁺ DV ⁺ FE ⁺ ID ⁺ HK ⁺ VRAAH ⁺ G ⁺ LDDSEVAETEEETTAS ⁺ KT ⁺ KA ⁺ KA ⁺ KK ⁺ EE ⁺ KAVETEEIELELED.....	387
DEKIGQGSEN ⁺ AK ⁺ KYL ⁺ AE ⁺ HP ⁺ EI ⁺ FE ⁺ ID ⁺ KQ ⁺ VR ⁺ SK ⁺ FL ⁺ IDGEEVSEQD ⁺ TEN ⁺ KK ⁺ DE...PK ⁺ KE ⁺ EA...VNEEVPLDLGDELEIE	385
DEKIGQGS ⁺ EN ⁺ AK ⁺ KYL ⁺ AE ⁺ HP ⁺ EI ⁺ FE ⁺ ID ⁺ KQ ⁺ VR ⁺ SK ⁺ FL ⁺ IDGEEVSEQD ⁺ TEN ⁺ KK ⁺ DE...PK ⁺ KE ⁺ EA...VNEEVPLDLGDELEIE	385
DEKIGQGS ⁺ EN ⁺ AK ⁺ KYL ⁺ AE ⁺ HP ⁺ EI ⁺ FE ⁺ ID ⁺ KQ ⁺ VR ⁺ SK ⁺ FL ⁺ IDGEEVSEQD ⁺ TEN ⁺ KK ⁺ DE...PK ⁺ KE ⁺ EA...VNEEVPLDLGDELEIE	385
DEKIGQGS ⁺ EN ⁺ AK ⁺ KYL ⁺ AD ⁺ NPEI ⁺ FE ⁺ ID ⁺ HQ ⁺ VR ⁺ VQ ⁺ FL ⁺ IDGEEIAES...KK ⁺ DE...AAQV ⁺ DS...VNEEVTLDLGDLEIE	381
DEKIGQGS ⁺ EN ⁺ AK ⁺ KYL ⁺ AD ⁺ NPEI ⁺ FE ⁺ ID ⁺ HQ ⁺ VR ⁺ VQ ⁺ FL ⁺ IDGEEAAES...KK ⁺ DE...VAQV ⁺ DS...VNEEVTLDLGDLEIE	381
GEKIGQGS ⁺ EN ⁺ AK ⁺ RYL ⁺ AD ⁺ HP ⁺ QL ⁺ FE ⁺ ID ⁺ DR ⁺ KVR ⁺ VK ⁺ FL ⁺ LEESEEEESAMAVASE.....E.....TDDLALDLNGIEIE	377
GEKIGQGS ⁺ EN ⁺ AK ⁺ KFL ⁺ AD ⁺ NPEI ⁺ FD ⁺ ID ⁺ HK ⁺ VR ⁺ VQ ⁺ GL ⁺ IEEDNTEEKQSSKEK.....ET ⁺ DE ⁺ K...ADKNLVLELDDTIELE	382
EERL ⁺ GQGR ⁺ EN ⁺ AK ⁺ QFL ⁺ KEN ⁺ KDIM ⁺ LM ⁺ IQEQ ⁺ IREHYGLD ⁺ NGGVVQQQAETQEEL.EFE.E.....	348
SERIGQGR ⁺ EN ⁺ AK ⁺ NYL ⁺ AD ⁺ HED ⁺ VE ⁺ DEV ⁺ RLK ⁺ VRAAY ⁺ GISD ⁺ VPEEDLPTTEDE.QI.NILPDDST...EE	355
DERIGQG ⁺ KE ⁺ KTI ⁺ AI ⁺ YIA ⁺ ERP ⁺ EME ⁺ QE ⁺ IT ⁺ DR ⁺ VMAA ⁺ IRAGNAGEAPALAPAA.....PE ⁺ AA...EA	363
DKKL ⁺ GQGR ⁺ EN ⁺ SA ⁺ FL ⁺ KEN ⁺ PEI ⁺ ADE ⁺ IT ⁺ KA ⁺ IQNS ⁺ MG ⁺ IEGMISGSEDDEG.....EE	343
DKKL ⁺ GQGR ⁺ EN ⁺ AK ⁺ ALL ⁺ KED ⁺ KAL ⁺ ADE ⁺ IT ⁺ LK ⁺ IKES ⁺ IG ⁺ SNEEIMPLPDEPL.....EEM...E	347
DKKL ⁺ GQGR ⁺ EN ⁺ AK ⁺ ALL ⁺ KED ⁺ KAL ⁺ ADE ⁺ IT ⁺ LK ⁺ IKES ⁺ IG ⁺ SNEEIMPLPDEPL.....EEM...E	347
DTKL ⁺ GQGR ⁺ DT ⁺ VKEL ⁺ IK ⁺ DN ⁺ PEL ⁺ AEE ⁺ LEAK ⁺ IREKLQAK.....	343
GT ⁺ KIGQG ⁺ KNN ⁺ AV ⁺ KVL ⁺ KEN ⁺ PEI ⁺ AD ⁺ KL ⁺ EQIR ⁺ NTL ⁺ ASNPNSVLKATASDSNDES...DVE ⁺ EF...SEEL	372
GAKIGQGD ⁺ KN ⁺ VR ⁺ VWL ⁺ KEN ⁺ PEI ⁺ SD ⁺ EID ⁺ AK ⁺ IRAL ⁺ NGVEMHITEGTQDET...DGE...RPEE	348
GAKIGQGD ⁺ KN ⁺ VR ⁺ VWL ⁺ KEN ⁺ PEI ⁺ SD ⁺ EID ⁺ AK ⁺ IRAL ⁺ NGVEMHITEGTQDET...DGE...RPEE	348
GAKIGQGD ⁺ KN ⁺ VR ⁺ VWL ⁺ KEN ⁺ PEI ⁺ SD ⁺ EID ⁺ AK ⁺ IRAL ⁺ NGVEMHITEGTQDET...DGE...RPEE	348
GERIGQGR ⁺ DN ⁺ CRE ⁺ FL ⁺ RENA ⁺ E ⁺ LARE ⁺ IE ⁺ EN ⁺ KV ⁺ REHL ⁺ GV ⁺ TP.MGAVTLAEE...VEE...D	352
GD ⁺ KIGQG ⁺ KAN ⁺ ACK ⁺ YL ⁺ KEN ⁺ PEI ⁺ AK ⁺ TL ⁺ DKKL ⁺ REMLLN ⁺ PEN ⁺ MLIAETSS...AAD...DVE.F...GAVPEEF	354
GDKIGQGG ⁺ KAN ⁺ ACK ⁺ YL ⁺ KEN ⁺ PEI ⁺ AK ⁺ TL ⁺ DKKL ⁺ REMLLN ⁺ PEN ⁺ MLIAETSS...AAD...DVE.F...GAVPEEF	354
GN ⁺ KIGQG ⁺ KNN ⁺ VIR ⁺ YLE ⁺ EN ⁺ PQ ⁺ ISTE ⁺ IE ⁺ AV ⁺ IREQLLT ⁺ TKASDQ ⁺ TAAHDET...EE...EP ⁺ DL...ES	349
GN ⁺ KIGQG ⁺ KNN ⁺ VIR ⁺ YFE ⁺ ENN ⁺ QIA ⁺ EIE ⁺ ERN ⁺ IREQLLT ⁺ TGTNGAVQI.ED...EE...EP ⁺ DL...LES	349
GN ⁺ KIGQGG ⁺ KANA ⁺ AK ⁺ YL ⁺ EN ⁺ PA ⁺ IGA ⁺ EIE ⁺ KQ ⁺ IREKLL ⁺ TAGAVAAAGKAAA...AEA...DADD...ADADAGY	355
GS ⁺ KIGQG ⁺ KANA ⁺ AK ⁺ YLE ⁺ EN ⁺ PEI ⁺ GQ ⁺ EIE ⁺ QQ ⁺ IRAKLL ⁺ V ⁺ TSANTKATPVA...EDL...ADADI	347

logo

		IEE	
NTDB id 600 KW2 RS01765 WP 011675413.1	...	387	
NTDB id 269 KZH43 RS08665 WP 001085462.1	IEE	388	
NTDB id 228 SPD RS09265 WP 001085462.1	IEE	388	
NTDB id 194 SPR RS08825 WP 001085462.1	IEE	388	
NTDB id 159 SP RS09750 WP 001085462.1	IEE	388	
NTDB id 499 SM12261 RS08315 WP 078228268.1	IEE	384	
NTDB id 527 SMSK321 RS11065 WP 001085454.1	IEE	384	
NTDB id 453 Spy49 1753c ACI62003.1	D..	378	
NTDB id 408 SMU RS09530 WP 002262392.1	D..	383	
NTDB id 115 BSU 16940 NP 389576.2	...	348	
NTDB id 625 LCA RS02525 WP 011374196.1	...	355	
NTDB id 1313 DR RS12030 WP 010888966.1	...	363	
NTDB id 1245 Cj1673c YP 002345041.1	...	343	
NTDB id 1214 NCTC11637 00359 SQJ03206.1	...	347	
NTDB id 1211 C694 RS00795 WP 000963128.1	...	347	
NTDB id 1334 RAOC RS04870 WP 004918236.1	...	343	
NTDB id 1389 A4U84 RS07560 WP 021112712.1	...	372	
NTDB id 1130 NGFG RS03960 WP 003688695.1	...	348	
NTDB id 1124 NGFG RS03960 WP 003688695.1	...	348	
NTDB id 1123 OK783 RS03845 WP 003688695.1	...	348	
NTDB id 1427 RS RS02760 WP 011000510.1	...	352	
NTDB id 1165 A1552VC RS01565 WP 000344154.1	...	354	
NTDB id 1142 GTF74 RS11550 WP 000344154.1	...	354	
NTDB id 1022 ACIAD RS06390 WP 004925639.1	...	349	
NTDB id 1074 ABD1 RS10195 WP 000344167.1	...	349	
NTDB id 40566 X969 RS04225 WP 013971339.1	...	355	
NTDB id 1205 PSJM300 05545 AFN77183.1	...	347	
consensus			

-  non conserved
-  similar
-  ≥ 50% conserved