

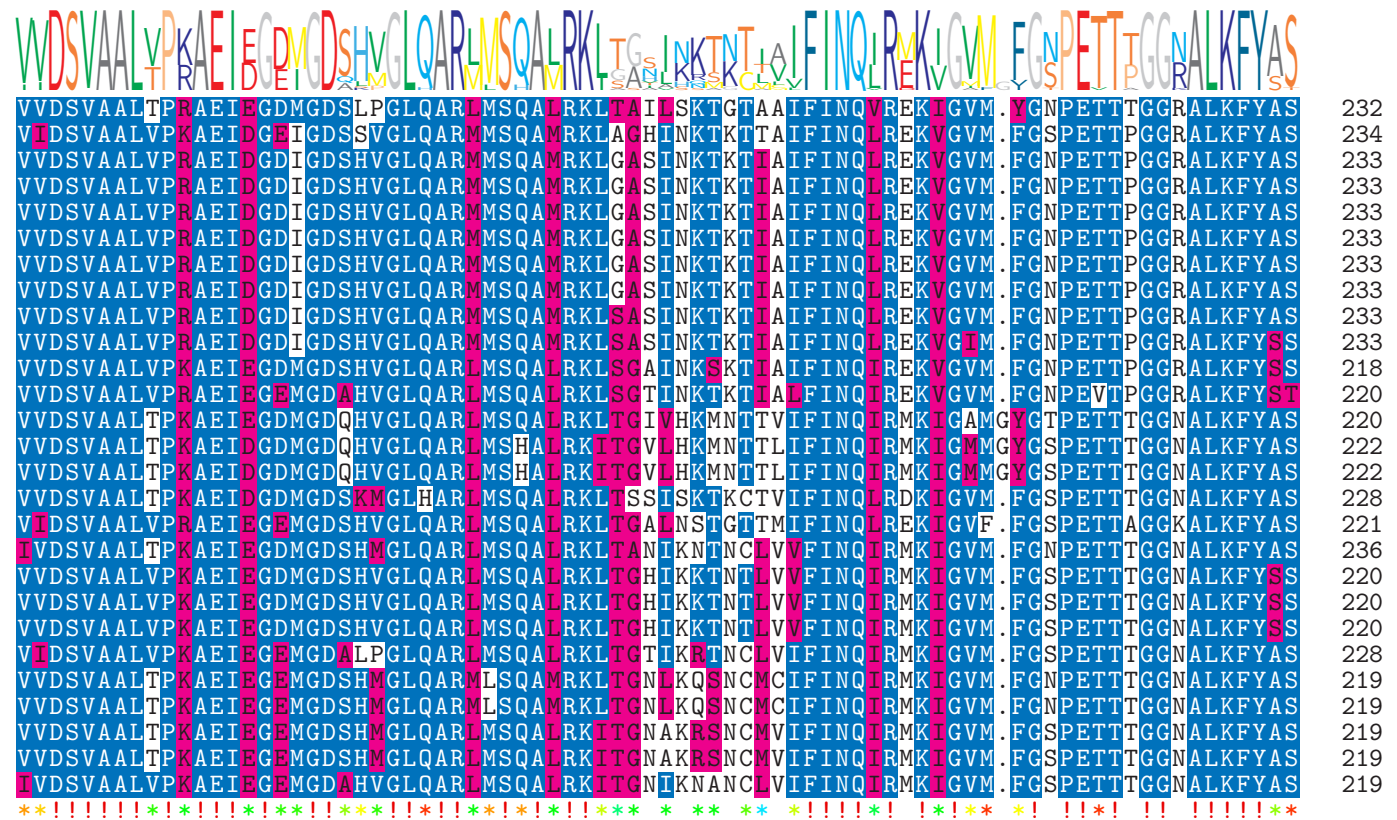
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NTDB id 1313 DR RS12030 WP 010888966.1	.MSK....DATKEISAPTDAK	ERSKATETAMSQIEKAFGKGSIMKLGAE	.SKLDVQVVSTGSLSLDLALGVGGIP	RGRI	73
NTDB id 600 KW2 RS01765 WP 011675413.1	MATKK...KTNFDDI.TKKYGA	ERDKALADALALIEKDFGKGSIMRLGEA	.ANQKVSVVSSGSLALDIALGAGGY	PKGRI	75
NTDB id 269 KZH43 RS08665 WP 001085462.1	.MAKK...PKKLEEI.SKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDIALGSGGY	PKGRI	74
NTDB id 228 SPD RS09265 WP 001085462.1	.MAKK...PKKLEEI.SKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDIALGSGGY	PKGRI	74
NTDB id 194 SPR RS08825 WP 001085462.1	.MAKK...PKKLEEI.SKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDIALGSGGY	PKGRI	74
NTDB id 159 SP RS09750 WP 001085462.1	.MAKK...PKKLEEI.SKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDIALGSGGY	PKGRI	74
NTDB id 499 SM12261 RS08315 WP 078228268.1	.MAKK...PKKLDEI.SKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDIALGSGGY	PKGRI	74
NTDB id 527 SMSK321 RS11065 WP 001085454.1	.MAKK...PKKLDEI.SKKFGA	EREKALNDALKLIEKDFGKGSIMRLGER	.AEQKVQVMSSGSLALDIALGSGGY	PKGRI	74
NTDB id 453 Spy49 1753c ACI62003.1	.MAKK...LKKNEEI.TKKFGD	ERRKALDDALKNIEKDFGKGAVMRLGER	.AEQKVQVMSSGSLALDIALGAGGY	PKGRI	74
NTDB id 408 SMU RS09530 WP 002262392.1	.MAKR...IKKTEEI.TKKFGD	ERKKALDDALKNIEKDFGKGAVMRLGER	.AEQKVQVMSSGSLALDIALGAGGY	PKGRI	74
NTDB id 115 BSU 16940 NP 389576.2	MSDRQAALDMALKQIEKQFGKGSIMKLGEK	.TDTRISTVPSGSLALDTALGIGGY	PRGRI		59
NTDB id 625 LCA RS02525 WP 011374196.1	MAKD	ERQAALDAALKKIEKNFGKGSIMRMGEK	.VDTQVSTVSSGSLALDEALGVGGY	PRGRI	61
NTDB id 1245 Cj1673c YP 002345041.1	MDDNKRKSLDAALKSLDKTFGKGTTILRLGDK	.EVEQIDSIGTGSVGLDLALGIGGV	PKGRI		60
NTDB id 1211 C694 RS00795 WP 000963128.1	MAIDEDKQKALSLAIKQIDKVF	GKGA LVRLGDK.QVEKIDSISTGSLGLDLALGIGGV	PKGRI		62
NTDB id 1214 NCTC11637 00359 SQJ03206.1	MAIDEDKQKALSLAIKQIDKVF	GKGA LVRLGDK.QVEKIDAISTGSLGLDLALGIGGV	PKGRI		62
NTDB id 1334 RAOC RS04870 WP 004918236.1	MAKTE.TTSPND	DKKKALAAVLEKLDKTYGKGTVMKLGD	GTVDHNIIEVIPSGSLGLDLALGVGGY	PKGRI	69
NTDB id 364746 FFA46 RS02280 WP 138275190.1	MPSPSDREKALDTALAQIDRQFGKGSVMRLGSD	.DRAPVEVIPTGSTIALDVALGTGGL	PKGRI		62
NTDB id 1389 A4U84 RS07560 WP 021112712.1	.MAENKKSSKNNTVV.KQVEPEQ	KKQALEAALAQIEKQFGKGSIMKL	GQT.EQLDIESVSTGSLGLDLALGIGGL	PMGRI	77
NTDB id 1130 NGFG RS03960 WP 003688695.1	MSDDKSKALAAALAQIEKSFGKGAIMKMDGSQQEEN	LEVISTGSLGLDLALGVGGL	PRGRI		61
NTDB id 1124 NGFG RS03960 WP 003688695.1	MSDDKSKALAAALAQIEKSFGKGAIMKMDGSQQEEN	LEVISTGSLGLDLALGVGGL	PRGRI		61
NTDB id 1123 OK783 RS03845 WP 003688695.1	MSDDKSKALAAALAQIEKSFGKGAIMKMDGSQQEEN	LEVISTGSLGLDLALGVGGL	PRGRI		61
NTDB id 1427 RS RS02760 WP 011000510.1	.MEDG.....KK.AASMSAEKKQKALAAALAQIEKQFGKGSIMKMGDA	.EVEPVQVVSTGSLGLDVALGVGGL	PRGRV		69
NTDB id 1165 A1552VC RS01565 WP 000344154.1	MDENKQKALAAALGQIEKQFGKGSIMRLGDN	.RAMDVETISTGSLSLDIALGAGGL	PMGRI		60
NTDB id 1142 GTF74 RS11550 WP 000344154.1	MDENKQKALAAALGQIEKQFGKGSIMRLGDN	.RAMDVETISTGSLSLDIALGAGGL	PMGRI		60
NTDB id 1074 ABD1 RS10195 WP 000344167.1	MDENKSKALQAALSQIEKQFGKNTVMRLGDN	.TVQAVEAVSTGSLTLDIALGIGGL	PKGRI		60
NTDB id 1022 ACIAD RS06390 WP 004925639.1	MDDNKS	KALQAALSQIEKQFGKNTVMRLGDN	.TVQAVEAVSTGSLTLDIALGIGGL	PKGRI	60
NTDB id 1205 PSJM300 05545 AFN77183.1	MDENKKRALAAALGQIEKQFGKGAVMRMGDH	.DRQAIP	SISTGSLGLDIALGIGGL	PKGRI	60
consensus	* * * * *	* * * * *	* * * * *	* * * * *	


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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 1245 Cj1673c YP 002345041.1
NTDB id 1211 C694 RS00795 WP 000963128.1
NTDB id 1214 NCTC11637 00359 SQJ03206.1
NTDB id 1334 RAOC RS04870 WP 004918236.1
NTDB id 364746 FFA46 RS02280 WP 138275190.1
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 1074 ABD1 RS10195 WP 000344167.1
NTDB id 1022 ACIAD RS06390 WP 004925639.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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NTDB id 1313 DR RS12030 WP 010888966.1	ERIGQGKEKTIA Y IAERP ME QEIRDRVMAAIRAGNAGEAPALAPAPAA.....PEAAEA.....	363
NTDB id 600 KW2 RS01765 WP 011675413.1	EKIGQGA E KAK N YLKEHQDV F DEIDHKVRAAHGLDDSEVAETEEETTASKT.KAKAKKEEKAVETEEIELELED.....	387
NTDB id 269 KZH43 RS08665 WP 001085462.1	EKIGQGSENAKKYLAEHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE.....PKKEEA..VNEEVPLDLGDELEIE	385
NTDB id 228 SPD RS09265 WP 001085462.1	EKIGQGS E NAKKYLAEHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE.....PKKEEA..VNEEVPLDLGDELEIE	385
NTDB id 194 SPR RS08825 WP 001085462.1	EKIGQGS E NAKKYLAEHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE.....PKKEEA..VNEEVPLDLGDELEIE	385
NTDB id 159 SP RS09750 WP 001085462.1	EKIGQGS E NAKKYLAEHPEIFDEIDKQVRSKFGLIDGEEVSEQDTENKKDE.....PKKEEA..VNEEVPLDLGDELEIE	385
NTDB id 499 SM12261 RS08315 WP 078228268.1	EKIGQGS E NAKKYLADNPEIFDEIDHQVRVQFGLIDGEEIAES....KKDE.....AAQVDS..VNEEVTLDLGDELEIE	381
NTDB id 527 SMSK321 RS11065 WP 001085454.1	EKIGQGS E NAKKYLADNPEVFAEIDHQVRVQFGLIDGEEAAES....KKDE.....VAQVDS..VNEEVTLDLGDELEIE	381
NTDB id 453 Spy49 1753c ACI62003.1	EKIGQGS E NAKRYLADHPQLFDEIDRKVRVKFGLLEEESEESAMAVASE.....E.....TDDLALDLNGIEIE	377
NTDB id 408 SMU RS09530 WP 002262392.1	EKIGQGS E NAKKFLADNPEIFDDIDHKVRVQYGLIEEDNTEEKQSSKEK.....ETDEK..ADKNLVLELDDTIELE	382
NTDB id 115 BSU 16940 NP 389576.2	ERLGQGR E NAKQFLKENKDIMLMIQEQIREHYGLDNNGVVQQAEETQEEL..EFE.E.....	348
NTDB id 625 LCA RS02525 WP 011374196.1	ERIGQGR E NAKNYLADHEDVEDEVRLKVRAAYGTS D VPEEDLPTTEDE.QI..NILPDDST.....EE.....	355
NTDB id 1245 Cj1673c YP 002345041.1	KKLGGGREN S KAFLKENPEIADEITKAIQNSMGIEGMISGSE..DDEGEE.....	343
NTDB id 1211 C694 RS00795 WP 000963128.1	KKLGQGR E NAKALLKEDKALADEITLKIKESIGSNEEIMPLP..DEPLEEME.....	347
NTDB id 1214 NCTC11637 00359 SQJ03206.1	KKLGQGR E NAKALLKEDKALADEITLKIKESIGSNEEIMPLP..DEPLEEME.....	347
NTDB id 1334 RAOC RS04870 WP 004918236.1	TKLGQGRD T VKELIKDNPELAELEAKIREKLQAK.....	343
NTDB id 364746 FFA46 RS02280 WP 138275190.1	EQLGQGG E NA R KYLIENGKVADEIEKKIKTKLGTIGVPRAVADLPAESAPDASTKAKAAS.....AG.....	357
NTDB id 1389 A4U84 RS07560 WP 021112712.1	TKIGQGG K NNAVKWLKENPEIADKLEQEIRNTLASNPNSVLKATASDSNDES DVVEEFSEE.....EL.....	372
NTDB id 1130 NGFG RS03960 WP 003688695.1	AKIGQGG K DNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQDETD.....GERPEE.....	348
NTDB id 1124 NGFG RS03960 WP 003688695.1	AKIGQGG K DNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQDETD.....GERPEE.....	348
NTDB id 1123 OK783 RS03845 WP 003688695.1	AKIGQGG K DNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQDETD.....GERPEE.....	348
NTDB id 1427 RS RS02760 WP 011000510.1	ERIGQGRD N CREFLRENAELAREIENKVREHLGVTPMGAV.TLAEVEVED.....	352
NTDB id 1165 A1552VC RS01565 WP 000344154.1	DKIGQGG K ANACKY L KENPEIAKTLDKKLREMLLNPNENMQLIAETSSAADDVEFGAVPEE.....F.....	354
NTDB id 1142 GTF74 RS11550 WP 000344154.1	DKIGQGG K ANACKY L KENPEIAKTLDKKLREMLLNPNENMQLIAETSSAADDVEFGAVPEE.....F.....	354
NTDB id 1074 ABD1 RS10195 WP 000344167.1	NKIGQGG K NNVIRYFEENNQIAEEIERNIREQLLTTGTNGAVQ.I.EDEEEPD..LLES.....	349
NTDB id 1022 ACIAD RS06390 WP 004925639.1	NKIGQGG K NNVIRYLEENPQISTEIEAVIREQLLTKASDQTAA.HDETEEEP..LLES.....	349
NTDB id 1205 PSJM300 05545 AFN77183.1	SKIGQGG K ANA A KYLEENPEIGQEIEQQIRAKLLVTSANTKATPVAEDLADAD..I.....	347
consensus	***!!****** ** ***** ***** ** *	

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NTDB id 1313 DR RS12030 WP 010888966.1	...		363
NTDB id 600 KW2 RS01765 WP 011675413.1	...		387
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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
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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
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NTDB id 1022 ACIAD RS06390 WP 004925639.1	...		349
NTDB id 1205 PSJM300 05545 AFN77183.1	...		347
consensus			

- X non conserved
- X similar
- X ≥ 50% conserved