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NTDB id 600 KW2 RS01765 WP 011675413.1	VVIVDSVAALVPKAEIDGDI	IGDSSVGLQARMSQAMRKL	AGHINKTKT	TAIFINQLREKVGVM	FGSPETTPGGRALKFYA	233
NTDB id 269 KZH43 RS08665 WP 001085462.1	VVVDSVAALVPRAEIDGDI	IGDSHVGLQARMSQAMRKL	GASINKTKT	TAIFINQLREKVGVM	FCNPETTPGGRALKFYA	232
NTDB id 228 SPD RS09265 WP 001085462.1	VVVDSVAALVPRAEIDGDI	IGDSHVGLQARMSQAMRKL	GASINKTKT	TAIFINQLREKVGVM	FCNPETTPGGRALKFYA	232
NTDB id 194 SPR RS08825 WP 001085462.1	VVVDSVAALVPRAEIDGDI	IGDSHVGLQARMSQAMRKL	GASINKTKT	TAIFINQLREKVGVM	FCNPETTPGGRALKFYA	232
NTDB id 159 SP RS09750 WP 001085462.1	VVVDSVAALVPRAEIDGDI	IGDSHVGLQARMSQAMRKL	GASINKTKT	TAIFINQLREKVGVM	FCNPETTPGGRALKFYA	232
NTDB id 527 SMSK321 RS11065 WP 001085454.1	VVVDSVAALVPRAEIDGDI	IGDSHVGLQARMSQAMRKL	GASINKTKT	TAIFINQLREKVGVM	FCNPETTPGGRALKFYA	232
NTDB id 499 SM12261 RS08315 WP 078228268.1	VVVDSVAALVPRAEIDGDI	IGDSHVGLQARMSQAMRKL	GASINKTKT	TAIFINQLREKVGVM	FCNPETTPGGRALKFYA	232
NTDB id 408 SMU RS09530 WP 002262392.1	VVVDSVAALVPRAEIDGDI	IGDSHVGLQARMSQAMRKL	SASINKTKT	TAIFINQLREKVGIM	FCNPETTPGGRALKFYS	232
NTDB id 453 Spy49 1753c ACI62003.1	VVVDSVAALVPRAEIDGDI	IGDSHVGLQARMSQAMRKL	SASINKTKT	TAIFINQLREKVGVM	FCNPETTPGGRALKFYA	232
NTDB id 115 BSU 16940 NP 389576.2	VVVDSVAALVPKAEIEGDMGDS	HVGLQARMSQALRKL	SGAINKSKT	TAIFINQLREKVGVM	FCNPETTPGGRALKFYS	217
NTDB id 625 LCA RS02525 WP 011374196.1	LVVDSVAALVPRAEIEGEMGDA	HVGLQARMSQALRKL	SGTINKTKT	TAIFINQLREKVGVM	FCNPETTPGGRALKFYS	219
NTDB id 1313 DR RS12030 WP 010888966.1	VVVDSVAALTPRAEIEGDMGDS	LPGLQARMSQALRKL	TAISKTGTA	IFINQLREKVGVM	YCNPETTTGGRALKFYA	231
NTDB id 1245 Cj1673c YP 002345041.1	IVVDSVAALTPKAEIEGDMGDQ	HVGLQARMSQALRKL	TGIVHKMNT	TVIFINQLRMKIGAMGY	CTPETTTGGNALKFYA	219
NTDB id 1211 C694 RS00795 WP 000963128.1	VVVDSVAALTPKAEIDGDMGDQ	HVGLQARMSHALRKL	ITGVLHKMNT	TLIFINQLRMKIGMNGY	GSPETTTGGNALKFYA	221
NTDB id 1214 NCTC11637 00359 SQJ03206.1	VVVDSVAALTPKAEIDGDMGDQ	HVGLQARMSHALRKL	ITGVLHKMNT	TLIFINQLRMKIGMNGY	GSPETTTGGNALKFYA	221
NTDB id 1334 RAOC RS04870 WP 004918236.1	VVVDSVAALTPKAEIDGDMGDS	KMGLHARMSQALRKL	TSSISKTKCT	VIFINQLRDKIGVM	FGSPETTTGGNALKFYA	227
NTDB id 1389 A4U84 RS07560 WP 021112712.1	ITVDSVAALTPKAEIEGDMGDS	HVGLQARMSQALRKL	TANIKNTNCT	VVIFINQLRMKIGVM	FCNPETTTGGNALKFYA	235
NTDB id 1130 NGFG RS03960 WP 003688695.1	VVVDSVAALVPKAEIEGDMGDS	HVGLQARMSQALRKL	TGHIKKTNTL	VVIFINQLRMKIGVM	FGSPETTTGGNALKFYS	219
NTDB id 1124 NGFG RS03960 WP 003688695.1	VVVDSVAALVPKAEIEGDMGDS	HVGLQARMSQALRKL	TGHIKKTNTL	VVIFINQLRMKIGVM	FGSPETTTGGNALKFYS	219
NTDB id 1123 OK783 RS03845 WP 003688695.1	VVVDSVAALVPKAEIEGDMGDS	HVGLQARMSQALRKL	TGHIKKTNTL	VVIFINQLRMKIGVM	FGSPETTTGGNALKFYS	219
NTDB id 1427 RS RS02760 WP 011000510.1	IVVDSVAALVPKAEIEGEMGDA	LPGLQARMSQALRKL	TGTIKR	TNCLVIFINQLRMKIGVM	FGSPETTTGGNALKFYA	227
NTDB id 188143 VSVS05 RS01050 WP 005593985.1	LVVDSVAALTPKAEIEGEMGDS	HVGLQARMSQAMRKL	TGNLQKSNCMC	IFINQLRMKIGVM	FCNPETTTGGNALKFYA	218
NTDB id 1165 A1552VC RS01565 WP 000344154.1	IVVDSVAALTPKAEIEGEMGDS	HVGLQARMSQAMRKL	TGNLQKSNCMC	IFINQLRMKIGVM	FCNPETTTGGNALKFYA	218
NTDB id 1142 GTF74 RS11550 WP 000344154.1	IVVDSVAALTPKAEIEGEMGDS	HVGLQARMSQAMRKL	TGNLQKSNCMC	IFINQLRMKIGVM	FCNPETTTGGNALKFYA	218
NTDB id 1205 PSJM300 05545 AFN77183.1	ITVDSVAALVPKAEIEGEMGDA	HVGLQARMSQALRKL	TGNLKNANCL	VIFINQLRMKIGVM	FGSPETTTGGNALKFYA	218
NTDB id 1074 ABD1 RS10195 WP 000344167.1	IVVDSVAALTPKAEIEGEMGDS	HVGLQARMSQALRKL	TGNAKRSNCMV	VIFINQLRMKIGVM	FGSPETTTGGNALKFYA	218
NTDB id 1022 ACIAD RS06390 WP 004925639.1	LVVDSVAALTPKAEIEGEMGDS	HVGLQARMSQALRKL	TGNAKRSNCMV	VIFINQLRMKIGVM	FGSPETTTGGNALKFYA	218
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	...	349
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consensus

 non conserved
 similar
 ≥ 50% conserved