```
logo
NTDB id 293473 DK843 RS15915 WP 114073772.1  ..............................................................................  0
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG  80
consensus                                    ******************************************************************************

logo
NTDB id 293473 DK843 RS15915 WP 114073772.1  ....................................MSWTERLKAGLAKTRDKLGKSLAGVFGGGKIDENLYEELETVLLT  45
NTDB id 1118 NGFG RS11455 WP 003696286.1     EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                    ************************************ !  !!! !!!!*!!!**!!!!!!!!!! ! ! !!!!!!!!!*!

logo
NTDB id 293473 DK843 RS15915 WP 114073772.1  ADMGMDATVHLLQDVRERVSLKGLKDASELKGALKDSLQDLIGPLEVPLNV.EGKKPFVIMMTGVNGAGKTTSIGKLAKY  124
NTDB id 1118 NGFG RS11455 WP 003696286.1     GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                    *!!!!*!! *!* !!! !!!!!!!!!* !!*!!!!* ! !!! !!! !! **! ! !!!!!* !*!!!!!!!!!!!!!!!!

logo
NTDB id 293473 DK843 RS15915 WP 114073772.1  YQSQGKSVLLAAGDTFRAAAREQLIAWGERNSVTVIAQQGGDSAAVCYDAIQAAVARGIDIVLADTAGRLPTQLHLMEEI  204
NTDB id 1118 NGFG RS11455 WP 003696286.1     FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                    *! !!!!!!!!!!!!!!!!!!!!! !!! !! !!!! !  !!!!!!!*!!*!!! !!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 293473 DK843 RS15915 WP 114073772.1  KKVKRVIQKALPDAPHEVVLVLDANIGQNTVNQVKAFDDALGLTGLILTKLDGTAKGGVIAAIAKQRPIPLRFVGVGESI  284
NTDB id 1118 NGFG RS11455 WP 003696286.1     KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                    !!!!!!*!!!*! !!!!***!!!!!!!!! !!!!!!!!!!!!!!!!!*!!!!!!!!!!**!!*!  !!*!*!**!!!! !

logo
NTDB id 293473 DK843 RS15915 WP 114073772.1  DDLRPFDSRDYIDALFE  301
NTDB id 1118 NGFG RS11455 WP 003696286.1     DDLRPFDARAFVDALLD  417
consensus                                    !!!!!!! ! **!!! *
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X non conserved

X similar

X ≥ 50% conserved