```
logo
NTDB id 261277 CXB49 RS18965 WP 101709811.1  MFSFFKKKTPPAAT....PQIETPAPAVPEPI...........PAAPEAIQPAAEPRH.......EDIAATVSEAPAPSA   58
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                    !!!!!**! **  ******!*  *!* !* **************  *! * * !!* ********! *  !! ! ! * *

logo
NTDB id 261277 CXB49 RS18965 WP 101709811.1  PPAA...........APAALVAAPAAPAEPASLKKMSWTDRLKAGLSKTRDRLGKQLAGLFGGGKIDEELYEELETVLLT  127
NTDB id 1118 NGFG RS11455 WP 003696286.1     EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                    **! *********** **  !* *  *  *   *!* !  !!! !! !*!!***! !!!*!!!! ! !*!!!!!!!!!*!

logo
NTDB id 261277 CXB49 RS18965 WP 101709811.1  ADIGIDATAHLLQDVRERVSLKGLKDSSELKGALKDSLQELIGPLQSPLNV.EGKKPFVIMMTGVNGAGKTTSIGKLAKY  206
NTDB id 1118 NGFG RS11455 WP 003696286.1     GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                    *!*!**!! *!* !!! !!!!!!!!!  !!*!!!!* ! *!! !!  !! **! ! !!!!!* !*!!!!!!!!!!!!!!!

logo
NTDB id 261277 CXB49 RS18965 WP 101709811.1  YQSQGKSVLLAAGDTFRAAAREQLIAWGERNNVTVIAQQGGDSAAVCYDAIQAAVARGIDIVLADTAGRLPTQLHLMEEI  286
NTDB id 1118 NGFG RS11455 WP 003696286.1     FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                    *! !!!!!!!!!!!!!!!!!!!!! !!! !!!!!!! !  !!!!!!!*!!*!!! !!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 261277 CXB49 RS18965 WP 101709811.1  KKVKRVIQKALPDAPHEVVLVLDANIGQNTVSQVKAFDDALGLTGLILTKLDGTAKGGVIAAIAKQRPIPLRFIGVGESI  366
NTDB id 1118 NGFG RS11455 WP 003696286.1     KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                    !!!!!!*!!!*! !!!!***!!!!!!!!! ! !!!!!!!!!!!!!!!!*!!!!!!!!!!**!!*!  !!*!*!*!!!!! !

logo
NTDB id 261277 CXB49 RS18965 WP 101709811.1  DDLRPFDSRDYIDALFE   383
NTDB id 1118 NGFG RS11455 WP 003696286.1     DDLRPFDARAFVDALLD   417
consensus                                    !!!!!!! ! **!!! *
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X non conserved

X similar

X $\geq$ 50% conserved