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logo

NTDB id 1103416 ACNH6H RS16065 WP 418319063.1  MFSFFKKRKAAAAEAAAKAAQQGPS..............AEELEAASINVSKRSA...............EPATEEPAH..   50
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                      !!!!!***!    *!   !  ! * ************ !**!     !  !  **************!*  ! ! ***

logo

NTDB id 1103416 ACNH6H RS16065 WP 418319063.1  ...DPFAFLAE...PLPGPSAEEISAQIAAEIAAREAWRKKLRGGLRKT....GSALSAVFTTTKVDEQLYEELEAALLM  120
NTDB id 1118 NGFG RS11455 WP 003696286.1       EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                      *****     !**** *!*    *   *!    ** *!  *!* !! !******  ! *!!    * ! !!!!!!  !*

logo

NTDB id 1103416 ACNH6H RS16065 WP 418319063.1  SDTGVKATQFLLDDLRRRVKQARETDPMKVKELLSEAIVDLLQPLEKALVVGQ.YSPTVMMVVGVNGAGKTTSIGKLTRY  199
NTDB id 1118 NGFG RS11455 WP 003696286.1       GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                       ! !* !! *!* !*! !!      !*  **  ! !!* !!* !!!!*!!** *  ! !*!* !*!!!!!!!!!!!! *!

logo

NTDB id 1103416 ACNH6H RS16065 WP 418319063.1  LSDHGQKVLLAAADTFRAAAREQLAVWAERSDAHSAGGARAAGLAPPAPVEFVSQTGGDPAAVAFDAVAAGRARRSDVVI  279
NTDB id 1118 NGFG RS11455 WP 003696286.1       FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNN................VTVISQTTGDSAAVCFDAVQAAKARGIDIVL  303
consensus                                         *!  !!!!!*!!!!!!!!!!!  !* !  ****************!  *!!! !!*!!!*!!!! !**!!  !*!*

logo

NTDB id 1103416 ACNH6H RS16065 WP 418319063.1  ADTAGRLPTQLHLMEELKKIKRVITKAQPDAPHEVLLVIDGNTGQNALSQVKAFDEALQLTGLIVTKLDGTAKGGVMAAI  359
NTDB id 1118 NGFG RS11455 WP 003696286.1       ADTAGRLPTQLHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAAL  383
consensus                                      !!!!!!!!!!!!!!!!!*!!*!!!* !! ! !!!!***!*!*! !!!!* !!!!!!*!! !!!!!!!!!!!!!!!!**!!*

logo

NTDB id 1103416 ACNH6H RS16065 WP 418319063.1  ALWSRERAAAKPGAGGVPIYFIGVGEQLEDLQTFNAREFAQALLG  404
NTDB id 1118 NGFG RS11455 WP 003696286.1       AS...........DRPVPVRYIGVGEGIDDLRPFDARAFVDALLD  417
consensus                                      ! ***********  *!!* *!!!!! **!! *! !! !  !!!
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X non conserved

X similar

X $\geq$ 50% conserved