logo
NTDB id 406663 GO999 RS14750 WP 211906342.1 MFSFWKKRKAEPQPAAEPAPPAAAPEAAQAPAP...AP.....VATPVPTPAPAP...VAAP.....VAVPDTPAAPA 62
NTDB id 1118 NGFG RS11455 WP 003696286.1 MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG 80
consensus !!!!****! * !! !*!** * * ************! *!* *!* **** ******* !** ! **
logo
NTDB id 406663 GO999 RS14750 WP 211906342.1 ............ALDMQADDIE.TVPTPPVVEQARKGWMSRLRSGLSKTS....KSLTTLFVGVKVDEALREELETALLM 125
NTDB id 1118 NGFG RS11455 WP 003696286.1 EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT 160
consensus ************ * ** * * * ** ! ** !! !!* !! !* ****!!! *! ! * ! !*!!!!! !*
logo
NTDB id 406663 GO999 RS14750 WP 211906342.1 ADAGVDATEYLLDELRRRVKAQRIETAEGVKTALRDLLIELLHPLEKTMVLGR.DQPTVIMIAGVNGAGKTTSIGKLCKH 204
NTDB id 1118 NGFG RS11455 WP 003696286.1 GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY 240
consensus *! !**!!!!!* **! !! * * ** !!** ! *!**!!!!**!!* * ! !!!*!!*!!!!!!!!!!!!*!*
logo
NTDB id 406663 GO999 RS14750 WP 211906342.1 FQTYGQSVLLAAGDTFRAAAREQLVIWGQRNNVTVVAQESGDPAAVIFDAVNAARARGIDIVMADTAGRLPTQLHLMEEL 284
NTDB id 1118 NGFG RS11455 WP 003696286.1 FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI 320
consensus !! ! !!!!!!!!!!!!!!!!!!! !! !!!!!!* ! *!!*!!!*!!!!*!!*!!!!!!!*!!!!!!!!!!!!!!!*
logo
NTDB id 406663 GO999 RS14750 WP 211906342.1 KKVRRVIGKAMATAPHETLLVIDANTGQNALAQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVGEQV 364
NTDB id 1118 NGFG RS11455 WP 003696286.1 KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI 400
consensus !!!*!!* !!** !!!! **!*!!! !!!!* !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*! !!!!! *!!!!! *
logo
NTDB id 406663 GO999 RS14750 WP 211906342.1 EDLQPFSAREFADALLG 381
NTDB id 1118 NGFG RS11455 WP 003696286.1 DDLRPFDARAFVDALLD 417
consensus *!! !! !! ! !!!!
X non conserved
X similar
X ≥ 50% conserved