```
logo
NTDB id 1239 Cj0634 YP 002344063.1          ...MASEILPDEFLELFKDLKKPPKKLHYKGNLSLL.....KQDKIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGVD   77
NTDB id 23002 HH RS05935 WP 011116067.1     ..MIADILSLRSLPKAFEILPNPPKKLFYCGKNEVISTLLDCEHKIAIVGTRKPNPYTKSFVATLASKIAQNNGVVISGGALGVD   83
NTDB id 1210 C694 RS01710 WP 001862983.1    MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKL....LDTPLKVAIIGTRRPTPYSKQHTITLARELAKNGAVIVSGGALGVD   81
consensus                                      **  **** ****!**!* !!!!!** !   **   **   !*!!*!*!****!*!* * *!!* **** ****!!!!!!!!

logo
NTDB id 1239 Cj0634 YP 002344063.1          ITASMAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSGSMQSAK  162
NTDB id 23002 HH RS05935 WP 011116067.1     IIAHTHAYPRTIMFSPASLEILYPKSNAEMIQKMMKECLVLSEYEKEYMPHKYSFLERNRLVIALSDKVIIPQADKQSGSMQSAR  168
NTDB id 1210 C694 RS01710 WP 001862983.1    IIAQENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSGSMSSAR  166
consensus                                   !*!   !*!*!!**** *!* *!!**! **!* *  **!*!!!!!****! ***!! !!!!!!!!!* !***!!!**!!!!*!!*

logo
NTDB id 1239 Cj0634 YP 002344063.1          LALELNKPLYVLPQRLGESTATNLLLKENKAKLICDFKEFVSEFAS......IDMNQDEFLEFCKKGVSVDEALKIYGQKVYEYE  241
NTDB id 23002 HH RS05935 WP 011116067.1     IALELGKPIFVLPHRIGESEGTNALLSTGEAQAIYDVNAFIESTFGT.....KIREDDDEVLLFCANNPSFEEAYERFGAKIYEYE  248
NTDB id 1210 C694 RS01710 WP 001862983.1    LAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKEMPEMEDEFLEYCAKNPSYEEAYLKFGDKLLEYE  251
consensus                                   *!***!!**!!!*!**!!**!! !!  * !**!**** !* *          *   *!!*!**!****!**!!* **! !**!!!

logo
NTDB id 1239 Cj0634 YP 002344063.1          LEGKISIEGLFIRVLA   257
NTDB id 23002 HH RS05935 WP 011116067.1     FEGKIIRESSFVRVP.   263
NTDB id 1210 C694 RS01710 WP 001862983.1    LLGKIKRINHIVVLA.   266
consensus                                   **!!! **  ****
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X non conserved

X similar

X ≥ 50% conserved