```
logo
NTDB id 293122 SALB1 RS01900 WP 109992319.1  ..................................................................................  0
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG  80
consensus                                    ********************************************************************************

logo
NTDB id 293122 SALB1 RS01900 WP 109992319.1  ......................MAKPDAPKTTKLKQPGFLTRMRASLNKGNSWLTYDLRNLLPGETVDQDALDELETQLLL  59
NTDB id 1118 NGFG RS11455 WP 003696286.1     EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                    *********************   **!* !!  * *!*  !**  ! !   *   !  * *!  *  !  *!!!! !*

logo
NTDB id 293122 SALB1 RS01900 WP 109992319.1  ADVGVDATGQVMHRLQTLYDGGQLRTSKDLGKSLREHLGNILAPCEAPLAIPDDIKPYVILMVGVNGVGKTTTIGKLSKK  139
NTDB id 1118 NGFG RS11455 WP 003696286.1     GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                    *!*!**!!  *!* *       !*   *!   !*!*!  ** !*! !! *!*   !*!!** !*!! !!!!*!!!! !

logo
NTDB id 293122 SALB1 RS01900 WP 109992319.1  LQDAGKSVMLAAGDTFRAAAVEQLTEWGERNNVPVVSQGSGADSASVVFDAYSRATARGVDVLIADTAGRLHTQGHLMEE  219
NTDB id 1118 NGFG RS11455 WP 003696286.1     FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTG.DSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEE  319
consensus                                     !  !!!!*!!!!!!!!!!! !!!  !! !!!!*!*!! *!*!!! !*!!!   ! !!!*!***!!!!!!! !! !!!!!

logo
NTDB id 293122 SALB1 RS01900 WP 109992319.1  LKKIKRVLAKQNADAPHEIMLVVDATTGGNALQQAVAFNDAVGLTGITVTKLDGSASGGVVFAICQKLGLPLRFIGVGET  299
NTDB id 1118 NGFG RS11455 WP 003696286.1     IKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEG  399
consensus                                    *!!*!!!! !  * !!!!!**!*!!  ! !!**!  !! !!*!!!!* !!!!!!*! !!** !**   **!*!*!!!!!

logo
NTDB id 293122 SALB1 RS01900 WP 109992319.1  VADLGVFEAEHFMDALFDRQA   320
NTDB id 1118 NGFG RS11455 WP 003696286.1     IDDLRPFDARAFVDALLD...   417
consensus                                    * !! *!*! *!*!!! !***
```

X non conserved

X similar

X $\geq$ 50% conserved