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logo                                                
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NTDB id 1184 GTF74 RS02335 WP 000649326.1      .........DANGLWTCSTT...................VTSEIAPKGCTAGATIN.......   153
NTDB id 1167 A1552VC RS11065 WP 000649326.1    .........DANGLWTCSTT...................VTSEIAPKGCTAGATIN.......   153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1     .........DANGLWTCSTT...................VTSEIAPKGCTAGATIN.......   153
NTDB id 1038 HON27 RS01510 WP 000993725.1      .........STDGAWTCVSD...................VPTEFLPKGCTAGTPTEGAITSL   149
NTDB id 1192 PAKAF RS23870 WP 024947914.1      .........AADGLWKCTSD...................QDEQFIPKGCSK............   150
NTDB id 1197 PSJM300 03940 AFN76866.1          .........TADGEWSCAST...................LDAKYKPGKCS.............   139
NTDB id 1191 AAC28468.1 484..945( )            .......ASGQISTWTCAPG................AANGIETRRLPTSCQ*............   153
NTDB id 1021731 AB1723 RS10680 WP 367380155.1  .......DNNGSISWACAGD................GTKVAAKYLPKACR.............   142
NTDB id 1119 NGFG RS11465 WP 017147169.1       .......RENGSVKWFCGQPVTRADDDTVADA..KDGKEIDTKHLPSTCRDKASDAK.....   166
NTDB id 1109 OK783 RS10670 WP 157147416.1      .......RQDGSVKWFCGQPVTRTGDNDDTVA..DANNAIDTKHLPSTCRDESSAT.......   164
NTDB id 1188 NELON RS01755 WP 003772458.1      AVP.AATSTGAIDWACASAAKETADKRFGTGSGVAVGTLPAKYAPSECK.............   188
NTDB id 1424 RS RS02795 WP 011000517.1         ALPVSSAPSGPIMWTCYAQNKAQAASSVAPS...GTMSLAAKYAPAECR.............   168
NTDB id 1002 ACIAD RS15100 WP 004923779.1      .......YASGAVTWKCTTT...................S.DKKYVPSECRG............   147
NTDB id 1274 LPP RS09530 WP 011214168.1        .......QANGDVTWTCTGG...................TLLAKYRPASCRP............   136
NTDB id 1258 GCO85 RS09745 WP 014844212.1      .......QANGDVTWTCTGG...................TLLAKYRPASCRP............   136
consensus                                                 *   !*!                   *  **  !  !
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X non conserved

X similar

X ≥ 50% conserved