```
logo

AT10112 QIP66197.1 MK544943:1-270                MAATKKETRLNIRIDPELKSAAQIVANDMGIDLTAAVTMFMTKMVKDHALPFTPTSL..PVETLQALKEAKH...    70
AT10111 QIP66199.1 MK544944:1-270                MAATKKETRLNIRVDPELKSAAQIVANDMGIDLTAAVTMFMTKMVKDHALPFTPTSL..PVETLQALKEAKH...    70
AT10048 WP 020862962.1 NZ CP001668:893772-894044 ....MKKVNVNIKMDPEIKKQASLLFKEFGMTMSSAINLFVKTAVEQNDIPFEYNREITNKETLEAFKEGERLLA    71
AT6256 WP 012576473.1 NC 011593:c15720-15442     ..MSGTTTTVNVRMNAELKKQAEELFGELGMNLTTAVNIFIRQSLRQGGIPFNVSLDEPNRETVSAMLEAKRIAR    73
AT10095 WP 005924829.1 NC 013316:c2056188-2055910 ..MAGNTTNISIRMDADLKAQADALFTELGMNLTTAFNIFVRQSLREGGIPFEVRLEQPNKETVAAMLEAERIAK    73
consensus                                           * * * *********! *! *** **!* ***!***!**  *** **!!  * *  **!!* !**!****

logo

AT10112 QIP66197.1 MK544943:1-270                PELLKKYSTPDDMWRDLNV    89
AT10111 QIP66199.1 MK544944:1-270                PELLKKYSTPDDMWRDLNV    89
AT10048 WP 020862962.1 NZ CP001668:893772-894044 DKNAKRYSSFKEILDSLDK    90
AT6256 WP 012576473.1 NC 011593:c15720-15442     DGNVKGYRDLNALFKDLDA    92
AT10095 WP 005924829.1 NC 013316:c2056188-2055910 DPSVKGYNDLDELFADLKR    92
consensus                                        *  *!*!** *****!
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X non conserved

X similar

X ≥ 50% conserved