

logo

NTDB id 65617 HP1WT RS00225 WP 001045839.1

NTDB id 1218 CAA10657.1 2413..3543()

consensus

MNKLKGAIVFYGGFATITISLYYHQPKAPLNNQPSLLNDDEVKYPLQDYTFTQNPQPTNTESSKDATIKALQEQKAAALKAL

MNKLKGAIVFYGGFATITISLYYHQPKAPLNNQPSLLNDDEVKYPLQDYTFTQNPQPTNTESSKDATIKALQEQKAAALKAL

!!!!!!!!!!*!!!!!!!!!!*!!

8585

logo

NTDB id 65617 HP1WT RS00225 WP 001045839.1

NTDB id 1218 CAA10657.1 2413..3543()

consensus

NSKEMNHSKEETFKNPPIDLKANTNPPKKDFSSKQWDLAARITPFKQSPKNGYEENLIFPMDNPKGIDGFTNLKEKDIAITNENKL

NSKEMNHSKEETFKNPPIDLKANTNPPKKDFSSKQWDLAARITPFKQSPKNGYEENLIFPMDNPKGIDGFTNLKEKDIAITNENKL

!!!!!!!!*!!!!!!!!!!***!!!!!!!!!!!!!!!!!!*!!

170168

logo

NTDB id 65617 HP1WT RS00225 WP 001045839.1

NTDB id 1218 CAA10657.1 2413..3543()

consensus

LRTITADKMIPAFLLITPISSQIAGKVIAQVESDIFAHMGKAVLIPKGSKVIGYYSNNNMGEYRLDIVWSRIITPHGINIMLTNA

LRTITADKMIPAFLLITPISSQIAGKVIAQVESDIFAHMGKAVLIPKGSKVIGYYSNNNMGEYRLDIVWSRIITPHGINIMLTNA

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255253

logo

NTDB id 65617 HP1WT RS00225 WP 001045839.1

NTDB id 1218 CAA10657.1 2413..3543()

consensus

KGADIKGYNGLVGELIERNFQRYGVPLLLSTLTNGLLIGITSALNNRGNKEGATNFFGDYLLMQLMRQSGMGINQVNVQILRDKS

KGADIKGYNGLVGELIERNFQRYGVPLLLSTLTNGLLIGITSALNNRGNKEGATNFFGDYLLMQLMRQSGMGINQVNVQILRDKS

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340338

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logo

NTDB id 65617 HP1WT RS00225 WP 001045839.1

NTDB id 1218 CAA10657.1 2413..3543()

consensus

KIAPIVVIREGSRVFISPNTDIFFPIPRENEVIAEFLK

KIAPIVVIREGSRVFISPNTDIFFPIPRENEVIAEFLK

!!

378376

X non conserved

X similar

X ≥ 50% conserved