

NTDB id	125760 DP15 RS00385 WP 002984519.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEGLYEAESGYDILLLDMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id	258 KZH43 RS03475 WP 000590640.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEGLYEAESGVYDLILLDMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id	217 SPD RS03775 WP 000590640.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEGLYEAESGVYDLILLDMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id	183 SPR RS03560 WP 000590640.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEGLYEAESGVYDLILLDMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id	147 SP RS03905 WP 000590640.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEGLYEAESGVYDLILLDMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id	362 SMU RS05200 WP 002262214.1	..MIKILLVEDDLGLSNSV.FDFLDDFADVMQVFDGEGLYEAESGVYDLILLDMLPEKNGFQVLKELREKGITTPVLIMTAKE	82
NTDB id	413 AAK55817.1 141..845()	..MKKILIVDEKPIISDIIFKNMTKEGYEVTAFNGREALQEFAEQPDIIILDMLPBLDGLEVAKTIRKT.SSVPILMLSAD	82
NTDB id	374 SMU RS06885 WP 002262930.1	..MKKILIVDEKPIISDIIFKNLAKEGYDTITAFDGREALSKYEENPDIIILDMLPBLDGLEVAKEVRKN.SHVPIIMLSAD	82
NTDB id	606 V4T04 RS01910 WP 003130756.1	MTSKKILIVDEKPNLARFVSLELEHEGYATEIKDNGRSGLEATSKDYDILLLDMLPBLDGFEVARRLRKE.KDTPIMMTARD	84
NTDB id	471 HSISS4 RS01430 WP 002883757.1	.MSKRILIVEDEKNLARFVSLELQHEGYDVVTADNREGLEMALEKFDLILLDMLPEMDGFEVTRRLQCE.KDTPIMMTARD	83
consensus		*****!****!* **** * *****!*****!*****!*****!*****!*****!*****!*****!*****!	

[illegible][illegible]

- X non conserved
- X similar
- X >50% conserved