

[illegible][illegible]

NTDB id	1111 NGFG RS09220 WP 003689814.1	MTLPLIWLGLIFNLDGGFVPLQSAVLGAVAGYSSLWLLCAVVKLLTGKTMGMNGDFKLIAALGAWLGISAPVLI FVSS	235
NTDB id	1059530 ACFSOV RS22370 WP 002888947.1	LTLPLIWAGILFNINLEVYLIALPD AVA GAMAAGLYALSWSVYWLFRLLTGKEALGYGDFKLLAALGAWCGWQVLPQVILLAS	224
NTDB id	1080 FDQ49 RS18305 WP 022575844.1	F TLPLAALGLGINTFNIYTSPNSAIWG YLIGFLCLWIIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLI V LLS	239
NTDB id	1061 ABD1 RS18470 WP 001152280.1	F TLPLAALGLGINTFNIYTSPNSAIWG YLIGFLCLWIIVYYLFKVITGKEGMGYGDFKLLAALGAWMGPLMLPLI V LLS	239
NTDB id	1170 A1552VC RS11080 WP 000418747.1	LTLPLIWWGIALLLGFSFVSLSDAVI GAMAGYLSLWSIYWLFKLLTGTGEGMGYGDFKLLAALGAWLGWQQLPVIVLLS	236
NTDB id	1403 DSB67 RS12675 WP 010643256.1	L TLPLMW TGIALLAGISPSVSLQDSII GAMAGYLSLWSVYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQS LPM I ILLS	236
consensus		* ! ! ! ! * ! ! ! ! * ! ! ! ! * ! ! ! ! * ! ! ! ! *	

NTDB id	1111	NGFG	RS09220	WP	003689814.1	LIGLVAAIVMRV...AKGRHFAGFPALT VSGWITFTANDSVWRANWNWLTHPVR...	286
NTDB id	1059530	ACFSOV	RS22370	WP	002888947.1	ASGLVWTLLQRLWTRQSLQQPLAFGPWLALAGGSIFLWQQMV.....	266
NTDB id	1080	FDQ49	RS18305	WP	022575844.1	LLGAIITIGIILLKLRND..NQPFAGFPGYTAIAGWVAFWLGDDQI...MKIYVLGG.....	286
NTDB id	1061	ABD1	RS18470	WP	001152280.1	LLGAIIIGIILLKLRND..NQPFAGFPGYTAIAGWVAFWLGDDQI...MKIYVLGG.....	286
NTDB id	1170	A1552VC	RS11080	WP	000418747.1	VVGVIIFGLIQRLRQQKKKGIDMAFFPGPYLAIAGWFA LLWGDKV...IDWYFTTWVGQPL	291
NTDB id	1403	DSB67	RS12675	WP	010643256.1	VVGVIIFGLIQRLRKQKGIEKAFFPGPYLAIAGWVSLT WGHQI...LNWYFTSILGV..	289
consensus						**!*!!** ** * * *	*! !*****! * * ***** *

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