

NTDB id 443 SPYM3 RS00325 WP 011054099.1	MLEHFGGKVKVLRLEKRLISREDLCGDESELSVRQLARIELGQSIPSLSKVIFIAKALNVSVGYLTDGADLLELPKRYKELKYLILR	85
NTDB id 70556 FGCD RS00710 WP 129555155.1	MLEHFGGEKVRGLRLEKRLISREDLCGDEAELSVRQLARIELGQSIPSLAKVIFIAKALNVSVGYLTDGANLLELPKRYKELKYLILR	85
NTDB id 451 SPYM18 RS00325 WP 002986681.1	MLEHFGGKVKVLRLEKRLISREDLCGDESELSVRQLARIELGQSIPSLSKVIFIAKALNVSVGYLTDGADLLELPKRYKELKYLILR	85
NTDB id 45 SMU RS00375 WP 002263400.1	MLKDFGKKIKSLRLEKRLTKEAVCLDESQLSTRQLTRIESGQSTPTLNKAVYIAGRLGVTLGYLTDGENVELPSRYKELKYLILR	85
NTDB id 555 STRINF RS01865 WP 174221231.1	MREEKEISREEFCEGDETELSVRQLARIELNQSIPNLKASFIANRLGVKLGTLTDGDSLLELPKRYKELKYLILR	74
consensus	*****	

[illegible][illegible]

NTDB id 443 SPYM3 RS00325 WP 011054099.1	WKLSTFYFLKDIRNARNYFEQSSILFTQMTGDTYLVQKLRGEWNKDIHYI.	303
NTDB id 70556 FGCD RS00710 WP 129555155.1	WKYFLIKQKNRKQAEEESYTKACLFQAQITGDKYLEEQLIKIEWEKDICAY.	303
NTDB id 451 SPYM18 RS00325 WP 002986681.1	WKYFLIKQKDIKAEQSFMKACLFQAMTADQYLENKLIQIEWEKDVKSY.	303
NTDB id 45 SMU RS00375 WP 002263400.1	WKHHLFVEKDYAEAKKSYDAAILFQAQLTENINLRENLEKEWQKDSQNGT	304
NTDB id 555 STRINF RS01865 WP 174221231.1	WKYYLKLKHDYFALAEQSFSTNATLFAIRLVGDTYLENKKLKEEWQLDIDAVE	292
consensus	! ! ! ! ! * * * * ! ! ! ! ! * * * * ! ! ! ! ! * * * * ! ! ! ! !	

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