

```

NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1252 GCO85 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 301310 DPR03 RS01160 WP 004810078.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus

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consensus
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390
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390
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393
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logo

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NTDB id 1402 DSB67 RS12670 WP 010643257.1
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Position	1	2	3	4	5	6	7	8	9	10
Sequence	LVT	AM	YLP	IF	N	LM	SV	LG	.	.
407	LVT	AM	YLP	IF	N	LM	SV	LG	.	.
408	LVV	AM	YLP	IF	N	LM	SV	LG	.	.
410	LVV	AM	YLP	L	F	N	LG	N	V	A
410	LLV	AM	YLP	L	F	N	LG	N	V	A*
406	LVV	AM	YLP	IF	K	LG	S	A	V	.
405	LII	AM	YLP	IF	Q	M	GS	V	V	.
408	LV	I	AM	YLP	IF	Q	M	GS	V	V
409	LV	I	AM	YLP	IF	Q	M	GS	V	V
408	LV	I	AM	YLP	IF	Q	M	GS	V	V

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

-  non conserved
-  similar
-  $\geq 50\%$ conserved