

[illegible]

NTDB	id	1184	GTF74	RS02335	WP	000649326.1	GFTR...	L	G	T	V	E	D	M	.	G	.	D	G	K	I	V	I	A	P	.	T	A	S	G	A	L	G	G	.	T	I	K	Y	T	F	D	A	...	G	V	V	S	S	S	K	I	Q	L	A	R	D	...	A	N	G	L	W	131		
NTDB	id	1167	A1552VC	RS11065	WP	000649326.1	GFTR...	L	G	T	V	E	D	M	.	G	.	D	G	K	I	V	I	A	P	.	T	A	S	G	A	L	G	G	.	T	I	K	Y	T	F	D	A	...	G	V	V	S	S	S	K	I	Q	L	A	R	D	...	A	N	G	L	W	131		
NTDB	id	1160	ABDM36	RS02330	WP	000649326.1	GFTR...	L	G	T	V	E	D	M	.	G	.	D	G	K	I	V	I	A	P	.	T	A	S	G	A	L	G	G	.	T	I	K	Y	T	F	D	A	...	G	V	V	S	S	S	K	I	Q	L	A	R	D	...	A	N	G	L	W	131		
NTDB	id	1292	VP	RS12235	WP	005479693.1	...	T	L	A	Q	L	G	V	D	V	A	...	N	D	L	G	A	I	D	S	A	...	L	A	G	S	P	T	L	T	F	T	...	D	A	D	S	S	I	G	G	I	L	T	F	T	R	D	...	A	A	T	G	W	124					
NTDB	id	1038	HON27	RS01510	WP	000993725.1	...	G	F	T	A	A	T	...	S	S	V	C	A	T	S	I	D	.	A	F	.	T	E	T	D	G	A	K	I	T	.	C	T	L	K	G	...	N	P	K	I	A	T	K	I	V	E	L	T	R	S	...	T	D	G	A	W	121		
NTDB	id	1192	PAKAF	RS23870	WP	024947914.1	ATKKEVPL	G	V	A	A	D	A	...	N	K	L	G	T	I	A	.	L	K	P	...	D	P	A	D	G	T	A	D	I	T	L	T	F	T	M	G	G	A	P	K	N	K	G	K	I	I	T	L	T	R	T	...	A	A	D	G	L	W	133	
NTDB	id	1204	PSJM300	03935	AFN	76865.1	...	G	F	I	G	I	G	A	S	T	...	N	.	Y	C	T	V	T	.	L	I	...	A	T	A	T	G	S	...	I	T	.	C	T	T	Q	N	G	M	P	D	R	F	N	G	K	T	I	T	I	A	R	T	...	T	D	G	L	W	125
NTDB	id	1197	PSJM300	03940	AFN	76866.1	...	G	Y	I	G	I	T	D	S	T	...	S	.	Y	C	D	V	D	.	L	D	...	T	A	A	D	G	H	...	I	E	.	C	T	A	K	G	G	N	A	K	F	D	G	K	T	I	T	L	N	R	T	...	A	D	G	E	W	123	
NTDB	id	1383	A4U84	RS08530	WP	005711781.1	...	K	N	G	...	V	R	.	A	D	K	S	D	V	P	A	K	.	L	K	S	...	I	S	V	A	S	G	V	I	T	V	A	G	...	G	N	I	D	G	Y	G	T	M	P	K	...	F	A	N	N	I	T	W	131					
NTDB	id	1354	HI	0299	AAC21963.1	...	K	N	G	...	I	A	.	A	D	I	K	T	...	A	K	G	V	.	A	S	...	V	I	T	Q	S	G	G	I	T	V	K	G	N	...	G	T	L	A	N	M	E	Y	I	L	Q	A	K	...	G	N	A	A	G	V	T	W	127		
NTDB	id	1365	NTHI	RS01955	WP	011271973.1	...	K	N	G	...	I	A	.	A	D	I	T	...	A	K	G	V	.	K	S	...	V	T	T	S	N	G	A	I	T	V	K	G	D	...	G	T	L	A	N	M	E	Y	I	L	Q	A	T	...	G	N	A	A	T	G	V	T	W	127	
NTDB	id	1191	AAC28468.1	484..945()	...	G	N	...	V	I	F	D	S	A	N	G</																																																		

logo

TCSTT	VTSEIAP	PKGCT	TAGATIN
TCSTT	VTSEIAP	PKGCT	TAGATIN
TCSTT	VTSEIAP	PKGCT	TAGATIN
SCLPT	APAIA	VDGCS	
TCVSD	VPTEFLP	PKGCT	AGTPTTEGAITS
KCTISD	QDEQFIP	PKGCS	
DCTISD	LDSTRYK	PGKCL	
SCAST	LDACYK	PGKCS	
ATSCQ	GADISL	PANFCG	QKATP
TTTICK	GTDASL	PANFCGS	VTK
TTTICK	GTDASL	PANFCGS	VTQ
TCAPG	AANGT	ETRRLP	PTSCQ*
FCGQPVTRADDDTVADAKDGKE	IDTKHLP	STCRD	KASDAK
FCGQPVTRTGDNDTVDANNA	IDTKHLP	STCR	DESSAT
TCYAQNKAQAASS.VAPSGTMS	LAAKYAP	PAE	CR
VCISA	I	NDALL	PATCR
KCTTT	S	DKKYVP	SECRG
TCIGG	TLLAKYR	PAS	CRP
TCTGG	TLLAKYR	PAS	CRP

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