

logo

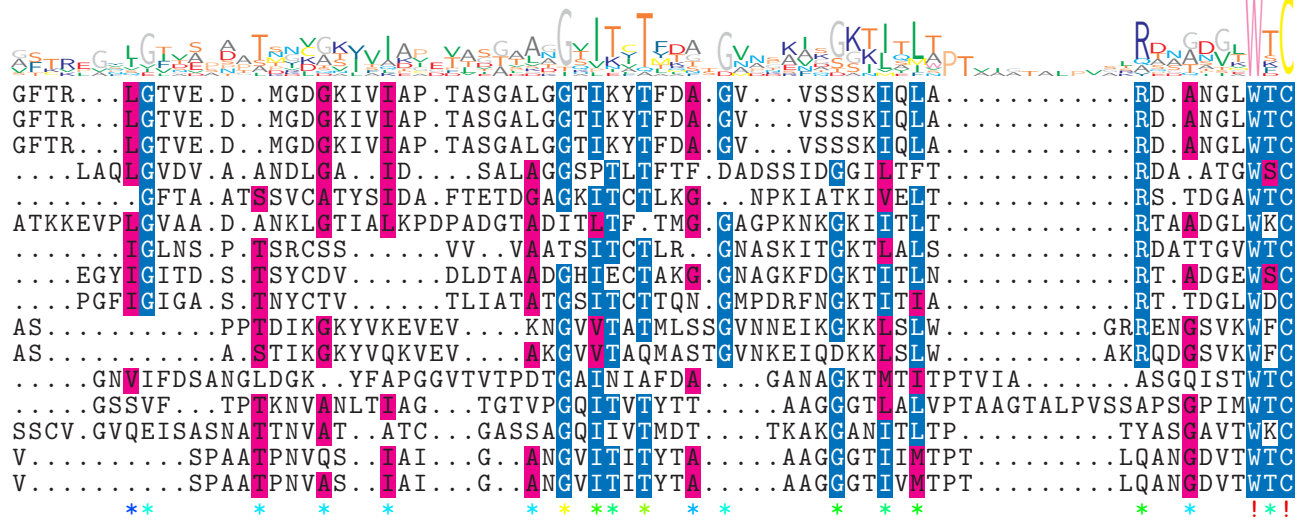
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NTDB id 1167 A1552VC RS11065 WP 000649326.1
NTDB id 1160 ABDM36 RS02330 WP 000649326.1
NTDB id 1292 VP RS12235 WP 005479693.1
NTDB id 1038 HON27 RS01510 WP 000993725.1
NTDB id 1192 PAKAF RS23870 WP 024947914.1
NTDB id 1000719 ABFU39 RS05165 WP 228442818.1
NTDB id 1197 PSJM300 03940 AFN76866.1
NTDB id 1204 PSJM300 03935 AFN76865.1
NTDB id 1119 NGFG RS11465 WP 017147169.1
NTDB id 1109 OK783 RS10670 WP 157147416.1
NTDB id 1191 AAC28468.1 484..945()
NTDB id 1424 RS RS02795 WP 011000517.1
NTDB id 1002 ACIAD RS15100 WP 004923779.1
NTDB id 1258 GCO85 RS09745 WP 014844212.1
NTDB id 1274 LPP RS09530 WP 011214168.1
consensus



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consensus



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logo

NTDB id 1184	GTF74	RS02335	WP 000649326.1	SI		TVTSEIAPKG	CTAGATIN		153
NTDB id 1167	A1552VC	RS11065	WP 000649326.1	ST		TVTSEIAPKG	CTAGATIN		153
NTDB id 1160	ABDM36	RS02330	WP 000649326.1	SI		TVTSEIAPKG	CTAGATIN		153
NTDB id 1292	VP	RS12235	WP 005479693.1	LPTAP			AIADVGC		139
NTDB id 1038	HON27	RS01510	WP 000993725.1	VS		DVPTEFL	PKGCTAGTPTEGAITS		149
NTDB id 1192	PAKAF	RS23870	WP 024947914.1	TS		DQDEQFIP	PKGCSK		150
NTDB id 1000719	ABFU39	RS05165	WP 228442818.1	QTGGSG		GGLD	AKYKPAGCS		135
NTDB id 1197	PSJM300	03940	AFN76866.1	AS		TLDAKYK	PGKCS		139
NTDB id 1204	PSJM300	03935	AFN76865.1	TS		DLDSRYK	PGKCL		141
NTDB id 1119	NGFG	RS11465	WP 017147169.1	GQPVTRADDDTVADAKDGKEIDTKHLPSTCRDKASDAK					166
NTDB id 1109	OK783	RS10670	WP 157147416.1	GQPVTRTGDNDDTVADANNAIDTKHLPSTCRDESSAT					164
NTDB id 1191	AAC28468.1	484..945()		APGAAN		GIETRRL	PTSCQ*		153
NTDB id 1424	RS	RS02795	WP 011000517.1	YAQNKAQAAS		SVAPSGTMS	LAAKYAPAECR		168
NTDB id 1002	ACIAD	RS15100	WP 004923779.1	TI		S	DKKYVPSECRG		147
NTDB id 1258	GC085	RS09745	WP 014844212.1	TGG		TLLAKYR	PASCRP		136
NTDB id 1274	LPP	RS09530	WP 011214168.1	TGG		TLLAKYR	PASCRP		136
consensus				*		*	**	* !	

- X

 non conserved
- X

 similar
- X

 ≥ 50% conserved