

logo

NTDB id 1384 A4U84 RS08525 WP 010787006.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
NTDB id 1468 ASU2 RS05090 WP 014991734.1		0
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 103485 A1R12 RS11730 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

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NTDB id 1384 A4U84 RS08525 WP 010787006.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
NTDB id 1468 ASU2 RS05090 WP 014991734.1		0
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
NTDB id 103485 A1R12 RS11730 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

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NTDB id 1384 A4U84 RS08525 WP 010787006.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAALLL	0
NTDB id 1468 ASU2 RS05090 WP 014991734.1		0
NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAALLL	240
NTDB id 103485 A1R12 RS11730 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

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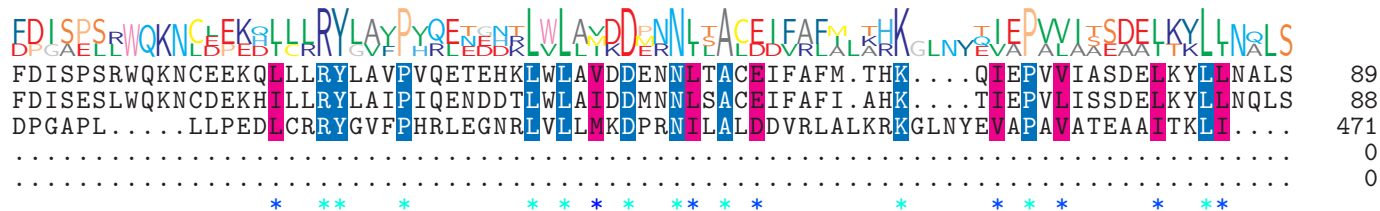
NTDB id 1384 A4U84 RS08525 WP 010787006.1	LRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	0
NTDB id 1468 ASU2 RS05090 WP 014991734.1		0
NTDB id 1028 TT RS08215 WP 011173991.1	LRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL	320
NTDB id 103485 A1R12 RS11730 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus		

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NTDB id 1384 A4U84 RS08525 WP 010787006.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGMSQYITICEQQTERI	14
NTDB id 1468 ASU2 RS05090 WP 014991734.1		13
NTDB id 1028 TT RS08215 WP 011173991.1	KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGMSQYITICEQQTERI	400
NTDB id 103485 A1R12 RS11730 WP 003153083.1		0
NTDB id 96 BSU 24730 NP 390353.1		0
consensus	*	

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consensus



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NTDB id 1384 A4U84 RS08525 WP 010787006.1	EGCSVYCRoyRRTRRAGYYE	LLYGKNLDS	LYGKGRG	YQY	LR	SS	RQK	AL		
NTDB id 1468 ASU2 RS05090 WP 014991734.1	GCKPDPETMSNL	LSTSLAEIQGARLYKMGCEE	VFGTAR	ETATFAC	FAKQA	F	DK	STAYL	DYS	T
NTDB id 1028 TT RS08215 WP 011173991.1	EVKPDPET...	LRLRLISEAEIQGARLYKMGCE	ERC	GGT	..	GYKGRY	AIH	...	ELLV	DD
NTDB id 103485 A1R12 RS11730 WP 003153083.1	ENCSVYCRMQRTRRAGIYE	...	LLYGKNLT	QCFQ	EAKGEHAN	FQYQ	...	TLRRL	IRK	GIAL
NTDB id 96 BSU 24730 NP 390353.1	GCSSVYCRQSRNTRRASVYE	...	LLYGKNLQ	CCIQ	EAKGNHAN	YQYQ	...	TLRQI	IRK	GIAL
consensus										

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NTDB id 1384 A4U84 RS08525 WP 010787006.1	GYTTNNYE	EVDRVLG	YE	...		461
NTDB id 1468 ASU2 RS05090 WP 014991734.1	NIT..TVDEVKRVLG	GDG	...			461
NTDB id 1028 TT RS08215 WP 011173991.1	GKSATEIKEIARRKGMKTLREDGLYKALQG	ITTLEEV	LARTIE			889
NTDB id 103485 A1R12 RS11730 WP 003153083.1	GYVTTDNYD	...	RWVYHEAD	...		356
NTDB id 96 BSU 24730 NP 390353.1	GYLTNNYD	...	RWVYHEKD	...		356
consensus						

- [X] non conserved
- [X] similar
- [X] ≥ 50% conserved