

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1168 A1552VC RS11070 WP 000957200.1
NTDB id 344916 SynWH8101 RS00100 WP 130128060.1
NTDB id 1410 SGL RS13950 WP 010873519.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK 80
..... 0
..... 0
..... 0

logo

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NTDB id 1410 SGL RS13950 WP 010873519.1
consensus

ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLD~~TL~~SL~~EE~~VEDLTGLVVEPYQT~~TK~~SAFLYALAKHYPELGLPVPPPPSG
 ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLD~~TL~~SL~~EE~~VEDLTGLVVEPYQT~~TK~~SAFLYALAKHYPELGLPVPPPPSG 160
 0
 0

logo

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consensus

EGQKDLKLGELLQKGI	SREALEEALVEQEK	TGDLGRILVRKGL	PEEALYRALAEQK	GLEFLESTEGIV	PDPSAALLL	240
EGQKDLKLGELLQKGI	SREALEEALVEQEK	TGDLGRILVRKGL	PEEALYRALAEQK	GLEFLESTEGIV	PDPSAALLL	0
.....						0

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[illegible]

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[illegible]

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consensus



GYPVDPEEDPPDPG.....APLLLPEDLCRRYGVPFHRLEG...NRLVLLMKDPRNIALDDVRIALKRGLNVEVAPA 460
 GIPETDLSR.....YDYANLCQQLGLRELITRYDALPIAKQG...NLLLLAVSDPTLLQAEFFRFRFAT.....GLQVELA 124
 .MTVH...APAAAA.....QNPWHQQRLE.LELLLQVPVPERLDAADAQQLQEAR..... 46
 GVDSDVPEVAPIATQQMAELIGRFFPLDTCRRFKFPLAQQEGDPPSVLVAVVDPDNLAA...QDELNRILRVKGFELRRL 161

logo

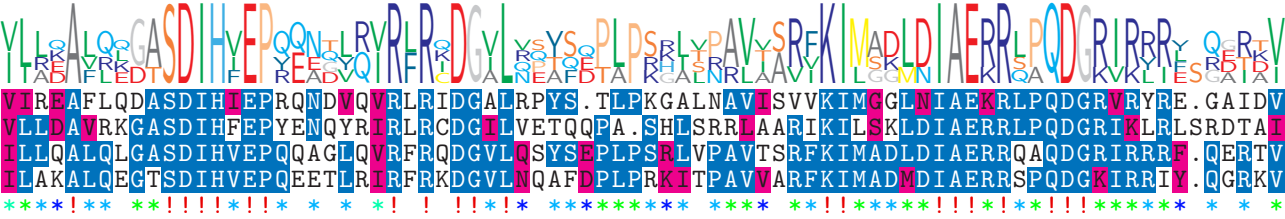
NTDB id 1028 TT RS08215 WP 011173991.1
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NTDB id 1410 SGL RS13950 WP 010873519.1
consensus



513
188
121
239

logo

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consensus



591
267
200
318

logo

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NTDB id 1410 SGL RS13950 WP 010873519.1
consensus



671
347
280
398

logo

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NTDB id 344916 SynWH8101 RS00100 WP 130128060.1
NTDB id 1410 SGL RS13950 WP 010873519.1
consensus



751
427
360
478

logo

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NTDB id 344916 SynWH8101 RS00100 WP 130128060.1
NTDB id 1410 SGL RS13950 WP 010873519.1
consensus



816
490
432
558

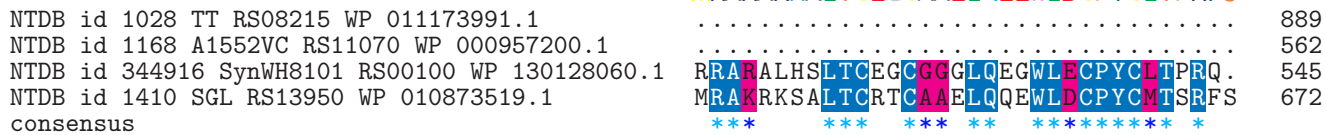
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NTDB id 1410 SGL RS13950 WP 010873519.1
consensus



889
562
512
638

logo



 non conserved
 similar
 ≥ 50% conserved