

```
NTDB id 419 DQL23 RS10745 WP 060553298.1
NTDB id 416 SGO RS10505 WP 012131078.1
NTDB id 404953 GOM48 RS09715 WP 235097586.1
NTDB id 486 SM12261 RS09375 WP 020902616.1
NTDB id 138 SP RS11430 WP 000364845.1
NTDB id 249 KZH43 RS10250 WP 000362880.1
NTDB id 208 SPD RS10925 WP 000362880.1
NTDB id 174 SPR RS10405 WP 000362880.1
NTDB id 514 SMSK321 RS10615 WP 001048129.1
consensus
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MDLFGFGTVIVHFLISHSYHFICK
MFLLFVIVVFFFTSVVRLIFNKKINRKELYVFAYTLLTEI
VLEFSFYLYLDGLGIERFLFPLG

.MILKLTIAAFFPTIIQVVSVEIFHKIMQSKKIDIKFGLFFSIVFGSITISSILIEIPSFGDVISSIFHYHFLFIQPLI	79
.MIFTIILSLFPVITIQVFSYDFFYKNINRKKKINIKLILFQWGLIMVISFLYSLFLLPDYWKIFRDIHFYFLLLIQPLI	79
MDFFAIIISVILHIFITSKSYNLICN....TT..RKEQVFFIFVTIVVEF...VVEFFFSYIYLSGL..GIEKFLYPFI	67
MDFLAVLVILHIFIISKSYLICK....NKINIKELCFRCGYTFLVES...VFELSFYFIYLDGL..GIEKFLFPLV	69
MDLLGFGTVIVHFLIISHSYRLICK....GRINRKELYVFGAYTLLTEI...VLEFSFYLYLDKI..GIERFLFPLG	69
MDLFGFGTVIVHFLIISHSYHFICK....GQINRKELVFGAYTLLTEI...VDFDPLYILYLDGL..GIERFLFPLG	69
MDLFGFGTVIVHFLIISHSYHFICK....GQINRKELVFGAYTLLTEI...VDFDPLYILYLDGL..GIERFLFPLG	69
MDLFGFGTVIVHFLIISHSYHFICK....GQINRKELVFGAYTLLTEI...VDFDPLYILYLDGL..GIERFLFPLG	69
MDLFGFGTVIVHFLIISHSYHFICK....GQINRKELVFGAYTLLTEI...VDFDPLYILYLDGL..GIERFLFPLG	69
MNLFVGGLVIVYFLIISHSYRLICK....GQINRKELYVFSAYTLLTEI...VLDFPSYLLHLDGL..GIVKFLFPLV	69

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

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[illegible]

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Position	Residue	Position	Residue	Position	Residue	Position	Residue	Position	Residue
1	Y	25	V	49	S	73	S	97	K
2	K	26	L	50	I	74	S	98	Q
3	E	27	L	51	V	75	F	99	K
4	D	28	L	52	F	76	A	100	H
5	E	29	L	53	S	77	M		
6	Y	30	L	54	S	78	N		
7	L	31	L	55	S	79	S		
8	P	32	L	56	S	80	H		
9	F	33	L	57	S	81	K		
10	V	34	L	58	S	82	V		
11	A	35	L	59	S	83	Q		
12	I	36	L	60	S	84	M		
13	N	37	L	61	S	85	E		
14	S	38	L	62	S	86	K		
15	Y	39	L	63	S	87	A		
16	T	40	L	64	S	88	I		
17	V	41	L	65	S	89	F		
18	L	42	L	66	S	90	L		
19	L	43	L	67	S	91	L		
20	N	44	L	68	S	92	L		
21	S	45	L	69	S	93	L		
22	Y	46	L	70	S	94	L		
23	I	47	L	71	S	95	L		
24	V	48	L	72	S	96	L		
25	L	49	L	73	S	97	L		
26	L	50	L	74	S	98	L		
27	L	51	L	75	S	99	L		
28	L	52	L	76	S	100	L		
29	L	53	L	77	S				
30	L	54	L	78	S				
31	L	55	L	79	S				
32	L	56	L	80	S				
33	L	57	L	81	S				
34	L	58	L	82	S				
35	L	59	L	83	S				
36	L	60	L	84	S				
37	L	61	L	85	S				
38	L	62	L	86	S				
39	L	63	L	87	S				
40	L	64	L	88	S				
41	L	65	L	89	S				
42	L	66	L	90	S				
43	L	67	L	91	S				
44	L	68	L	92	S				
45	L	69	L	93	S				
46	L	70	L	94	S				
47	L	71	L	95	S				
48	L	72	L	96	S				
49	L	73	L	97	S				
50	L	74	L	98	S				

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logo

[illegible]

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