

Probab=93.86 E-value=1.5 Score=36.69 Aligned_cols=105 Identities=19% Similarity=0.149 Sum_probs=0.0 Template_Neff=9.700

Q ss_pred		EeEeCccCceEEEEeCccCccEEcCCcCceeee	
Q SRAV	1864	RNVYKTDSDRYMVVSVDVMNYGGYSGGPVVTSDGEFLGIC	1903 (3835)
Q Consensus	1864	rnvkyktdsdrymvvsvdvmnyggysggpvvtsdgeflgic	1903 (3835)
	+.+ . . ++..+ +++ .	
T Consensus	189	g~~~~~-----G~SGgpv~~~~G~viGi~	224 (296)
T RPOA_LDVC/1418	189	GLSLDLLGKNSAFCFTK---CGDSGSPVVEDEGNLLGIH	224 (296)
T ss_pred		EeccccCCCeEeeeC---CCCCcEECCCCCEEEE	