

MuscleWS alignment of Uniref50 Sequence

File	Edit	Select	View	Annotations	Format	Colour	Calculate	Web Service
						Apply Colour To All Groups		
						Text Colour...		
						✓ None		
						Clustal		
						Blosum62		
						% Identity		
						Zappo		
						Taylor		
						Hydrophobic		
						Helix Propensity		
						Strand Propensity		
						Turn Propensity		
						Buried Index		
						Nucleotide		
						Purine/Pyrimidine		
						RNA Helices		
						T-Coffee Scores		
						User Defined...		
						By Conservation		
						Modify Conservation Threshold...		
						Above Identity Threshold		
						Modify Identity Threshold...		
						By Annotation...		