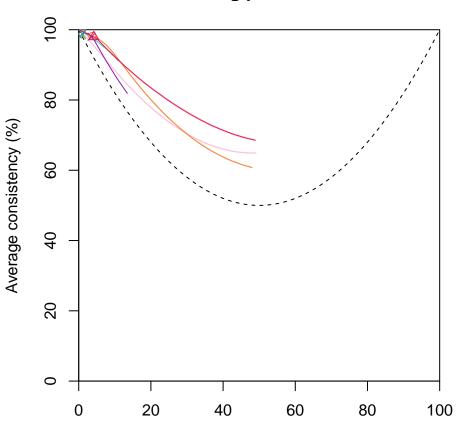
KEGG ortholog protein benchmark



Percentage of residues predicted as tandem repeat (%)