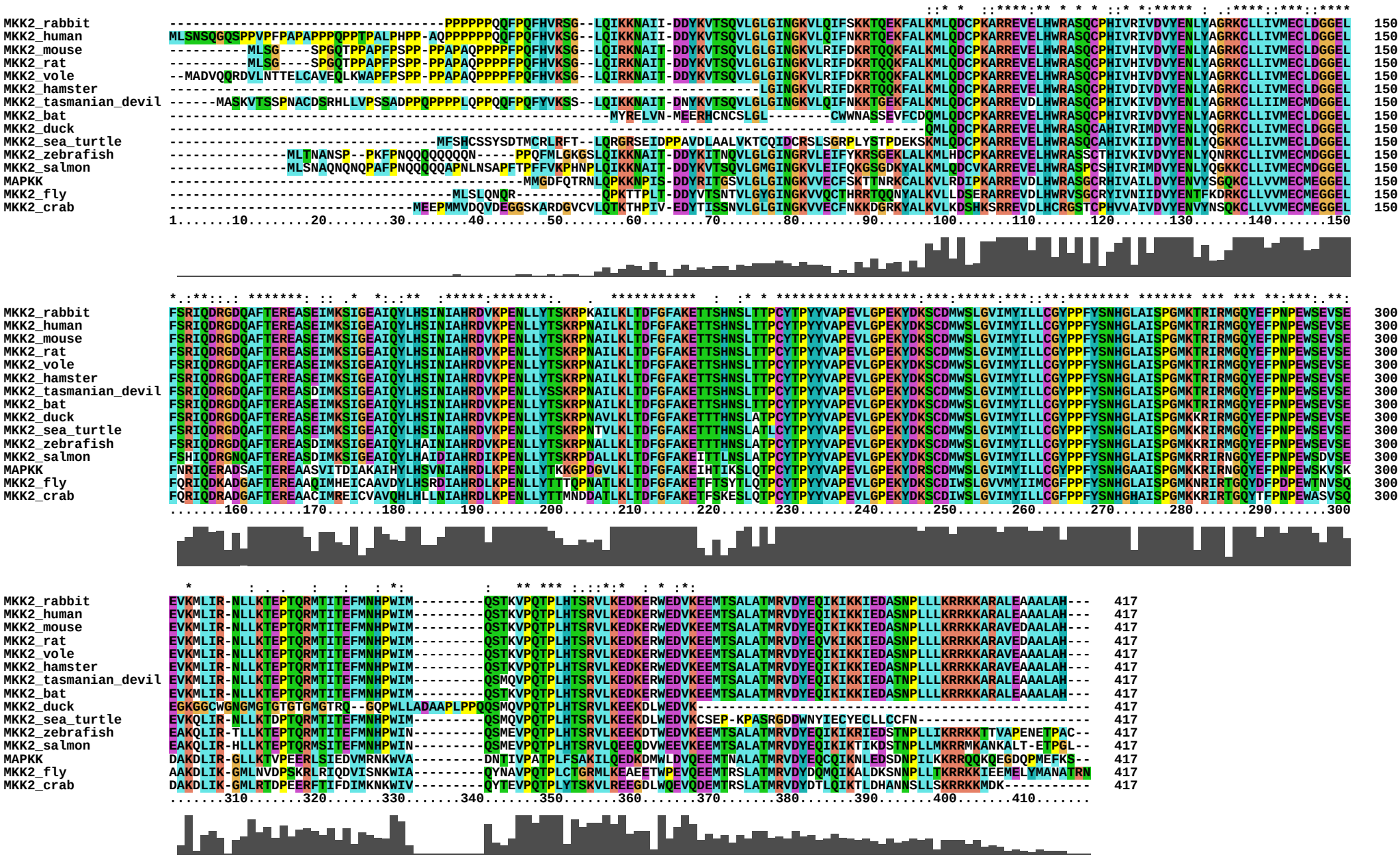CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT