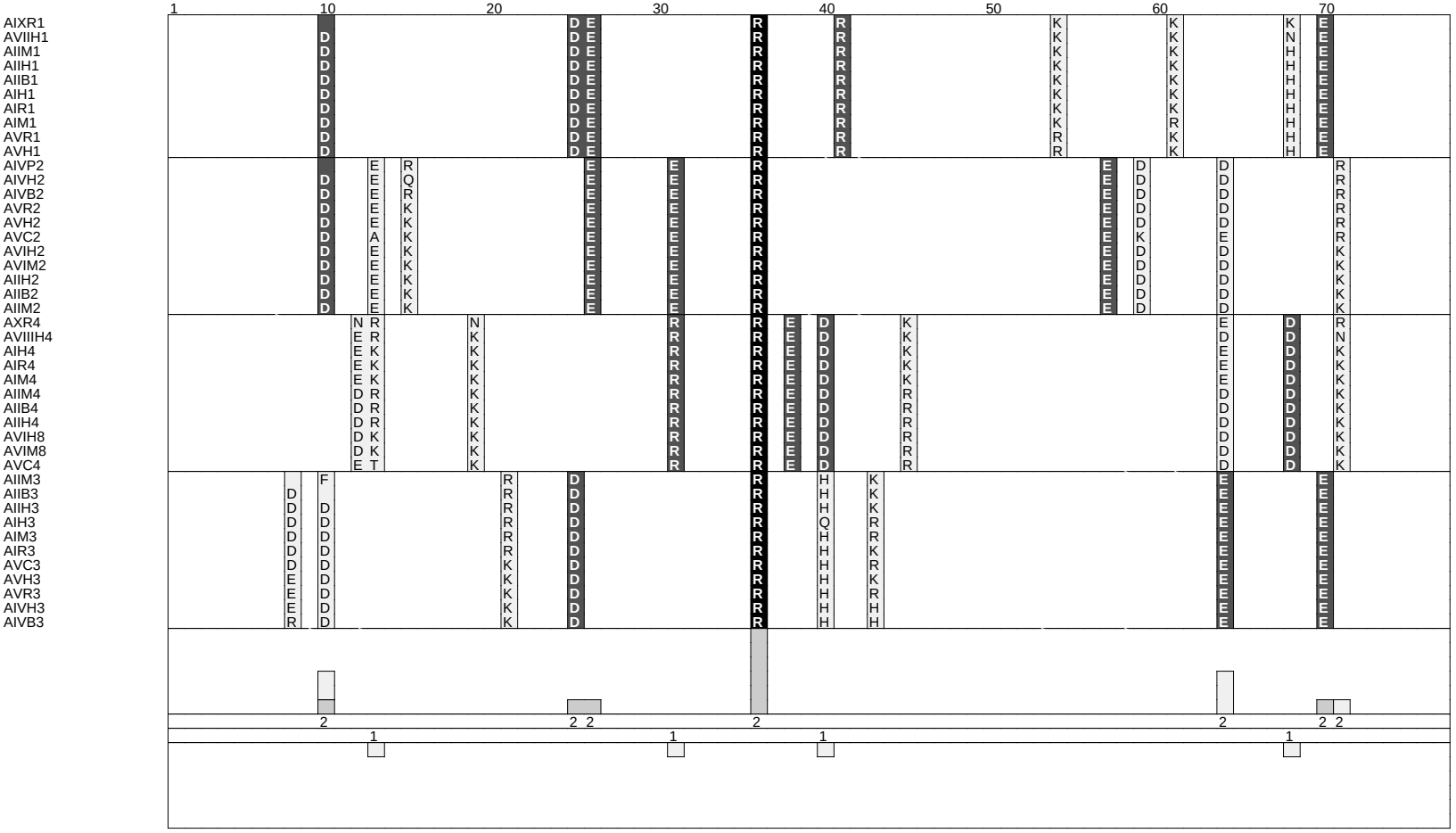

AMAS conservation plot of the sequence alignment newest

(Stringency is set to 66% with 3 property matrix /home/orac/cdl/bin/ch.pt)