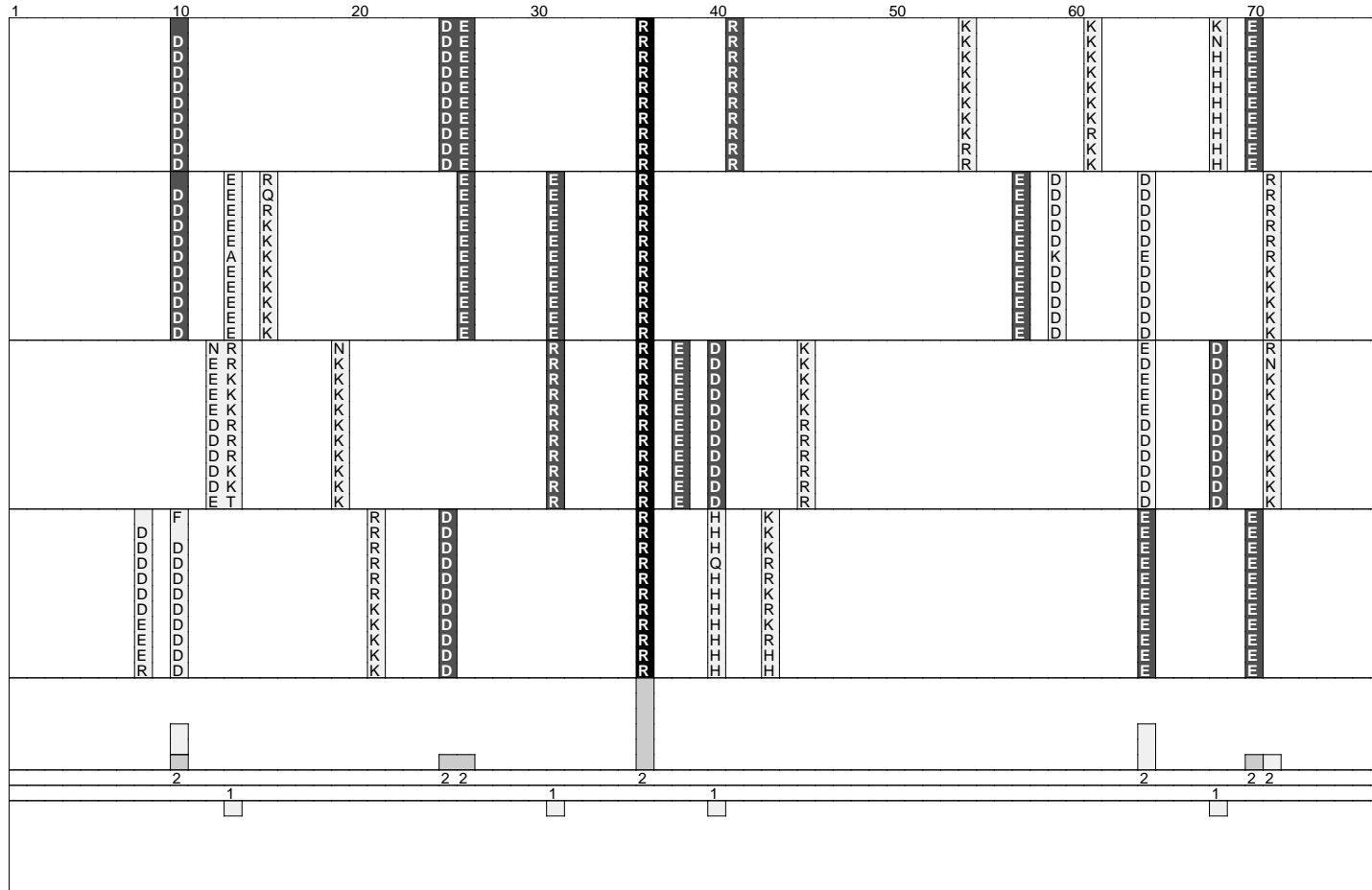


AIXR1
 AVIIH1
 AIIM1
 AIH1
 AIIB1
 AIR1
 AIM1
 AVR1
 AVH1
 AIVP2
 AIVH2
 AIVB2
 AVR2
 AVH2
 AVC2
 AVIH2
 AVIM2
 AIH2
 AIIB2
 AIIM2
 AXR4
 AVIIH4
 AIH4
 AIR4
 AIM4
 AIIM4
 AIIB4
 AIH4
 AVIH8
 AVIM8
 AVC4
 AIIM3
 AIIB3
 AIH3
 AIH3
 AIM3
 AIR3
 AVC3
 AVH3
 AVR3
 AIVH3
 AIVB3



AMAS conservation plot of the sequence alignment newest
 (Stringency is set to 66% with 3 property matrix: /home/orac/bdl/bin/ch.pt)