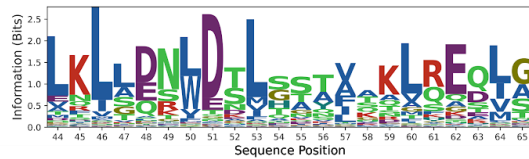
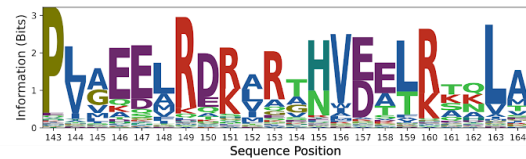


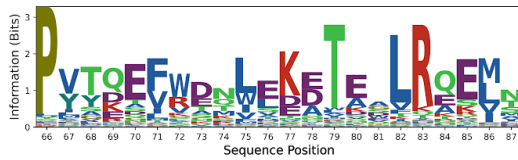
Repeat 1



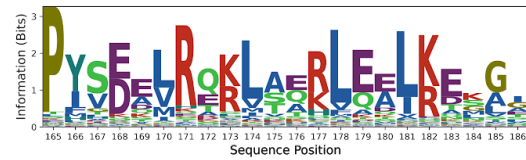
Repeat 6



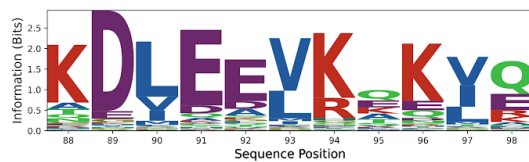
Repeat 2



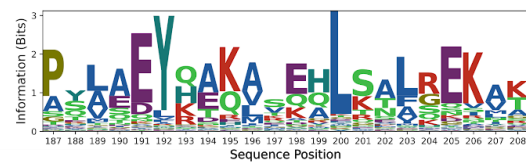
Repeat 7



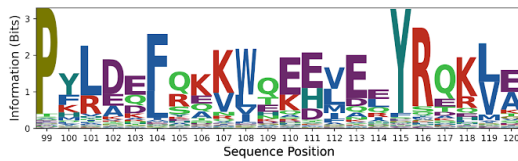
Repeat 3



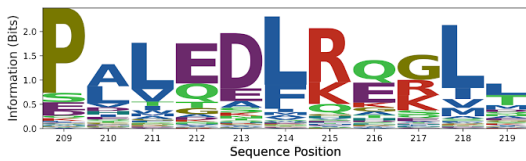
Repeat 8



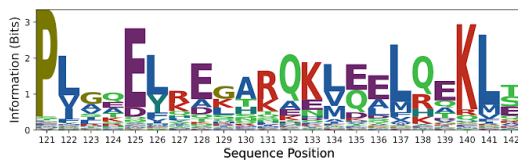
Repeat 4



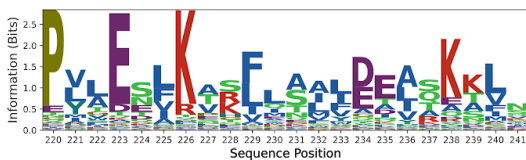
Repeat 9



Repeat 5



Repeat 10



### Supplementary Figure 2 Sequence logos for apoA-I tandem repeats

Sequence logos corresponding to each of the ten tandem repeats present in apoA-I (residues 48-243) were extracted from a multiple sequence alignment of mammalian sequence using the LogoMaker package. Position-specific conservation was calculated using information content (expressed in Bits).