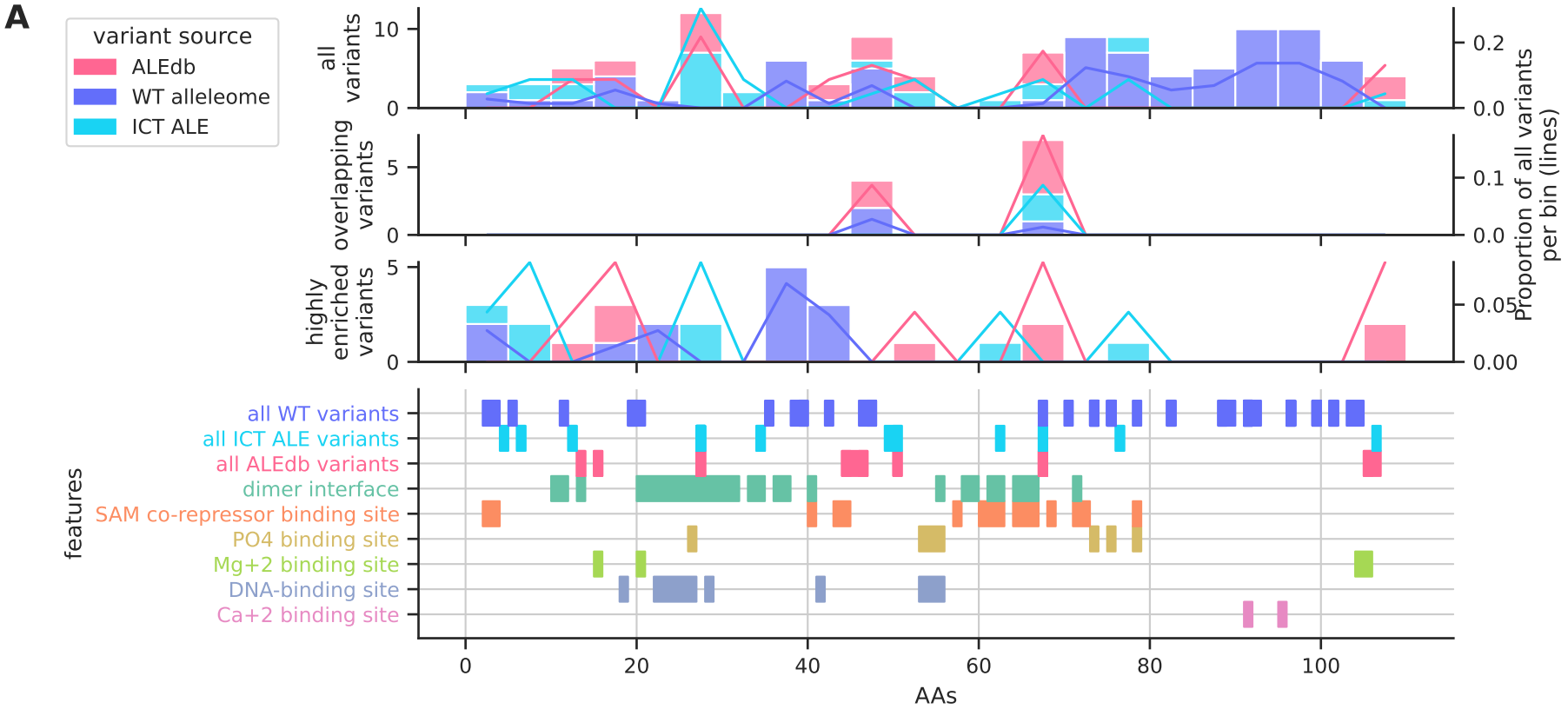
A
variant source
ALEdb
WT alleleome
ICT ALE
all variants
overlapping variants
highly enriched variants
Proportion of all variants per bin (lines)
features
all WT variants
all ICT ALE variants
all ALEdb variants
dimer interface
SAM co-repressor binding site
PO4 binding site
Mg+2 binding site
DNA-binding site
Ca+2 binding site
AAs