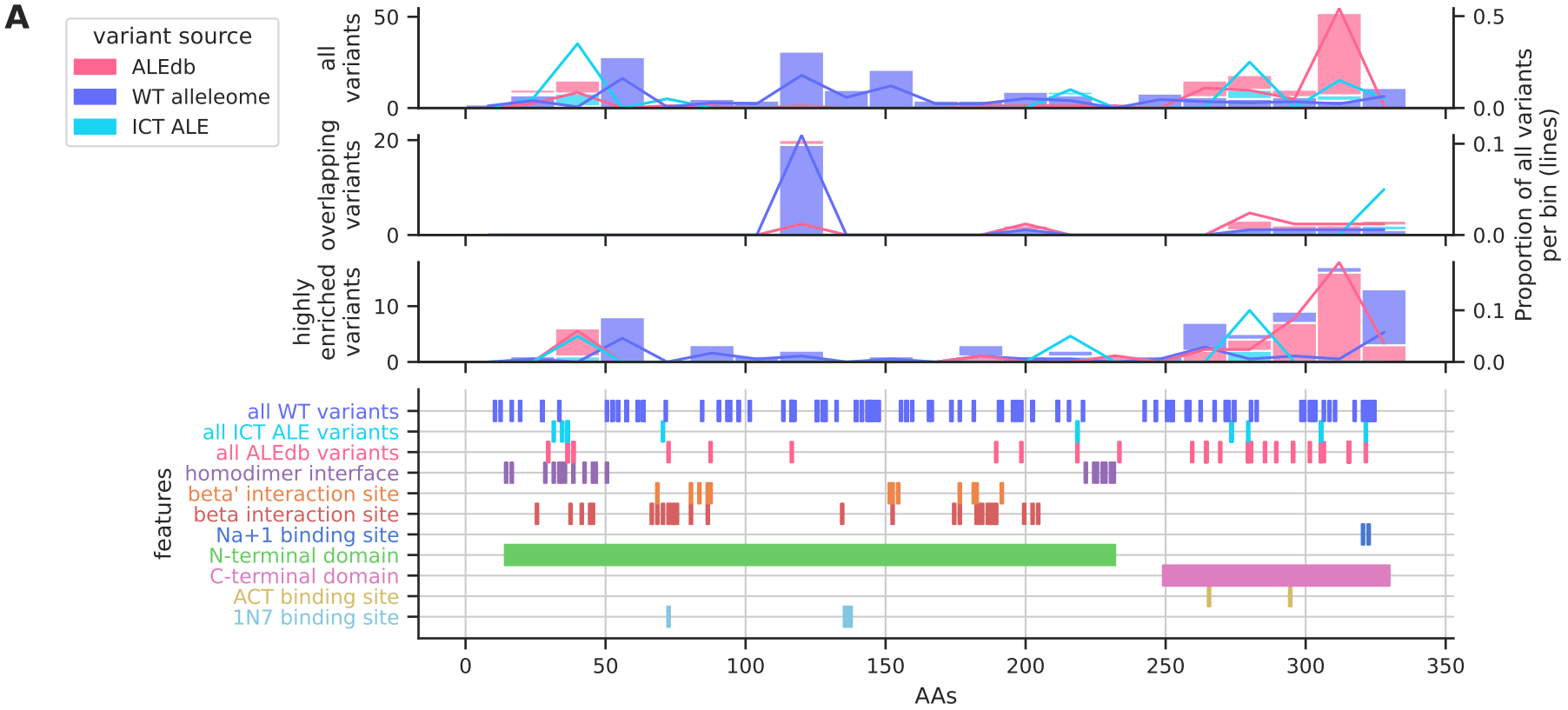
A
variant source
ALEdb
WT alleleome
ICT ALE
all variants
overlapping variants
highly enriched variants
Proportion of all variants per bin (lines)
features
all WT variants
all ICT ALE variants
all ALEdb variants
homodimer interface
beta' interaction site
beta interaction site
Na+1 binding site
N-terminal domain
C-terminal domain
ACT binding site
1N7 binding site
AAs