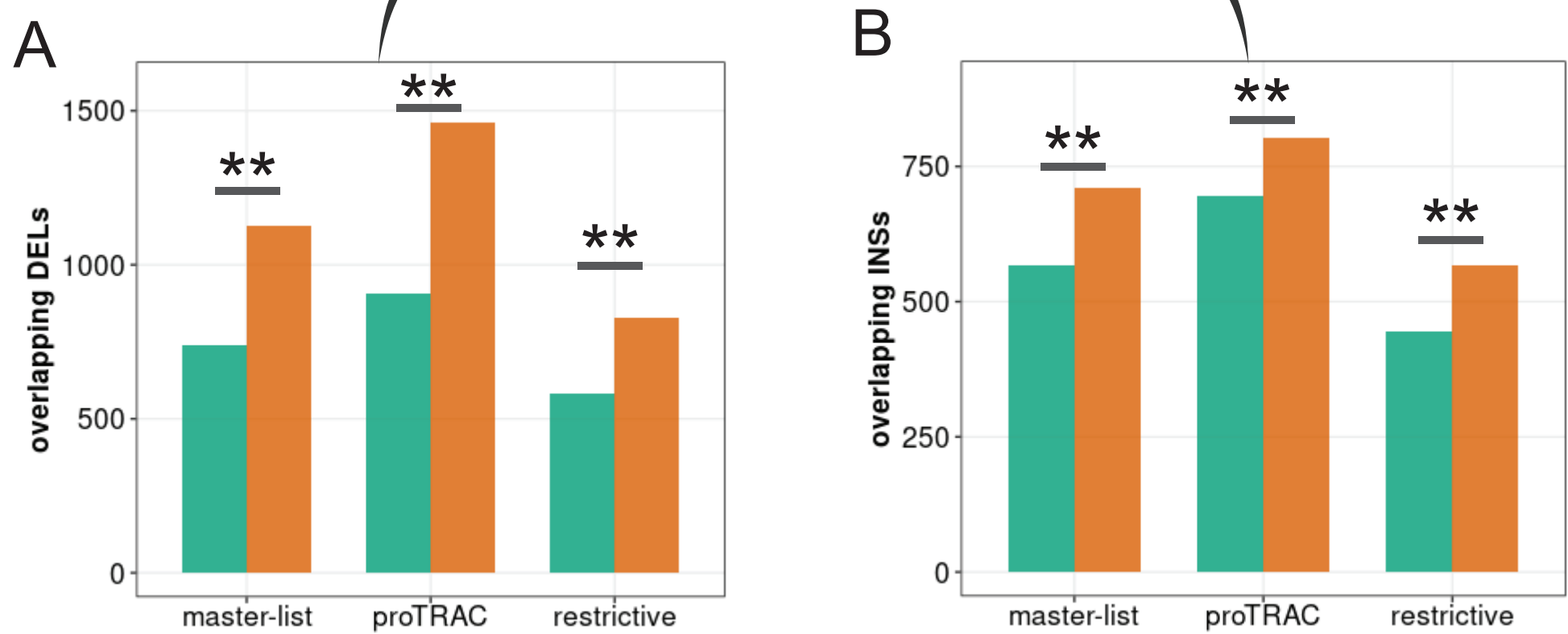
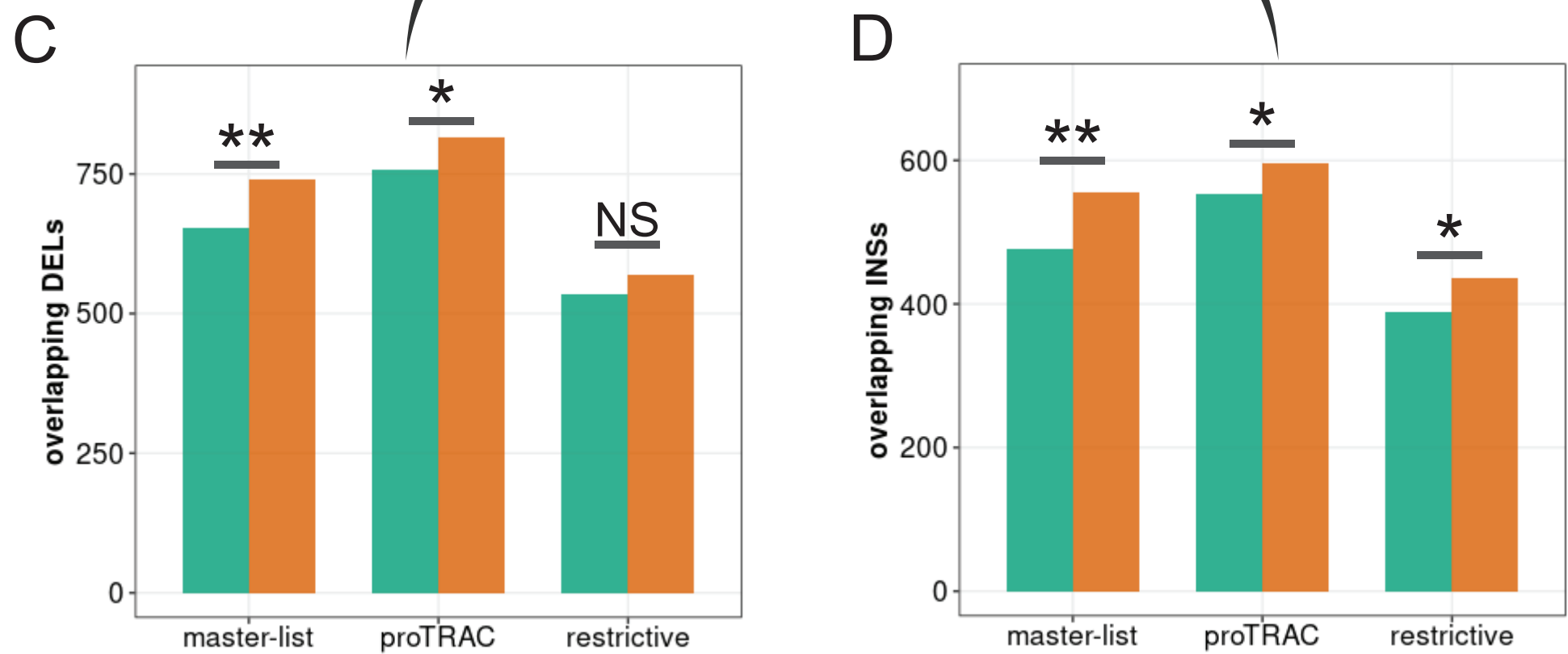


Supplementary Fig S7

whole genome



heterochromatin only
(Riddle et al, 2010)



mean_expected observed

Supplementary_Fig_S7. Comparisons of INDEL enrichment across whole genome and heterochromatin only to the piC regions from master-list. (A&B) Bar plots of observed and mean expected values of INDELs overlapping piCs and when piCs are randomly shuffled across whole genome. (C&D) Bar plots displaying observed and mean expected values of INDELs overlapping piCs and when piCs are shuffled only within heterochromatin boundaries. (p-value <0.05=*, <0.005= **)