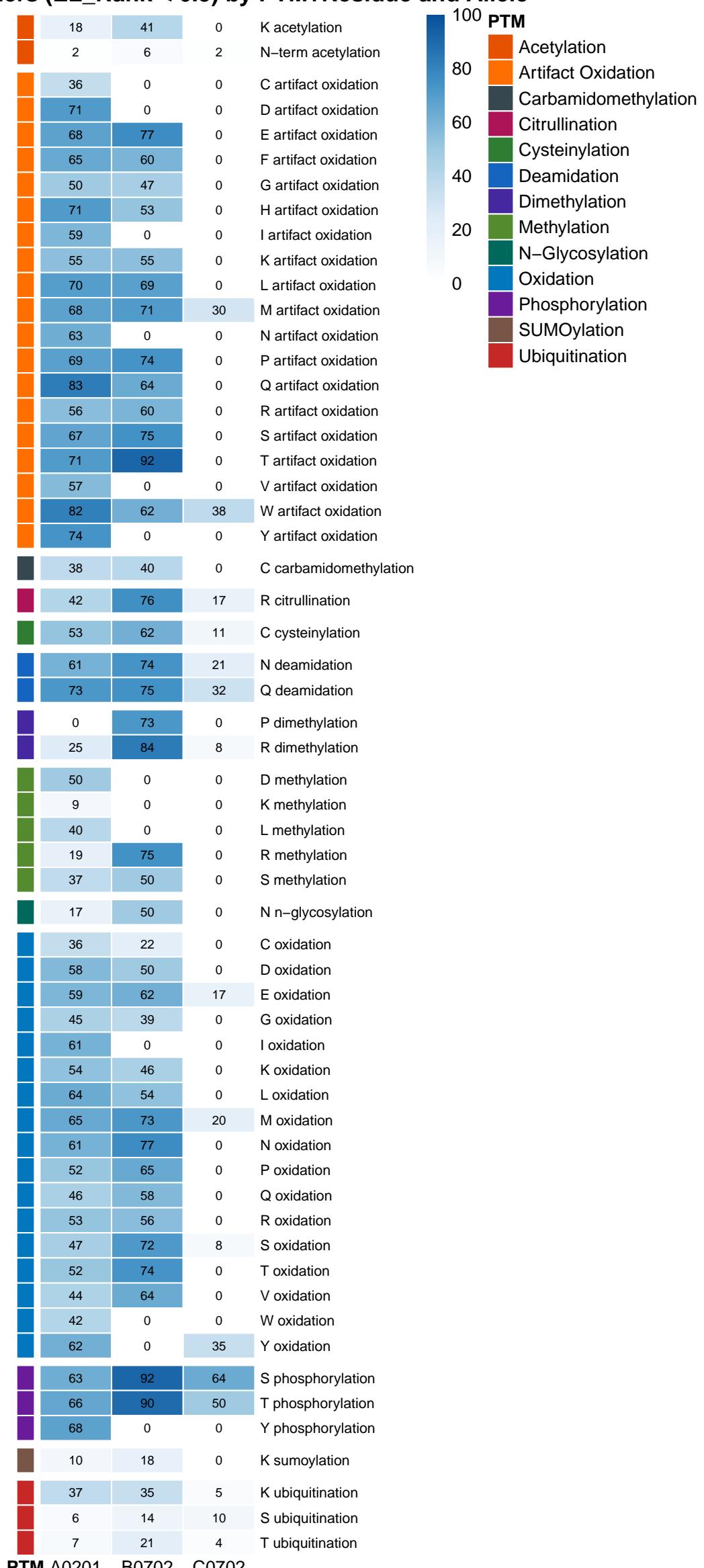


% Strong Binders (EL_Rank < 0.5) by PTM+Residue and Allele



PTM A0201 B0702 C0702