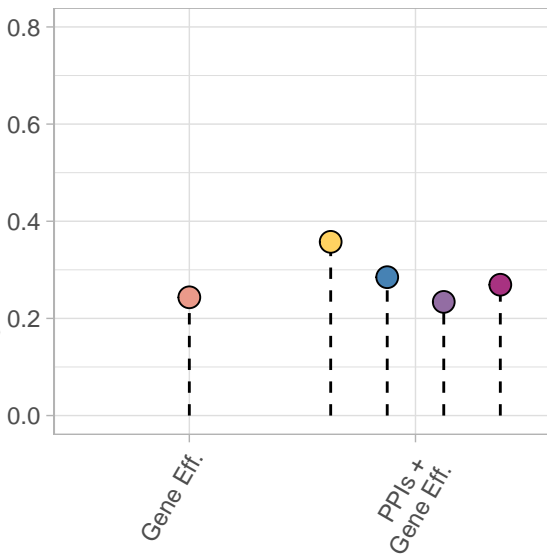
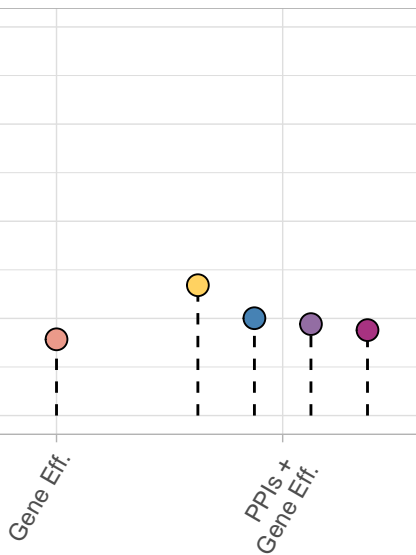


Cor. between Essentiality Scores  
and Phylogenetic Conservation Scores

sgRNA.input.dataset



shRNA.input.dataset



method    PRODE    MUFINN (Sum)    MUFINN (Max)    RWR    Average Gene Eff.