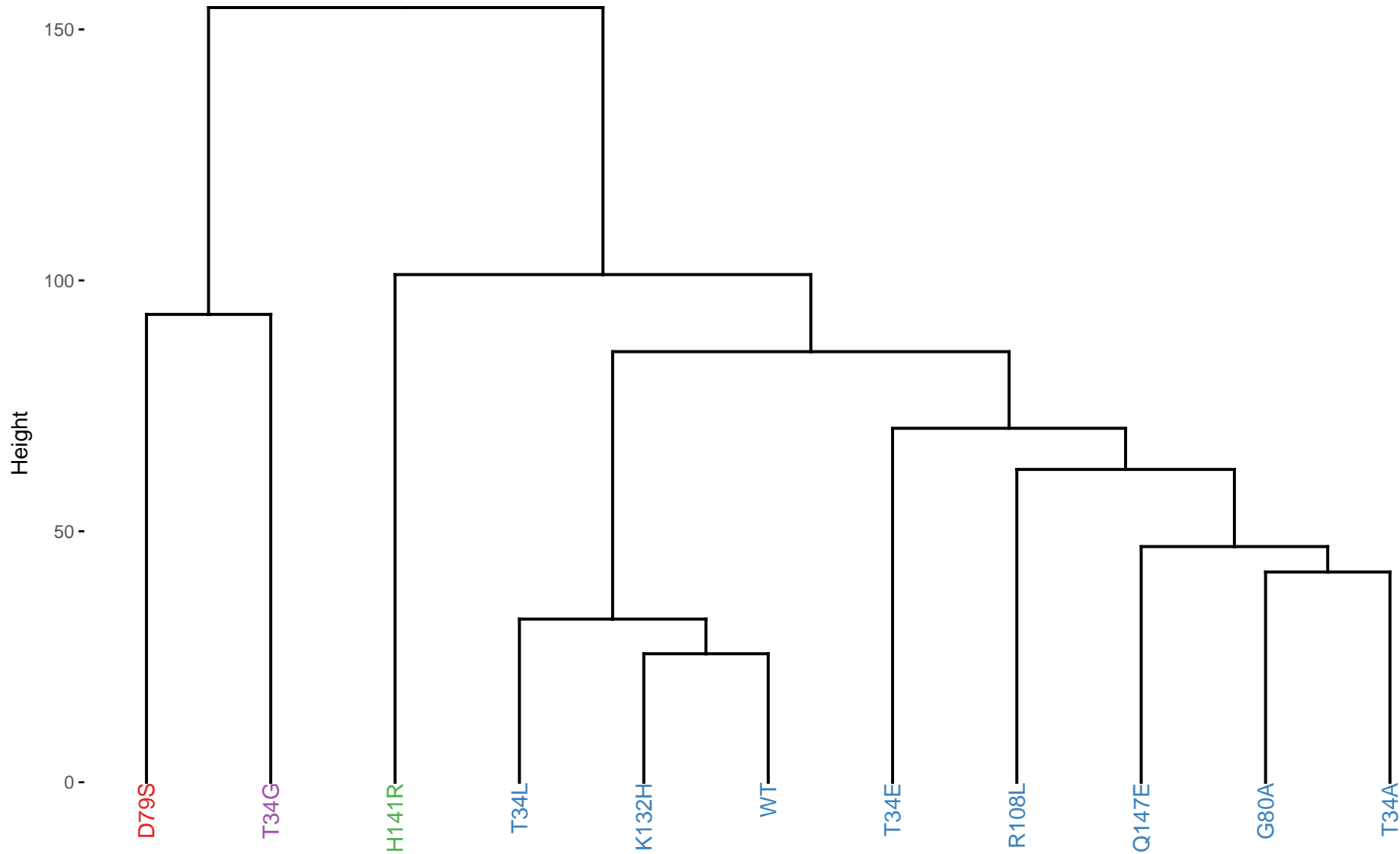


mutants clustered by E-MAP score, k = 4



mutants clustered by: (GAP_kcat GAP_Km GEF_kcat GEF_Km gamma2 SRM1 RNA1) -> k = 4

