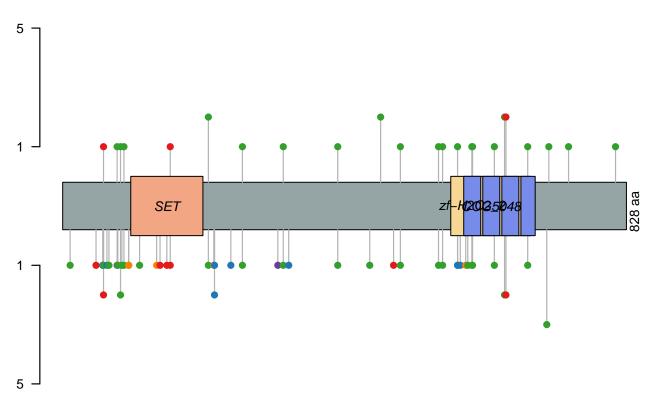
PRDM1: NM_001198



Reanalysis [4.91%; N = 958]

- Nonsense_MutationFrame_Shift_Del
- Missense_MutationSplice_Site
- Frame_Shift_Ins