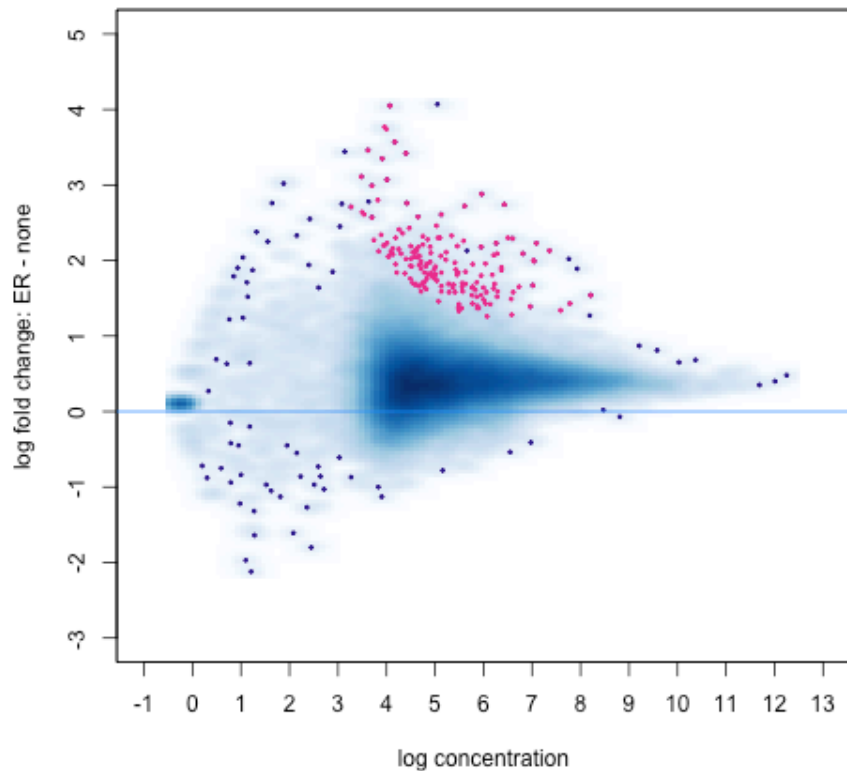


A**Binding Affinity: ER vs. none (166 FDR < 0.050)****B****VULCAN analysis of GRHL2 ChIP-Seq**