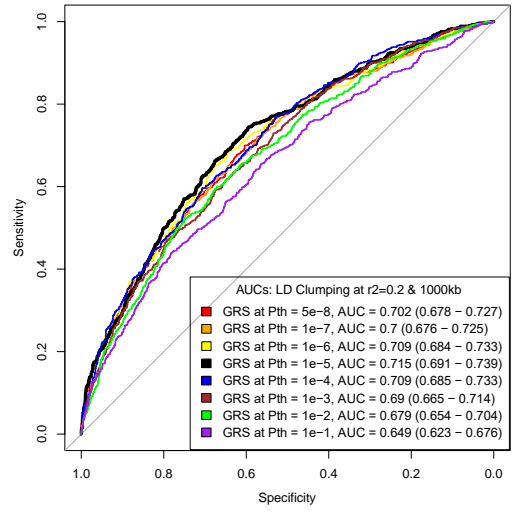
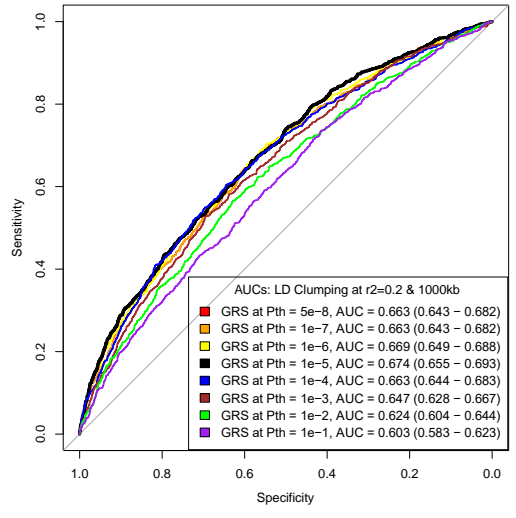


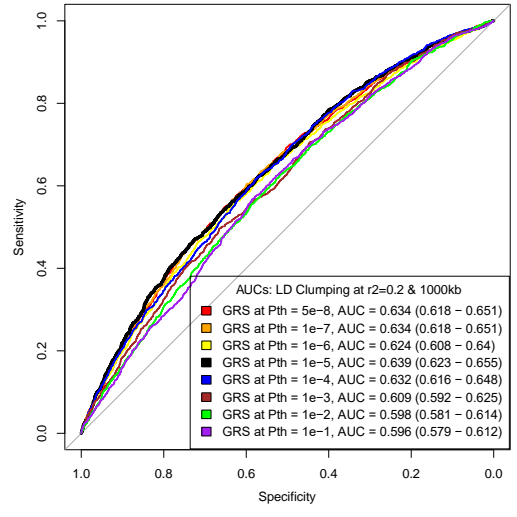
**A** SLE main cohort → SLEGEN cohort



**B** SLE main cohort → GENENTECH cohort



**C** EUR population → CHN population



**D** CHN population → EUR population

