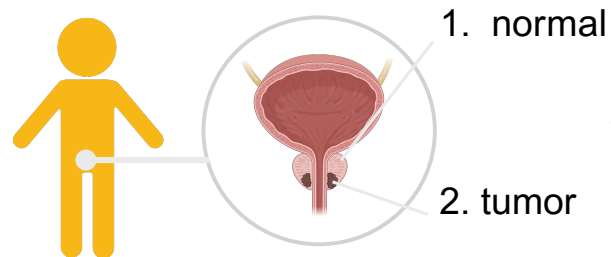
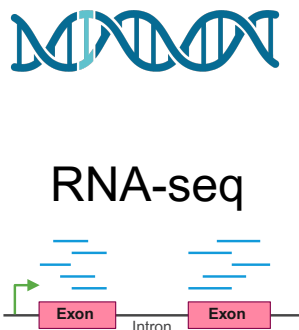


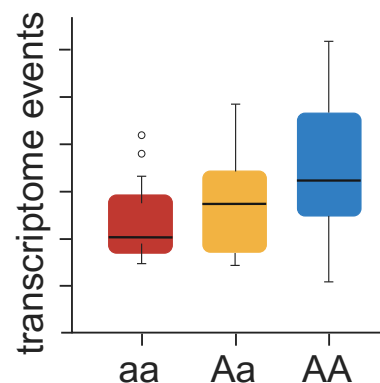
CPGEA (N=134)



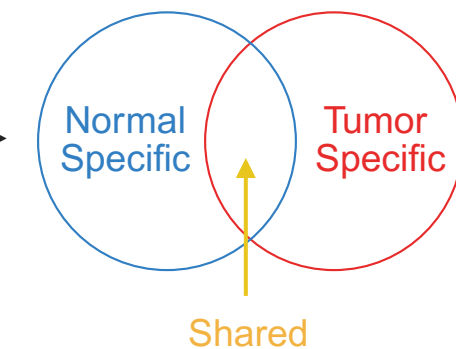
WGS



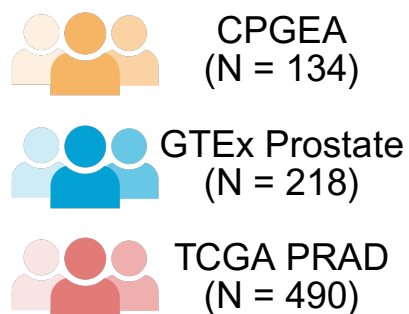
eQTL analysis



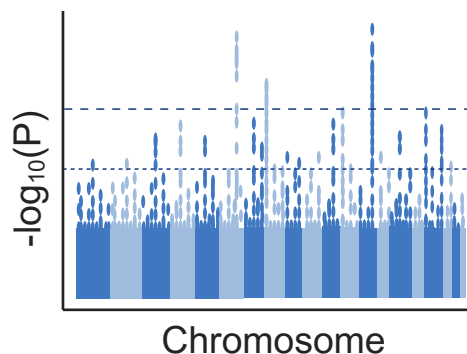
Comparison between normal and tumor



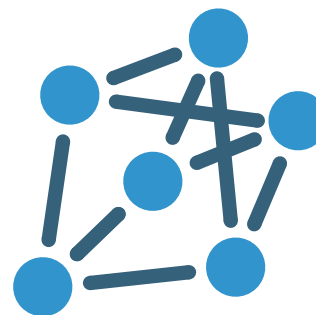
Multi-cohorts eQTL results



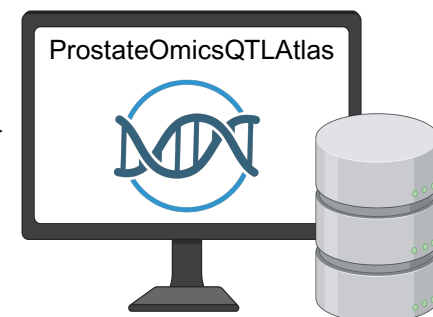
TWAS and colocalization analysis



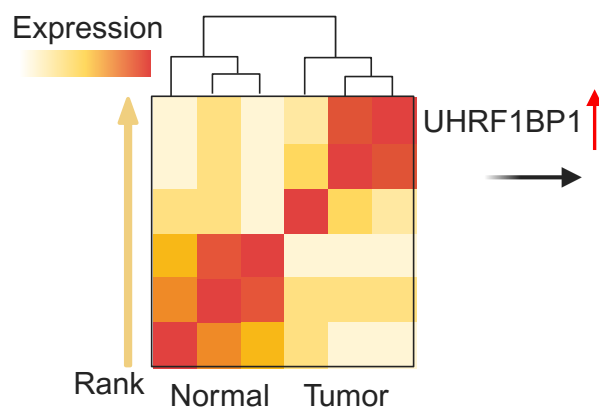
Risk genes network



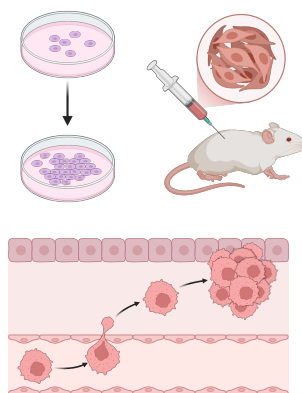
Database construction



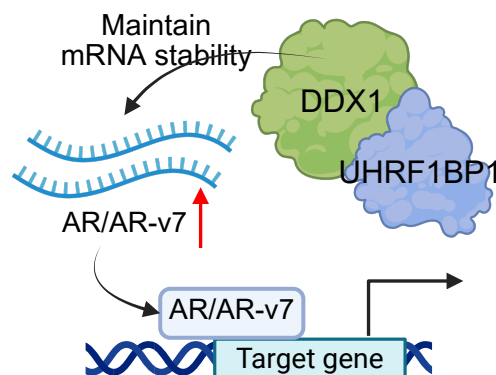
Key cross-population risk genes



Functional experiment



Functional mechanism



Clinical treatment

