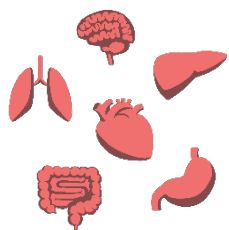


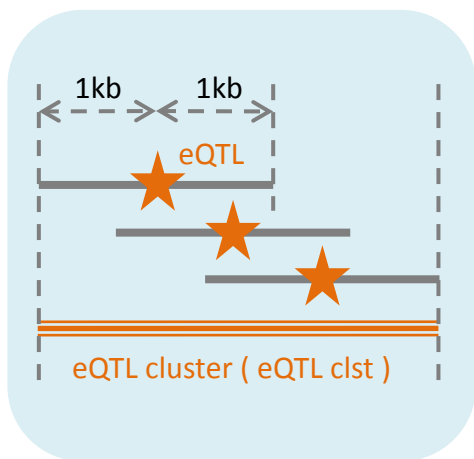
**A**

### GTEx eQTL & Single-cell Chromatin Accessibility Atlas

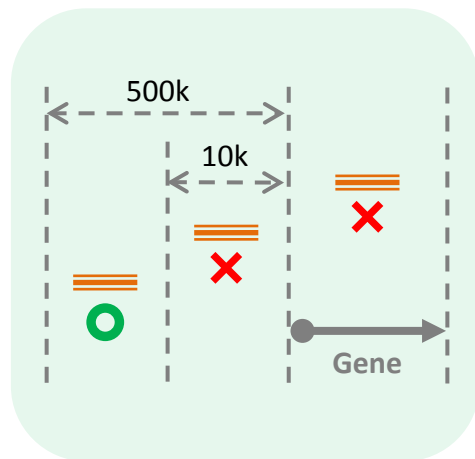


26 Tissues  
158 Cell Types

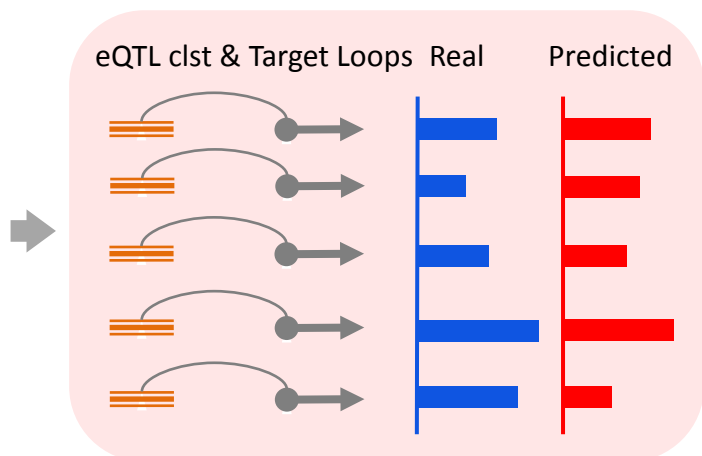
### Definition of eQTL Cluster



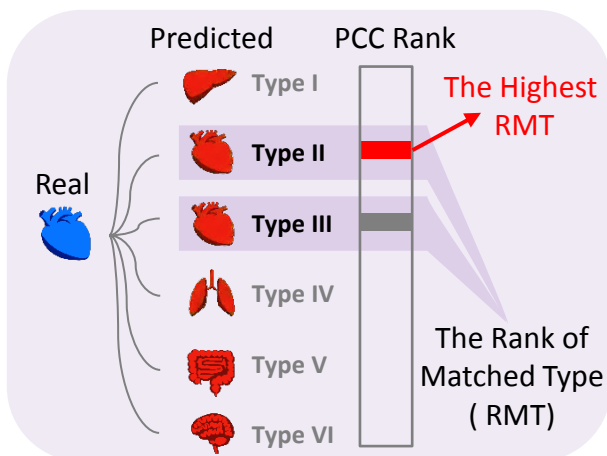
### Filtering of eQTL Cluster



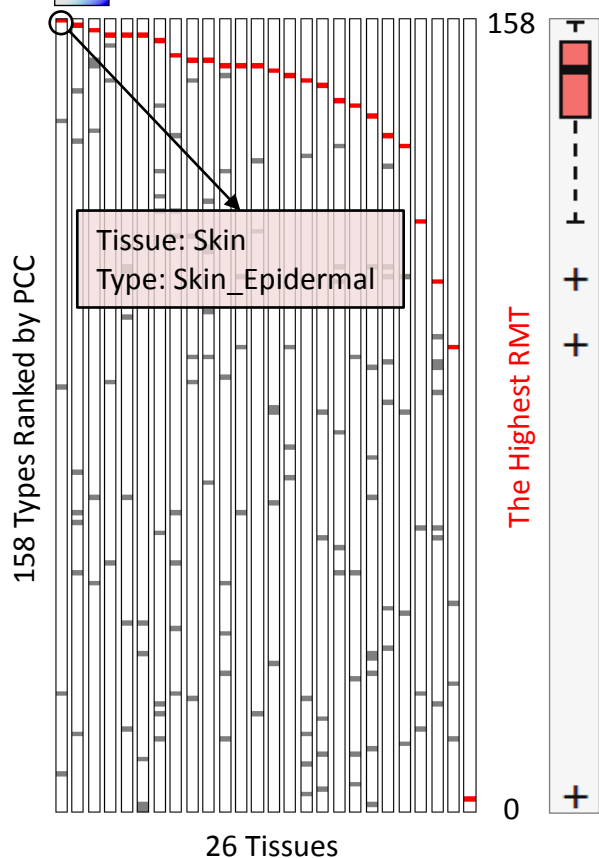
### Calculation of Real & Predicted Signal



### Comparison between Real & Predicted Signal

**B**

ILS ( $r=0$ )

**C**

The Highest RMT

