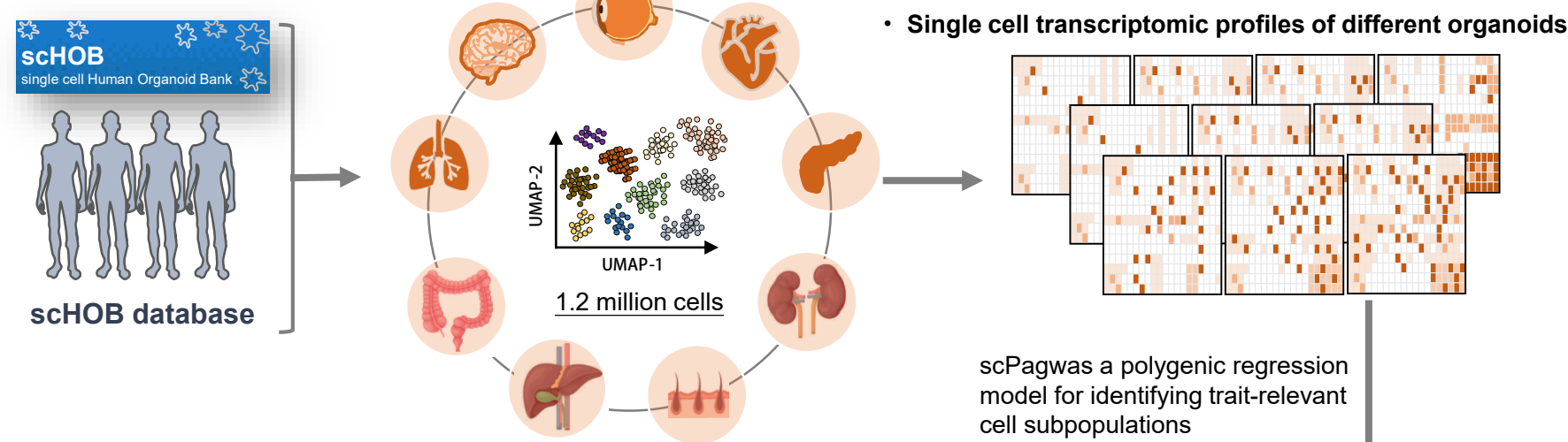
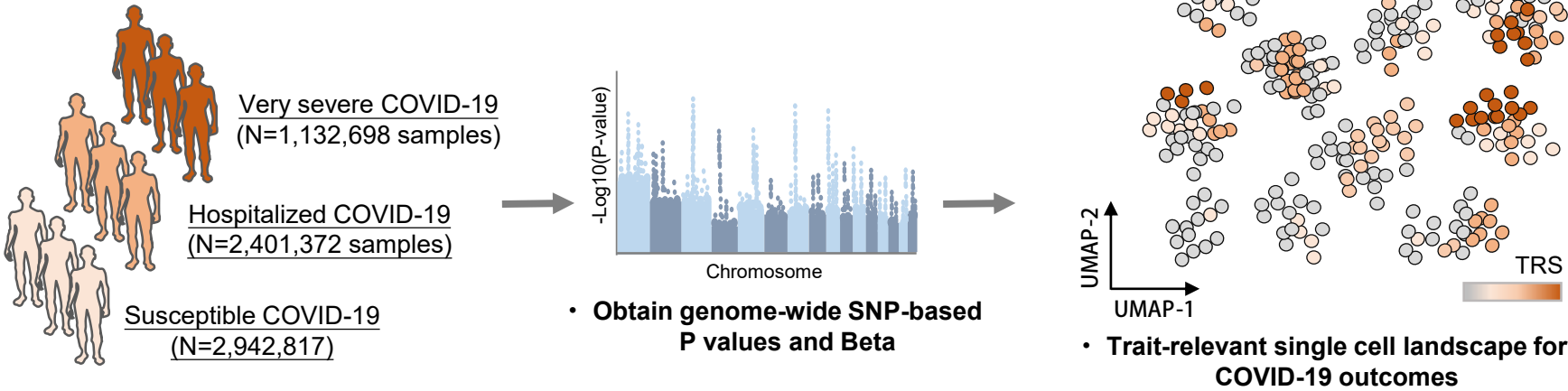


A

• Human organoids scRNA-seq datasets

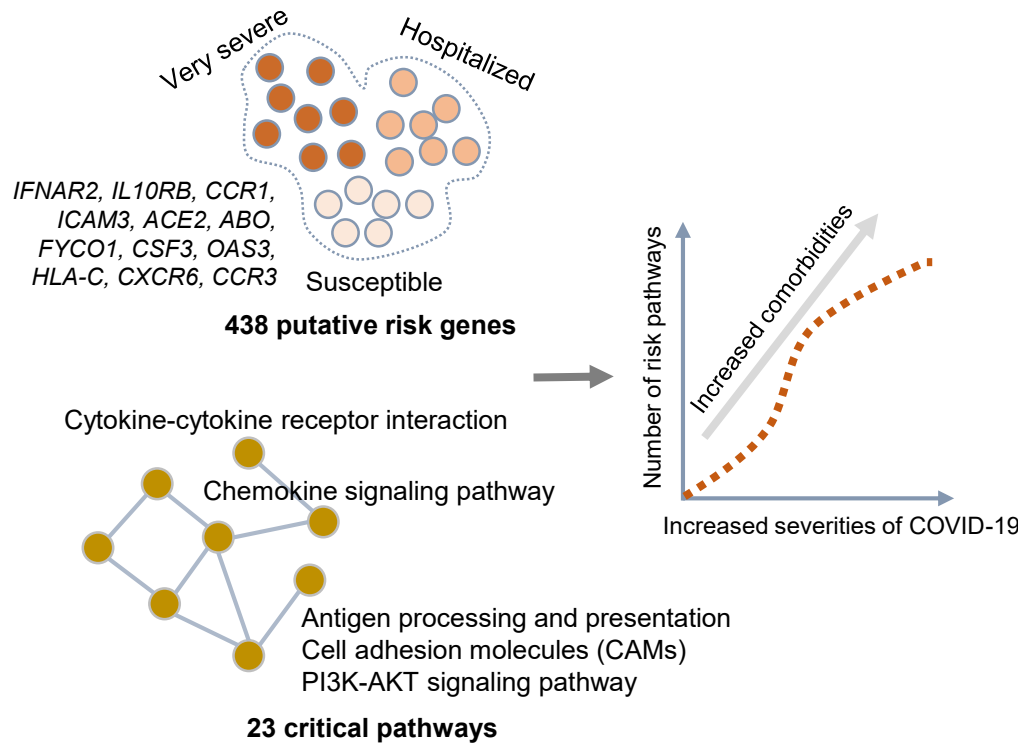


• GWAS summary statistics on COVID-19 outcomes



B

• Identifying 438 putative risk genes for COVID-19 severities



C

• Prioritizing druggable genes and interacting drugs for treating COVID-19 and related complications

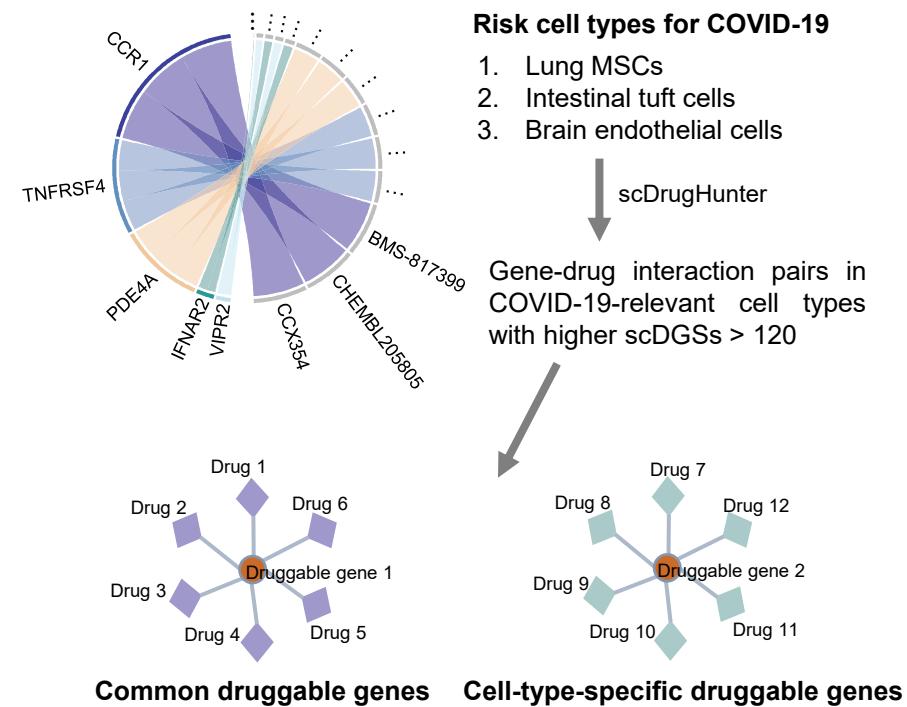


Figure 1