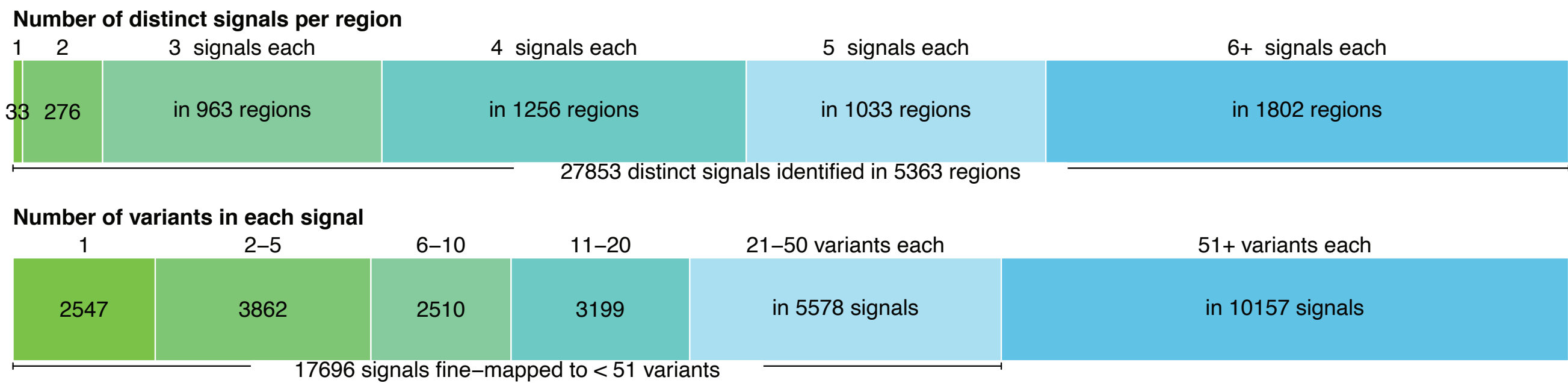


A. Summary of fine-mapping across all biomarkers



B. Number of fine-mapped variants per trait C. Allelic series at SLCO1B1/SLCO1B3 D. Pleiotropy at HGFAC in common and rare variants

