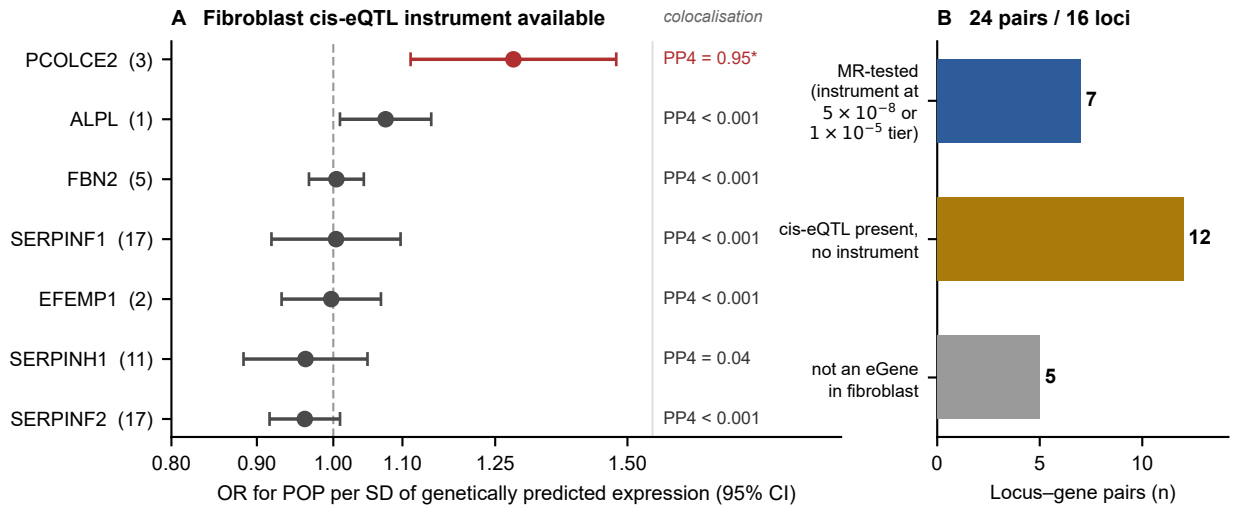


** BH-adjusted p < 0.05. PP4 = posterior probability of a shared causal variant (coloc.abf, $p_{12} = 10^{-5}$). Every testable gene was single-instrument ($r^2 = 1.00$ among candidate instruments). The screen relies on fibroblast cis-eQTL instruments only and cannot assess effects mediated by non-fibroblast cell types (macrophages, smooth muscle, vascular cells).*