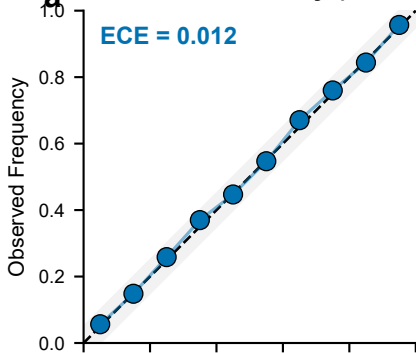
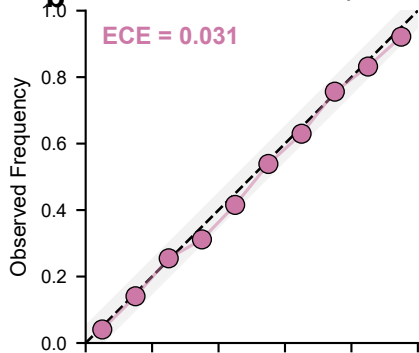


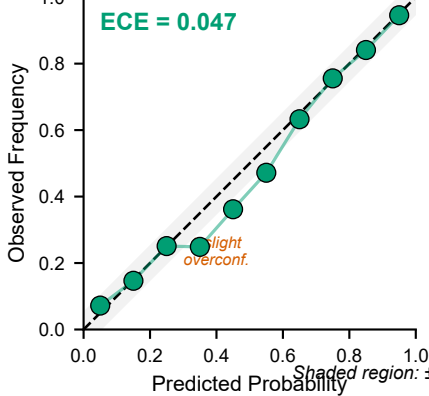
a Final Gene Probability (n=5,692)



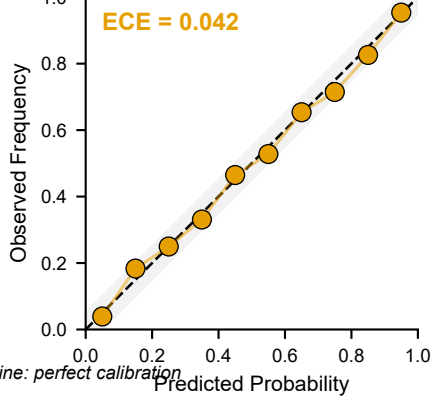
b Variant PIP from SuSiE (n=7,500)



c cCRE-Gene Linking (n=3,015)



d Gene-Tissue Colocalization (n=4,234)



Shaded region: ± 0.05 tolerance band | Dashed line: perfect calibration