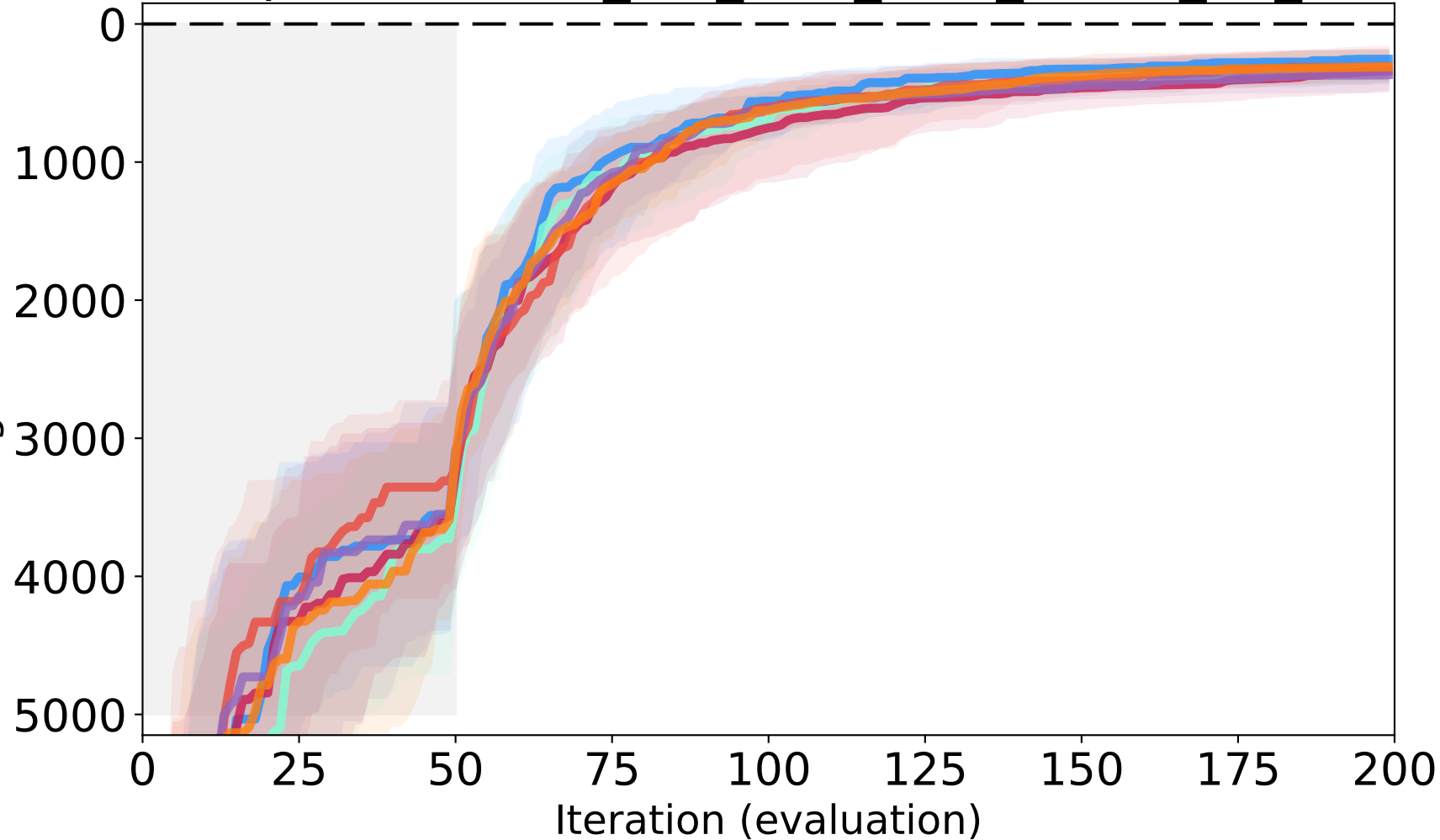


Comparison for 3_YRI_CEU_CHB_struct_25_Gut

Distance to optimum value
of log-likelihood



- LogEI + Matern52
- PI + RBF
- PI + Auto
- LogEI + RBF
- PI + Matern52
- LogEI + Auto
- Initial design