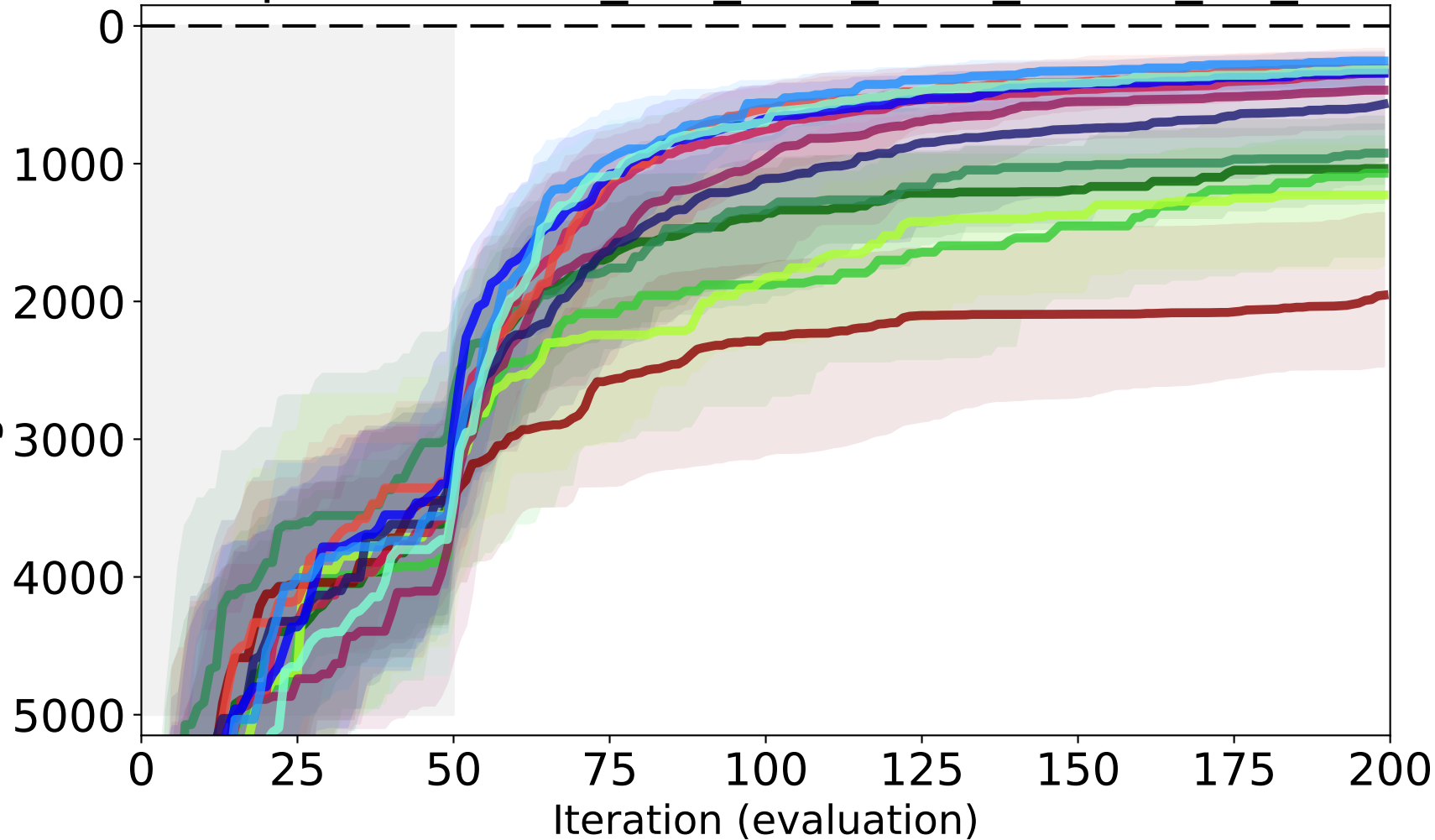


Comparison for 3_YRI_CEU_CHB_struct_25_Gut

Distance to optimum value
of log-likelihood



- LogEI + Matern52
- PI + RBF
- LogEI + RBF
- PI + Matern52
- LogEI + Matern32
- PI + Matern32
- LogEI + Exponential
- EI + Matern32
- EI + Exponential
- EI + Matern52
- EI + RBF
- PI + Exponential
- Initial design