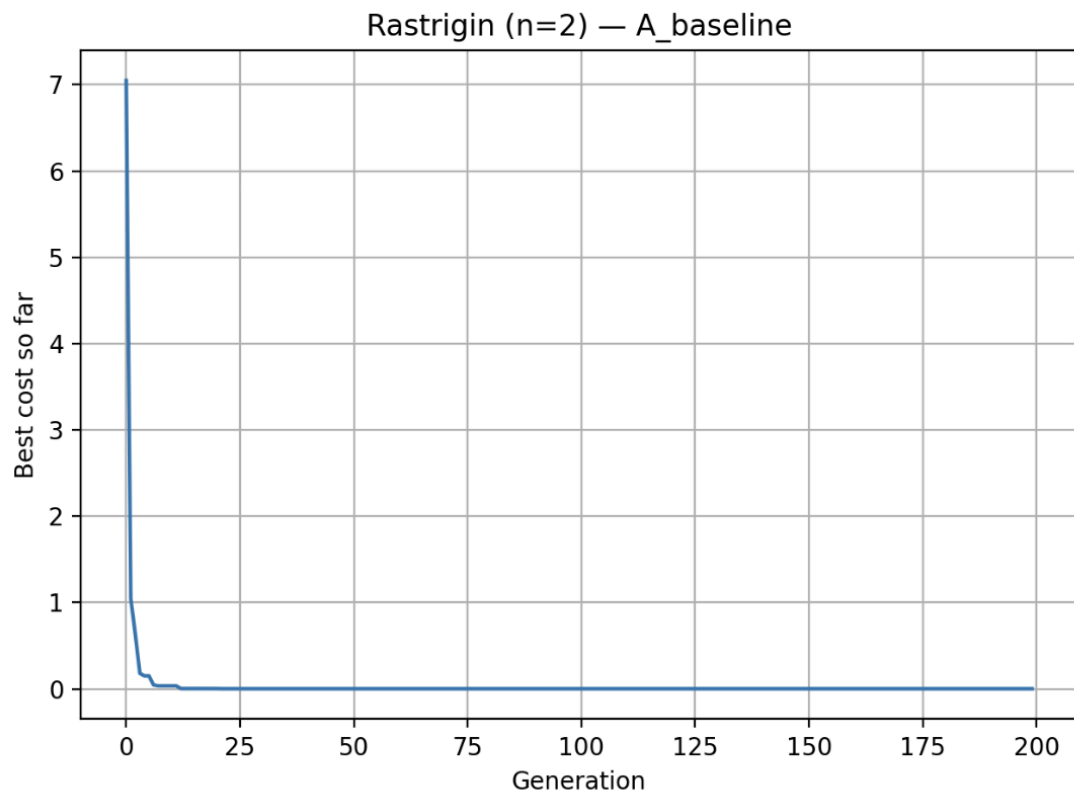
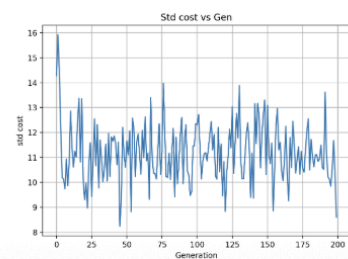
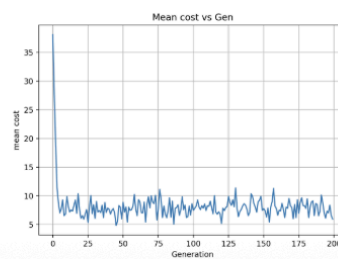
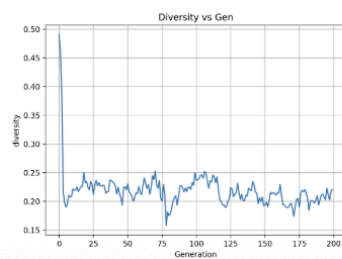


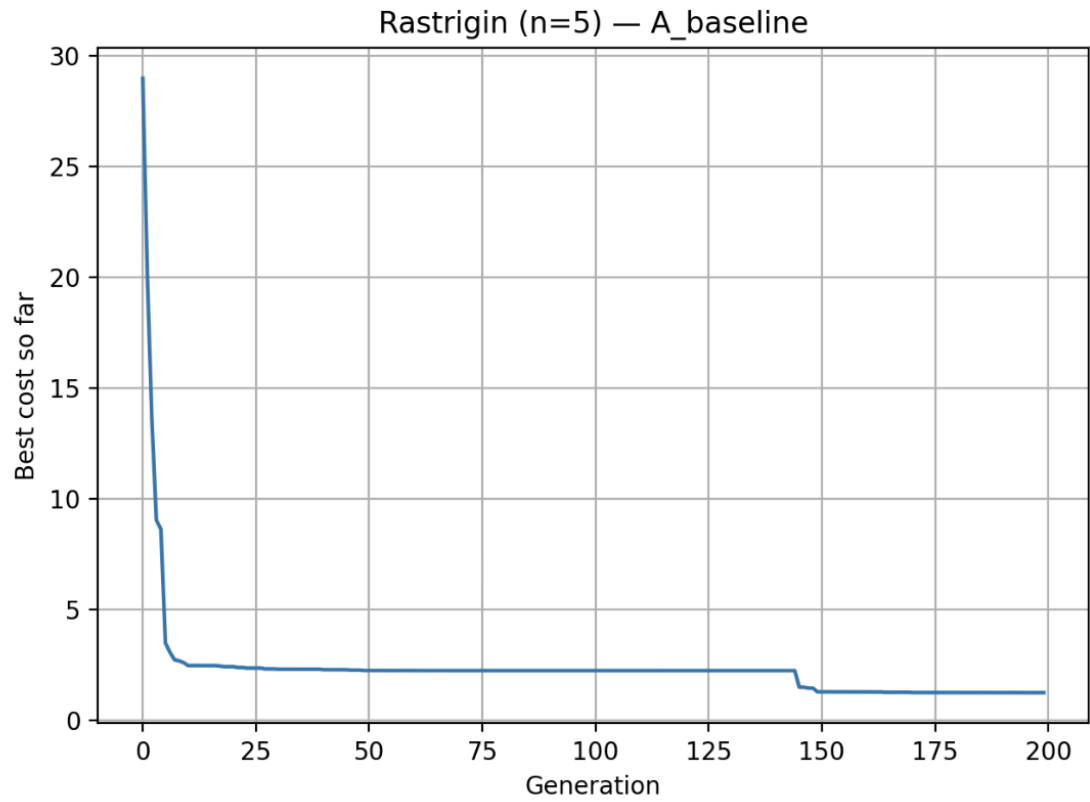
Acto 1 — Qué problema resolvemos y por qué GA

Single Run

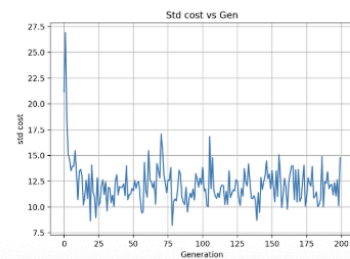
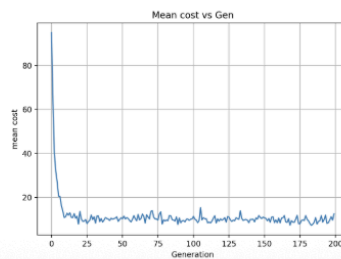
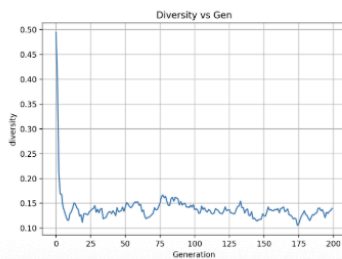


Population diagnostics

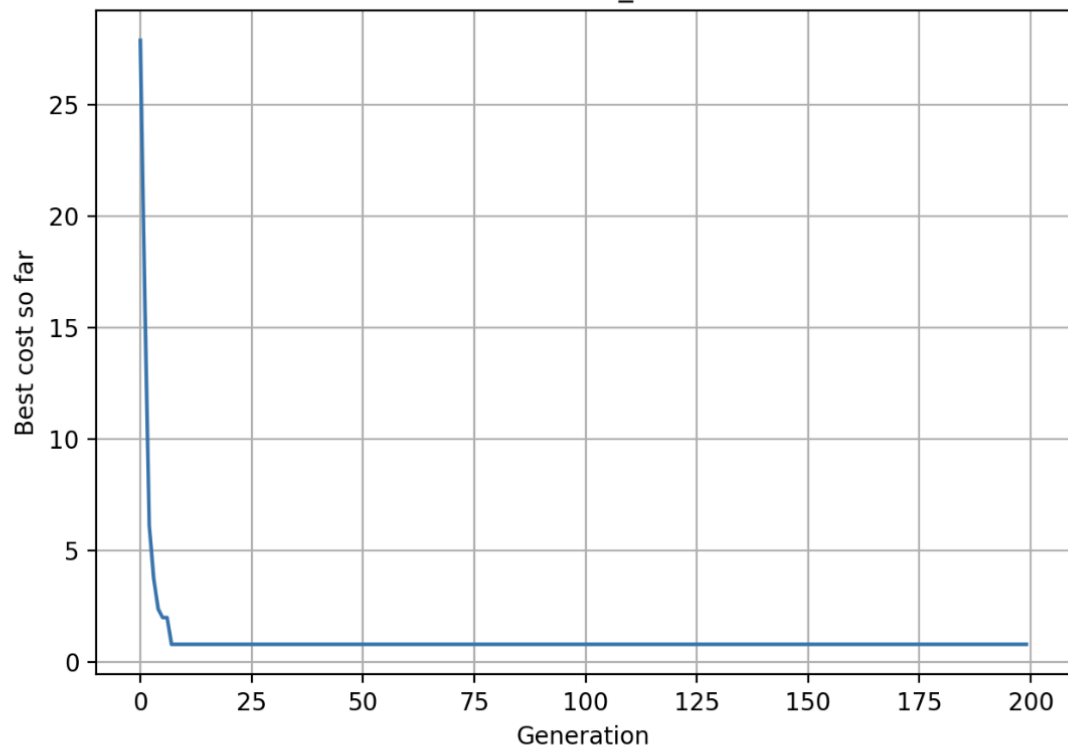




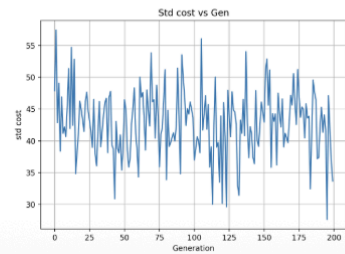
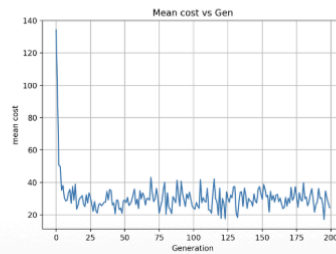
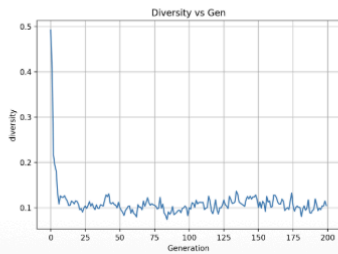
Population diagnostics

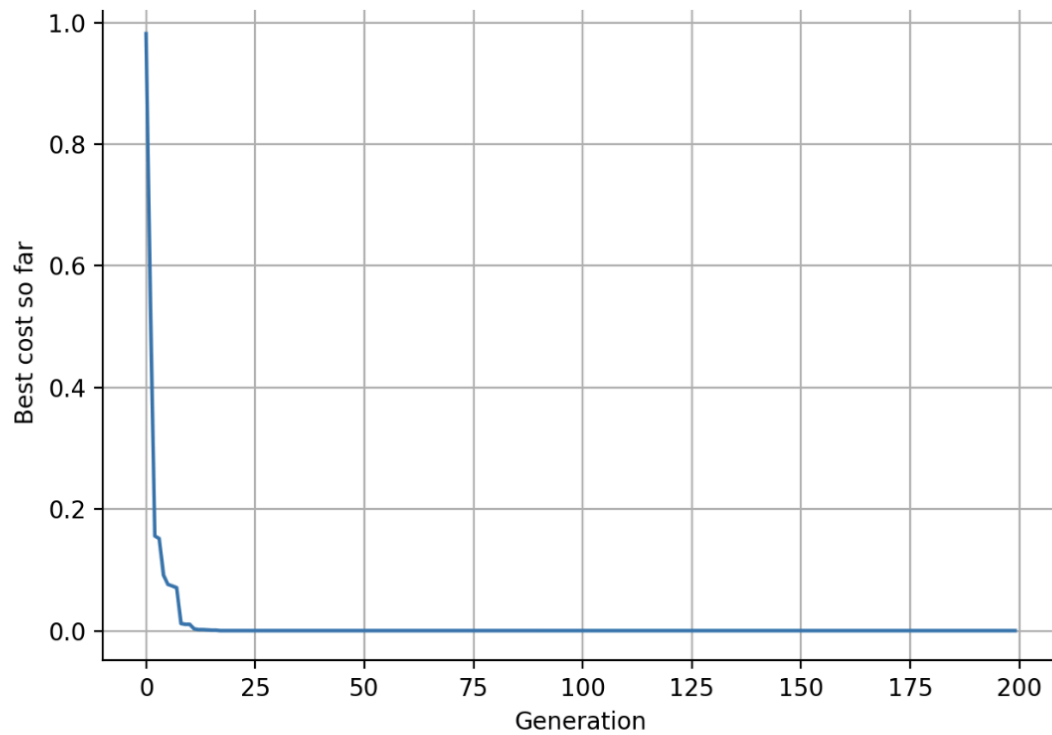


Bukin N.6 — A_baseline

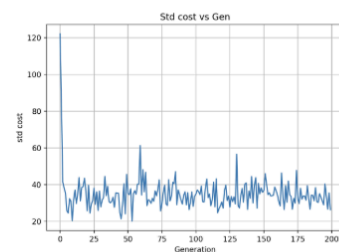
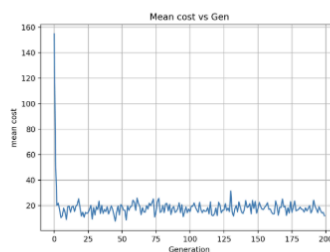
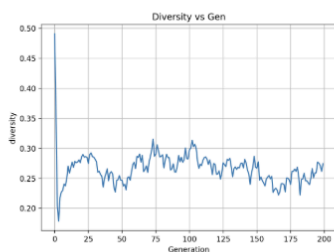


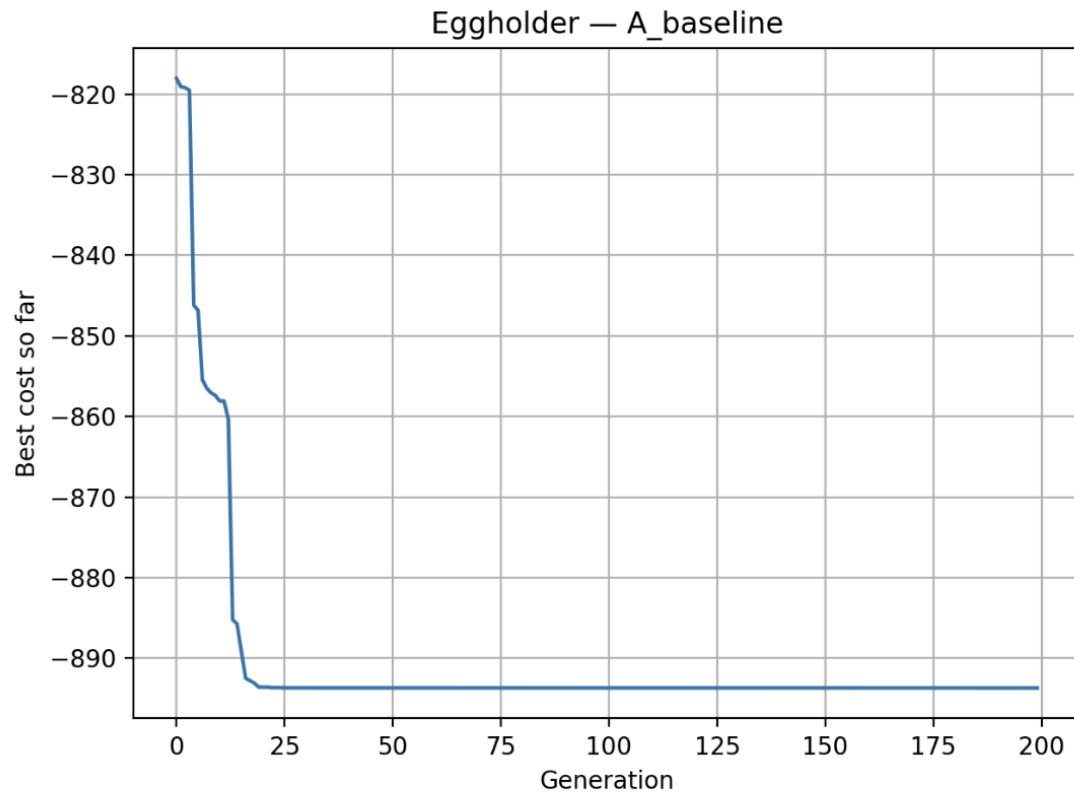
Population diagnostics



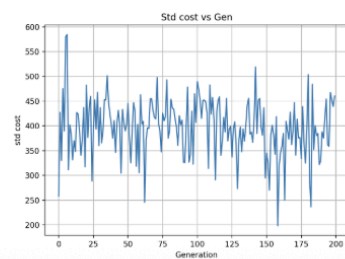
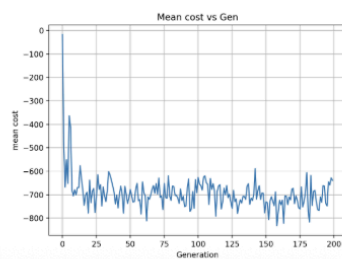
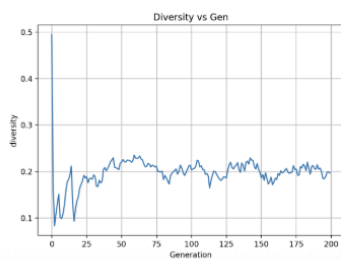


Population diagnostics



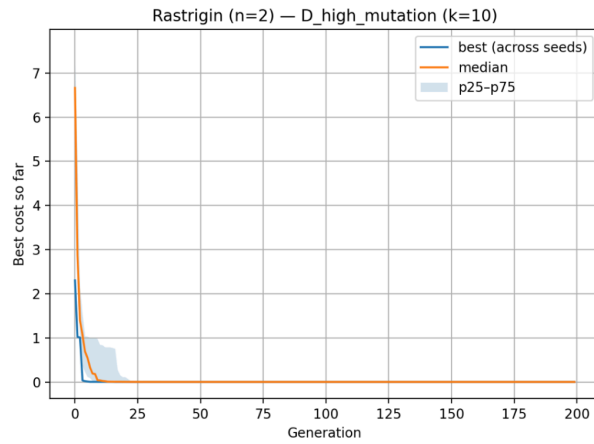


Population diagnostics

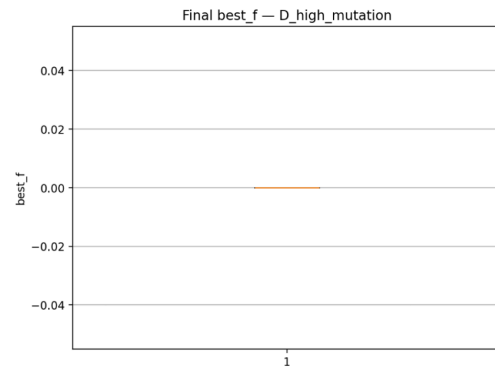


Mode: **Multi-seed (Scenario)**

Convergence bands (best-so-far across seeds)



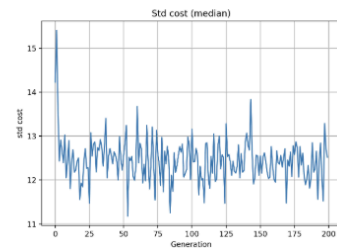
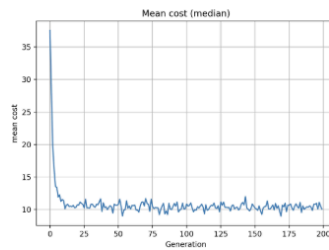
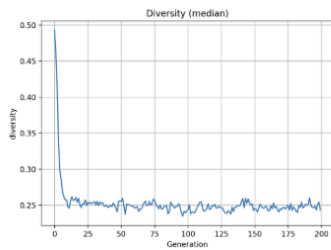
Final best_f distribution



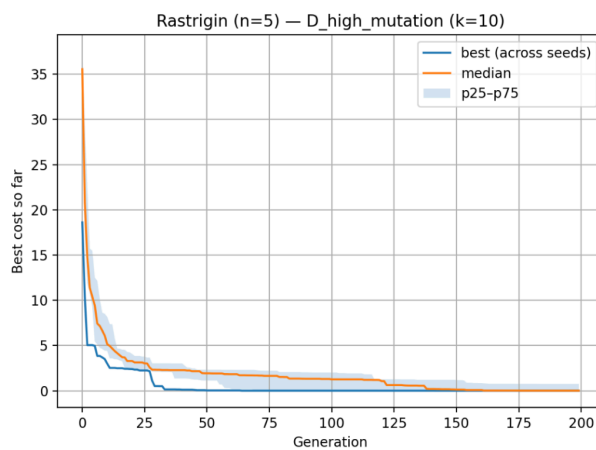
Robustness metrics

{

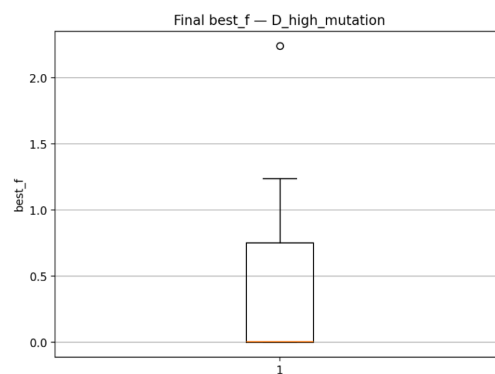
Population diagnostics (median across seeds)



Convergence bands (best-so-far across seeds)



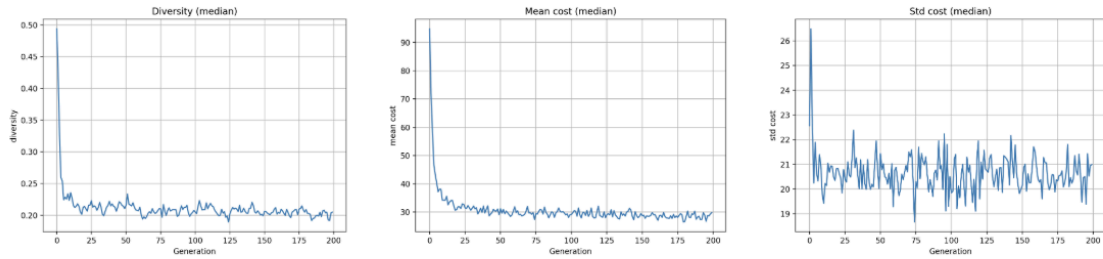
Final best_f distribution



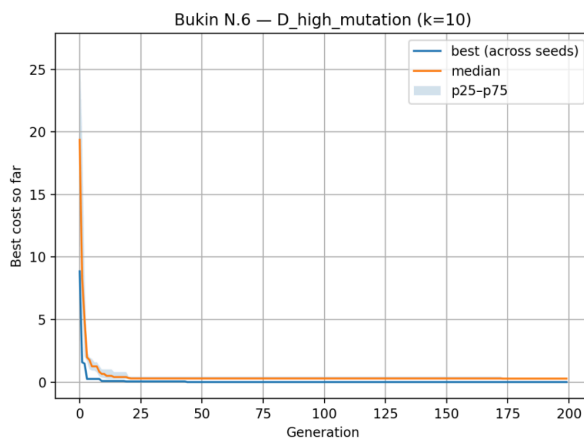
Robustness metrics

{

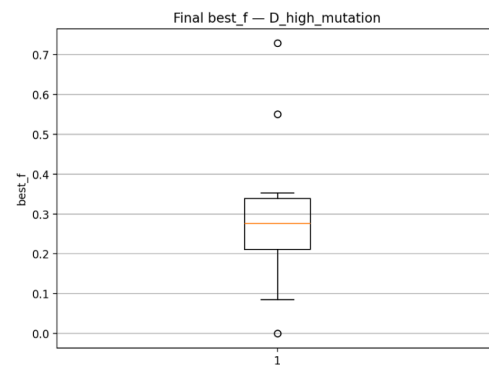
Population diagnostics (median across seeds)



Convergence bands (best-so-far across seeds)



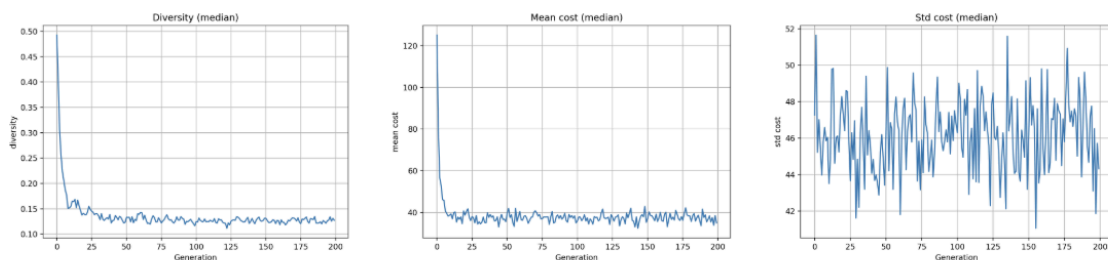
Final best_f distribution



Robustness metrics

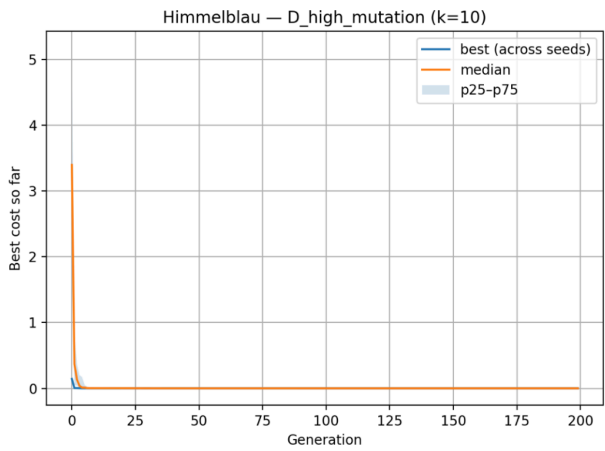
▼ {

Population diagnostics (median across seeds)

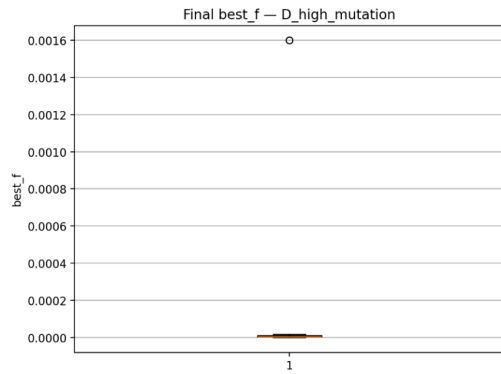




Convergence bands (best-so-far across seeds)



Final best_f distribution

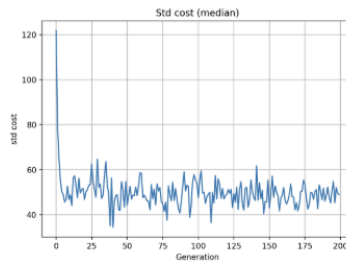
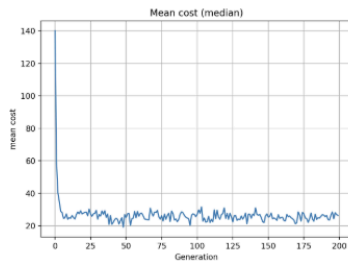
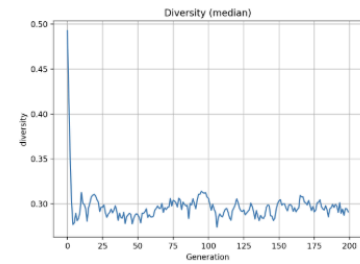


Robustness metrics

▼ {



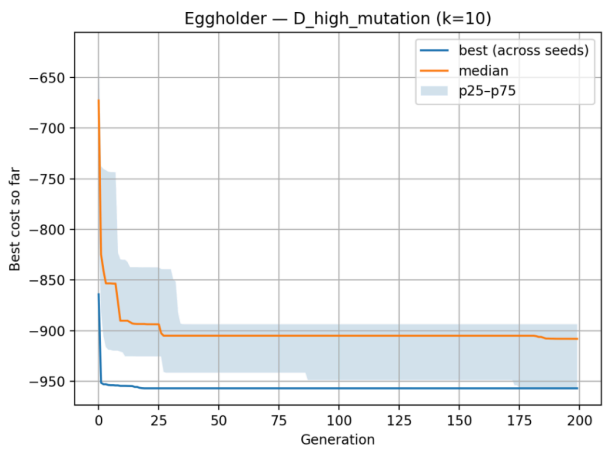
Population diagnostics (median across seeds)



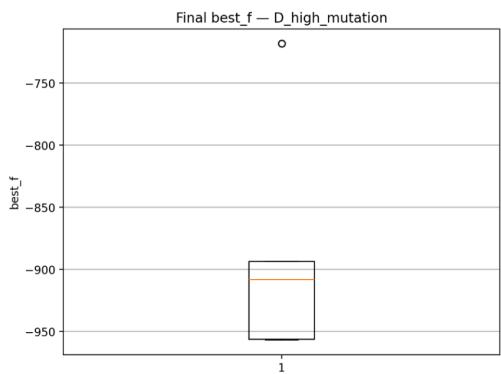
copy



Convergence bands (best-so-far across seeds)



Final best_f distribution

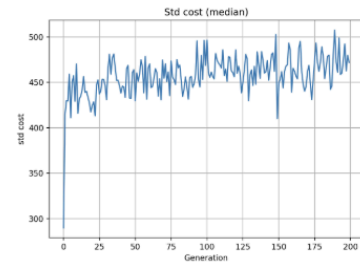
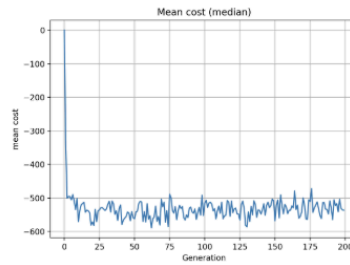
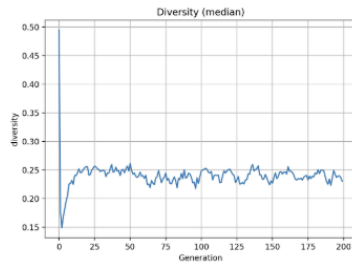


Robustness metrics

▼ {



Population diagnostics (median across seeds)

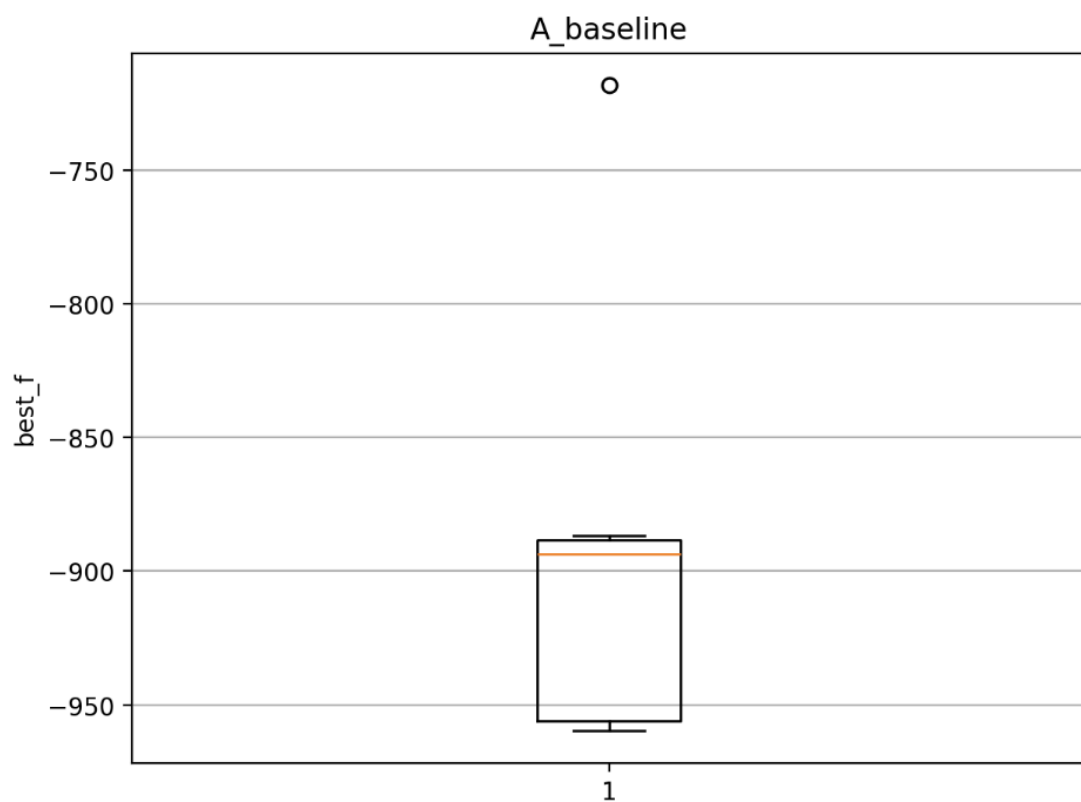


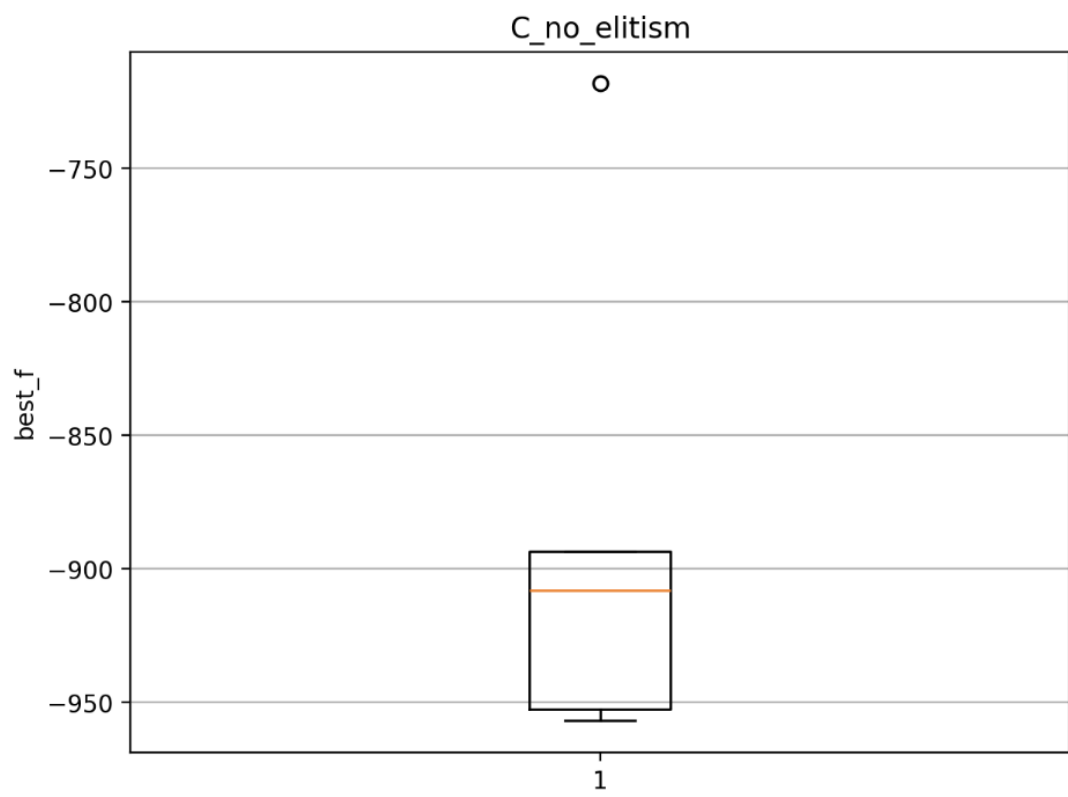
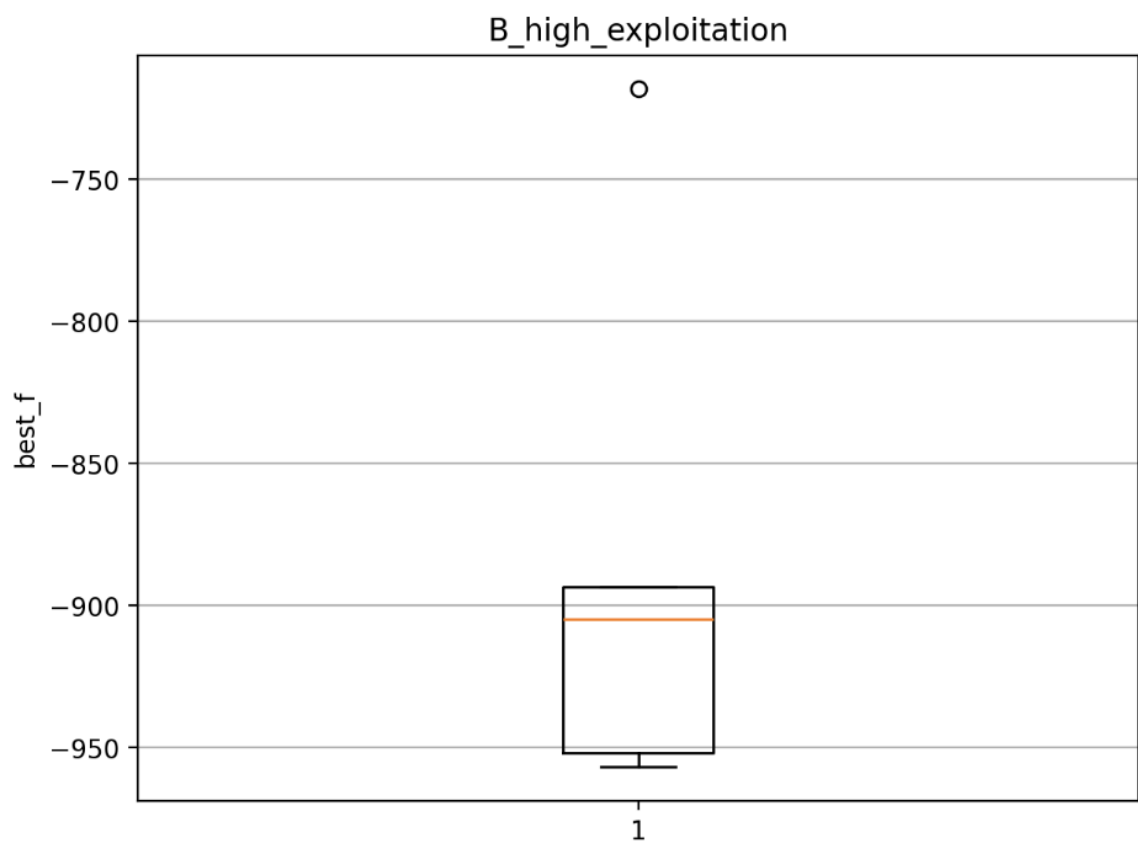
Mode: Compare Scenarios

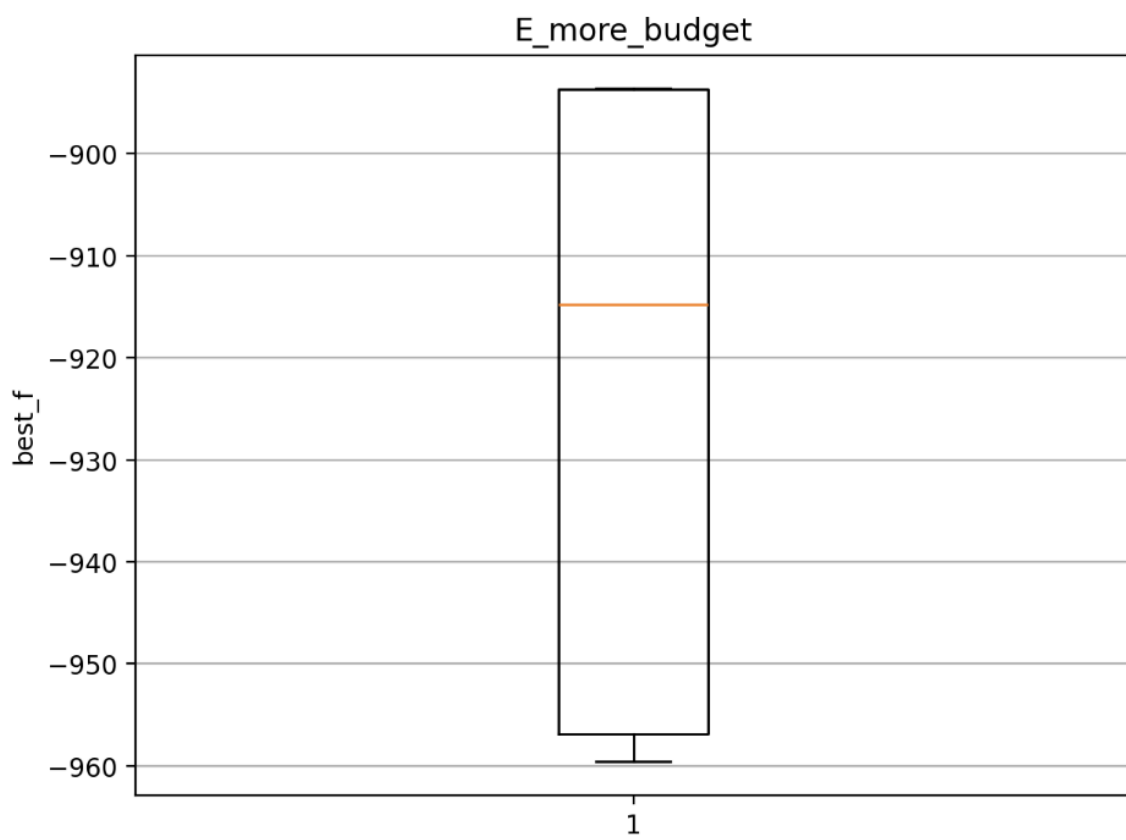
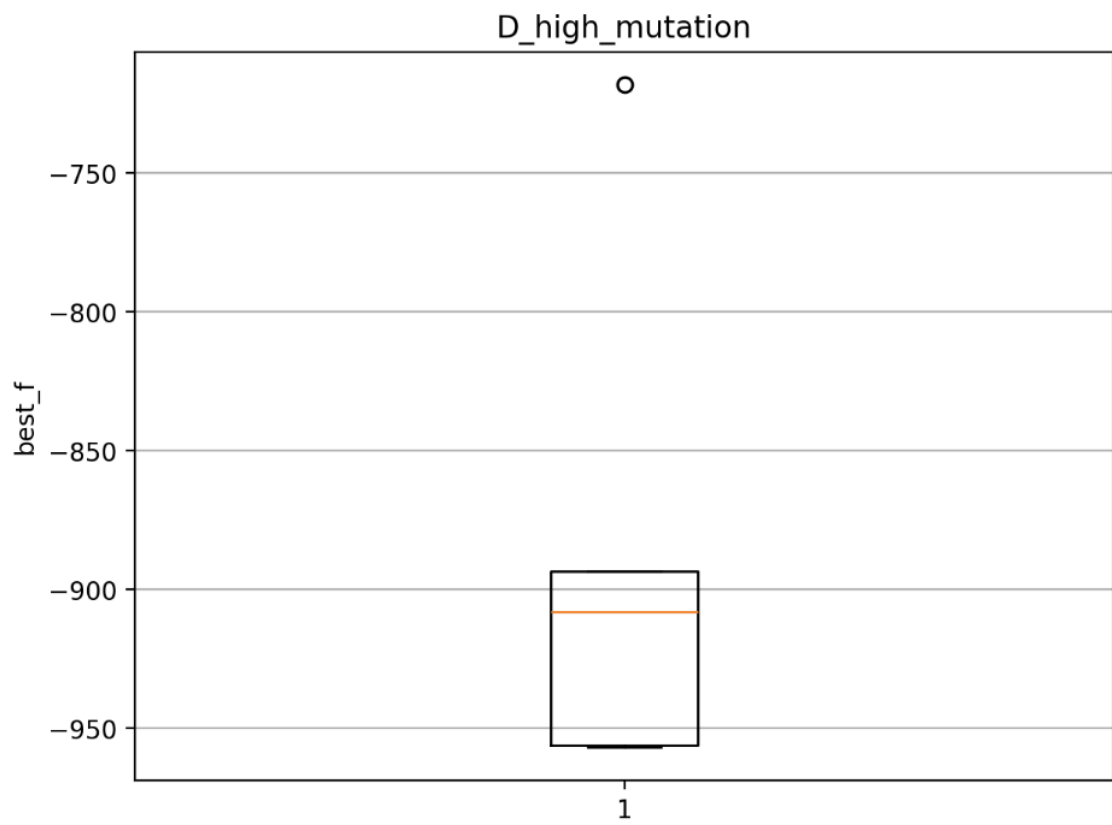
EGGHOLDER



Boxplots (final best_f) — one per scenario





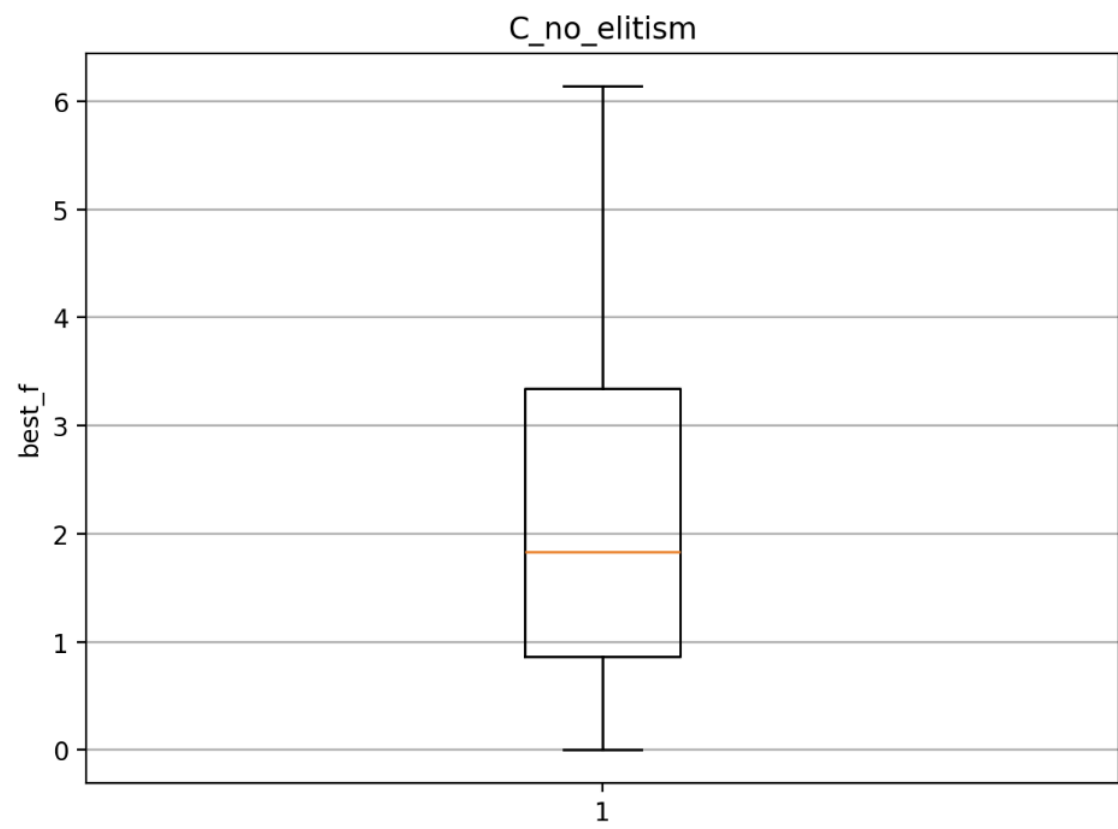
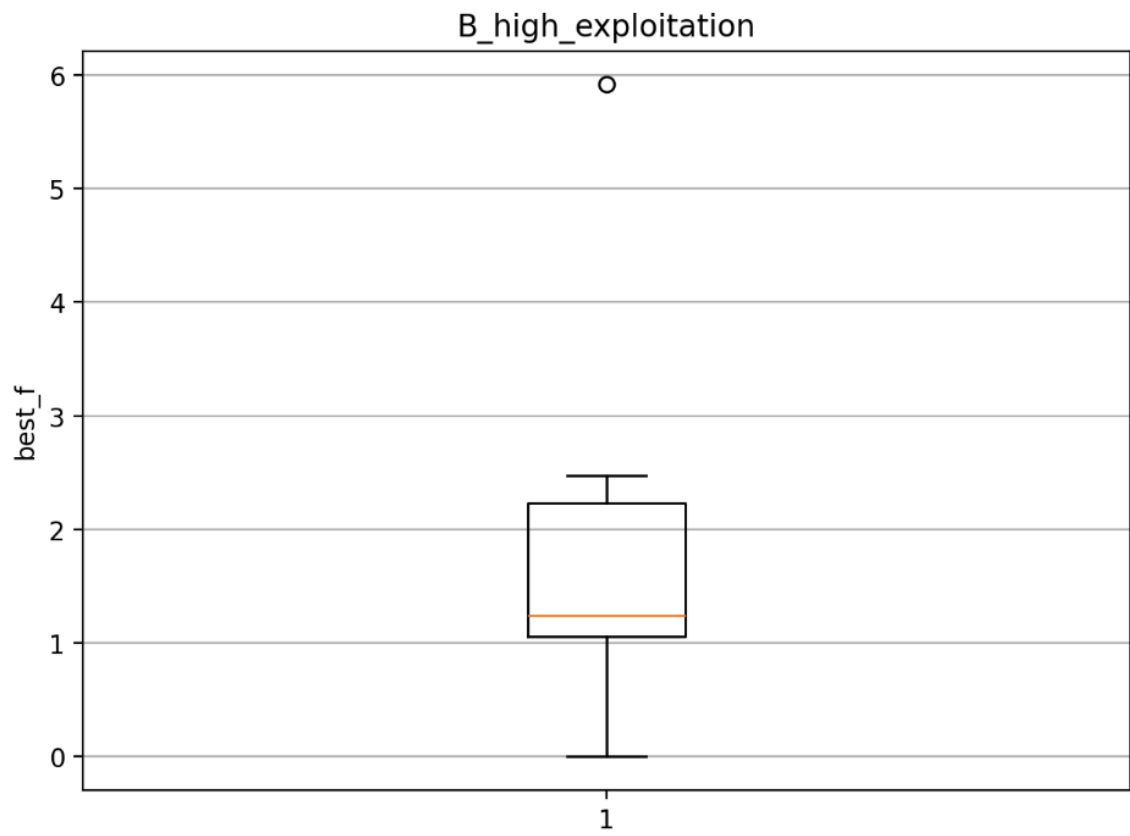


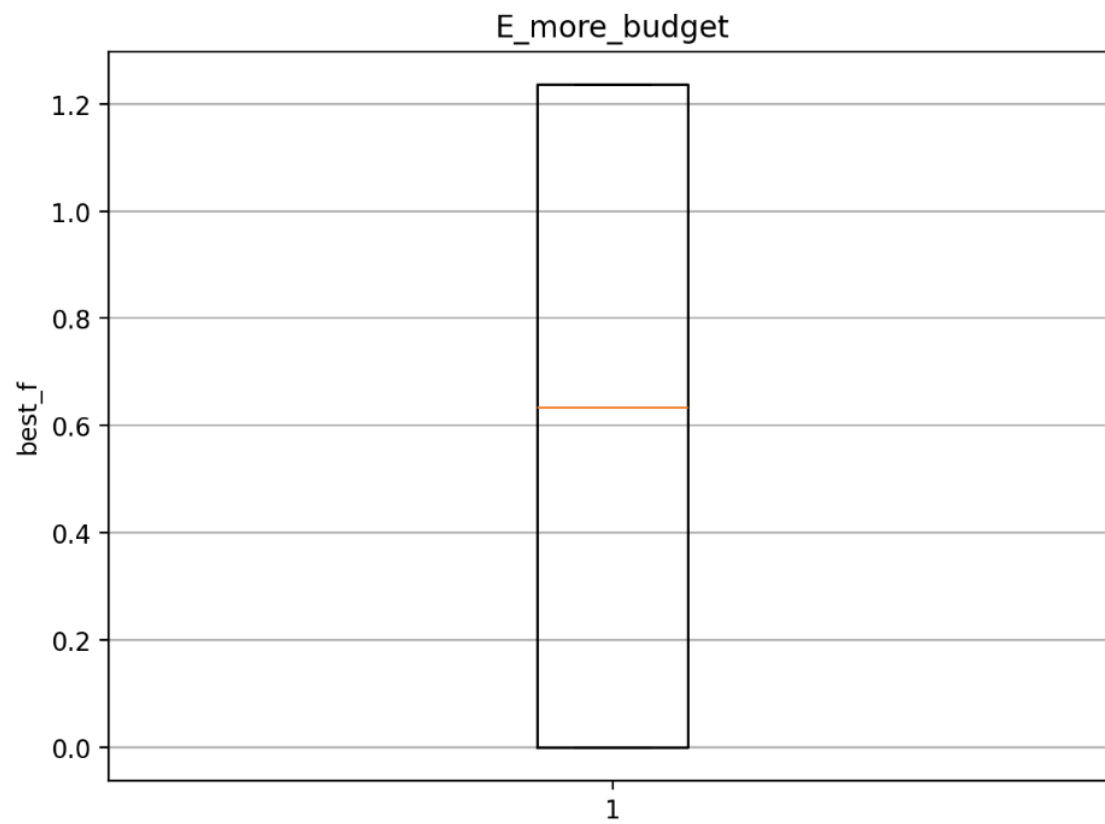
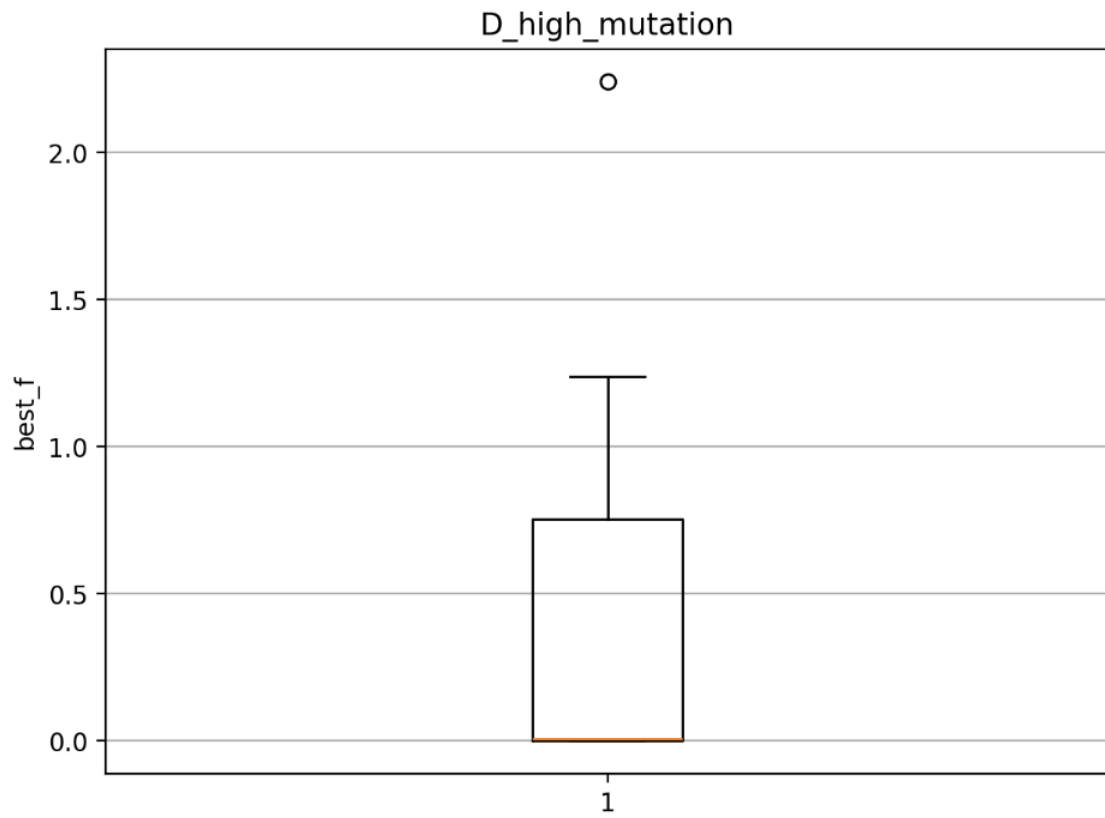
RASTRING(n=5)



Boxplots (final best_f) — one per scenario



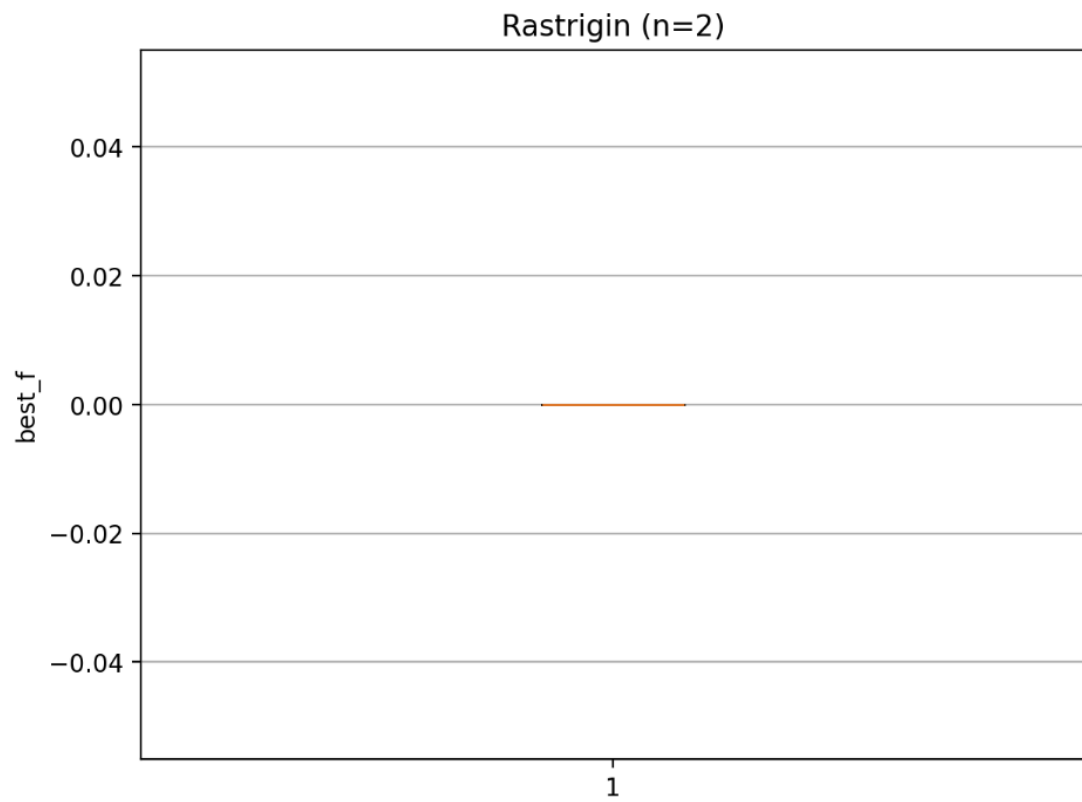


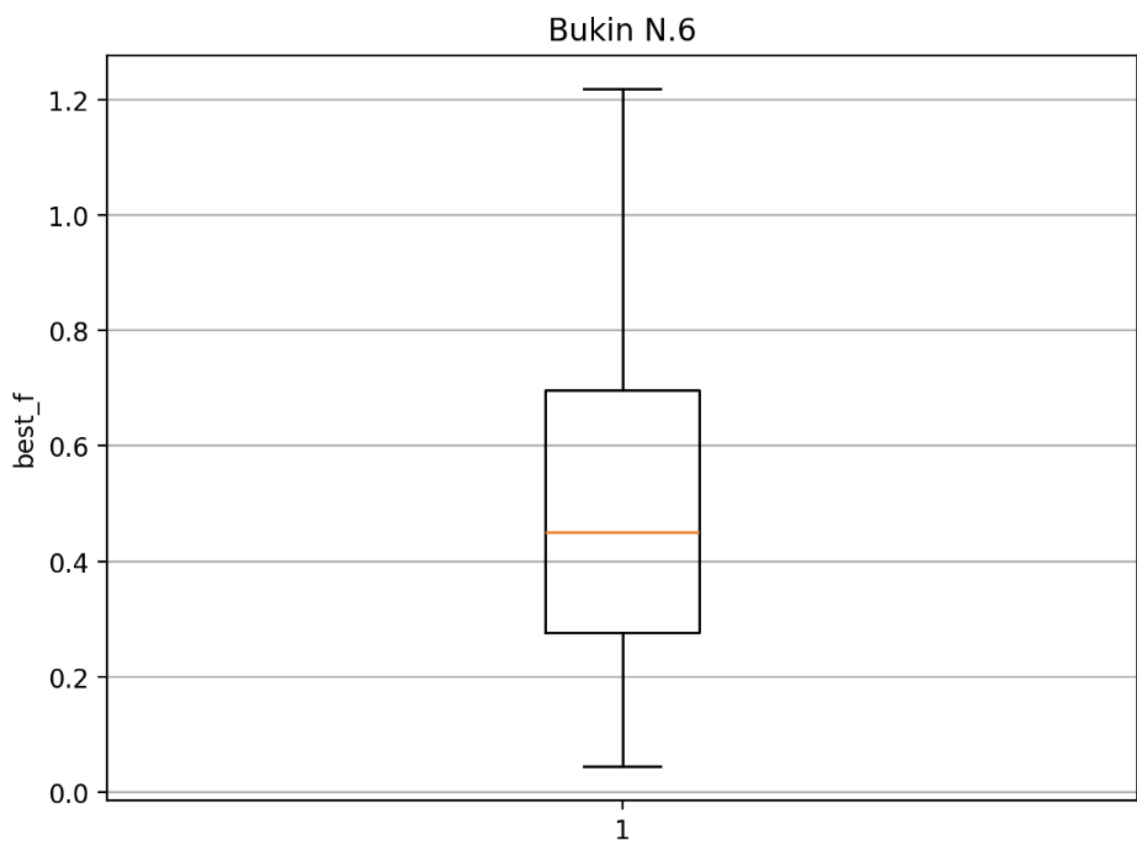
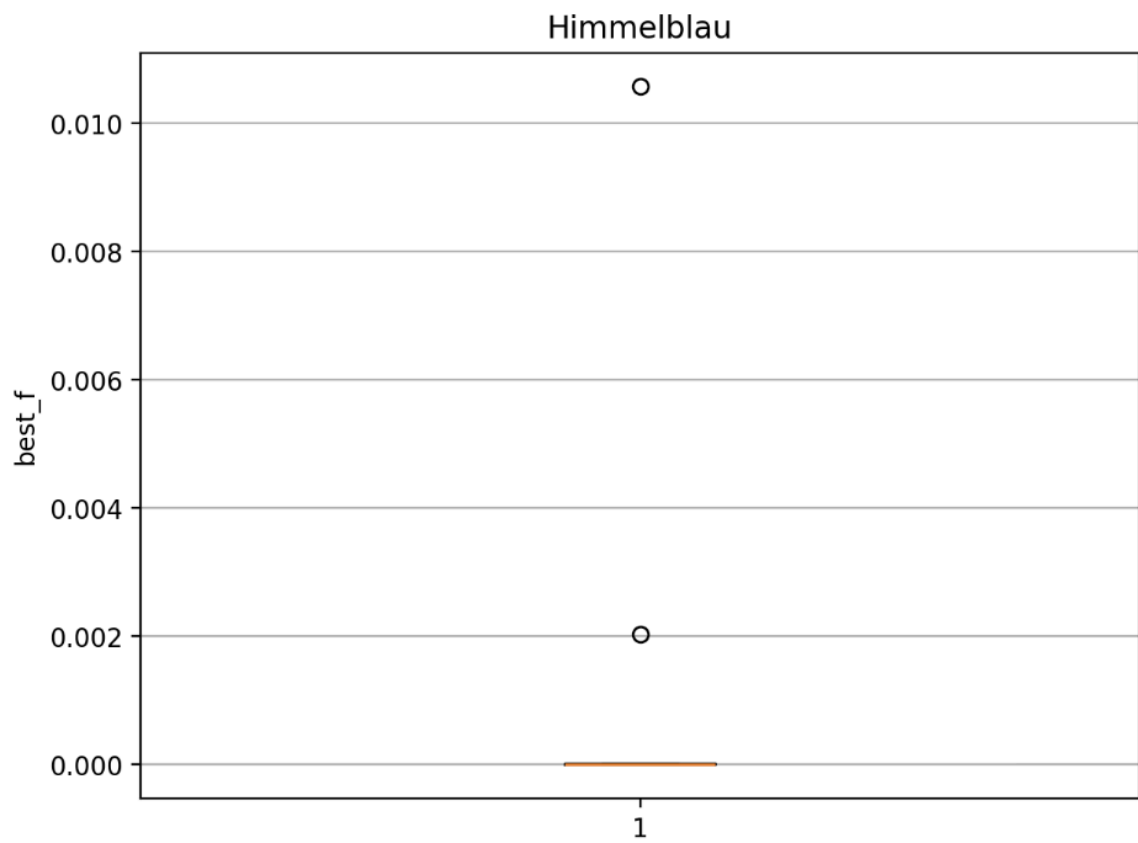


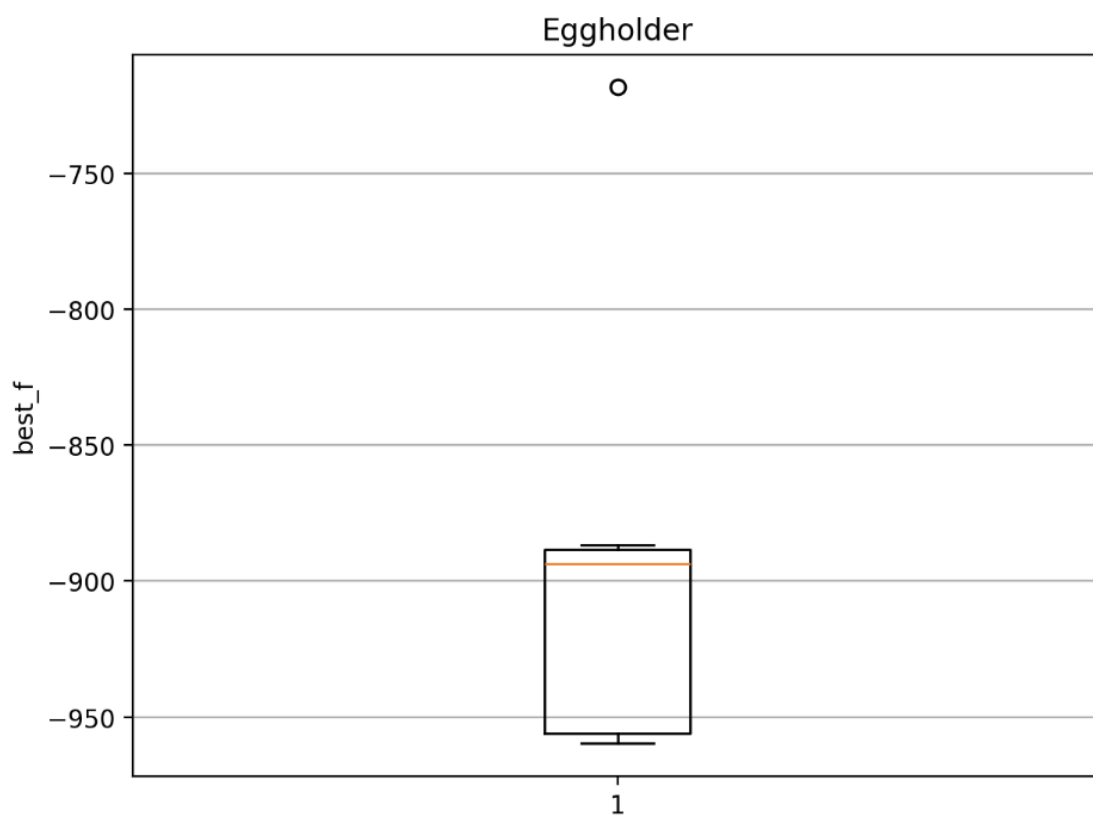
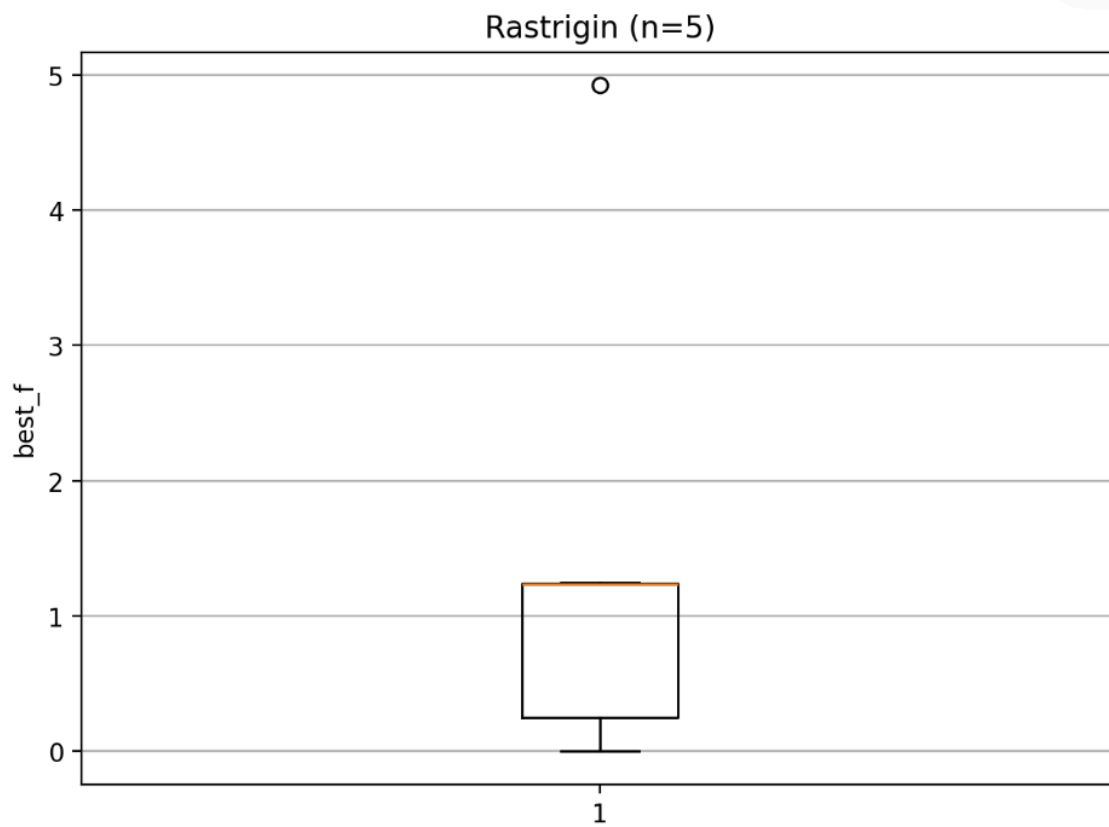
Mode: Compare Functions

a_baseline

Boxplots (final best_f) — per function

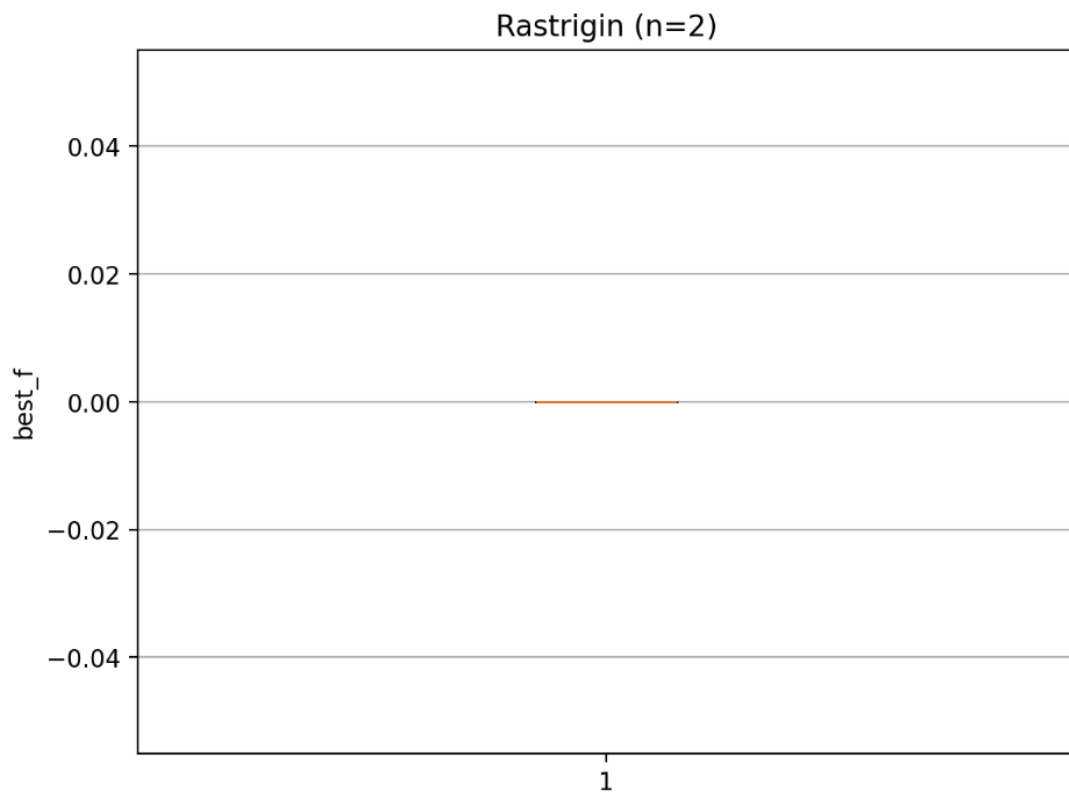


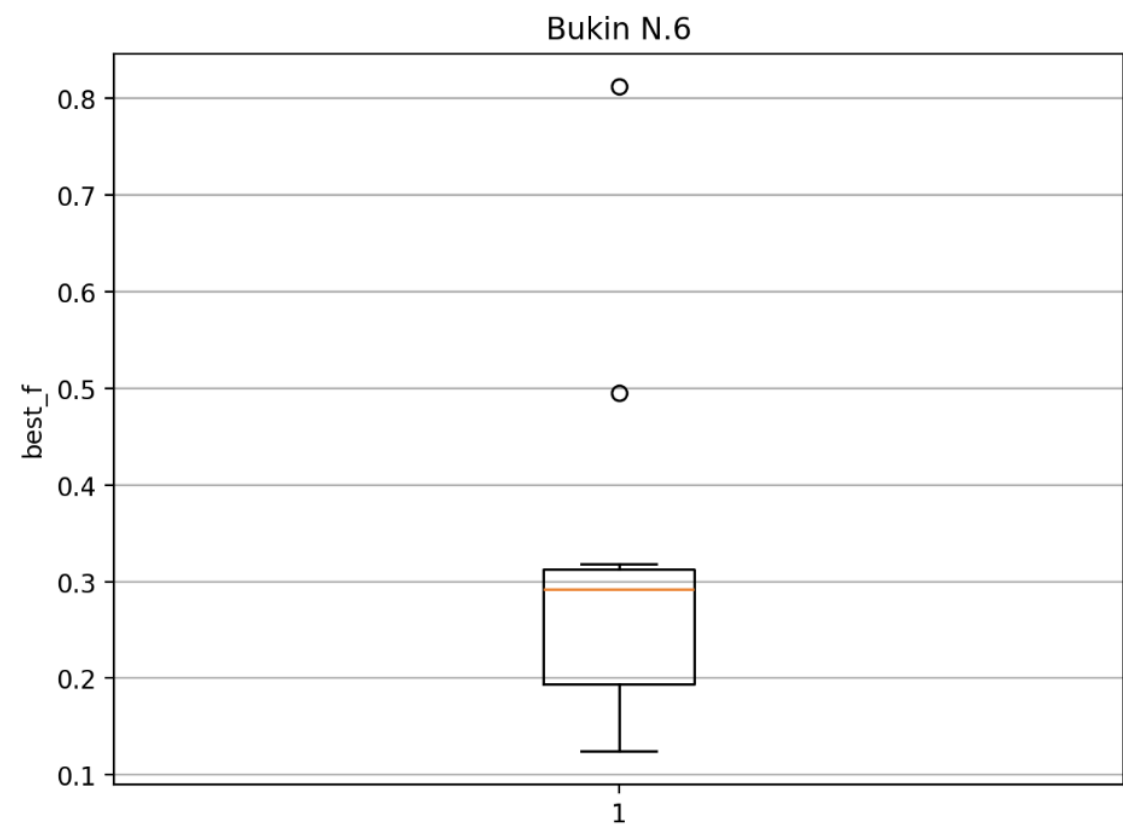
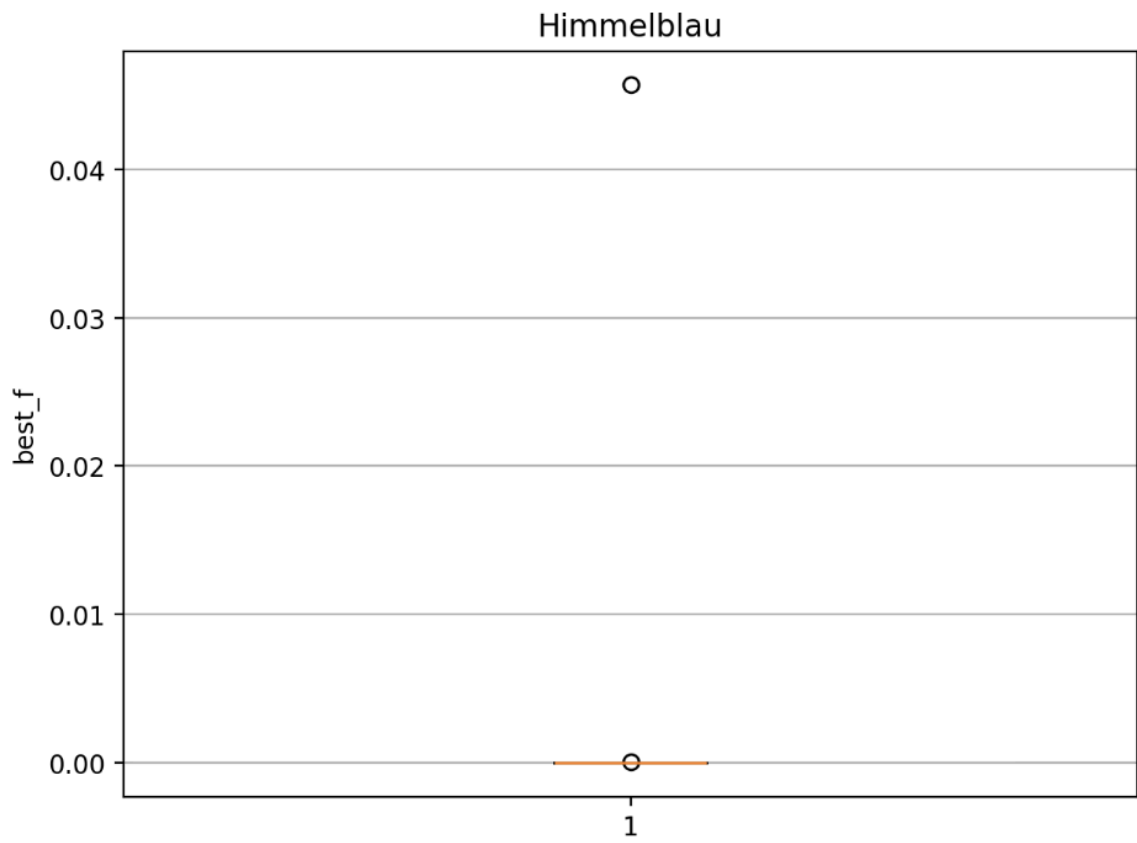


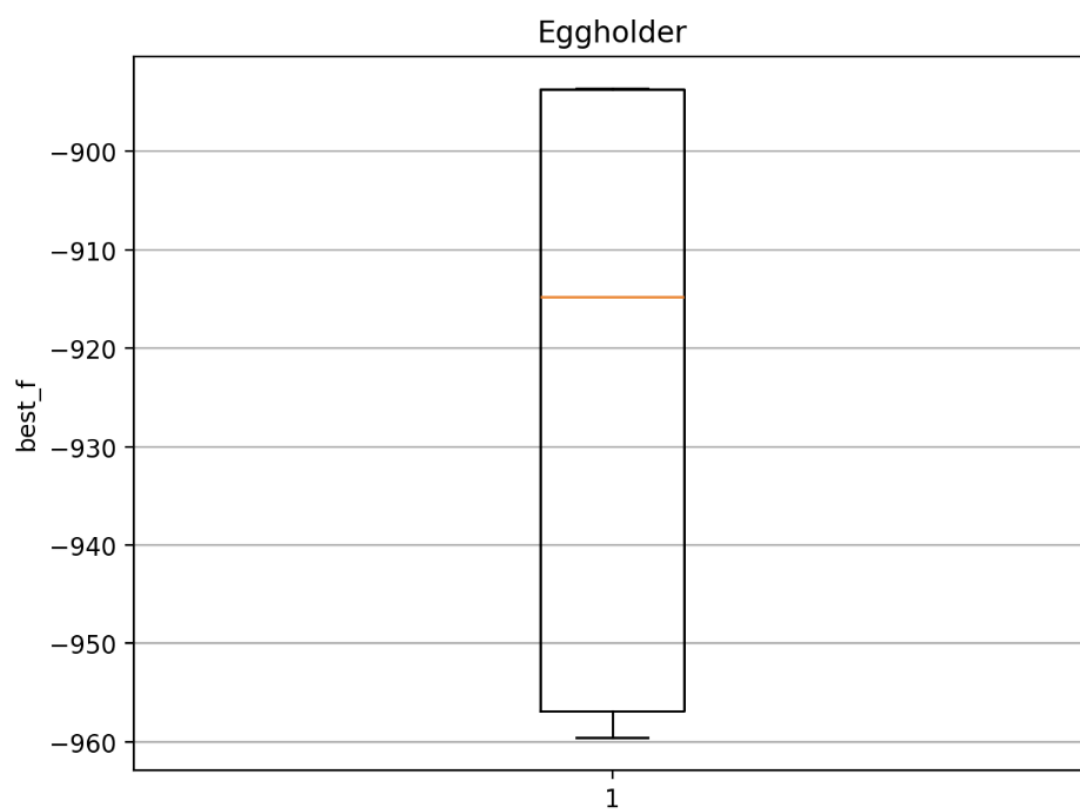
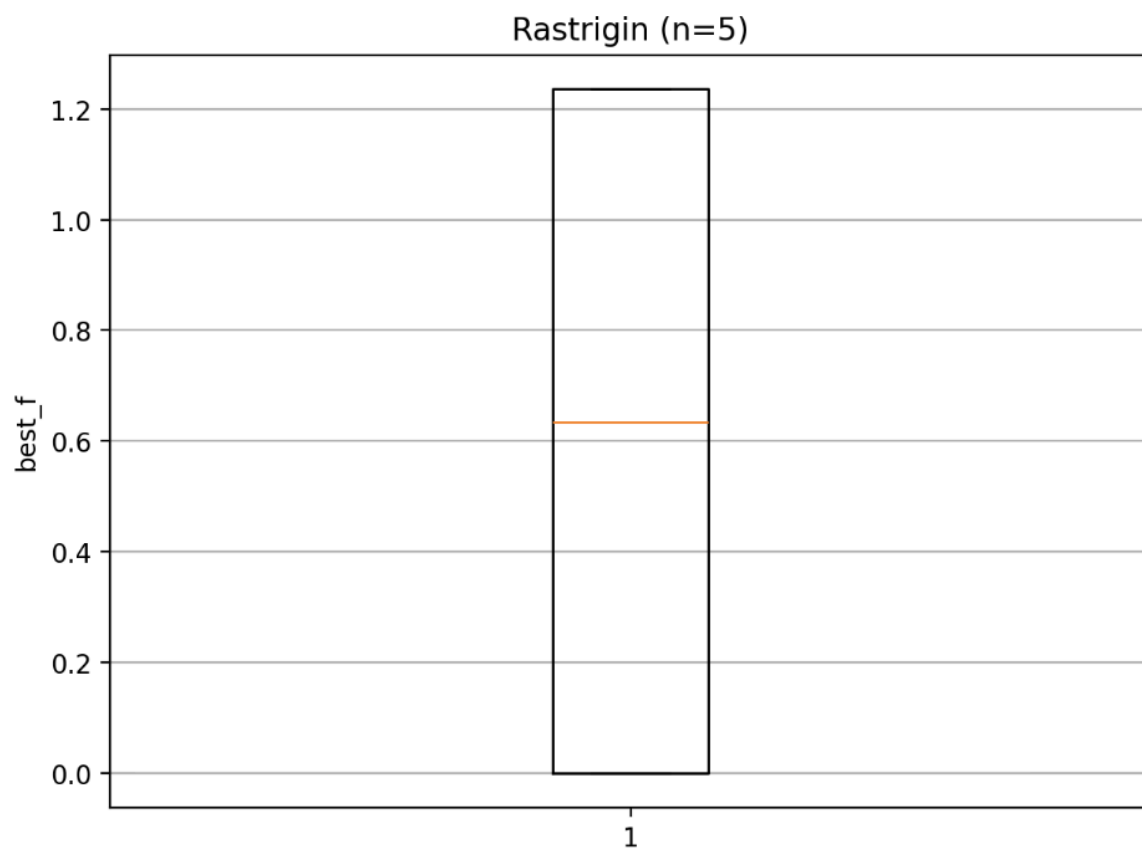


e-morebudget

Boxplots (final best_f) — per function



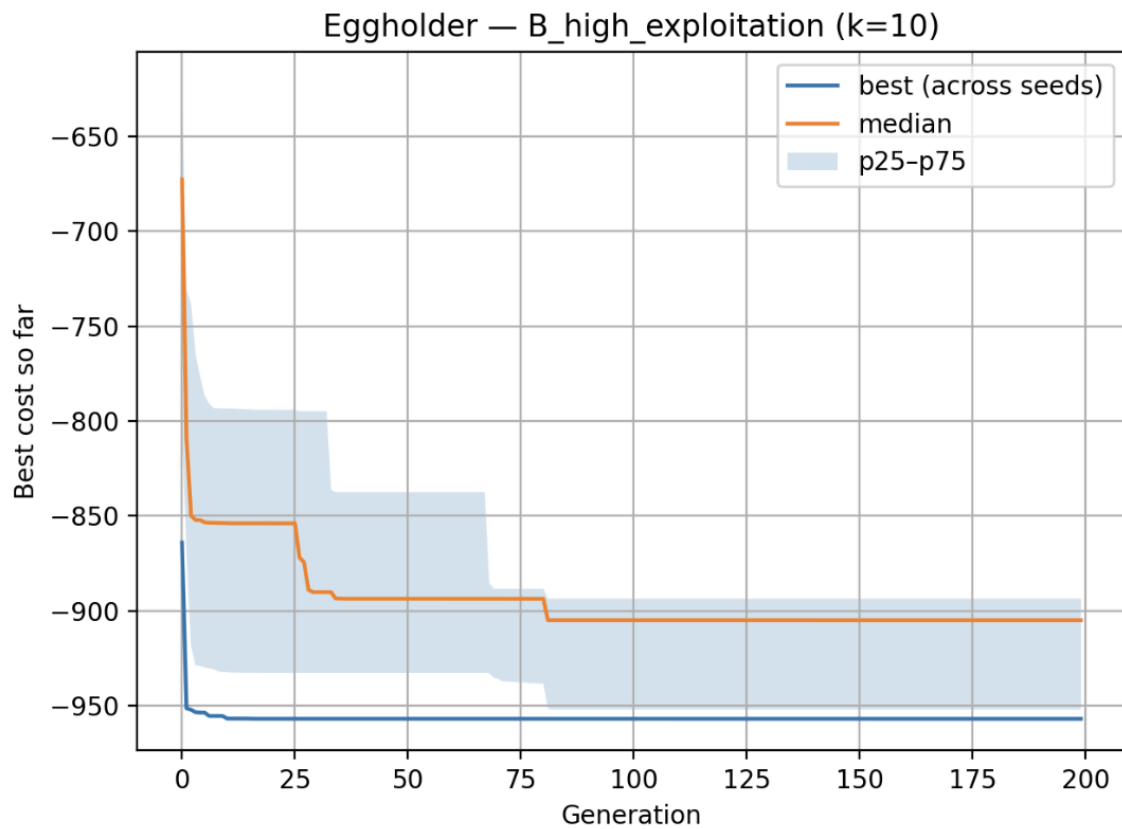




----- escenarios -----

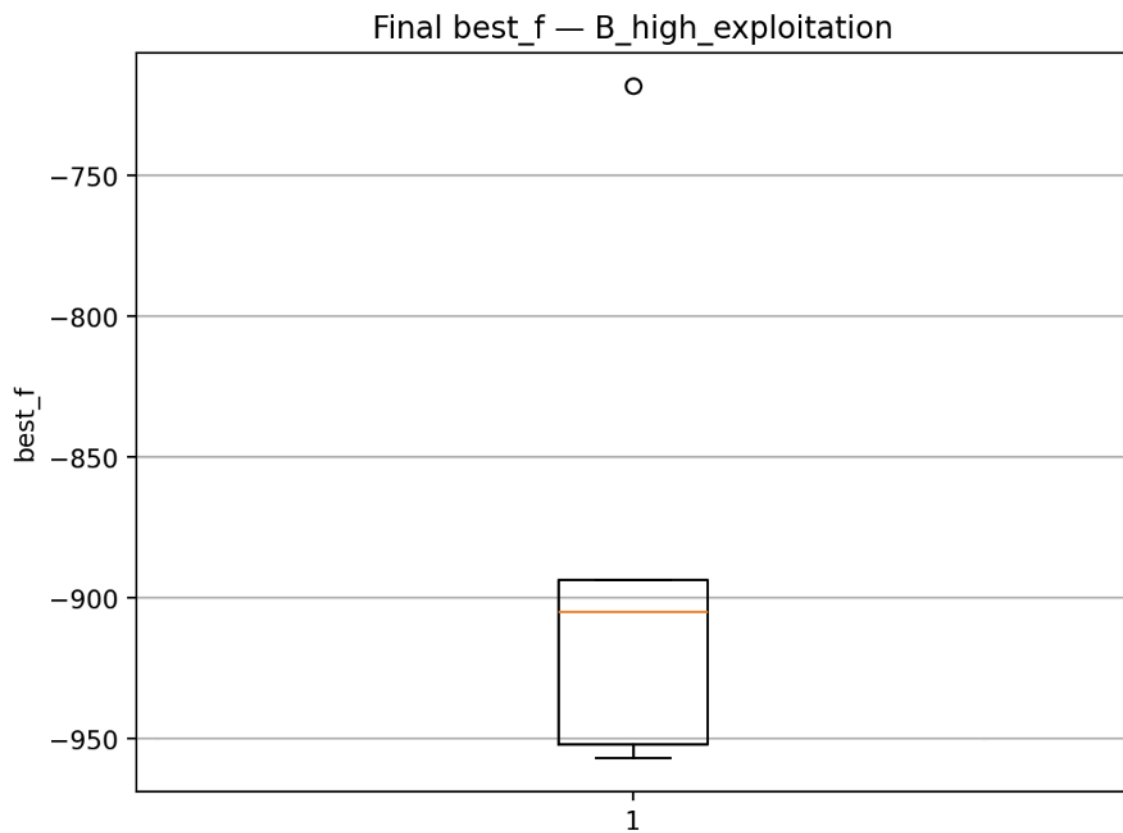
Multi-seed B y D en Eggholder para mostrar diversity y explicar por qué

Convergence bands (best-so-far across seeds)

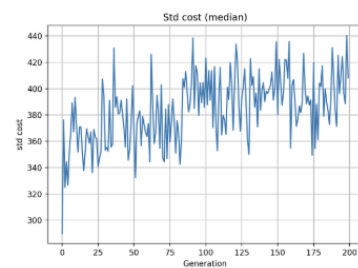
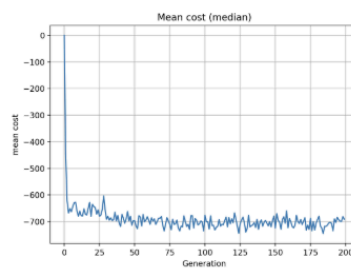
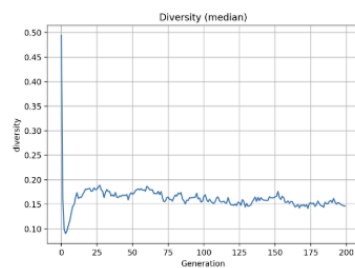




Final best_f distribution

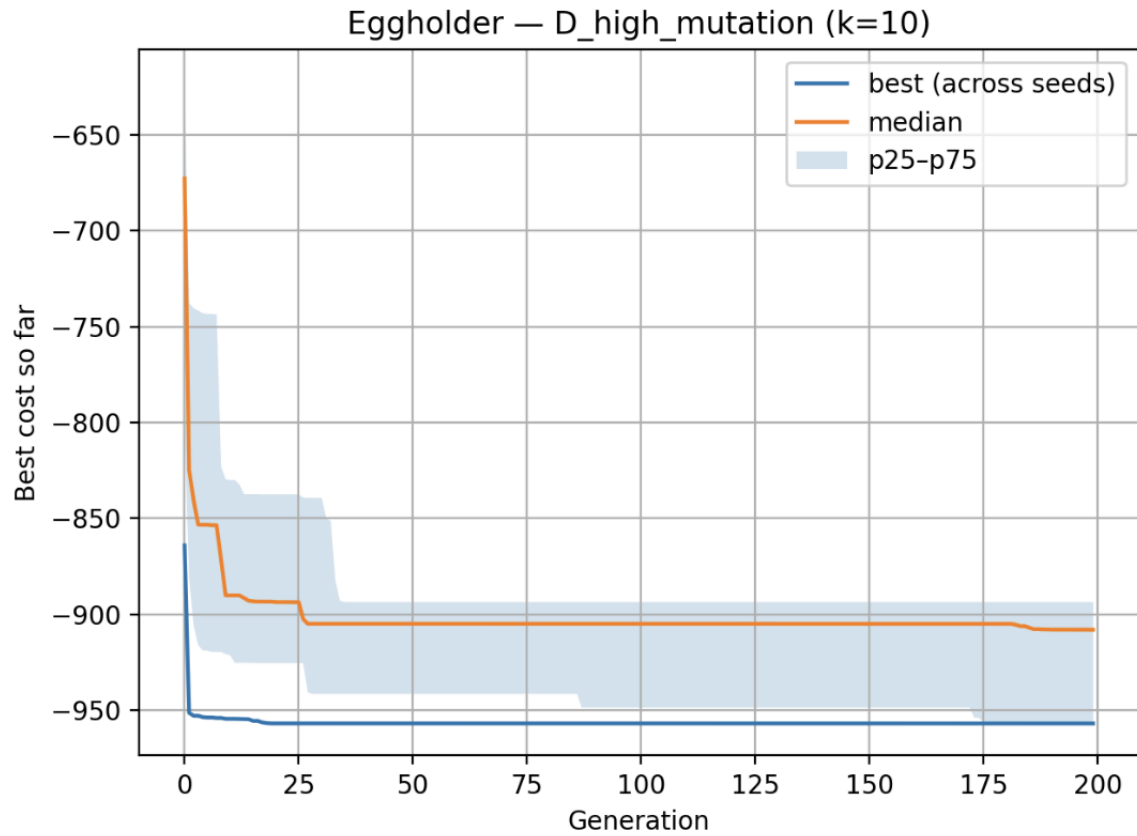


Population diagnostics (median across seeds)



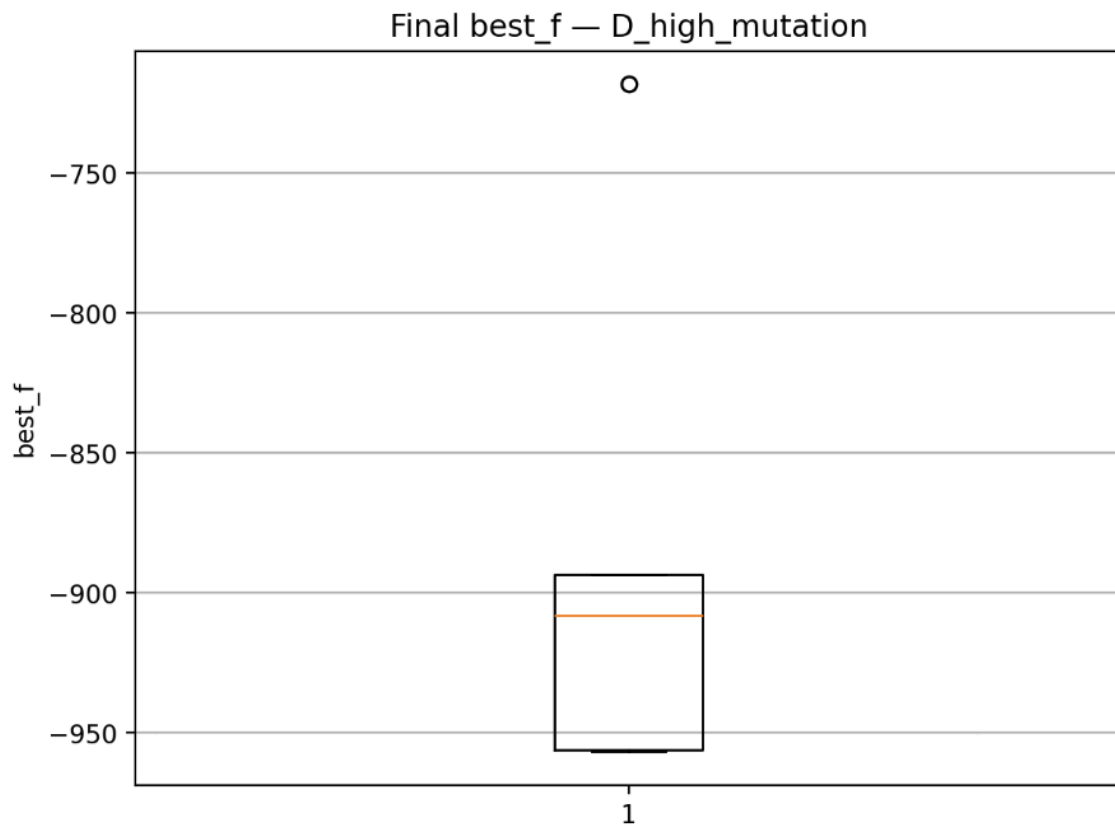


Convergence bands (best-so-far across seeds)





Final best_f distribution



Population diagnostics (median across seeds)

