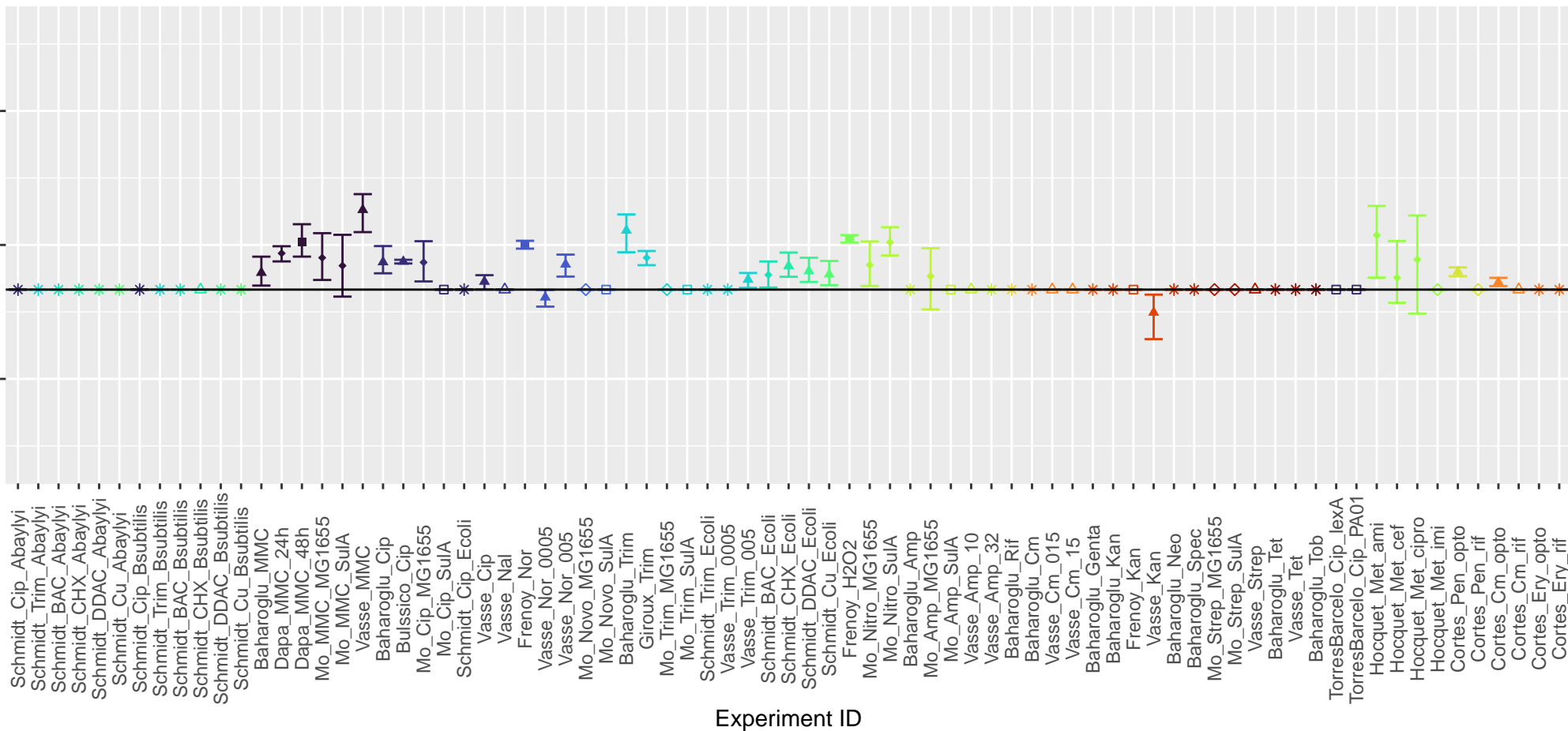


Fold-change population-wide mutation rate

Estimates from the homogeneous model with lowest AIC, if S significantly different from UT condition



Selected model

- △ N0
- ◇ N1
- N2
- ▲ HOM0
- ◆ HOM1
- HOM2
- * S/UT n.s. different

Antimicrobial (abbr)

- | | |
|-------|-------|
| MMC | Amp |
| Cip | Pen |
| Nal | Rif |
| Nor | Cm |
| Novo | Ery |
| Trim | Genta |
| BAC | Kan |
| CHX | Neo |
| DDAC | Spec |
| Cu | Strep |
| H2O2 | Tet |
| Met | Tob |
| Nitro | |