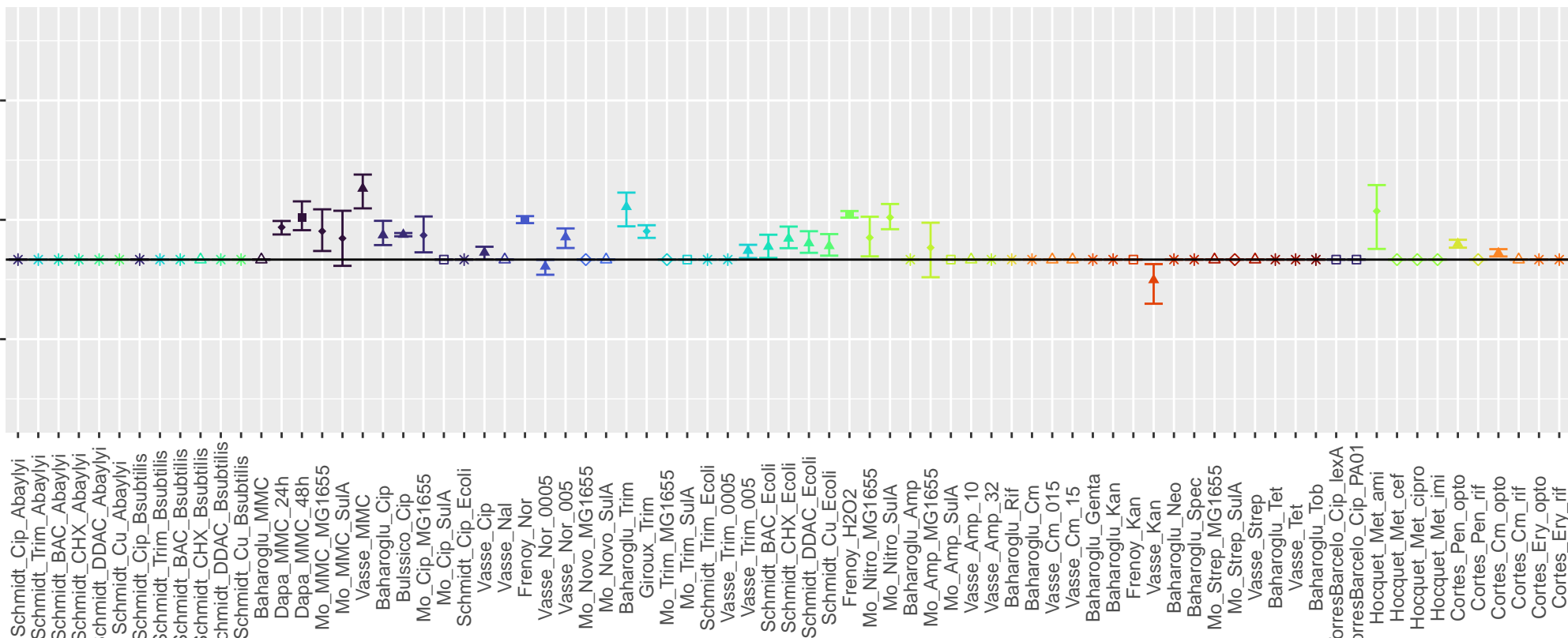


Fold-change population-wide mutation rate

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



Experiment ID

Selected model

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

Antimicrobial (abbr)

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