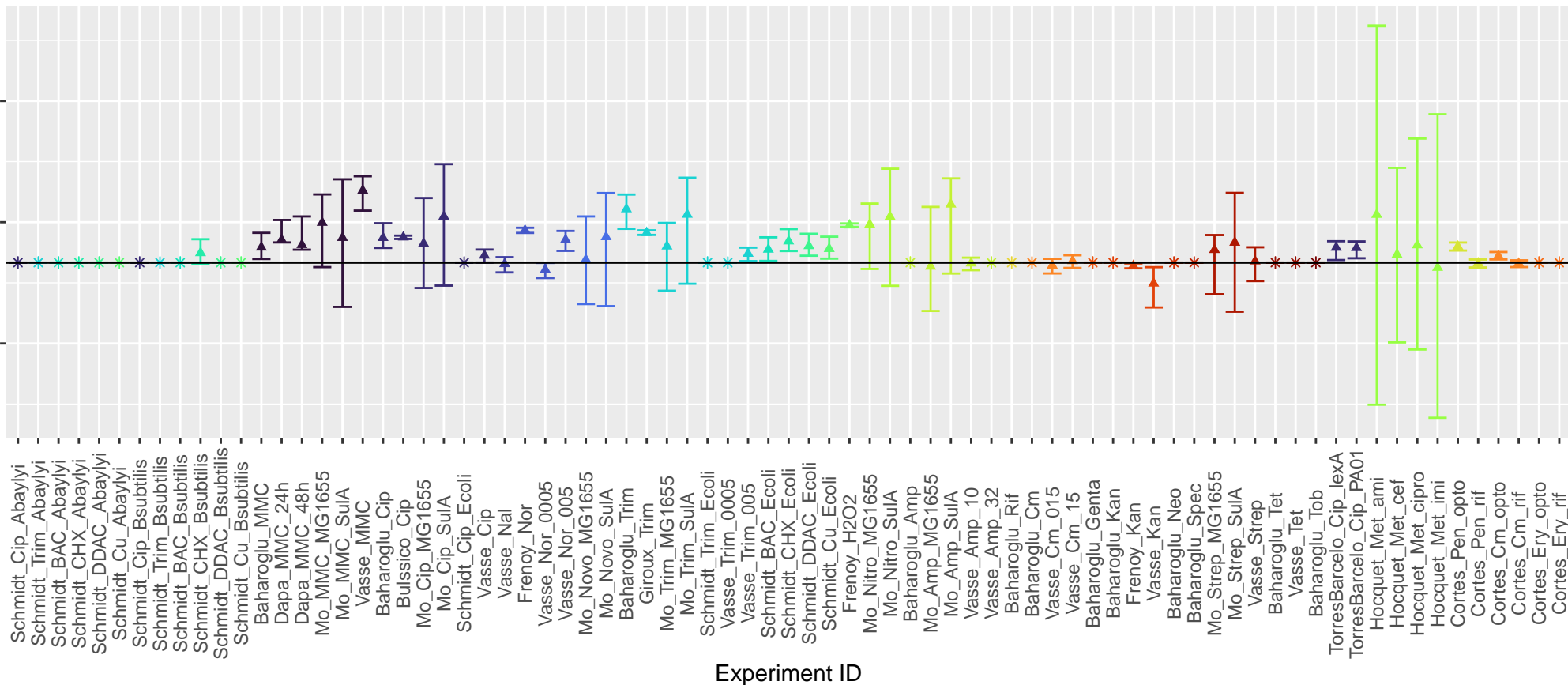


Fold-change population-wide mutation rate

Estimated using model HOM0, if S significantly different from UT condition



GoF test resulted in

- ▲ HOM0
- \* 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 |       |