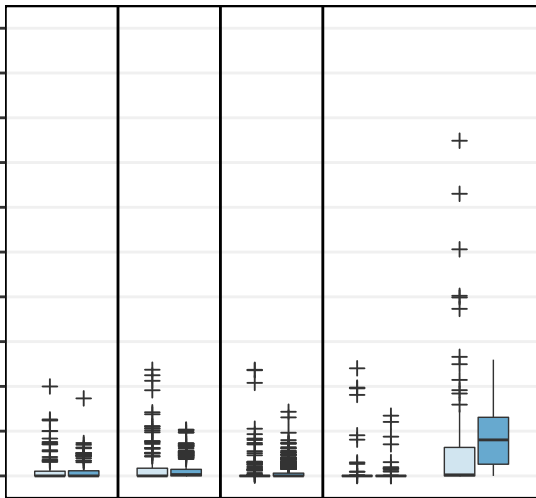


Highest incorrect cluster estimate



*C. jejuni/coli*

*E. coli*

*K. pneumoniae*

*S. epidermidis*

14 groups

82 groups

82 groups

3 groups

11 groups



mSWEEP single



mSWEEP mixture