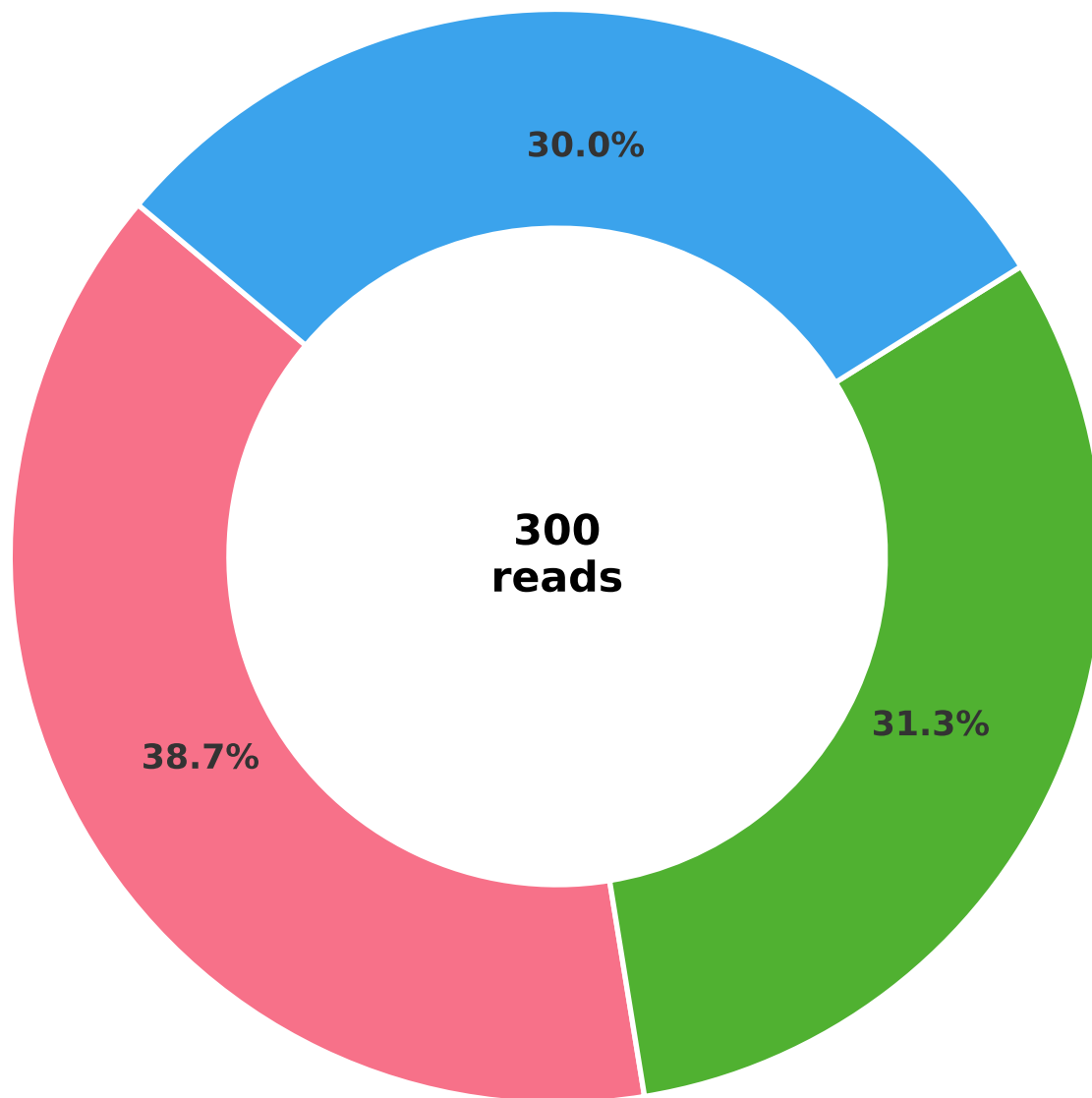


Microbial Community Composition



- Staphylococcus aureus (38.7%)
- Klebsiella pneumoniae (31.3%)
- Escherichia coli (30.0%)