Figure: Box plots showing the Number of Blast Hits per Genes per Megabase comparing MSS and ResCap sequencing protocols across different blast analyses.

All Blast Hits with Evalue lower than 10^{-100} (Blastn against ResCap DataBase)

Blast Hits with Identity lower than 95%

Blast Hits with Identity higher than 95%

Known (Identity Higher than 95%) (Blastp against Uniprot Bacteria Database)

Novelty (Identity Lower or equal than 95%)
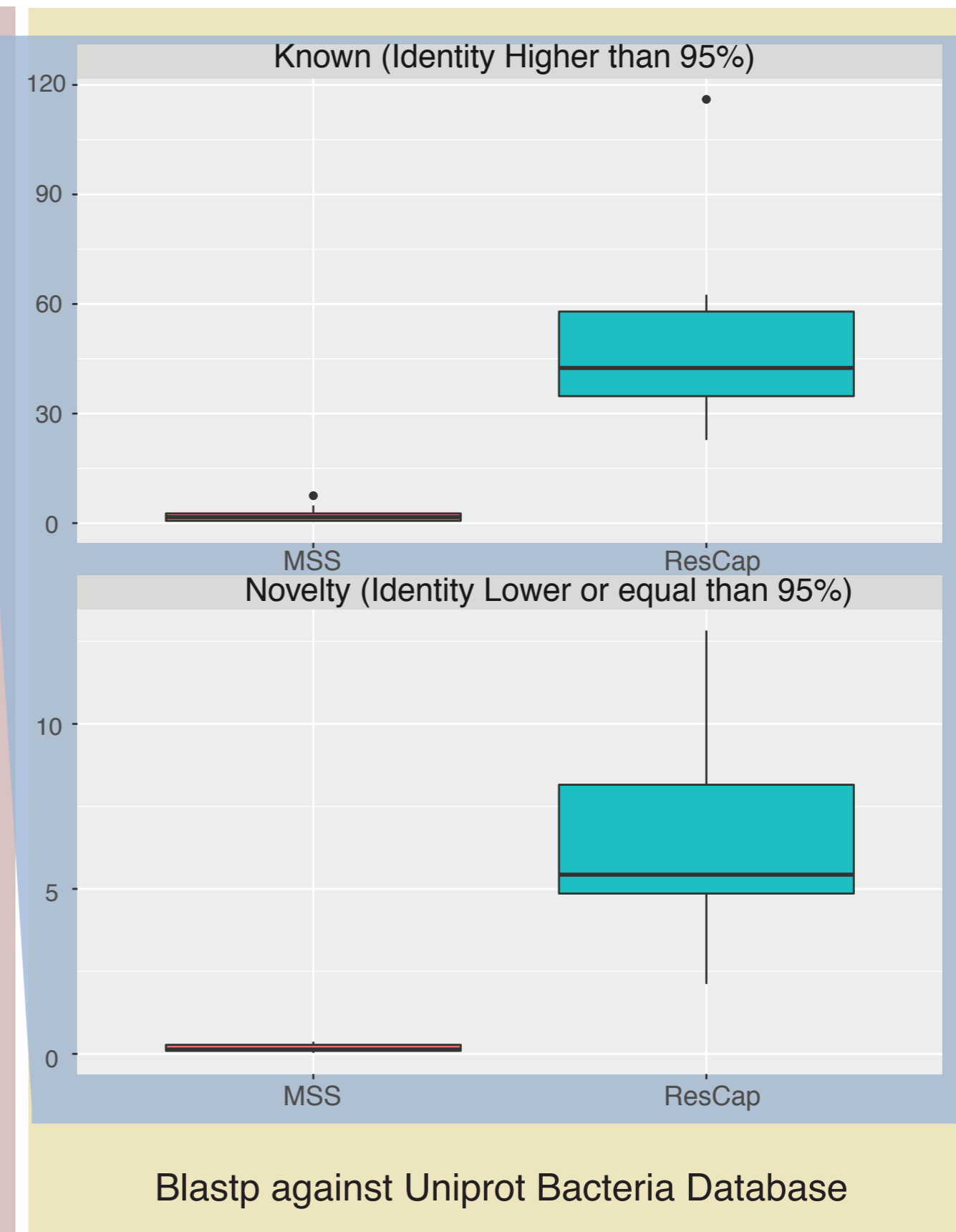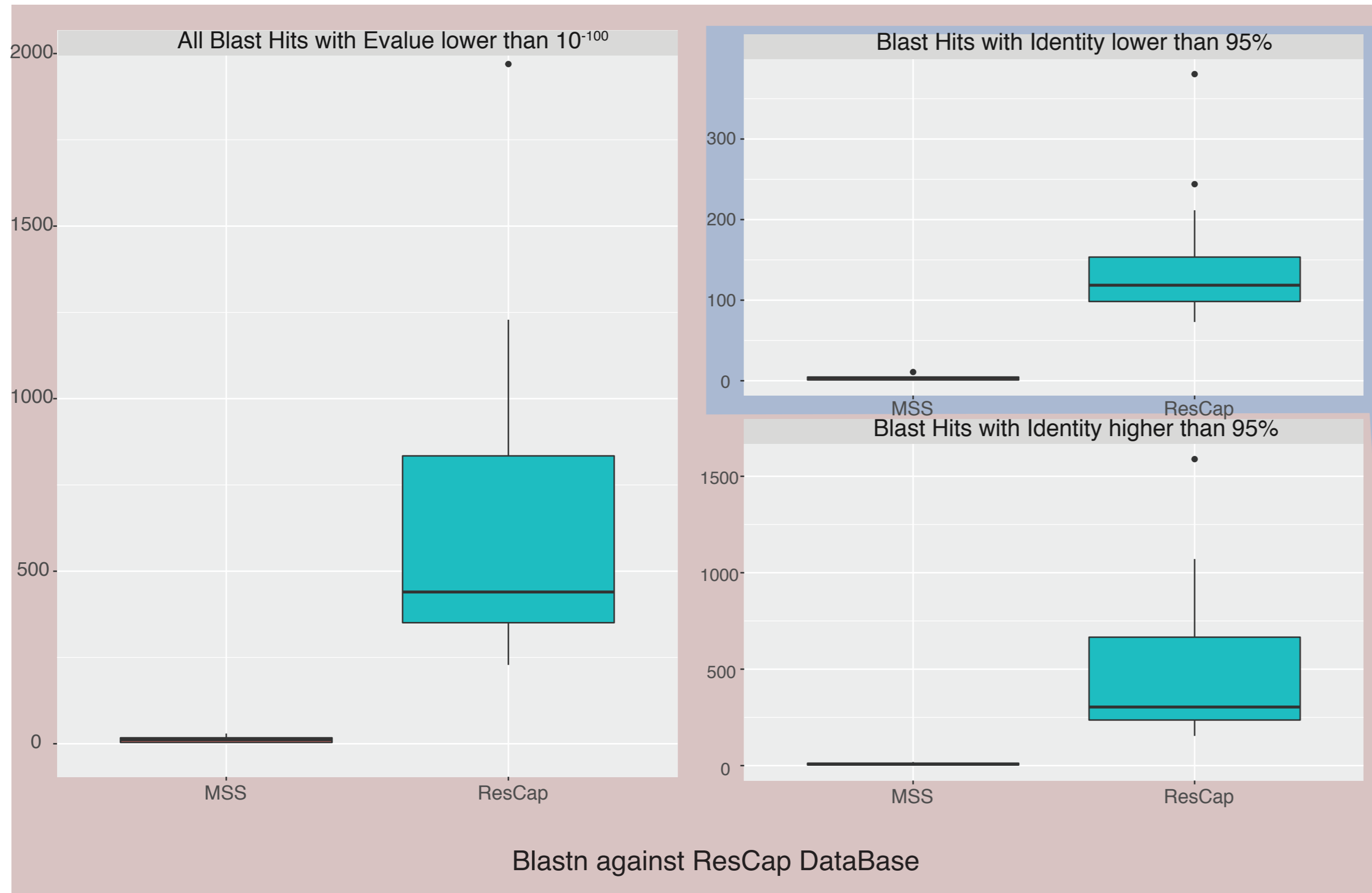
Sequencing Protocol
- MSS
- ResCap