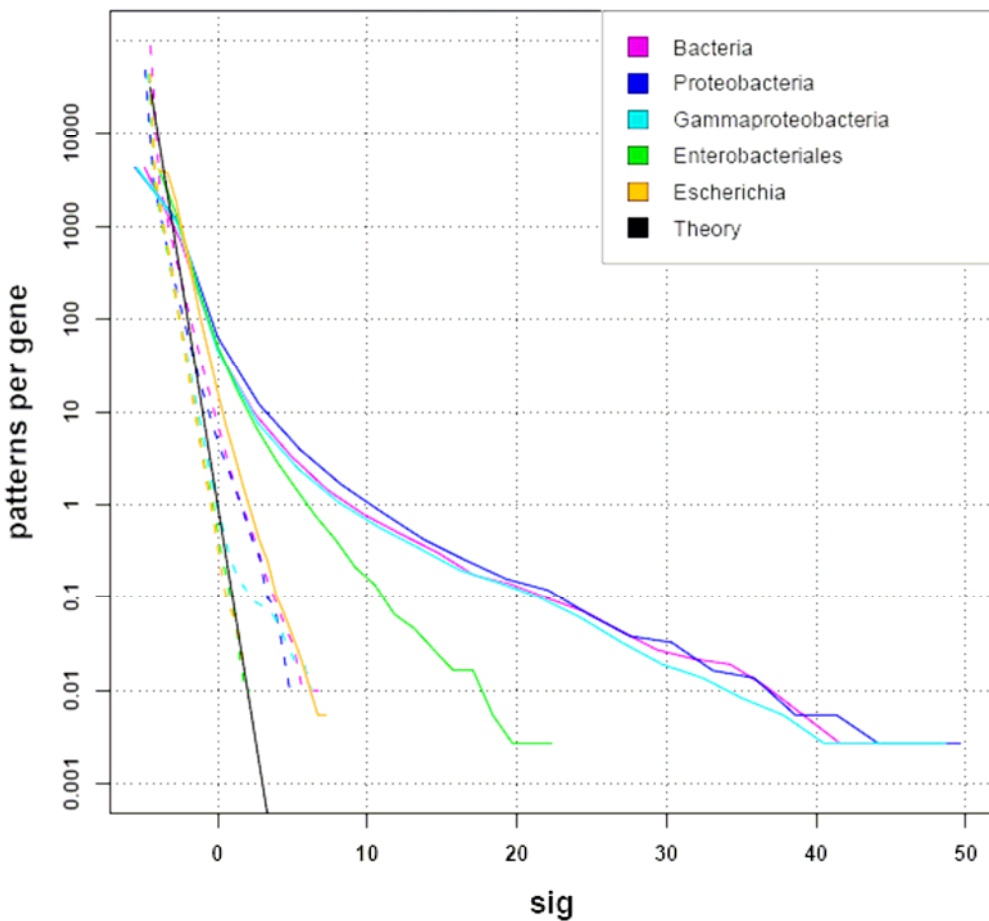


**A**

Significant patterns per gene

**B**

Fraction of sequence sets with at least one significant pattern

