

Clever title here

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Stats 744: Final Project

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The Organism

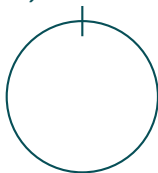
Bacteria:

- *Escherichia coli*
- *Bacillus subtilis*
- *Streptomyces*
- *Sinorhizobium meliloti*

insert pic of bacteria here!
probs sino bc its beautiful

Biological Background

1) Circular



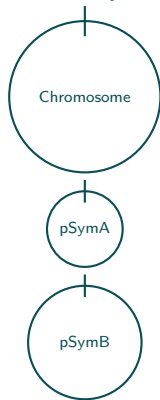
Escherichia coli
Bacillus subtilis

2) Linear



Streptomyces

3) Multi-repliconic



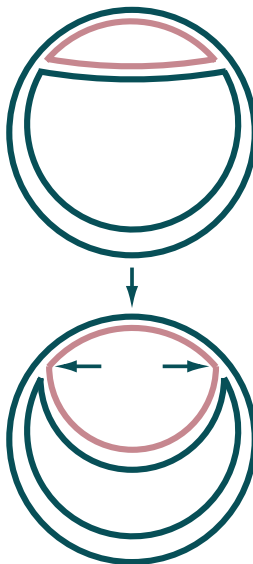
Sinorhizobium meliloti

Bacterial Genome Properties

Bidirectional Replication

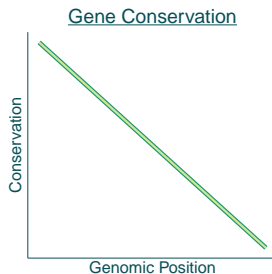
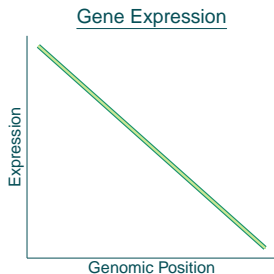
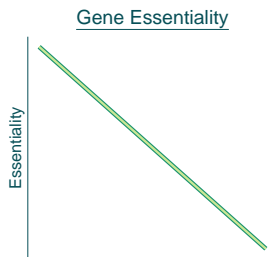
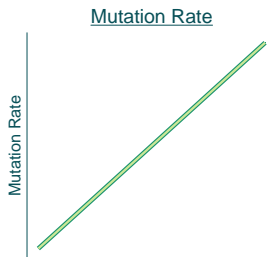
- Gene dosage
- Sequence composition
- Codon bias
- Replication errors
- Strand differences

– new DNA
– old DNA



Different **replicons** and **positions** of bacterial genomes can have **varying molecular trends**

Spatial Molecular Trends



Couturier et al. 2006, Cooper et al. 2010, Sharp et al. 2005, Morrow et al. 2012, Cooper and Rocha 2006

Selection

Measuring what evolutionary forces are acting on genes

- **dN** = non-synonymous substitution rate
 - change in amino acid sequence
 - can alter function of the protein
- **dS** = synonymous substitution rate
 - no change in amino acid sequence
 - will not alter function of protein
- $\omega = (dN/dS)$
 - $\omega > 1 \rightarrow$ positive selection
 - beneficial to the organism and will likely be maintained over time
 - $\omega = 1 \rightarrow$ neutral selection
 - neither beneficial nor deleterious to the organism
 - $\omega < 1 \rightarrow$ negative selection
 - deleterious to the organism and will likely not be maintained over time

Gene Expression

- *E. coli*, *B. subtilis*, *Streptomyces*, and *S. meliloti*
- Gene expression + distance from the origin of replication
- **Expectation:** Near the origin or replication = Higher Expression

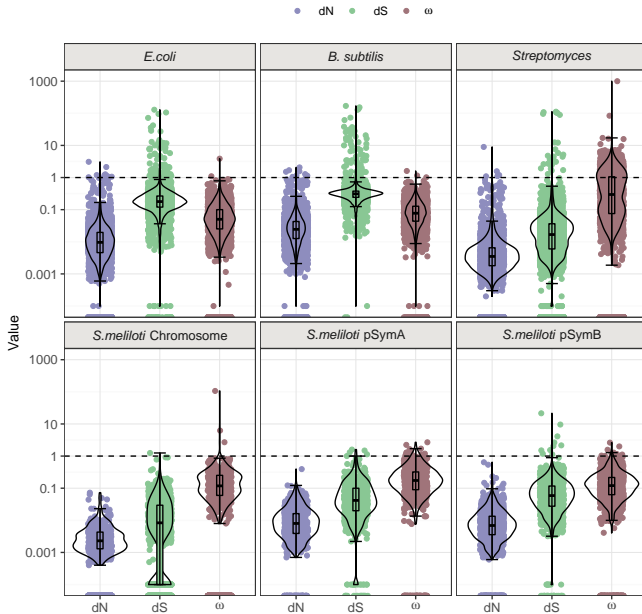
Selection

- *E. coli*, *B. subtilis*, *Streptomyces*, and *S. meliloti*
- dN, dS, and ω + distance from the origin of replication
- $\omega > 1$ (positive selection), $\omega = 1$ (neutral), $\omega < 1$ (negative selection)
- **Expectations:**
 1. dS > dN
 2. Most genes are under neutral or negative selection
 3. Genes that are under positive selection should most likely be accessory genes
 - Core genes are near the origin of replication
 - Accessory genes are usually near the terminus

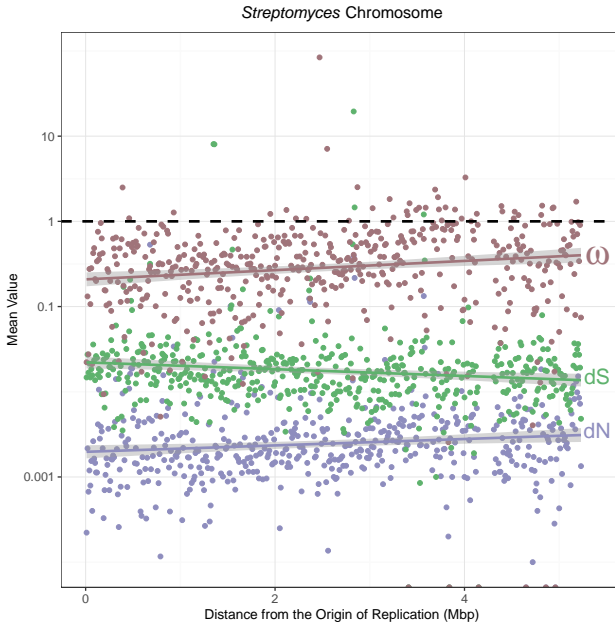
Gene Expression

Insert gene expression graphs here!!!!

Selection Summary



Streptomyces Selection



Summary

- ✓ Gene expression decreases (slightly) with increasing distance from the origin of replication
 - But we see more of a wave-like pattern!
- ✓ $dS > dN$
- ✓ Most genes are under neutral or negative selection ($\omega \leq 1$)
- ✓ In *Streptomyces*, genes under positive selection ($\omega > 1$) were near the terminus and most likely accessory genome