

Discussion Club 2018-09-04:

Soil-persistent *E. coli* and Mobile elements

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September 4, 2018

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Background

What does 10 years look like to *E. coli*?

Mobile Genetic Elements

In Closing

Background



- *E. coli* has been found to persist stably in the soil
- Isolates were cultured from lysimeter leachate
- Strains were sequenced, resulting in 149 soil-persistent *E. coli* genome



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- ◊ What virulence factors are harboured by these strains?
- ◊ What can we infer about adaptation from these?
- ◊ Can we differentiate soil-persistent *E. coli* from recent contamination?

What does 10 years look like to *E. coli*?



$$(5.9 / 2) * 365.25 * 10$$

$\approx 10\text{k}$ generations

Bååth 1998

(assuming generation time roughly equals half of turnover rate)



- o 32 days
- o 22 days
- o 8 weeks
- o 114 days

Detection of *Escherichia coli* in sequenced soil



Escherichia coli approximately .092% prevalence in soil



Highly diverse soil communities and environmental pressures favour rapid adaptation over incremental changes



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Adaptations will occur more rapidly because of mobile elements rather than mutations to core functions

Mobile Genetic Elements

Insertion Sequence (IS) (Transposon)

Short, mobile sequence capable of jumping



- Size: 750bp – 1.5Kb
- Detection: inverted tandem repeat lollipop with a transposase and promoter.
- Maintenance: genome replication
- Mobility: encoded transposase

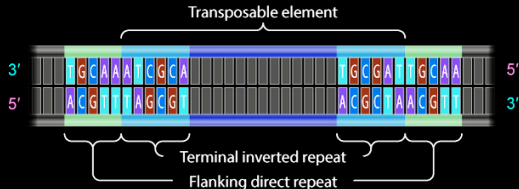


Figure: Insertion Sequence

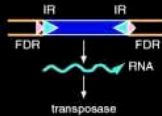
A DNA lollipop containing a transposase and promoter, and other accessory genes

- Size: $< 3\text{Kb}$
- Detection: inverted tandem repeat, transposase, bonus genes
- Maintenance: genome replication
- Mobility: encoded transposase

A DNA fragment with IS sequences on each end

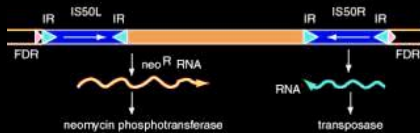
- Size: $< 5\text{Kb}$
- Detection: double IS, promoters, and accessory genes
- Maintenance: genome replication
- Mobility: encoded transposase

Insertion sequences

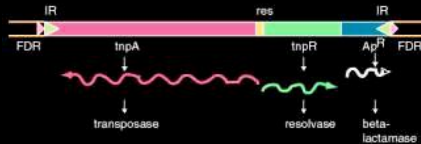


Transposons

Composite transposons, e.g. Tn5



Transposons lacking terminal ISs, e.g. TnA



Mini-plasmids often containing AMR gene cassettes

- Size: < 10Kb
- Detection: integrase, recombination site (attI), promoter, and (optionally) a cassette of resistance genes
- Maintenance: integration into host genome
- Mobility: recombination, or transfer when circularized



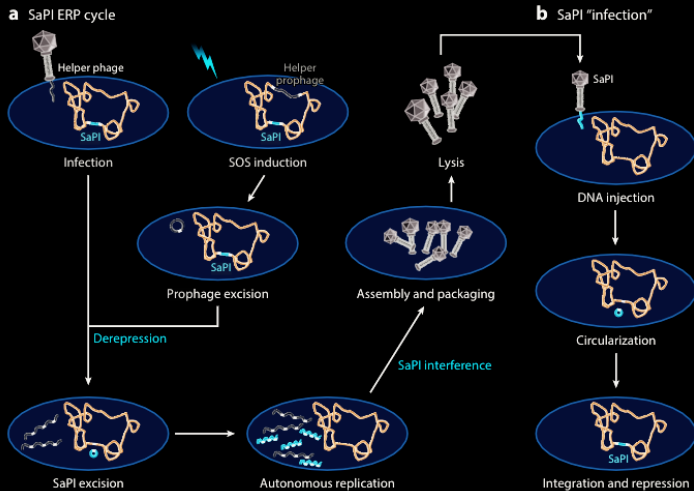
- Size: $>10\text{Kb}$
- Detection: GC Skew, phylogenetic analysis of ORFs
- Maintenance: genome replication
- Mobility: various



Phage parasites

- Size: 5 – 15Kb
- Detection: Look for phage proteins along with phage inhibitors
- Maintenance: lysogeny
- Mobility: phage-like particles

Phage-inducible Chromosomal Islands





Virus integrated into the host genome

- Size: ~ 50Kb
- Detection: Look for phage integrases, tail and capsule proteins
- Maintenance: genome replication
- Mobility: lysogenic and lytic phases



Freely replicating DNA not required for survival

- Size: $<1\text{Kb} - 1\text{Mb}$
- Detection: run a gel; look for circular contigs
- Maintenance: self-replicating or integrating
- Mobility: conjugation (directly or indirectly)

In Closing



- Develop mobile pangenome tool for detecting "recent" adaptations
- Characterize trends in MGEs in environmental vs enteric *E. coli*



- o <https://www.nature.com/scitable/topicpage/transposons-the-jumping-genes-518>
- o <https://www.nature.com/scitable/topicpage/transposons-the-jumping-genes-518>
- o https://www.researchgate.net/publication/283707425_The_Phage-Inducible_Chromosomal_Islands_A_Family_of_Highly_Evolved_Molecular_Parasites
- o <https://www.sciencedirect.com/science/article/pii/S0043135416302226>
- o http://www.bx.psu.edu/~ross/workmg/TranspositionCh9_files/



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- ☐ Dr. Florence Abram
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- ☐ Functional Environmental
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Questions?