

# GRC Annual Report 2017-2018: Soil-persistent *E. coli*

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# Outline



Background

Research

Publication Plans

Software

Publications

Other

# Background

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# Project Overview



- *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

# Research Questions



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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?

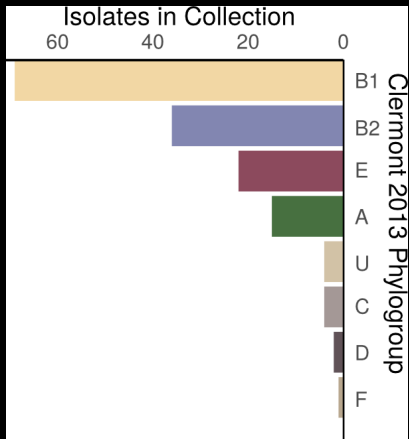


# Research

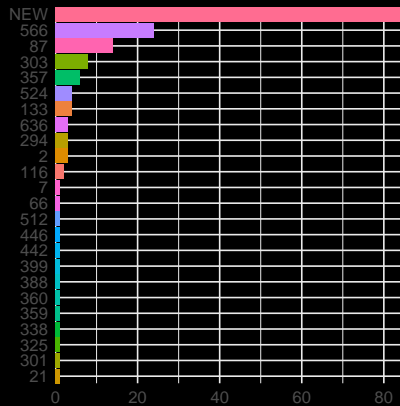
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# Phylogroups

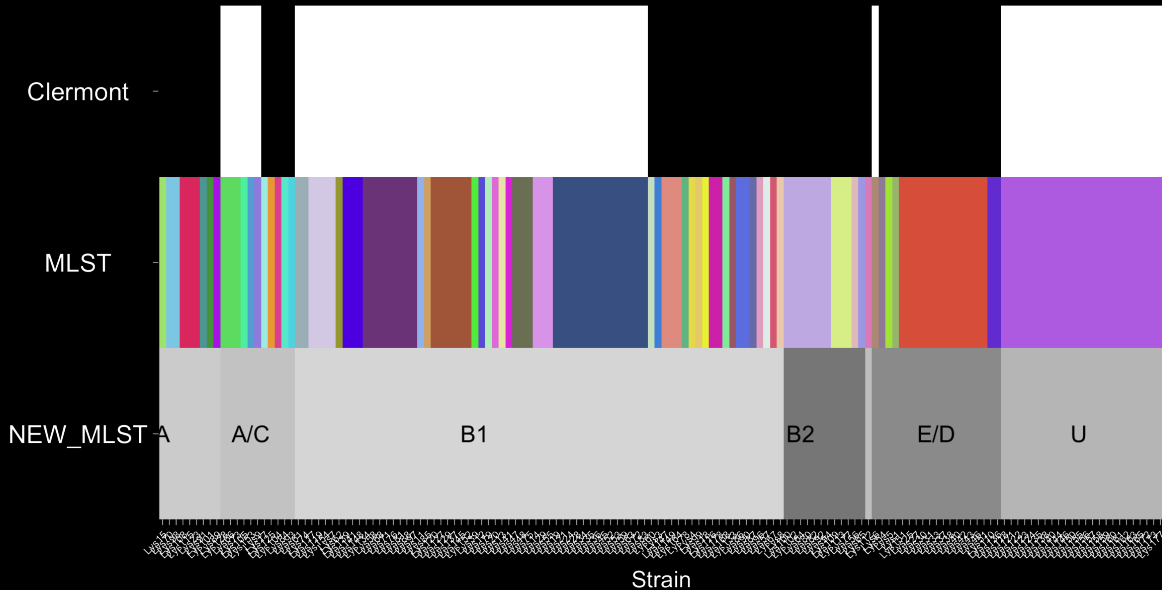
Clermont 2013



Achtman 7 gene MLST



# Phylogroups



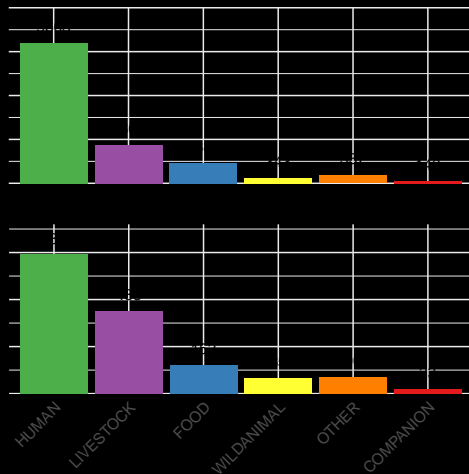
# Phylogenetics Trees



# Pangenome Analysis



- One isolate from each  
Achtman 7 MLST
- Total: 1193



# Pangenome Analysis



	N	Core	total
Soil	149	2662	21,662
Enterobase	1193	1822	79,288
All	1342	1806	83,868

# Detecting differential presence/absence



- Statistically compare traits to a pangenome



# Publication Plans

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## Focus: Genomic characterization of Soil persistent *E. coli*

- o Phylogenetic diversity
- o Pangenome
- o rRNA copy number survey
- o *Correlations with growth-rate phenotypes*
- o Survey of Cold-Shock Proteins
- o Plasmid Survey



## Focus: Comparison of Soil-Persistent Strains to Enteric/Strains

- Pangenome of Soil and Enteric strains
- Sub-pangnomes by phylogroup, plasmid profile, etc
- Secondary metabolite production
- Regulatory regions



## Potential Foci:

- ◊ Virulence
- ◊ AMR
- ◊ Plasmids/Prophages/Other mobile elements

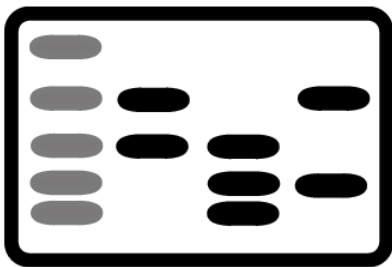
# Software

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[github.com/nickp60/clermontpcr](https://github.com/nickp60/clermontpcr)

build passing coverage 93% License MIT pypi package 0.1.1



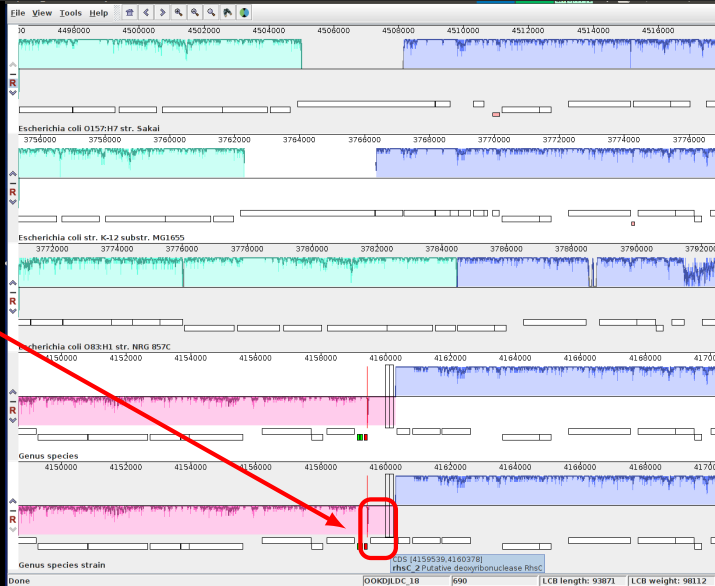
**E** **B2** **C**

Clermont PCR typing tool

# annofilt: Assessing Assemblies



Partial



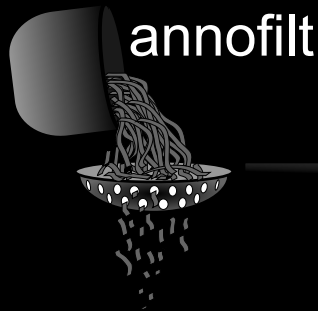


`nickp60.github.io/annofilt/`

1. Select trusted complete genomes
2. Create reference pangenome
3. Find genes next to contig borders
4. Blast against pangenome
5. Reject hits  $< 90\%$  of CDS length

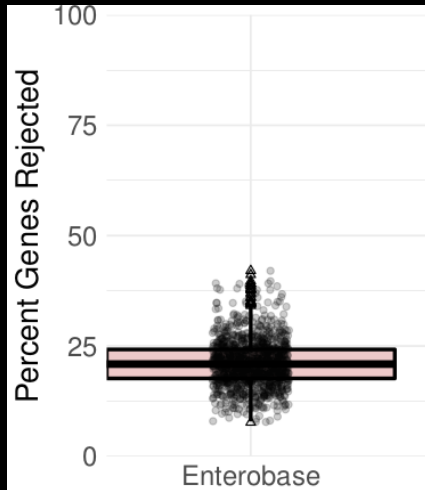
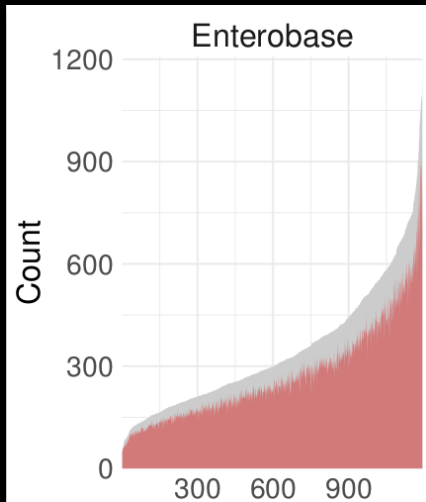
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# annofilt performance



# Publications

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# Articles



## Published:

- Dessì, et al. "Thermophilic versus mesophilic dark fermentation in xylose-fed fluidised bed reactors: Biohydrogen production and active microbial community" International Journal of Hydrogen Energy, 43(11), 10.1016/j.ijhydene.2018.01.158. 2018
- Waters**, et al. "riboSeed: leveraging prokaryotic genomic architecture to assemble across ribosomal regions" Nucleic Acids Research, 10.1093/nar/gky212. 2018

## In Preparation:

- Nolan, et al. "Pathogen survival in anaerobic co-digestion of slurry with organic waste" Frontiers

## Submitted:

- Somorin, et al. "Loss of Curli in Soil-Persistent *Escherichia coli* is Mediated by a c-di-GMP Signalling Defect and suggests biofilm-independent niche specialisation" Frontiers

Other

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# Taught Modules



- *Probability (fa16)*
- *Probabalistic Models for Bio (sp17)*

# Supervisor Assessed Modules



- o Visit JHI
- o Journal Club
- o Organize Meeting
- o Present Poster
- o Present Talk
- o Write a Paper
- o Workshop Participation

# Assorted Activities



- Organized and co-led Software Carpentry Workshop
- Participated at an NCBI Biohackathon
- Presented talk at Environ 2017
- Presented poster and talk at SGM 2018
- Contributed to open-source projects (as needed)