

Phylogenetic Diversity of Coliform Isolates in USA

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Phylogenetic Classification

- group organisms together based on probable evolutionary relationships
- usually based on direct comparison of genetic material and gene products

Constructing a “phylogenetic tree” for mickey mouse

Finding differences in their appearance (phenotype) or genetic code (genotype)

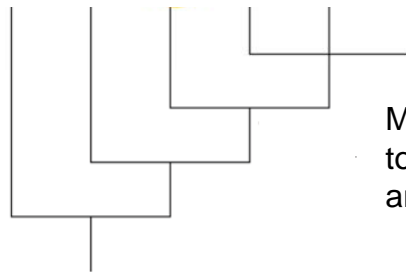


- Define characters: wearing shoes or not (1,0), color of skin (1,0), nose tilt (1,4)...
- Establish matrix:

M1	0	0	0	0	0	1
M2	1	0	1	0	1	1
M3	1	1	3	1	?	1
M4	1	1	3	?	?	1
M5	1	1	4	1	2	1
MX	1	1	2	1	2	1
- Calculate distance within the matrix
- Construct phylogenetic tree

From Professor Gustavo Caetano-Anolles

Constructing a phylogenetic tree based on phenotype



Mickey mouse X is more closer to M3, M4, and M5 than to M1 and M2

From Professor Gustavo Caetano-Anolles

Constructing a phylogenetic tree based on genotype

1. TTGACATGCCGGGAACCG
2. TTCGACATGCCGGTGGTAACGCCG
3. TTGACATCCTAGGAACGCCG
4. TTCGACATGCTAGGGAACCCG

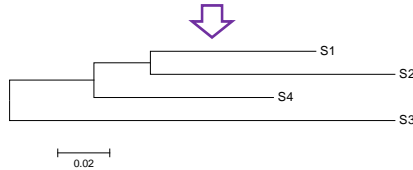
Characters: genetic code

1. TT-GACATGCCGG--GG-AA--CCG
2. TTCGACATGCCGGT-GGTAACGCCG
3. TT-GACAT-CC--TAGG-AAAGGCCG
4. TTCGACATG-C-TAGGGAACCCG

Establish matrix: alignment

	S1	S2	S3	S4
S1	0.00			
S2	0.16	0.00		
S3	0.26	0.30	0.00	
S4	0.16	0.18	0.25	0.00

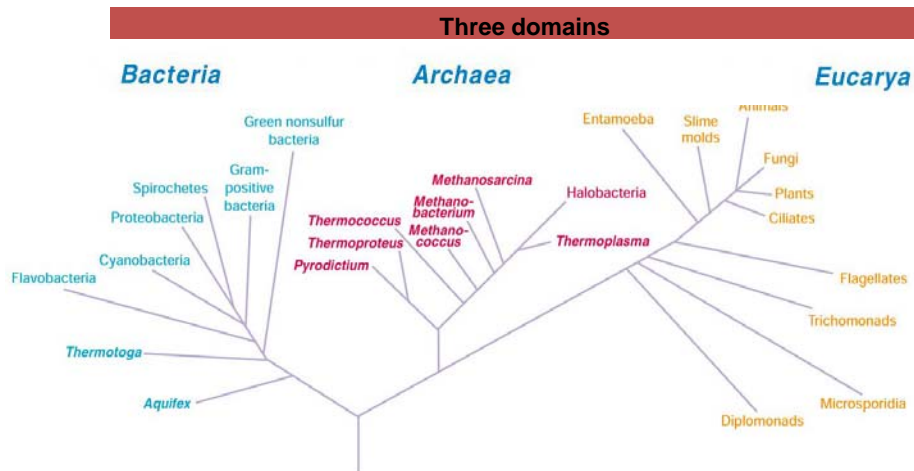
Calculate distance within the matrix



Construct phylogenetic tree

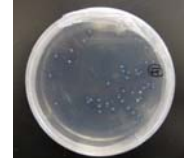
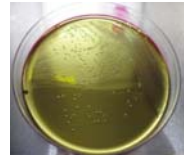
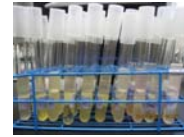
Phylogenetic systems

1. Direct comparison of genetic material and gene products such as rRNA and proteins
2. Three domains (Carl Woese 1970s)
3. Most accepted by microbiologists



EPA approved coliform tests

Technique	Medium	Working basis
MTF	LTB-BGLB broth	Lactose degradation and acid and gas production
	P-A broth-BGLB broth	Lactose degradation and acid and gas production
MF technique	m-Endo or LES-Endo-LTB, BGLB	Lactose degradation and metallic (golden) sheen production
	MI medium	MUGal-IBDG
	M-coliBlue 24°	Fermentation to produce color change. Use of X-Gluc to detect E. coli
	Chromocult®	Salmon-Gal and X-Gluc
	Coliscan® C MF	Salmon-Gal and X-Gluc
P/A tests	Colilert®	ONPG-MUG
	Colilert-18®	ONPG-MUG
	Colisure®	CPRG-MUG
	ReadiCult®	X-Gal-MUG
	Coliforms 100	X-Gal-MUG
	E*Colite®	X-Gal-MUG
	Colitag™	ONPG-MUG

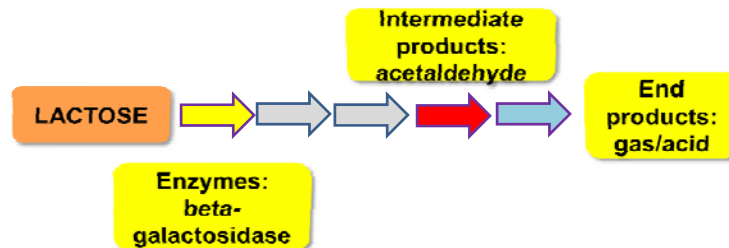
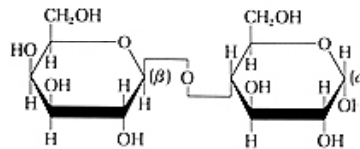


Use of lactose fermentation to detect coliforms

Lactose



Lactose is a disaccharide composed of D-galactose and D-glucose.



Leclerc, H. et al. 2001. Annu. Rev. Microbiol. 55:201-234.

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EPA approved analytical methods

Medium	Working basis
MI medium M-coliBlue 24® ★ Chromocult® Coliscan® C MF Colilert® Colilert-18® Colisure® ReadyCult® Coliforms 100 E*Colite® Colitag™	Enzymes: beta-galactosidase (total coliform) & beta-glucuronidase (<i>E. coli</i>)
M-Endo	Intermediate product (acetaldehyde)
Lauryl Tryptose Broth (LTB) Presence-Absence (P/A) broth	End product: acid and gas production

★ For detection of total coliform, it targets on end product

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What are coliforms?

Definition of Total Coliform

- **Original:**
 - “Gram-negative, facultative anaerobic rod-shaped bacteria that ferment lactose to produce acid and gas within 48 h at 35-37° C”.
- **Lactose Tryptose Broth:**
 - “all facultative anaerobic, Gram negative, **non-spore-forming**, rod shaped bacteria that ferment lactose with gas and acid formation within 48 h at 35°C”
- **Membrane test:**
 - “... that develop **red colonies with a metallic (golden) sheen** within 24h at 35°C on an Endo-type medium containing lactose.”
- **P/A (Beta-galactosidase) test:**
 - “A member of the *Enterobacteriaceae* which produces the enzyme **β -D-galactosidase**”
 - “a facultative, Gram-negative organism which grows at 35°C in the presence of bile salts, is **cytochrome oxidase negative** and produces the enzyme β -D-galactosidase”
 - “*Escherichia coli*: bacteria giving a positive coliform response and possessing the enzyme **β -D-glucuronidase**”.

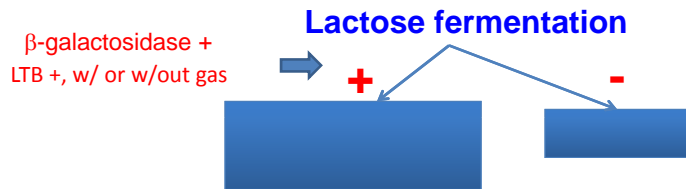
WaterRF report 4024

Classification by Bergey's Manual of Systematic Bacteriology, 2nd Edition

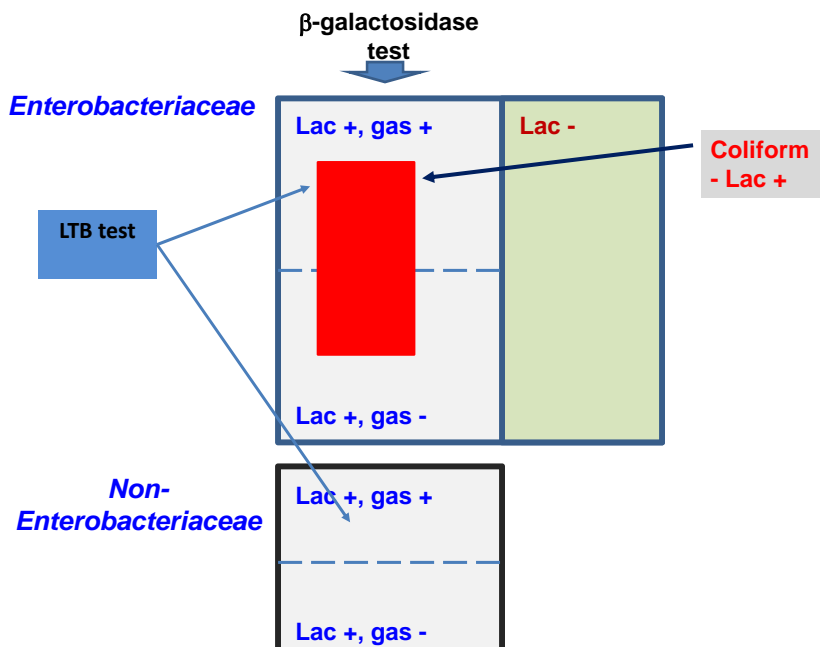
Facultatively Anaerobic Gram-Negative Rods

- Family:
 - *Enterobacteriaceae* (44 genera and 176 names species → 60 genera and 160 species in the next edition?)
 - *Arsenophonus, Budvicia, Buttiauxella, Cedecea, Citrobacter, Edwardsiella, Enterobacter, Erwinia, Escherichia, Ewingella, Hafnia, Klebsiella, Kluyvera, Leclercia, Leminorella, Moellerella, Morganella, Obesumbacterium, Pantoea, Pragia, Proteus, Providencia, Rahnella, Salmonella, Serratia, Shigella, Tatumella, Xenorhabdus, Yersinia, Yokenella*
 - Classification
 - Morphological traits
 - Phenotypic traits
 - Biochemical tests including substrate utilization, antibiotics resistant, etc...
 - Genetic traits
 - 16S rRNA gene analysis (species cutoff, 97% similar or higher)

Family *Enterobacteriaceae* – Phenotypic classification



Combining phylogeny with phenotypes in TC testing

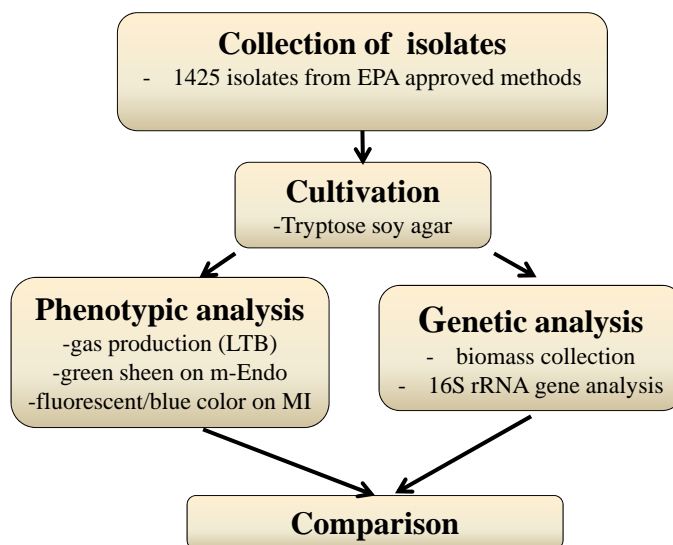


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Experiment objectives

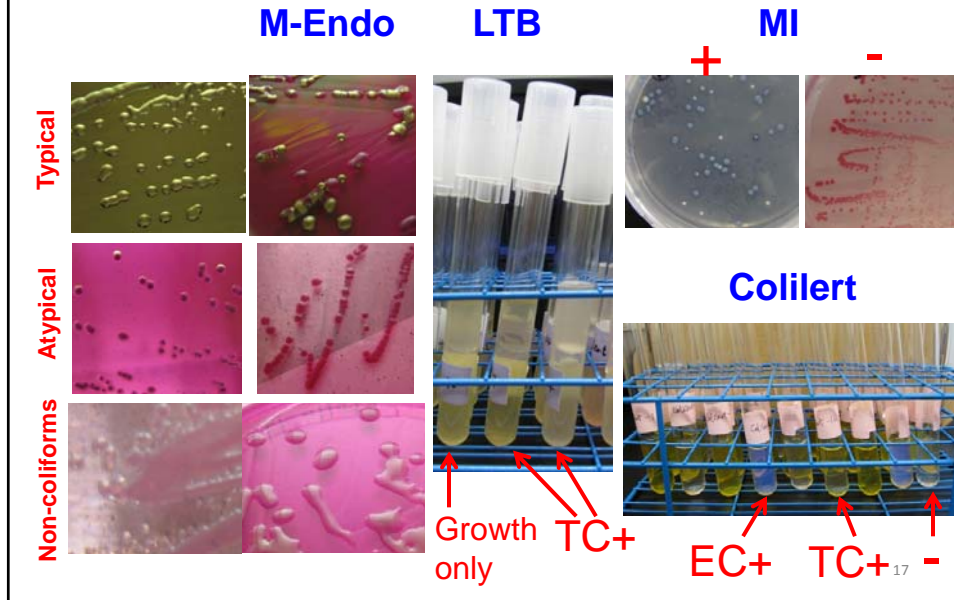
- What are coliforms from the view of phylogeny?
- How can phylogeny information of coliforms be related to current coliform tests?
 - True coliforms
 - False '+' and false '-'

Experimental procedure

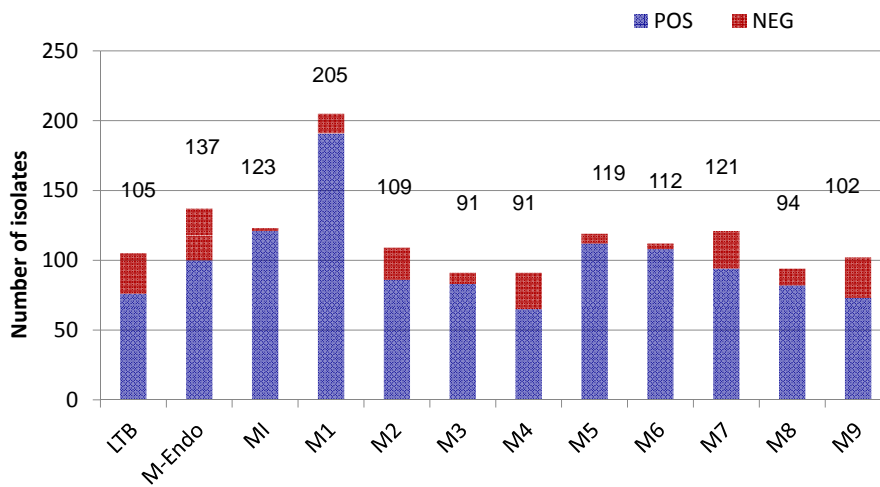


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Responses on different media



1425 bacterial isolates obtained from 12 different coliform tests



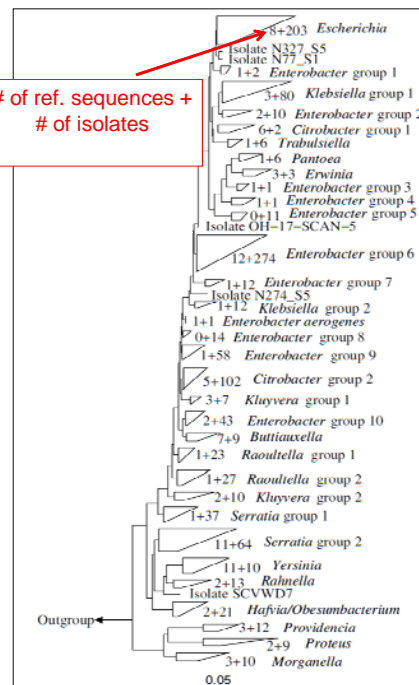
Criteria used for forming phylogenetic clusters

- ① 16S rRNA gene sequences that can form a cohesive cluster with known coliform bacteria at the **species** level within the family *Enterobacteriaceae*
- ② To group sequences that form a cohesive cluster with reference sequences at a known **genus** level.
 - *Escherichia* cluster
 - *Buttiauxella* cluster.
- ③ For sequences from a same genus that could not form a cohesive group, **more than one sub-cluster** was formed within each genus.
 - *Enterobacter*, *Citrobacter*, *Klebsiella*, *Raoultella*, *Serratia*, and *Kluyvera*
 - sequence similarity: 98.1% to 99.5%, higher than 'species' definition in microbial taxonomy.

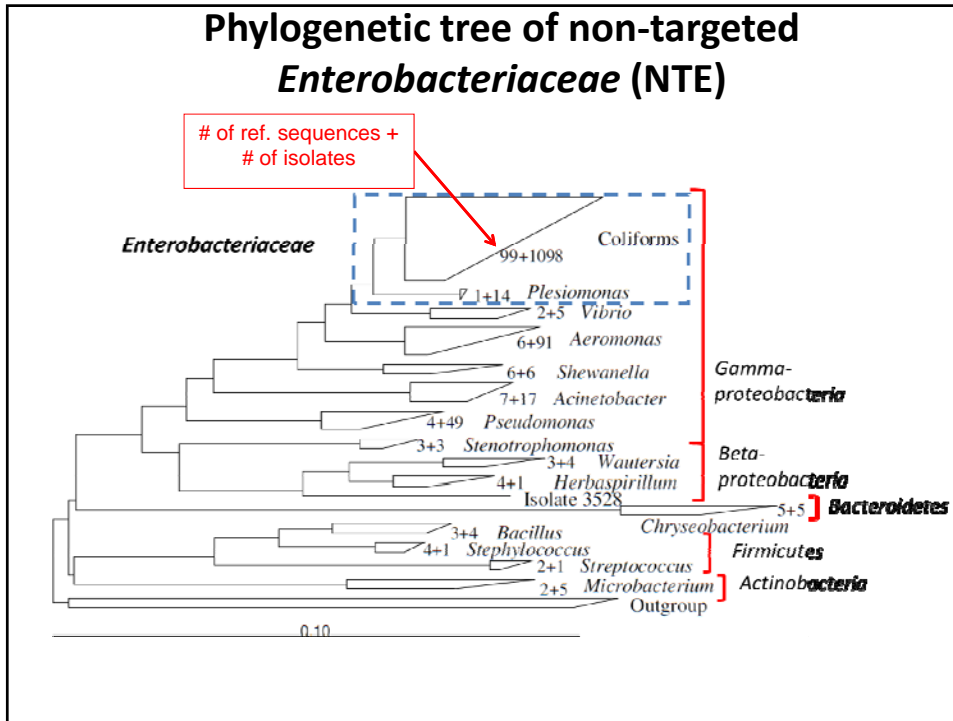
Phylogenetic tree of targeted *Enterobacteriaceae* (TE)

- **32** groups are formed.
- Some genera contain more than 1 sub-cluster.

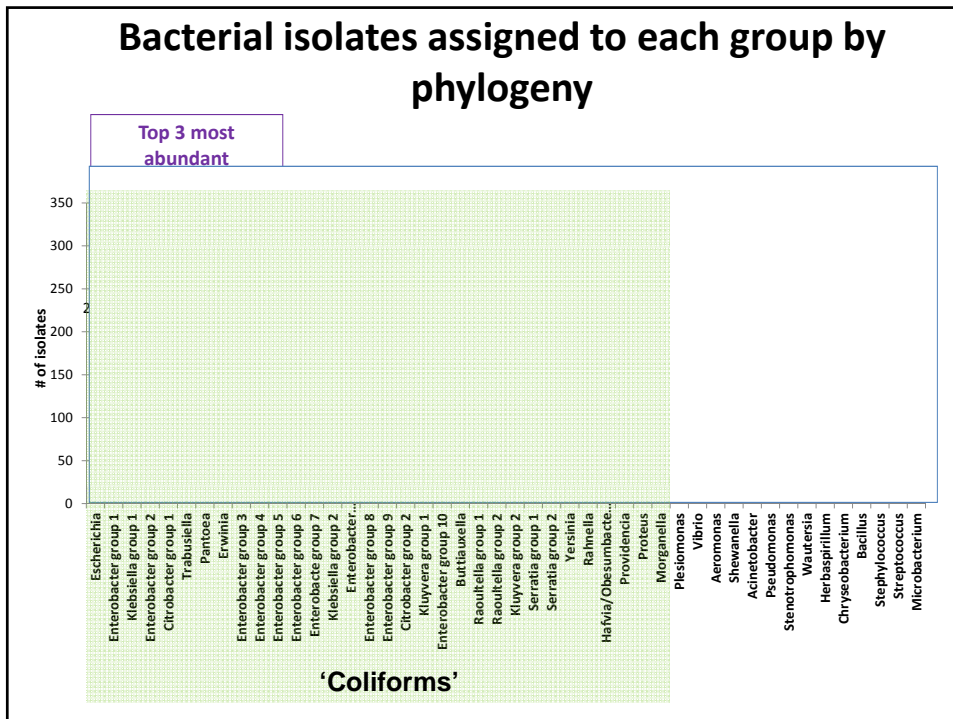
Genera	Group #
<i>Enterobacter</i>	10
<i>Klebsiella</i>	2
<i>Citrobacter</i>	2
<i>Kluyvera</i>	2
<i>Raoultella</i>	2
<i>Serratia</i>	2



Phylogenetic tree of non-targeted *Enterobacteriaceae* (NTE)



Bacterial isolates assigned to each group by phylogeny




Principles	Methods	POS (%)		NEG (%)	
		TE	NTE	TE	NTE
Enzymes	M1				
	M4				
	M5				
	M1				
	M2				
	M3				
	M7				
	M8				
	M9				
Intermediate product	m-Endo				
End product	LTB				
	M6				

Membrane-based
 Liquid-based

Summary

- 'coliforms':
 - at least 32 phylogenetic clusters.
 - Dominant phylogenetic clusters include for example *Enterobacter*, *Klebsiella*, *Pantoea*, and *Serratia*.
- 'non-coliforms':
 - at least 14 phylogenetic groups within and outside of the *Enterobacteriaceae*.
- Phylogenetic clustering together with coliform tests provides an effective way to identify coliform isolates that cause false-positive or false-negative results with different coliform testing methods.
 - False '+' rate: 0 – 67%
 - false '-' rate: 12 – 100%

Acknowledgement

- Water samples and coliform isolates provided by anonymous water utilities and testing laboratories
- WaterRF #4300 
The logo for the Water Research Foundation, featuring a stylized globe with water droplets and the text "WATER RESEARCH FOUNDATION" and "ADVANCING THE SCIENCE OF WATER".
- WaterRF #4371