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## Concomitant Antibiotic and Mercury Resistance among Gastrointestinal Microflora of Feral Brook Trout, *Salvelinus fontinalis*

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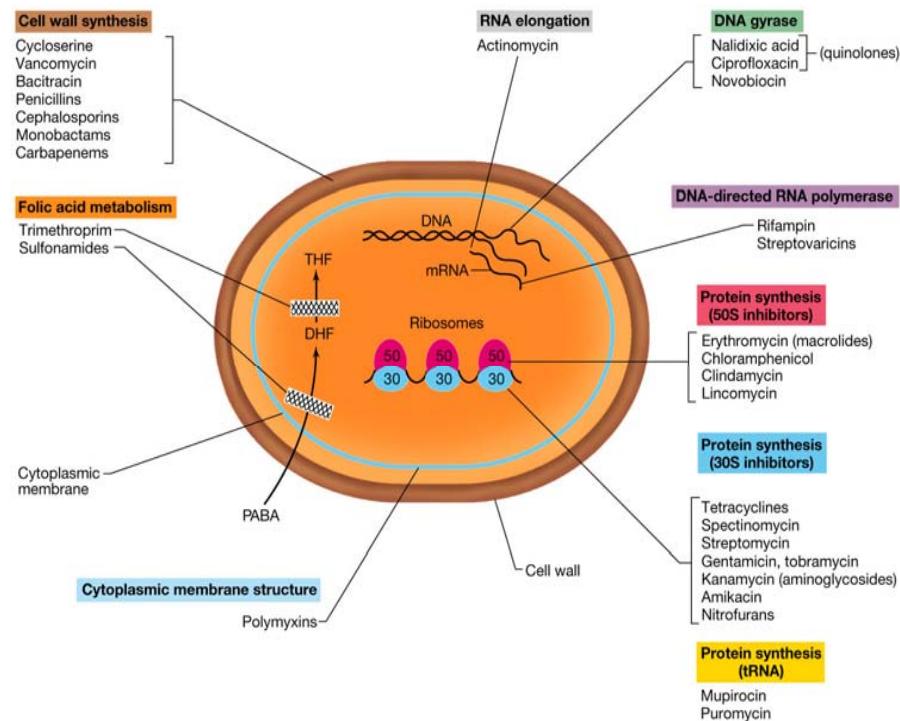
# Concomitant Antibiotic and Mercury Resistance among Gastrointestinal Microflora of Feral Brook Trout, *Salvelinus fontinalis*



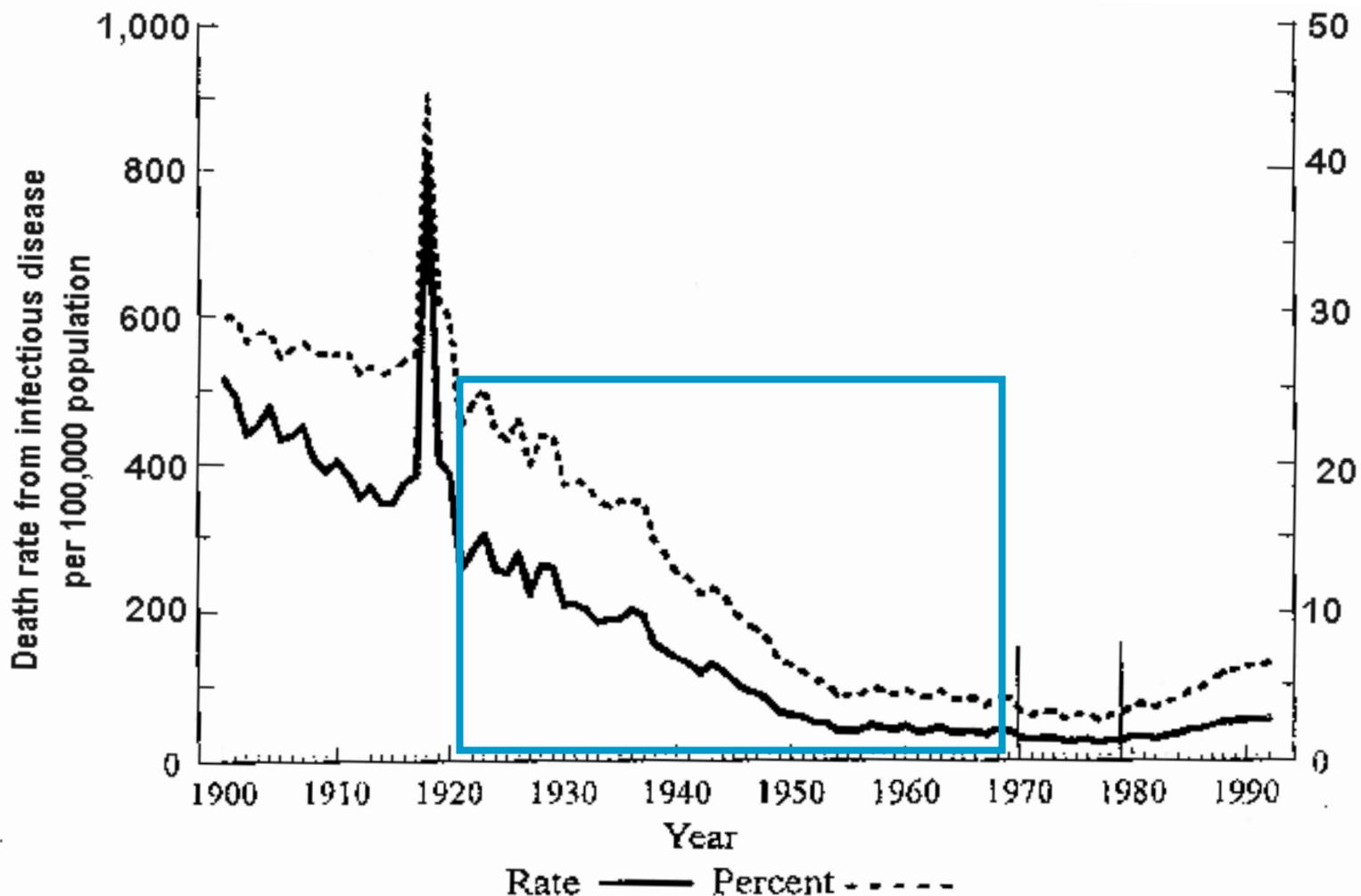
Matt Meredith  
Colby College  
Undergraduate Research Symposium  
May 5<sup>th</sup>, 2006

# Antibiotics (Ab)

- “Low molecular weight compounds produced by microorganisms to inhibit the growth of other microorganisms”
- Selective toxicity
- Confer competitive advantage
- Antimicrobial = synthetic Ab

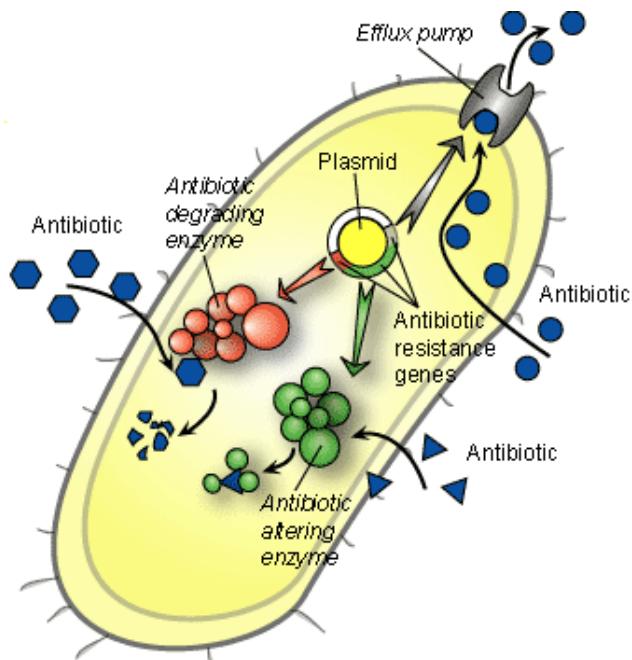


# US Infectious Disease Mortality in the 20<sup>th</sup> Century

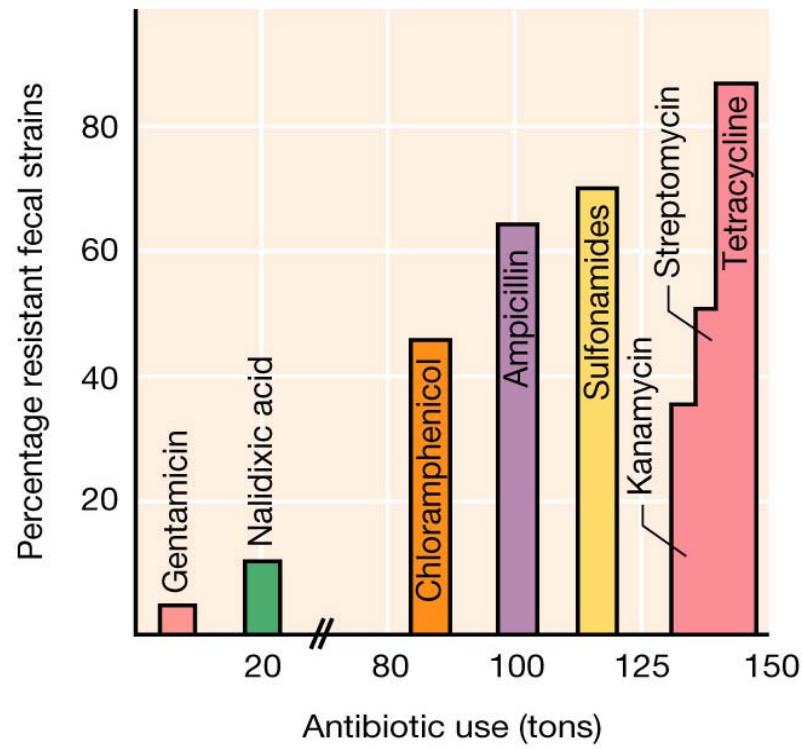


# Antibiotic Resistance ( $\text{Ab}^r$ )

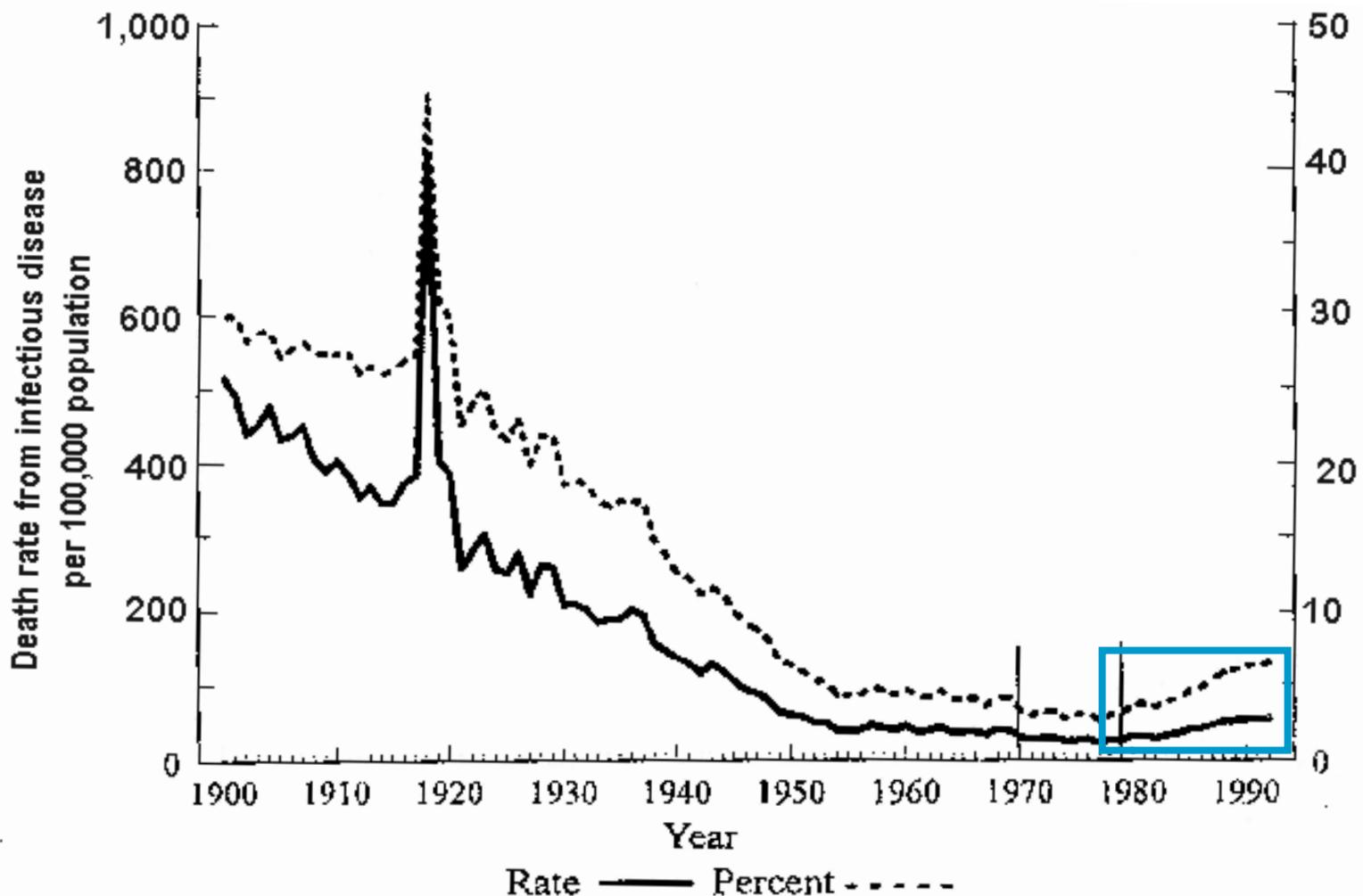
- $\text{Ab}^r$  gene origin in  $\text{Ab}^-$  producing organisms to prevent self-destruction
- Selected by excessive and imprudent use of Ab



# Ab Selection for Abr

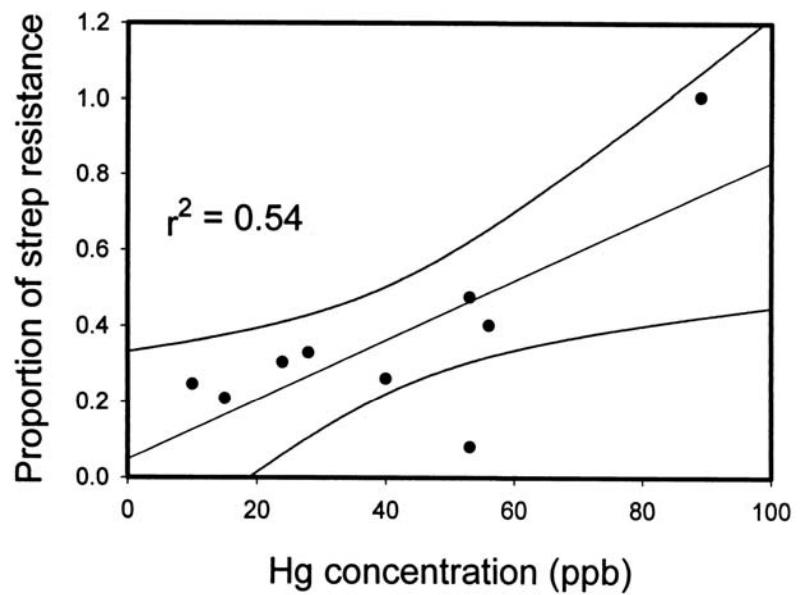


# US Infectious Disease Mortality in the 20<sup>th</sup> Century



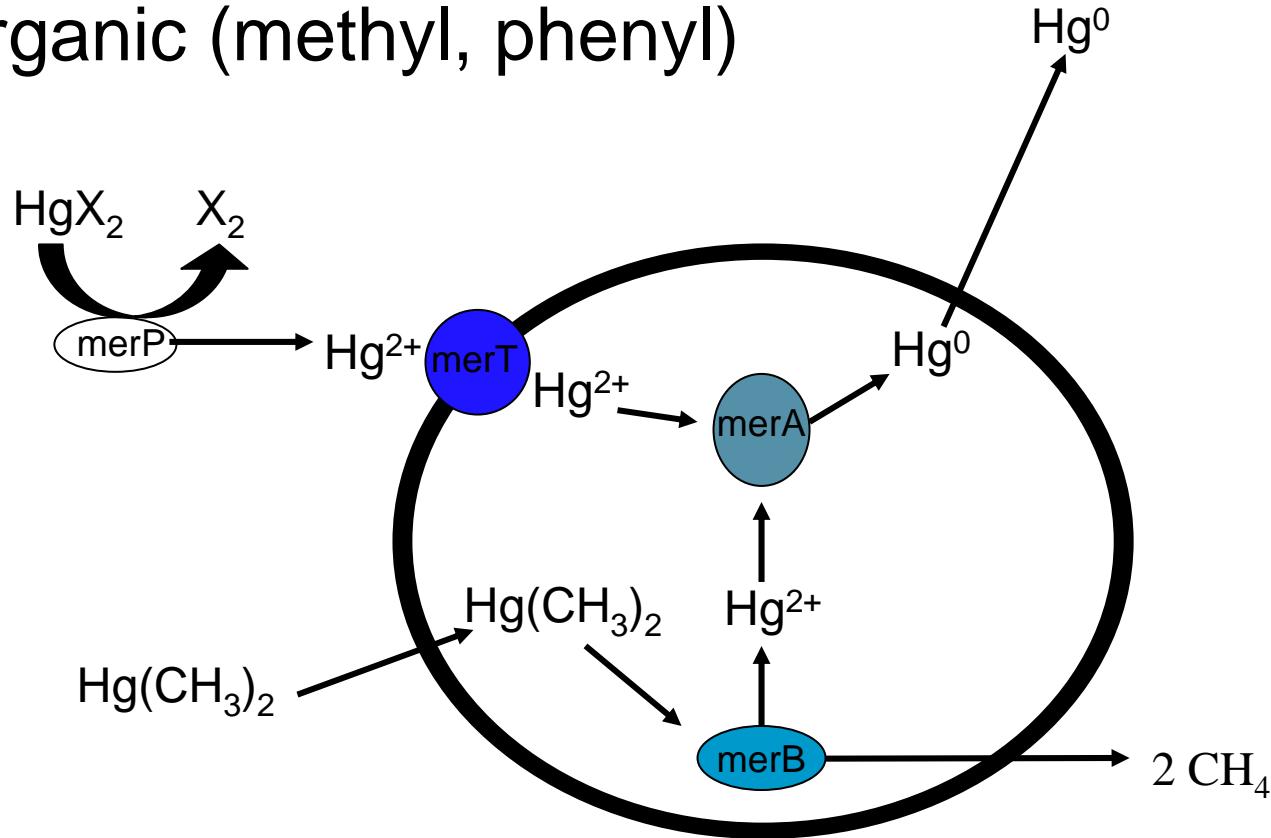
# Ab<sup>r</sup> and Mercury Resistance (Hg<sup>r</sup>)

- Hg as indirect selective agent for Ab<sup>r</sup>
- Ab<sup>r</sup> and Hg<sup>r</sup> proximate on mobile genetic element?

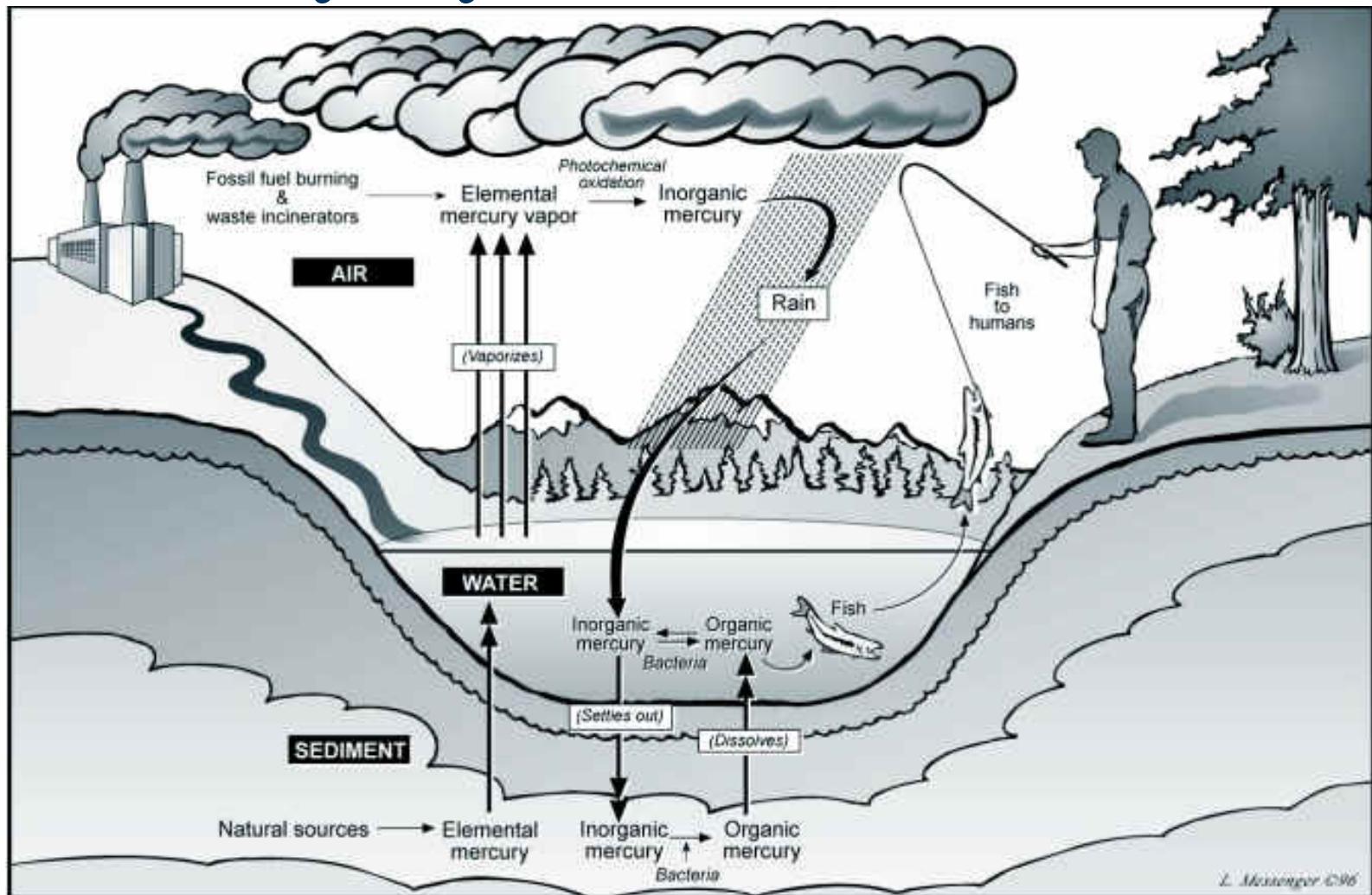


# Forms of Hg and Hg<sup>r</sup>

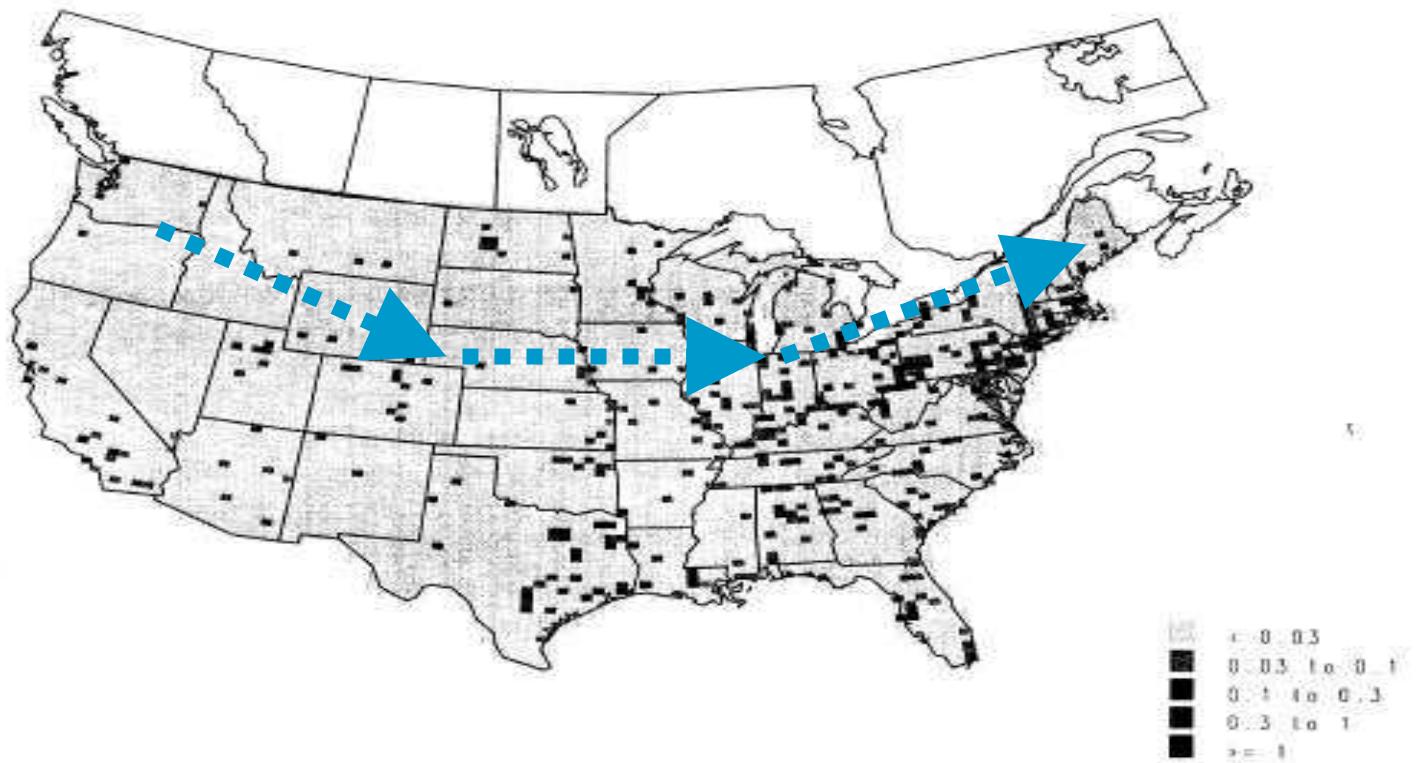
- Elemental ( $\text{Hg}^0$ )
- Ionic ( $\text{Hg}^{2+}$ )
- Organic (methyl, phenyl)



# Mercury Cycle



# Mercury in Maine



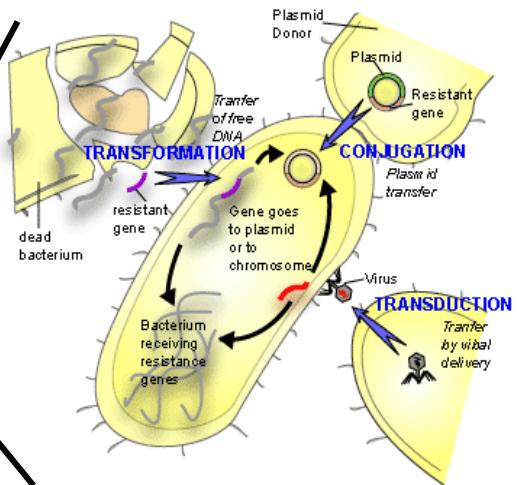
# Lake Kennebago, Maine

- $1.5 \text{ ng l}^{-1}$  total Hg
- Pristine
- Oligotrophic
- No Ab-usage



# Fish GI tract system

- Commensal populations stable
- Commensal gene reservoir stable
- GI bacterial commensals exposed to Hg
- Simultaneous selection for Hg<sup>r</sup> and Ab<sup>r</sup>?
  - Feral brook trout: *Salvelinus fontinalis*



# Isolate Isolation

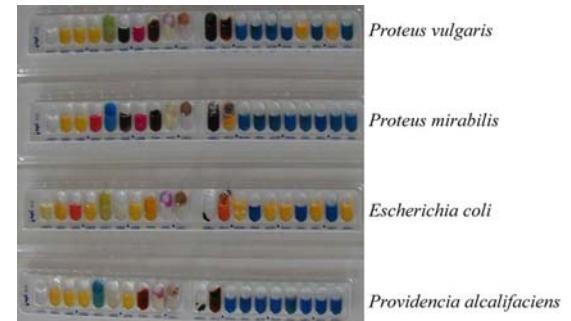
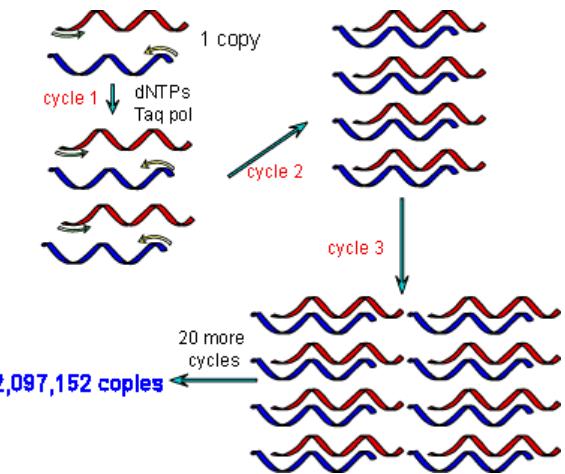
- FBT dissection, GI tract extraction
- GI tract/ingesta dilution in PO<sub>4</sub> buffer
- Serial dilution on TSA plating media containing 0 and 25 µM HgCl<sub>2</sub>
- Incubation at 22°C

# 29 Bacterial Isolates

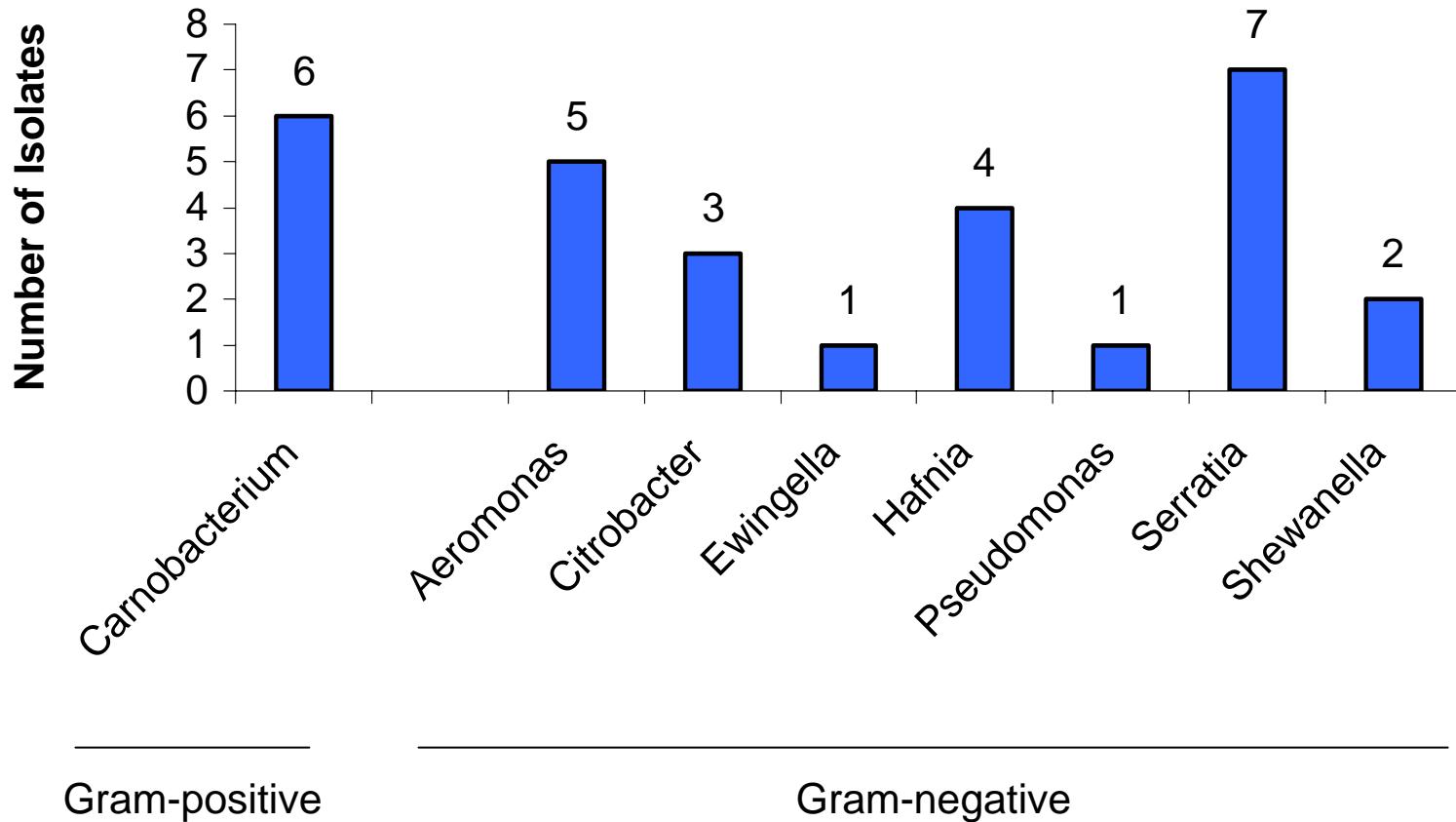
- August 2004: 3 fish – 6 FBT1
- September 2005: 4 fish – 9 FBT3
- October 2005: 2 fish – 14 FBT4

# Isolate Identification

- Genomic DNA extraction
- 16S rDNA PCR
- 16S rDNA sequencing
- Blastn database search
- API 20E strip

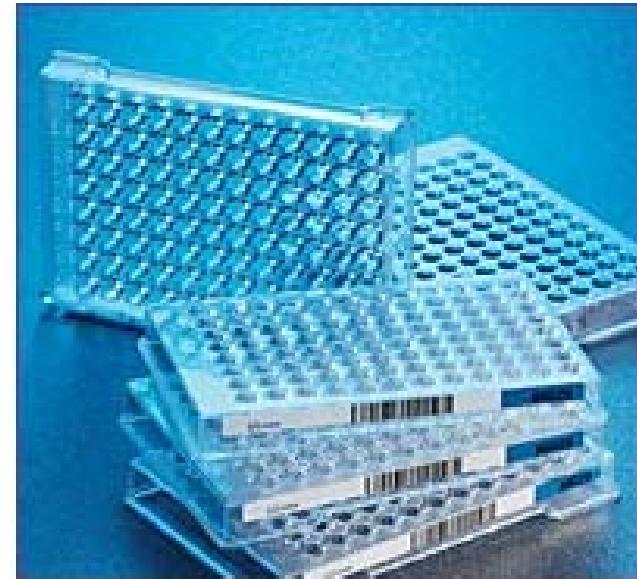


# 8 Genera



# Phenotypic Characterization of Ab<sup>r</sup>

- Ab MIC panel:
  - GPN2F (Gram-positive)
  - MJ (Gram-negative)



# Antibiogram (GPN2F)

Isolate	Genus	AMP 0.12- 16	PEN 0.06- 8	OXA+ 0.25- 8	FAZ 2- 16	AXO 8- 64	GEN 2- 16	CIP 0.5- 2	LEVO 0.25- 8	GAT 1- 8	VAN 1- 32	ERY 0.25- 8	CLI 0.5- 4	SYN 0.12- 4	LZD 0.5- 8
FBT1-19	<i>Carnobacterium</i>	>16	>8	>8	16	>64	>16	2	2		2	0.5	>4	2	2
FBT1-22	<i>Carnobacterium</i>	8	4	>8	8	>64	>16		1			0.5	>4	2	1
FBT3-9	<i>Carnobacterium</i>	0.5	1	>8	16	>64	>16	2	2	2	2		>4	1	
FBT3-14	<i>Carnobacterium</i>	1	2	>8		>64	>16		0.5		2	0.5	>4	1	
FBT4-1	<i>Carnobacterium</i>	>16	>8	>8	4	>64	>16	>2	2			0.5	>4	2	2
FBT4-18	<i>Carnobacterium</i>	4	8	>8	4	>64	>16			2		0.5	>4	2	1

AMP, ampicillin; AXO, ceftriaxone; CIP, ciprofloxacin; CLI, clindamycin; ERY, erythromycin; FAZ, cefazolin; GAT, gatifloxacin; GEN, gentamicin; LEVO, levofloxacin; LZD, linezolid; OXA+, oxacillin+2%NaCl; PEN, penicillin; SYN, quinupristin/dalfopristin; VAN, vancomycin.

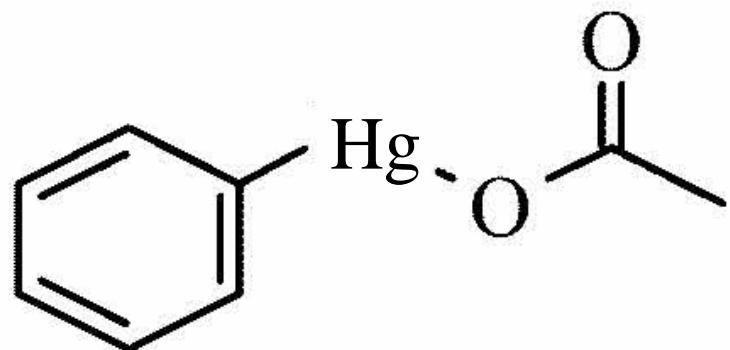
# Antibiogram (MJ)

Isolate	Genus	AMP 0.5- 16	A/S 8/4- 16/8	MEZ 4- 64	TIM 4/2- 64/2	AUG 0.5/0.25- 16/8	PIP 4- 64	FAZ 1- 16	CEP 8- 16	FOX 2- 16	TAZ 1- 16	AXO 4- 32	FUR 2- 16	AMI 4- 32	GEN 0.25- 8	TET 0.25- 8	CIP 0.06- 2	OFL 0.25- 4	LOM 0.5- 4	COT 0.5/9.5- 2/38	NIT 32- 64	FIS 0- 256
FBT1-8	<i>Serratia</i>	>16	16/8	16	8/2	16/8	8	>16	>16	>16		>16		1	>8	0.12	0.5	1		>64		
FBT1-15	<i>Ewingella</i>	>16		8		>16/8		>16	>16	8		>16		0.5	2					64		
FBT1-21	<i>Serratia</i>	>16	>16/8	32	8/2	>16/8	16	>16	>16	16		>32	>16	0.5	4					64	>256	
FBT1-23	<i>Serratia</i>	>16	>16/8	16	16/2	>16/8	16	>16	>16	16		>32	>16	0.5	8					64		
FBT3-1	<i>Serratia</i>	>16	>16/8	>64	8/2	16/8	32	>16	>16	16		>32	>16	0.5	8					64		
FBT3-2	<i>Aeromonas</i>	>16	>16/8		16/2	16/8		8						1							>256	
FBT3-5	<i>Aeromonas</i>	>16	>16/8		>64/2	16/8	8	>16	>16	16				8	2						>256	
FBT3-6	<i>Shewanella</i>					2/1		16							1	2					>256	
FBT3-8	<i>Serratia</i>	>16	>16/8	>64	8/2	16/8	>64	>16	>16	16		>32	>16	0.5	8					64		
FBT3-16	<i>Serratia</i>	>16	>16/8	64	16/2	>16/8	>64	>16	>16	16		>32	>16	0.5	8					64		
FBT3-17	<i>Hafnia</i>	>16	>16/8	8	16/2	>16/8	8	>16	>16	4	8	16		0.5	>8					>256		
FBT4-2	<i>Hafnia</i>	4		32	8/2	4/2	16	16	>16		4		4	1	>8					>2/38		
FBT4-3	<i>Aeromonas</i>	>16	>16/8	16	64/2	8/4		>16	>16					2	0.5					>256		
FBT4-5	<i>Aeromonas</i>	>16	>16/8		8/2	8/4		8						16	4	0.5						
FBT4-6	<i>Citrobacter</i>	>16		16/8	8/2	>16/8	8	>16	>16	>16			8	1	2				0.5			
FBT4-9	<i>Shewanella</i>	8			8/2	16/8		>16	>16					1	1	0.12				>256		
FBT4-10	<i>Citrobacter</i>	4				4/2		2	16	4					2					>256		
FBT4-11	<i>Hafnia</i>	>16	>16/8	8	32/2	>16/8	8	>16	>16	8	2	>16		0.5	>8					>256		
FBT4-12	<i>Citrobacter</i>	8				8/4		8	>16	4		4		0.5	4	0.12				>256		
FBT4-19	<i>Serratia</i>	>16	>16/8	64	16/2	>16/8	64	>16	>16	16		>32	>16	0.5	4					64	>256	
FBT4-21	<i>Aeromonas</i>	>16	>16/8	32	64/2	>16/8	16	>16	>16	8				8	2					>256		
FBT4-22	<i>Hafnia</i>	>16	>16/8	8	32/2	>16/8	8	>16	>16	4	8		16	0.5	>8					>256		
FBT4-23	<i>Pseudomonas</i>	>16	>16/8	>64	>64/2	>16/8	32	>16	>16	>16	>16	>32	>16	1	8	0.5	2	4	>2/38	>64	>256	

AMI, amikacin; AMP, ampicillin; A/S, ampicillin/sulbactam; AUG, amoxicillin/clavulanic acid; AXO, ceftriaxone; CEP, cephalothin; CIP, ciprofloxacin; COT, trimethoprim/sulfamethoxazole; FAZ, cefazolin; FIS, sulfisoxazole; FOX, cefoxitin; FUR, cefuroxime(parenteral); GEN, gentamicin; LOM, lomefloxacin; MEZ, mezlocillin; NIT, nitrofurantoin; OFL, ofloxacin; PIP, piperacillin; TAZ, ceftazidime; TET, tetracycline; TIM, ticarcillin/ clavulanic acid.

# Phenotypic Characterization of Hg<sup>r</sup>

- TSA plating media
- HgCl<sub>2</sub> (0-1000 µM) and phenylmercuric acetate (PMA) (0-16 µM)
- Inoculated with droplet of bacterial suspension
- Incubation at 22°C



# Hg MICs

- HgCl<sub>2</sub> : 100 µM (70%)  
250 µM (30%)
- PMA : 8 µM (10%)  
16 µM (20%)  
>16 µM (70%)

# Isolate Exposure to Hg

- Lake Kennebago water :  $1.5 \text{ ng l}^{-1}$  total Hg
- *S. fontinalis* muscle tissue :  $395 \text{ ng g}^{-1}$
- *S. fontinalis* diet (e.g. crawfish)

# Hg-susceptible isolates?

- Population counts on 0 and 25 µM HgCl<sub>2</sub>
- TSA plating media inoculated with GI/ingesta-PO<sub>4</sub> buffer suspension
- Triplicate
- Incubation at 22°C
- No significant difference between culturable bacterial growth on 0 and 25 µM HgCl<sub>2</sub> by paired Student's *t* test (P=0.088, df=5)

# Genotypic Characterization of Hg<sup>r</sup>

- *merA* PCR amplification
- Two primer sets: long and short
- Long set: Amplification in 12 isolates
- Short set: Amplification and sequencing in 5 isolates

# *merA* Amplification

## Gram Negative Isolates

Isolate	Genus
FBT1-8	<i>Serratia</i>
FBT1-15	<i>Ewingella</i>
FBT1-21	<i>Serratia</i>
FBT1-23	<i>Serratia</i>
FBT3-1	<i>Serratia</i>
FBT3-2	<i>Aeromonas</i>
FBT3-5	<i>Aeromonas</i>
FBT3-6	<i>Shewanella</i>
FBT3-8	<i>Serratia</i>
FBT3-16	<i>Serratia</i>
FBT3-17	<i>Hafnia</i>
FBT4-2	<i>Hafnia</i>
FBT4-3	<i>Aeromonas</i>
FBT4-5	<i>Aeromonas</i>
FBT4-6	<i>Citrobacter</i>
FBT4-9	<i>Shewanella</i>
FBT4-10	<i>Citrobacter</i>
FBT4-11	<i>Hafnia</i>
FBT4-12	<i>Citrobacter</i>
FBT4-19	<i>Serratia</i>
FBT4-21	<i>Aeromonas</i>
FBT4-22	<i>Hafnia</i>
FBT4-23	<i>Pseudomonas</i>

## Gram Positive Isolates

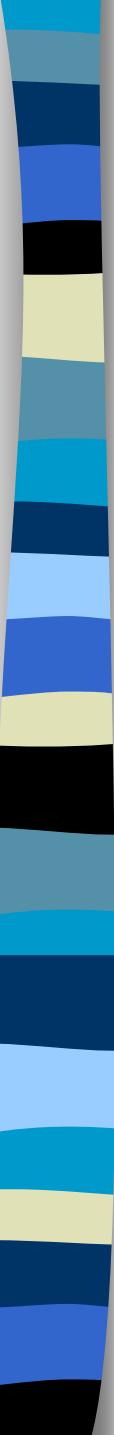
Isolate	Genus
FBT1-19	<i>Carnobacterium</i>
FBT1-22	<i>Carnobacterium</i>
FBT3-9	<i>Carnobacterium</i>
FBT3-14	<i>Carnobacterium</i>
FBT4-1	<i>Carnobacterium</i>
FBT4-18	<i>Carnobacterium</i>

Sequence  
Similarity

FBT1-21	86%
FBT1-23	85%
FBT3-1	84%
FBT3-8	91%
FBT3-16	89%

# Conclusions

- Presence of Ab<sup>r</sup> in absence of Ab-use
- Presence of Hg<sup>r</sup> due to Hg exposure
- Established relationship between Hg<sup>r</sup> and Ab<sup>r</sup>
- Hg likely indirect selective agent for Ab<sup>r</sup> determinants



# Acknowledgments

- Frank Fekete (Colby College)
- Justin Guay (Colby College)
- Erin Parry (Colby College)
- Russell Danner (Maine Department of Inland Fisheries and Wildlife)
- Tamar Barkay (Rutgers University)



# Questions?



Thank you.

# Ab<sup>r</sup> Prevalence

- Methicillin-resistant *Staphylococcus aureus* strains endemic in hospitals and are increasing in other settings such as locker rooms.
- Vancomycin-resistance *S. aureus* and enterococci emerged in 2002
- Incidence of sepsis has nearly tripled between 1979 to 2000.

*STAPHYLOCOCCUS AUREUS*  
VS. PENICILLIN



*ENTEROCOCCUS FAECIUM*  
VS. CIPROFLOXACIN (CIPRO)



*STREPTOCOCCUS PNEUMONIAE*  
VS. TETRACYCLINE



*STAPHYLOCOCCUS AUREUS*  
VS. METHICILLIN



*ENTEROCOCCUS FAECIUM*  
VS. AMPICILLIN



*STREPTOCOCCUS PNEUMONIAE*  
VS. PENICILLIN

