

### Why is horizontal gene transfer so important?

# Additional Save Lives Additional Save Lives

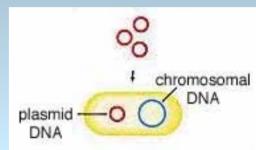
- Resistance acquired by mutation
  - Usually need multiple mutations in essential bacterial genes
  - One step mutations can make bacteria less fit, need compensating mutations
    (R. Lenski, Ciba Found. Symp. 1997. 207:131-140)
  - Many random mutations will be deleterious

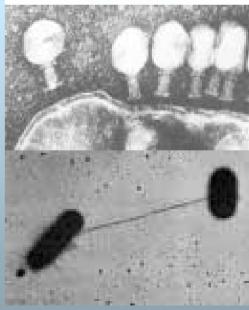
### Why is horizontal gene transfer so important?

- Resistance acquired by horizontal transfer
  - New gene unlikely to be deleterious
  - Promoter mutations can lead to expression

### Types of horizontal gene transfer events

- Narrow host range between closely related bacteria
  - Transformation
  - Phage transduction
- Broad host range cross species and genus lines
  - Conjugation can occur within an hour





### Conjugal gene transfer elements

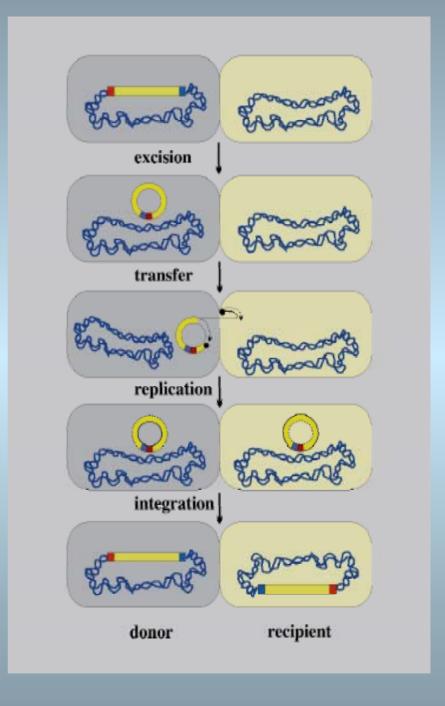
#### Plasmids

- Can acquire new resistance genes
  - Transposons
  - Integrons
- Bacteria acquire compensating mutations that lead to greater fitness

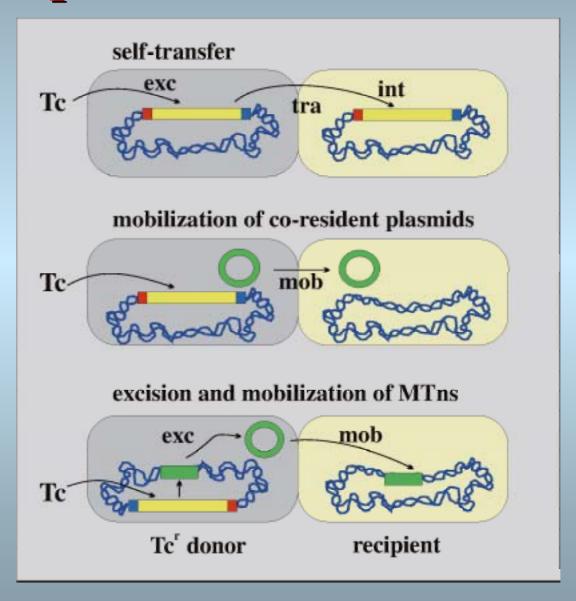
#### Conjugative transposons

- Can be as promiscuous as plasmids
- Stably integrated into the genome not easily lost

Steps involved in the intercellular transfer of a CTn



#### Transfer potential of Bacteroides CTns



### Stability of plasmids and conjugative transposons

#### Naturally occurring plasmids

- Usually have a low copy number enhances stability
- Often encode plasmid maintenance systems

#### Conjugative transposons

Integrated in the chromosome

### Stability of plasmids and conjugative transposons

- Inducible resistance genes and inducible transfer decrease fitness toll
  - Cessation of antibiotic treatment or changing antibiotics may not cause resistant strains to disappear
  - Even when incidence of resistance decreases, a
     "hard core" of strains persist

### Can resistance be prevented?

- End abuse and overuse of antibiotics
- Treating a patient with antibiotics means treating the patient's microflora
  - Can members of the microflora serve as reservoirs of resistance genes?



## Antibiotic resistance gene distribution in Community and Hospital Isolates of *Bacteroides* spp.

		Percentage of Isolates Carrying					
Source of Isolates	No.	tetQ	ermF	ermG	cfxA	CTn	NBU
Community (pre-1970)	69	32	0	0	0	42	51
Clinical (pre-1970)	23	22	9	0	4	30	35
Community (1996-1997)	102	81	15	8	3	84	74
Clinical (1980-1995)	87	86	30	18	14	85	84

#### Horizontal Gene Transfer

Bacillus spp.

ermG

Bacteroides spp.

Enterococcus spp. Staphylococcus spp. Streptococcus spp.

ermB

Bacteroides spp.

Enterococcus spp.
Staphylococcus spp.
Streptococcus spp.
Actinomyces spp.
Bifidobacterium spp.

tetM

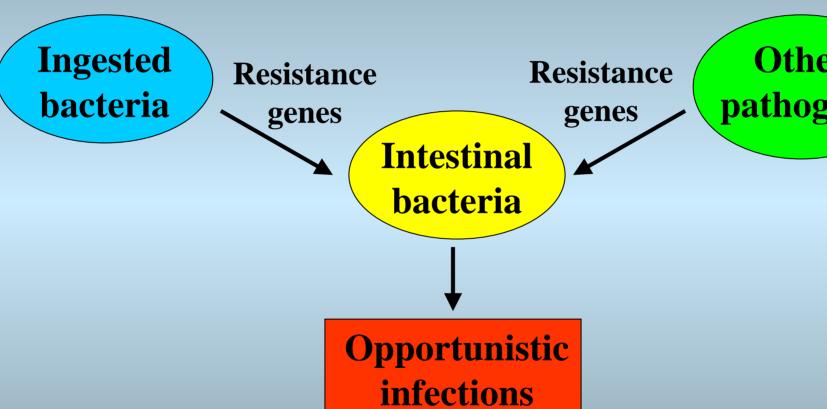
Campylobacter spp.
Fusobacterium nucleatum
Gardenella vaginalis
Haemophilus spp.
Neisseria spp.
Veillonella spp.

7

ermF tetQ

Bacteroides spp.

### Summary



Other pathogens