

# Chip Aquadro

- Mutation
- Drift
- Natural Selection
- Recombination
- Population Subdivision

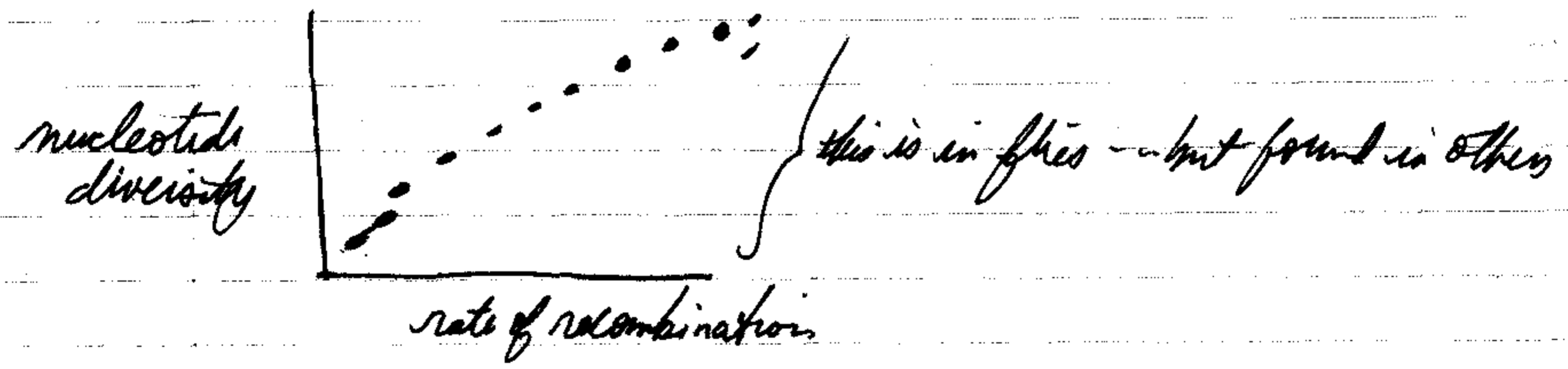
SPENCER

## D. melanogaster

- polyploid chromosomes allow genome map
- genetics dense

## Recombination

- rate per band varies a great deal
- ~~rate suppressed at telomeres/centromeres~~  
some
- rate suppressed at some telomeres
- rate suppressed at centromeres



## Why?

- ① less fit / constraint in high recomb. regions
- ② ~~recombination~~ recombination is mutagenic
- ③ selective sweeps (reduces variation)
- ④ deleterious mutations selected against  
reduces variation

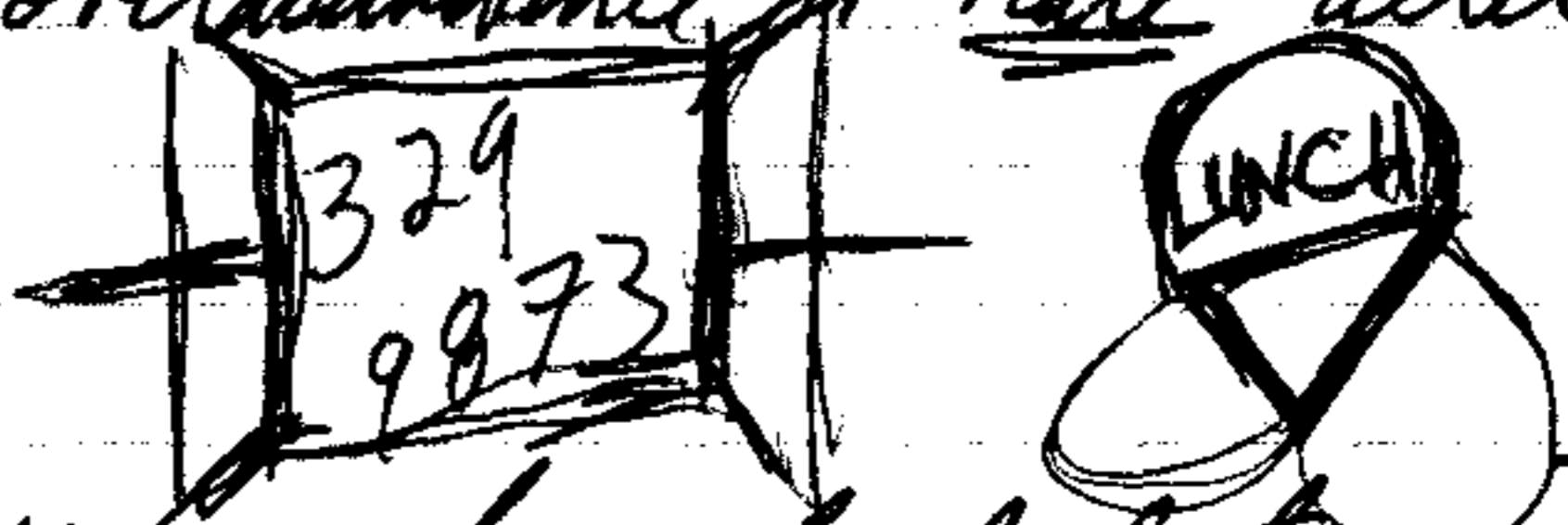
For 142

predict that there should be a <sup>'x'</sup> correlation  
between interspecies divergence  
and recombination

- this is not the case

$$\text{Heterozygosity} = 4Ne\mu$$

① selective sweeps (shorten coalescence time)  
should have overabundance of rare alleles



④ only chromosomal regions free of deleterious  
mutations persist in populations

DOES NOT  
VARIABILITY

AFFECT

~~MUTATION~~  
RECOMB.  
RATE

GABA

- mutations resistant to pesticides  
- levels of polymorphism v. low, around  
site of resistance mutations

IF THE  
PATTERN  
OF

RECOMB.  
CONSERVED.

Xanthine Dehydrogenase