

Correspondance between genotype and phenotype

Genotype

Phenotype

non-canalised/robust/buffered against internal or environmental noise

canalised/robust/buffered against genetic variations

- How stable are developmental processes when faced with internal/environmental/genetic variations ?
- What is the nature of genotypic changes that modify developmental processes?
- How do they propagate at the phenotypic level through the hierarchy of biological organisation?

Hierarchy of biological organization

DNA base
Molecule
my favorite module → Cell
Organism

... and some modularity inbetween all levels....

Modularity helps to describe morphological organization and developmental dynamics

How do these modules behave in evolution?

MODULAR ORGANIZATION OF DEVELOPMENT

- (1) Supracellular modules Organs
Interacting cells in development
- (2) Modules of interacting molecules

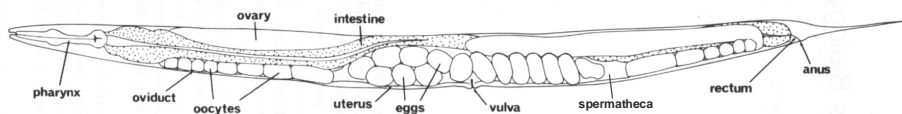
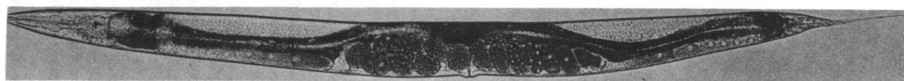
MODULAR ORGANIZATION IN EVOLUTION

- (3) Evolution of a supracellular network module - breakage
- (4) Evolution of modules within the organism is facilitated by modularity of gene organization
- (5) Morphological evolution and the body plan

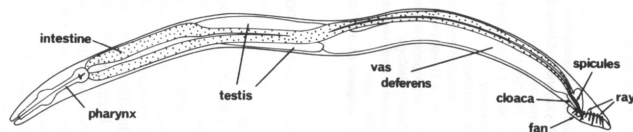
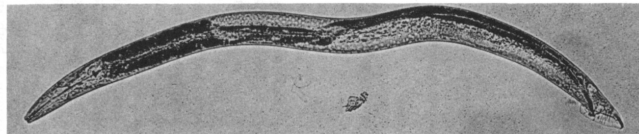
... and on nematode vulva evolution if time permits..

Caenorhabditis elegans

Hermaphrodite



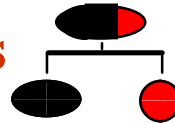
Male



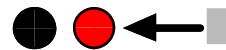
CELL SPECIFICATION MECHANISMS IN DEVELOPMENT

Segregation of determinants at mitosis

Visualization:
asymmetric cell size
segregation of determinant
(P granules, PAR-1, etc.)



Intercellular signaling

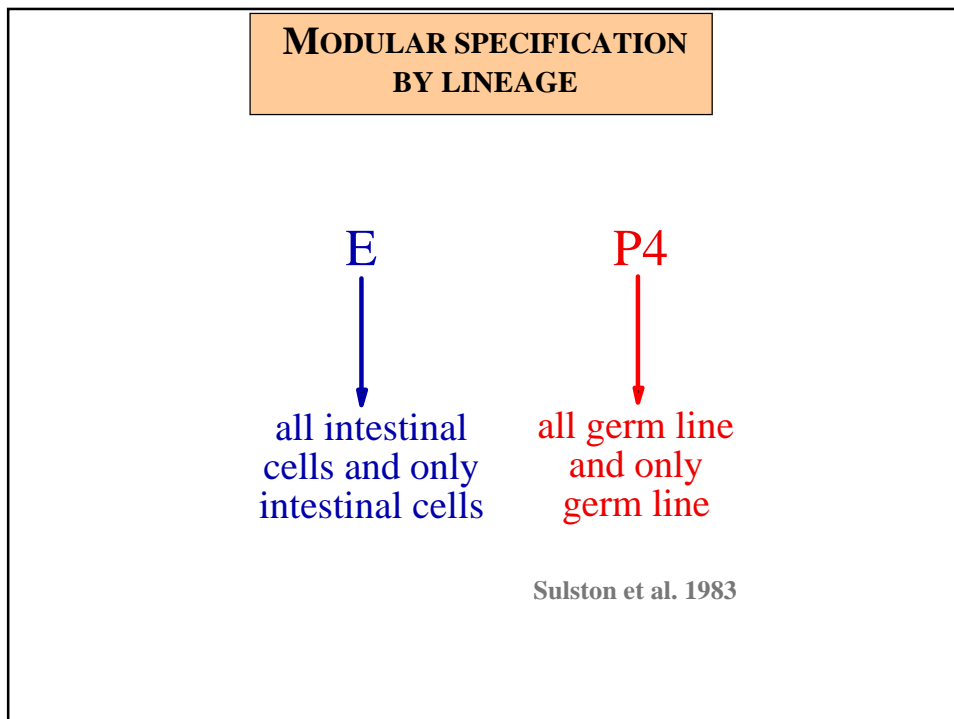
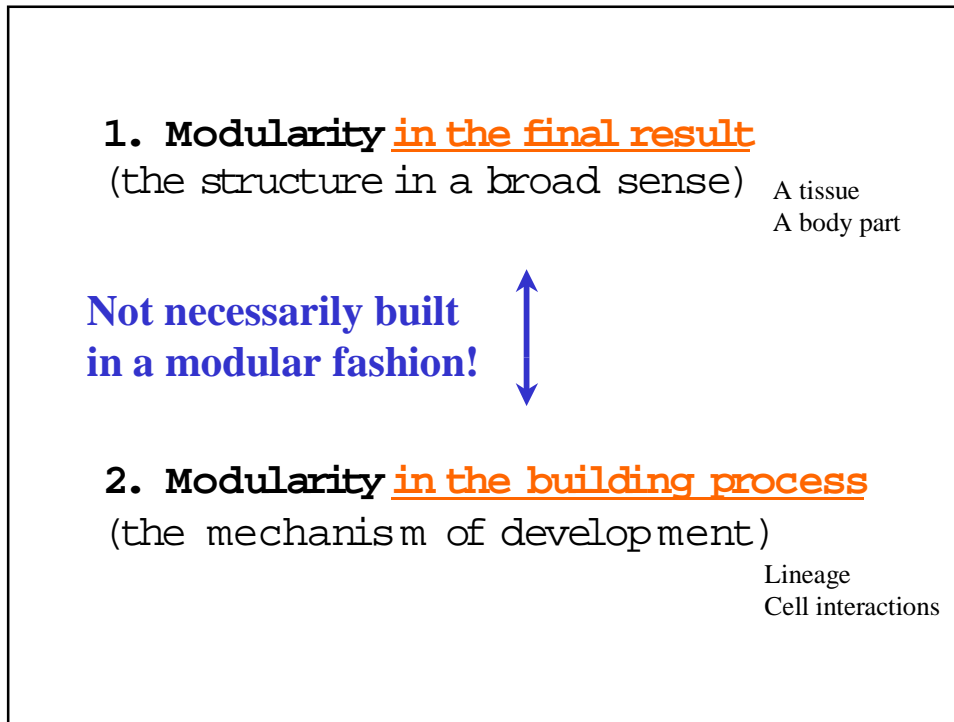


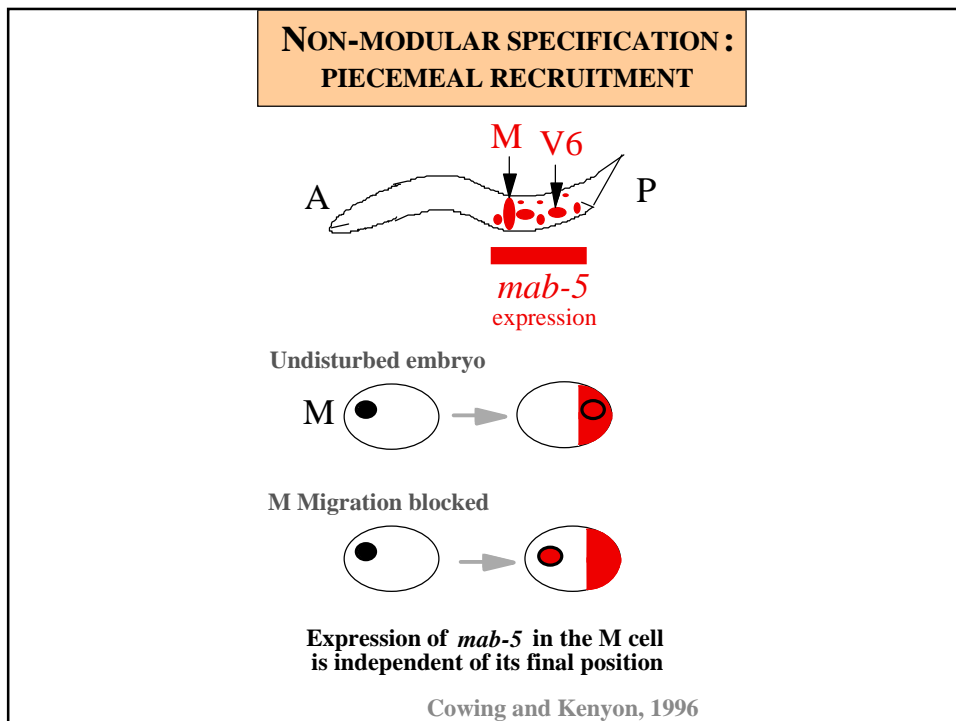
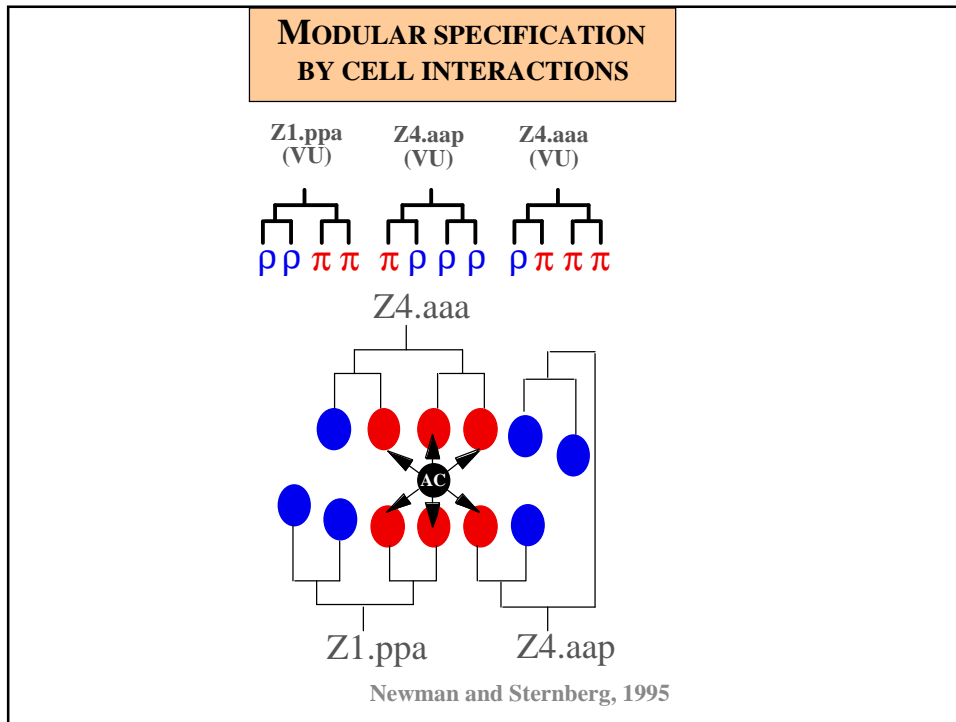
The specification of cell fates during
C. elegans development is the result of
asymmetric cell divisions
and **cell interactions**.



Its **cell lineage is invariant**
because of stereotyped asymmetric cell divisions,
but also because
the cell interactions are reproducible.

**Is there a supracellular level of description
of *C. elegans* development / morphology?**





Cell groups may be specified

A. in a **modular** fashion by:

- i. **shared lineage**
- ii. **shared cell interactions**

'Independent' cell interaction networks
Often used in the development
of coherent final structures (ex: limb bud)

B. in a **non-modular** fashion
by **piecemeal recruitment**

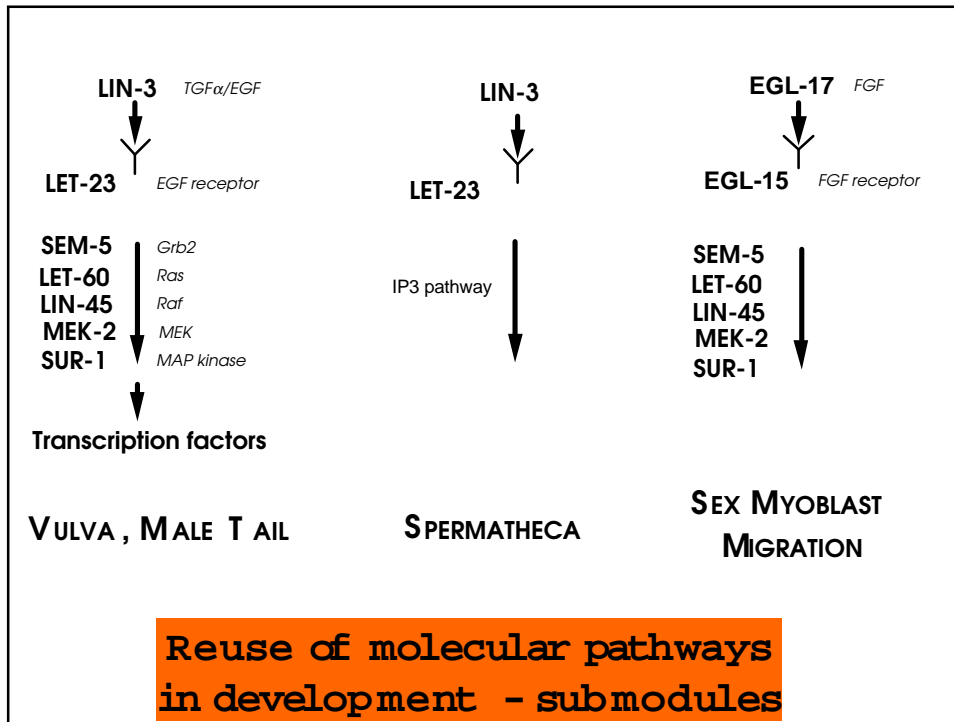
« perverse assignments »
(J. Sulston et al.)


MODULAR ORGANIZATION OF DEVELOPMENT

- (1) Supracellular modules
- (2) Modules of interacting molecules

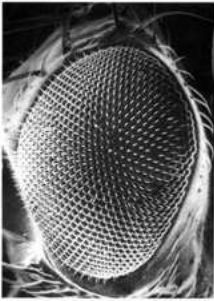
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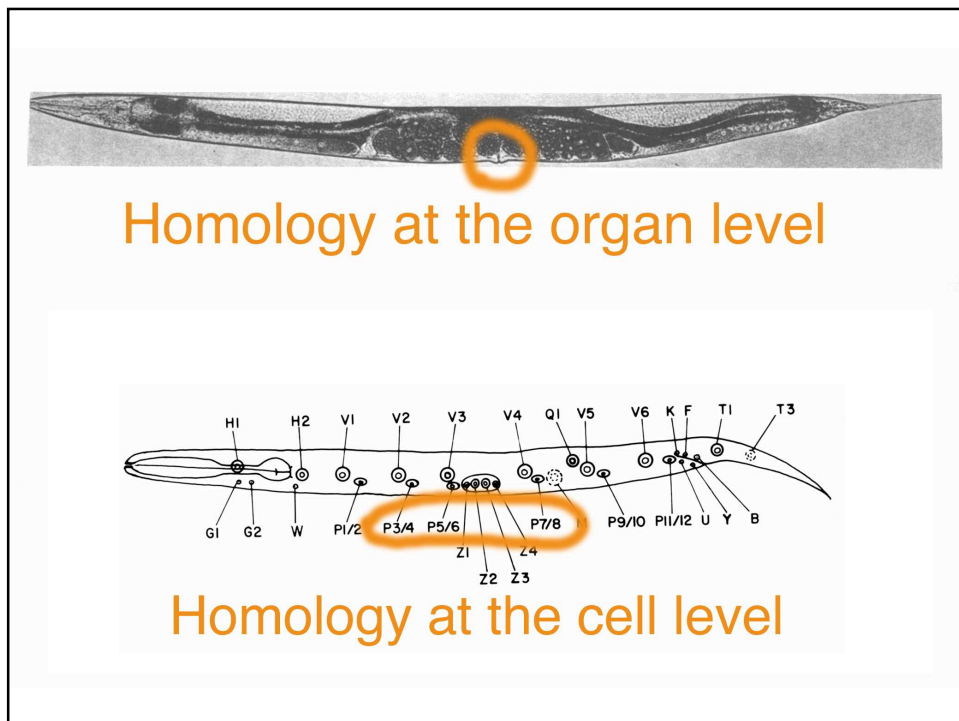
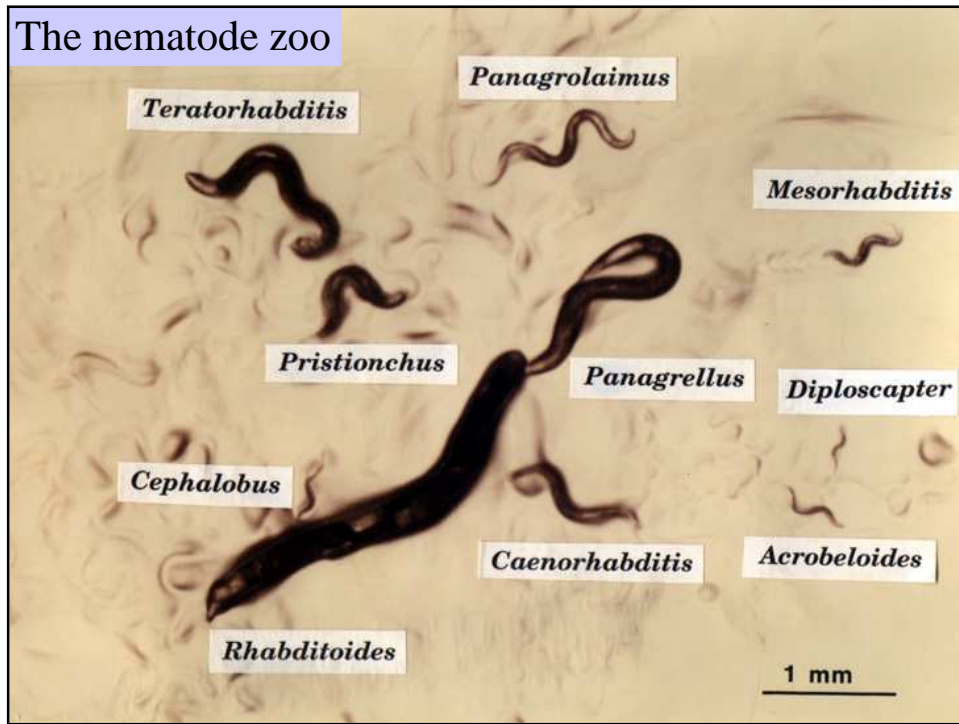
Caenorhabditis elegans
vulval induction
EGF-Ras-MAP kinase



Drosophila melanogaster
eye photoreceptor induction
EGF-Ras-MAP kinase

Molecular conservation
Reuse in non-homologous contexts

➔ Evolution of development of homologous structures



MODULAR ORGANIZATION OF DEVELOPMENT

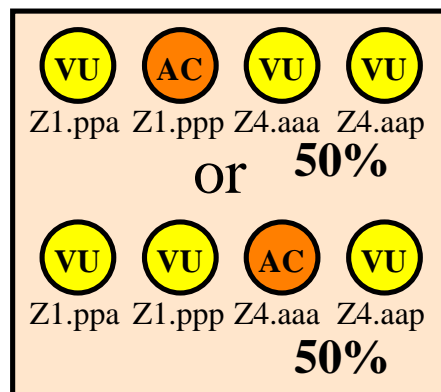
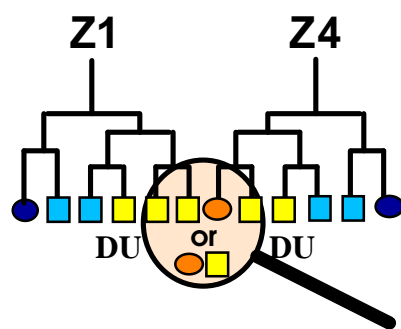
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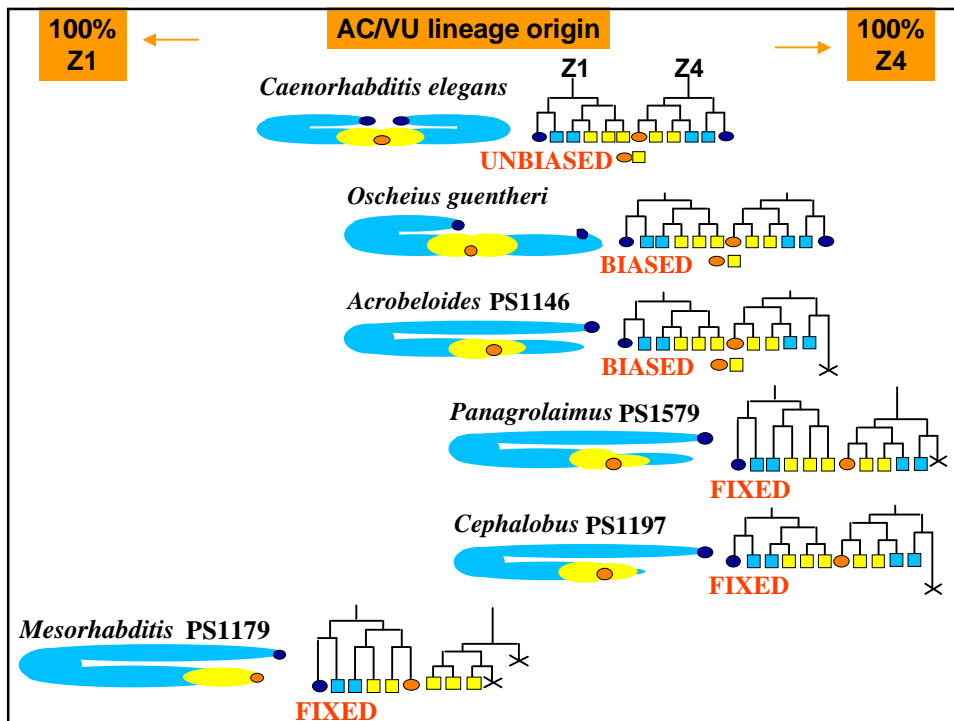
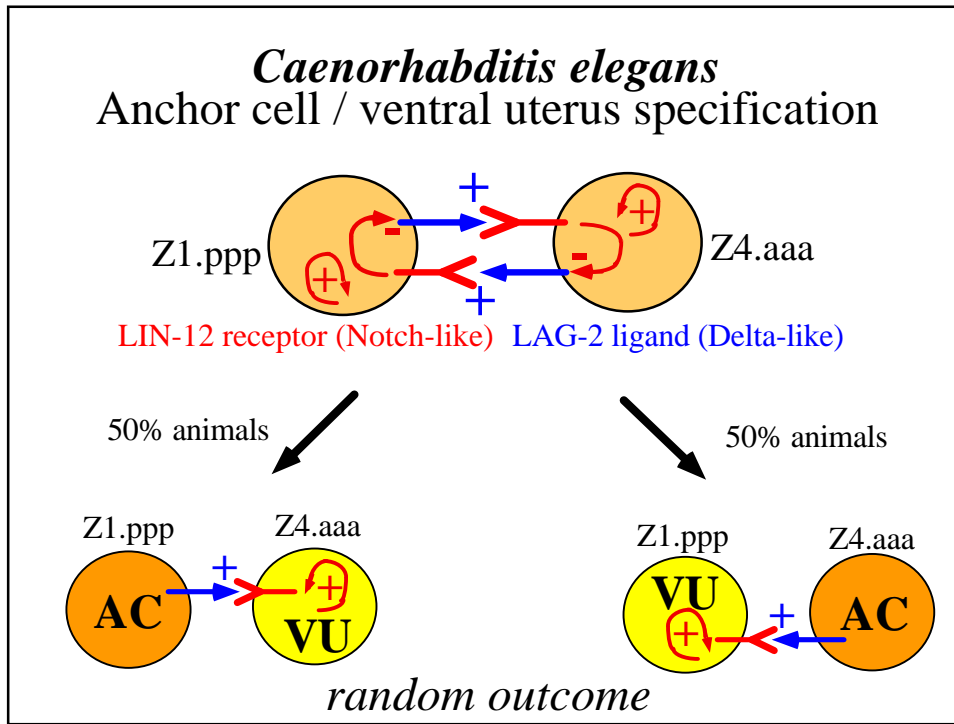
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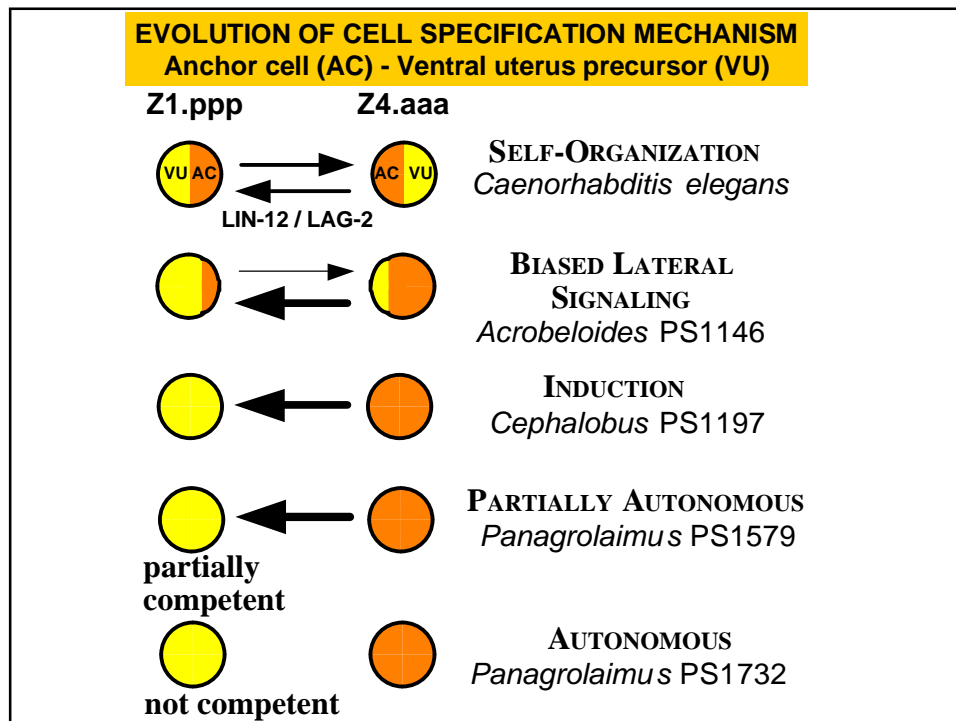
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Caenorhabditis elegans

The ventral uterus group

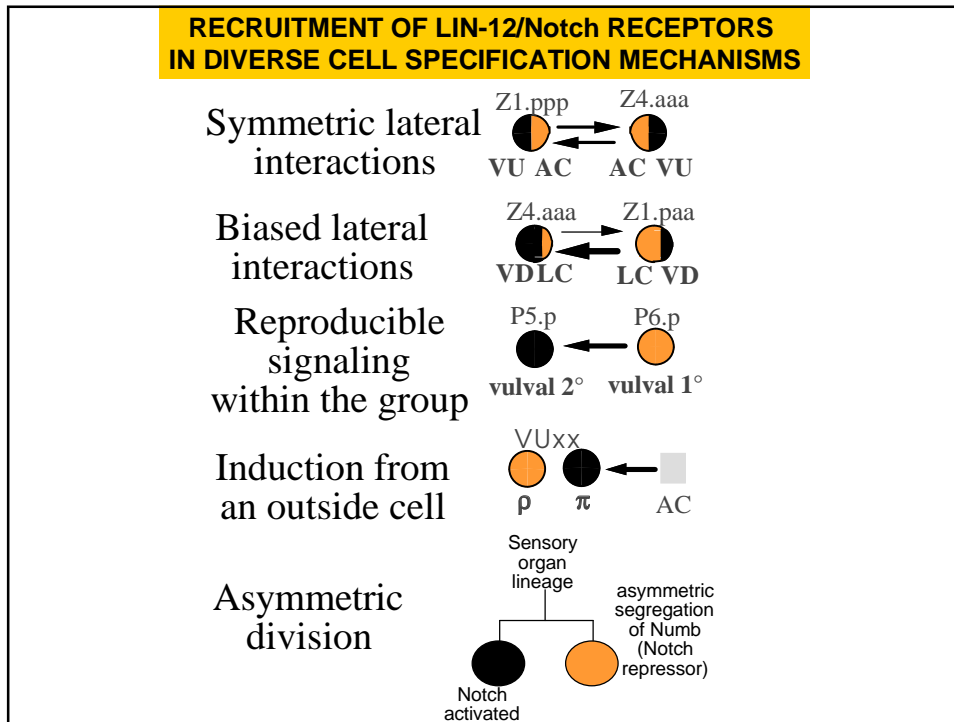






The developmental module AC/VU is useful to describe development in *C. elegans* but it is broken in evolution

'Intermediates' are found



CONSERVATION of MOLECULES

(e.g. gene expression pattern,
pathway involvement) does
not imply conservation of
SPECIFICATION MECHANISM

importance of details in the
molecular interaction network
ex. HOM-C genes!!!

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Problem:

Since **distinct developmental modules** use the **same molecular modules**, how can evolution occur in one developmental module without affecting the rest of the organism (pleiotropy)?

Note: Modularity in the dynamics of developmental modules is not sufficient for evolutionary independence between modules

Modularity in Gene Evolution

GENE DUPLICATION

Redundancy and subfunctionalization

EVOLUTION OF CODING REGIONS

Protein domains - shuffling,
phosphorylation sites in different tissues, etc.

EVOLUTION OF GENE EXPRESSION

Modularity in cis-regulatory regions

**Modularity of gene organization
(promoter, coding region)
might allow for
quasi-independent
evolution
of developmental modules.**

**What are the relative roles of
coding vs. regulatory sequence evolution?
In ubiquitous vs. regulatory molecules?**

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**MORPHOLOGICAL EVOLUTION
BY
DEVELOPMENTAL 'INNOVATION'**

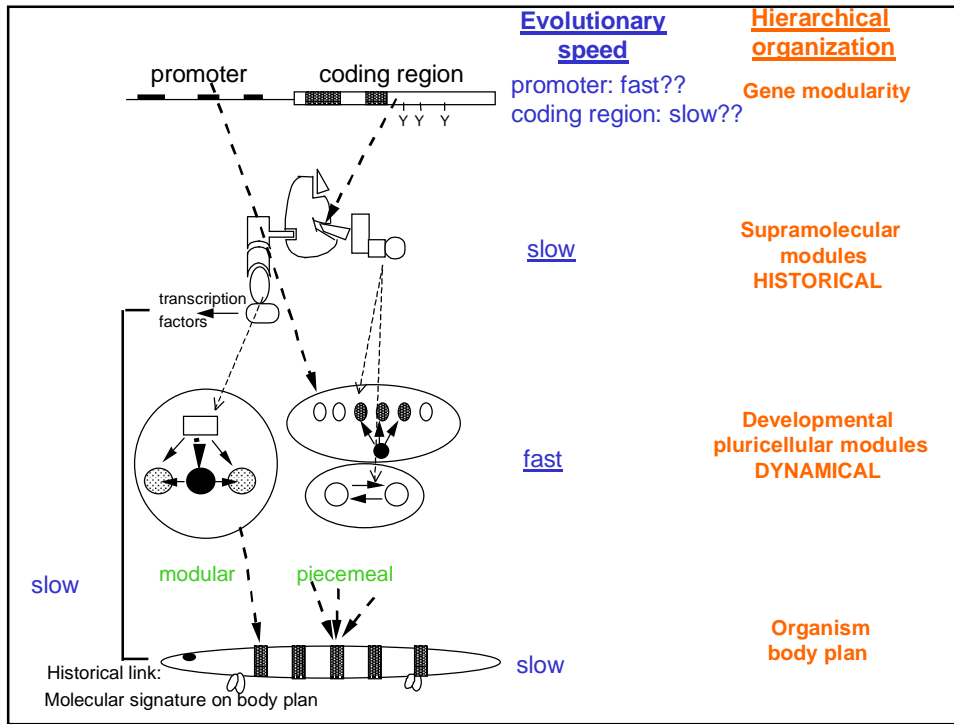
linkage/ breakage of (sub)modules
facilitated . by the evolutionary flexibility
in the dynamics of molecular modules
. by redundancy

nesting
reuse of the same molecular pathways

diversification by symmetry
breakage
facilitated by small asymmetries

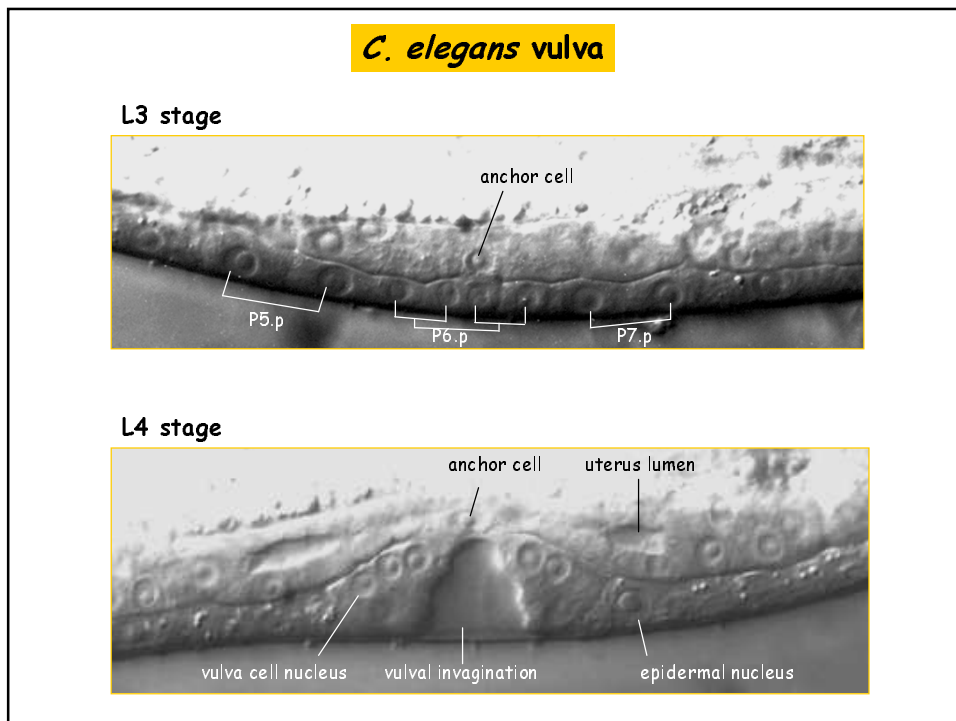
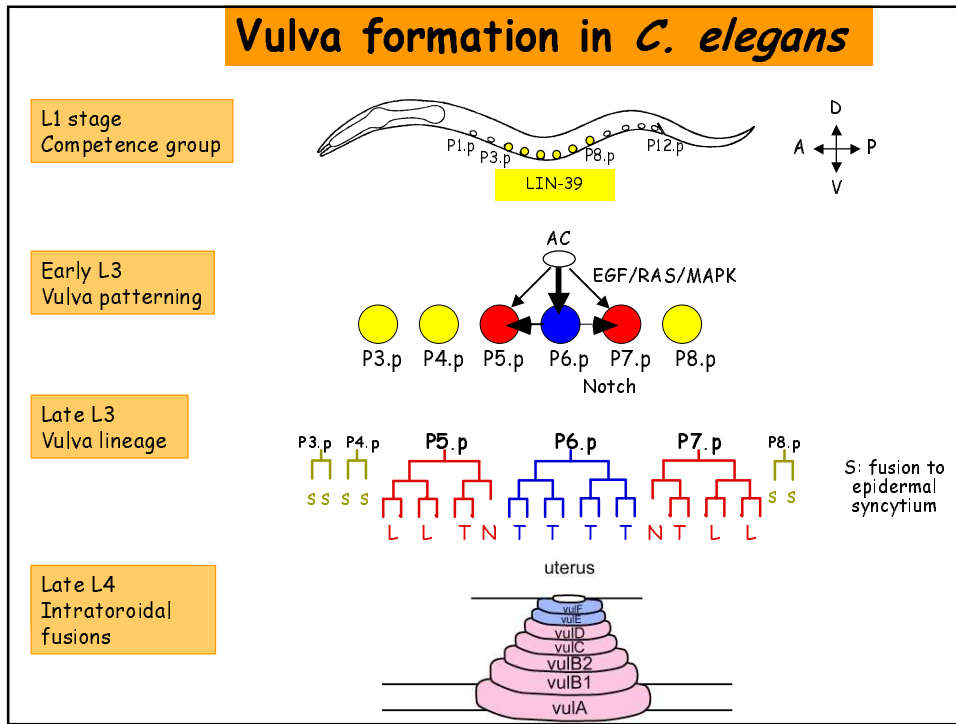
CONSTRAINTS ON BODY PLAN ?

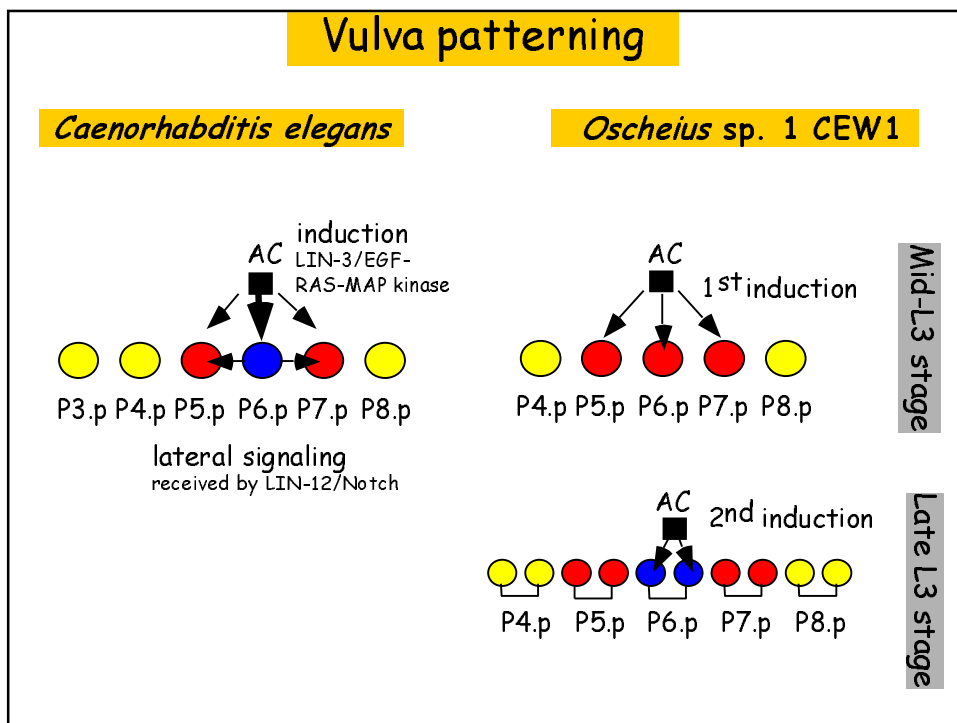
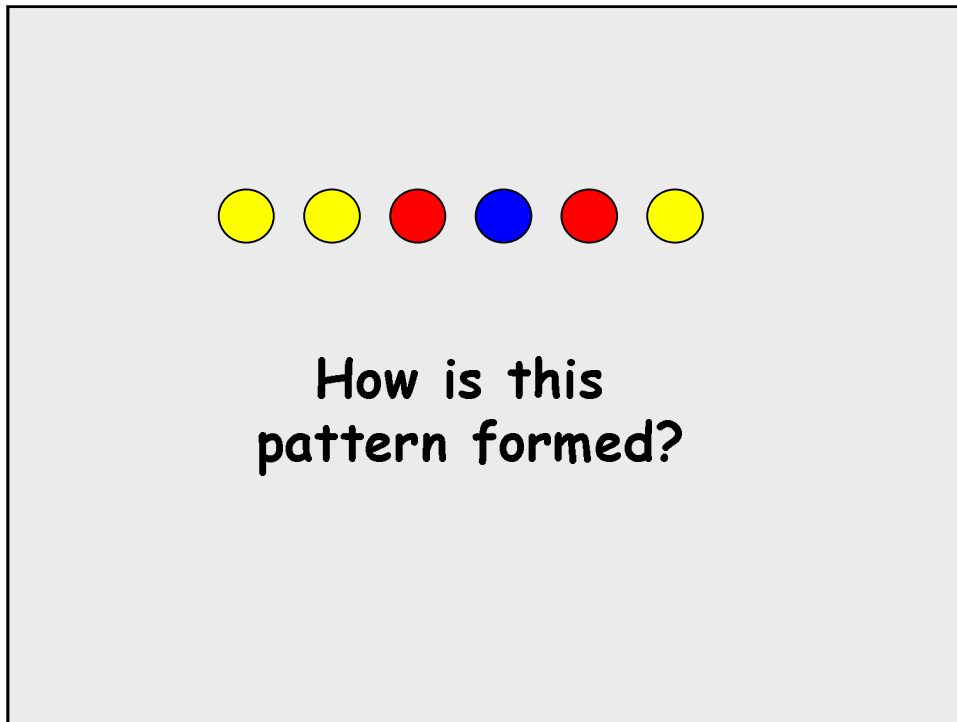
- not because early development is constrained
- not as much constrained in plants

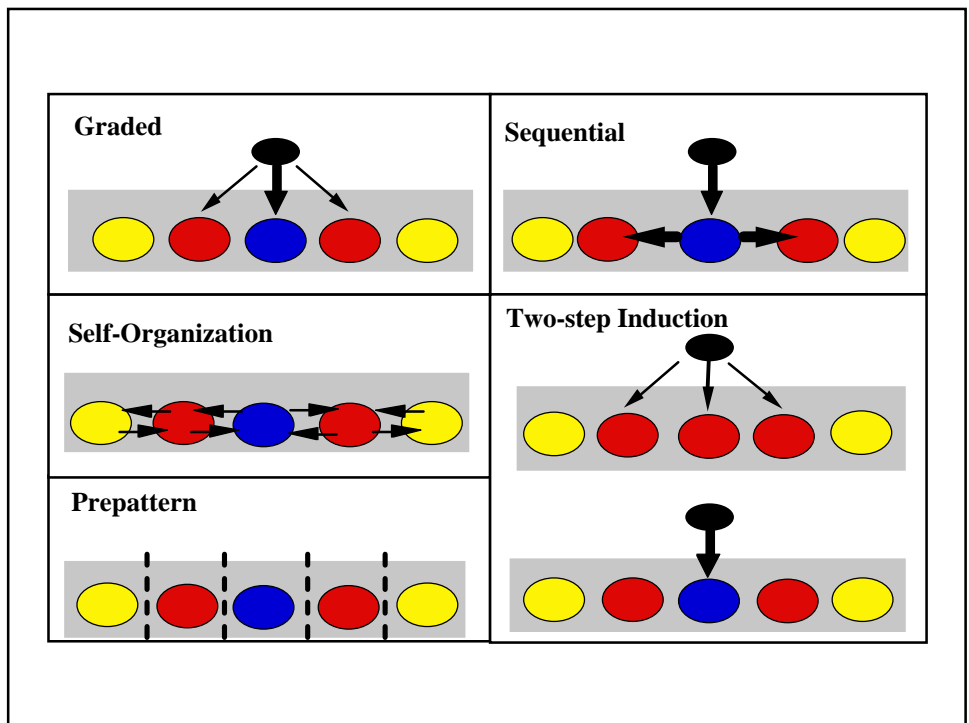
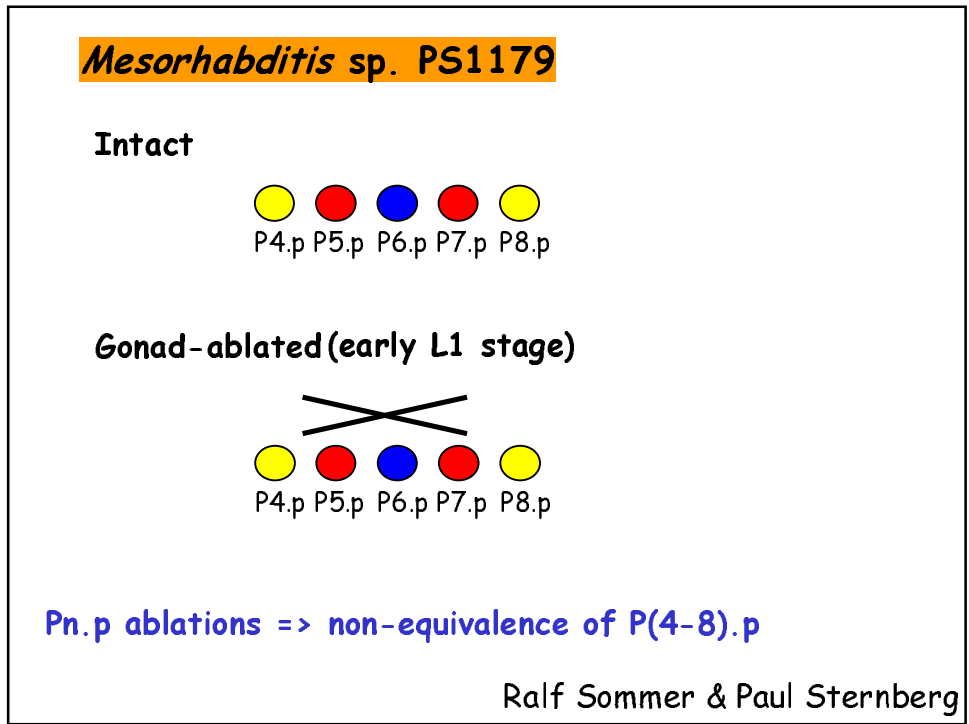


THE VULVA









The mechanisms
patterning the vulva
precursor cells vary
extensively among
these nematodes

(whereas the pattern
does not in most
cases).

Redundancy
between developmental mechanisms
ensures precision in a given species

and might in turn favor rapid evolution
of these mechanisms

(large changes in the weight of different
mechanisms may be quasi-neutral).

Microevolutionary example:
Panagrolaimus sp.
Degree of requirement for the gonadal induction
versus pre-patterning along the antero-posterior axis

Microevolutionary analysis

- Genetic dissection of phenotypic variation is possible

in hermaphroditic nematodes

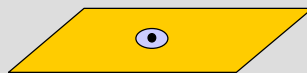
- Genetically homogeneous populations
 - studies of developmental variants/noise
 - recombinant inbred lines are easily driven to homozygosity

Microevolution within *C. elegans* or *Oscheius* sp. 1
(and then comparison of their evolvability)

Mutagenesis effect vs. Observed microevolutionary change
=> imprint of selection

Mutagenesis

Genotype



○ after mutagenesis

Phenotype



COMPARING THE VULVA SCREENS

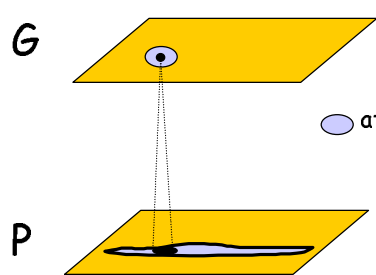
The spectrum of vulva mutations isolated is very different in the two species:

<p><i>C. elegans</i></p> <p>Competence Vulvaless Multivulva Lateral signaling</p>	<p><i>Oscheius</i> sp. 1 CEW1</p> <p>Centering and competence</p> <p>Hypoinduction of both steps Hyperinduction of 1st step Too few divisions Too many divisions</p>
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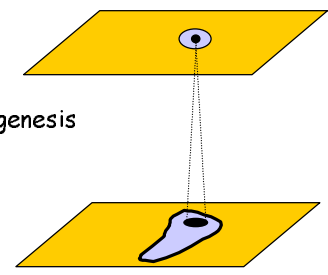
different sensitivities of the systems to one-gene perturbation

Mutagenicity and evolvability

species 1



species 2



○ after mutagenesis

Assessment by mutagenesis of the "reachable" phenotypic states from a given genotype
PHENOTYPIC NEIGHBOURHOOD
 Correlation with wild variants ?

Nonreductive variations

Genotype



different wild isolates
of the same species

Phenotype

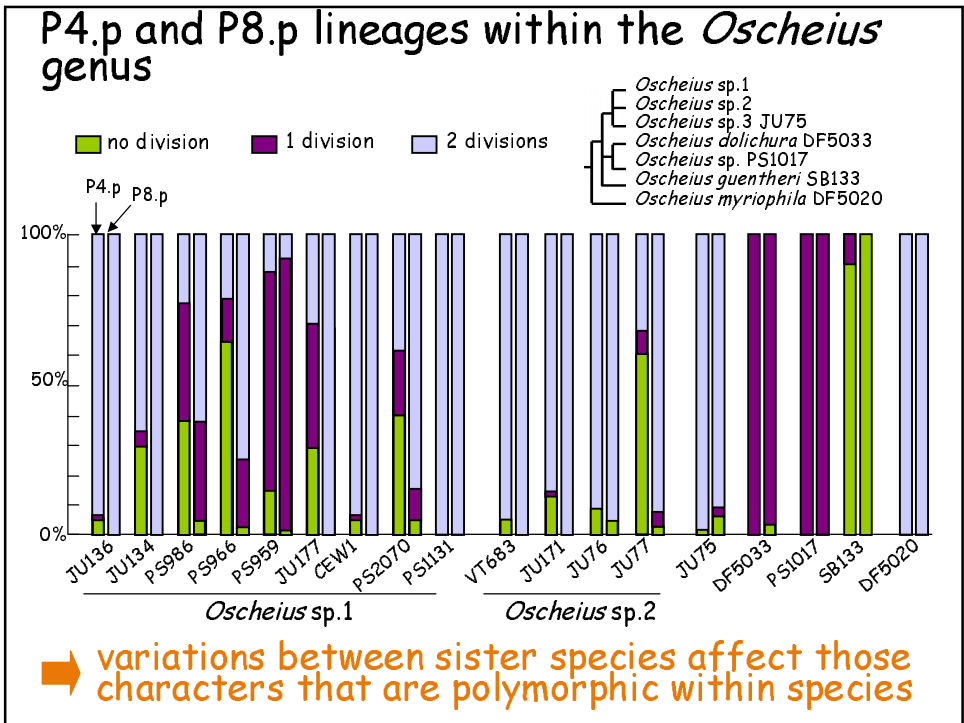
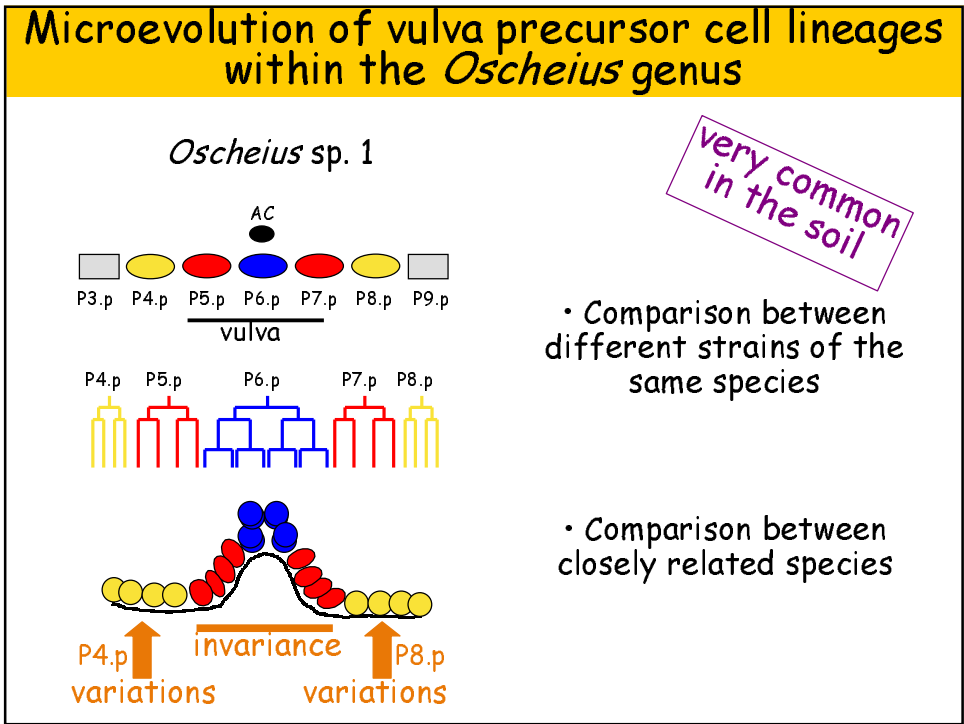


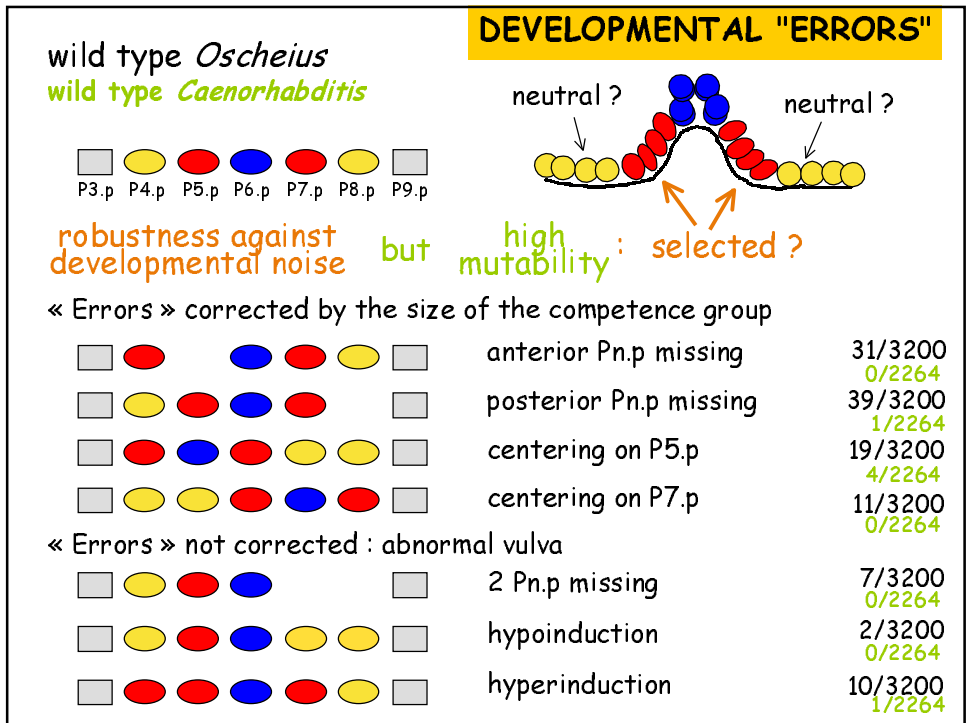
The cell lineage of a given species
of nematodes is globally invariant.

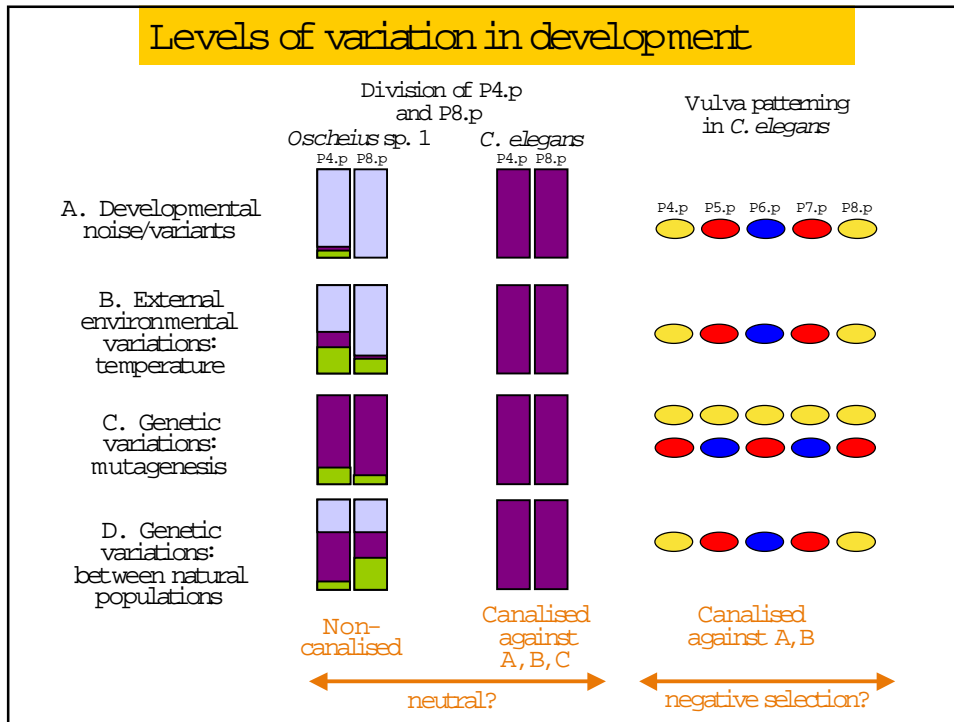
zygote

How can it evolve given this constancy?

Are inter-species variations the
consequences of fixation of intra-species
polymorphisms?









Thanks to:

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