

Sex chromosome evolution in organisms with female heterogamety and holokinetic structure of chromosomes: moths and butterflies

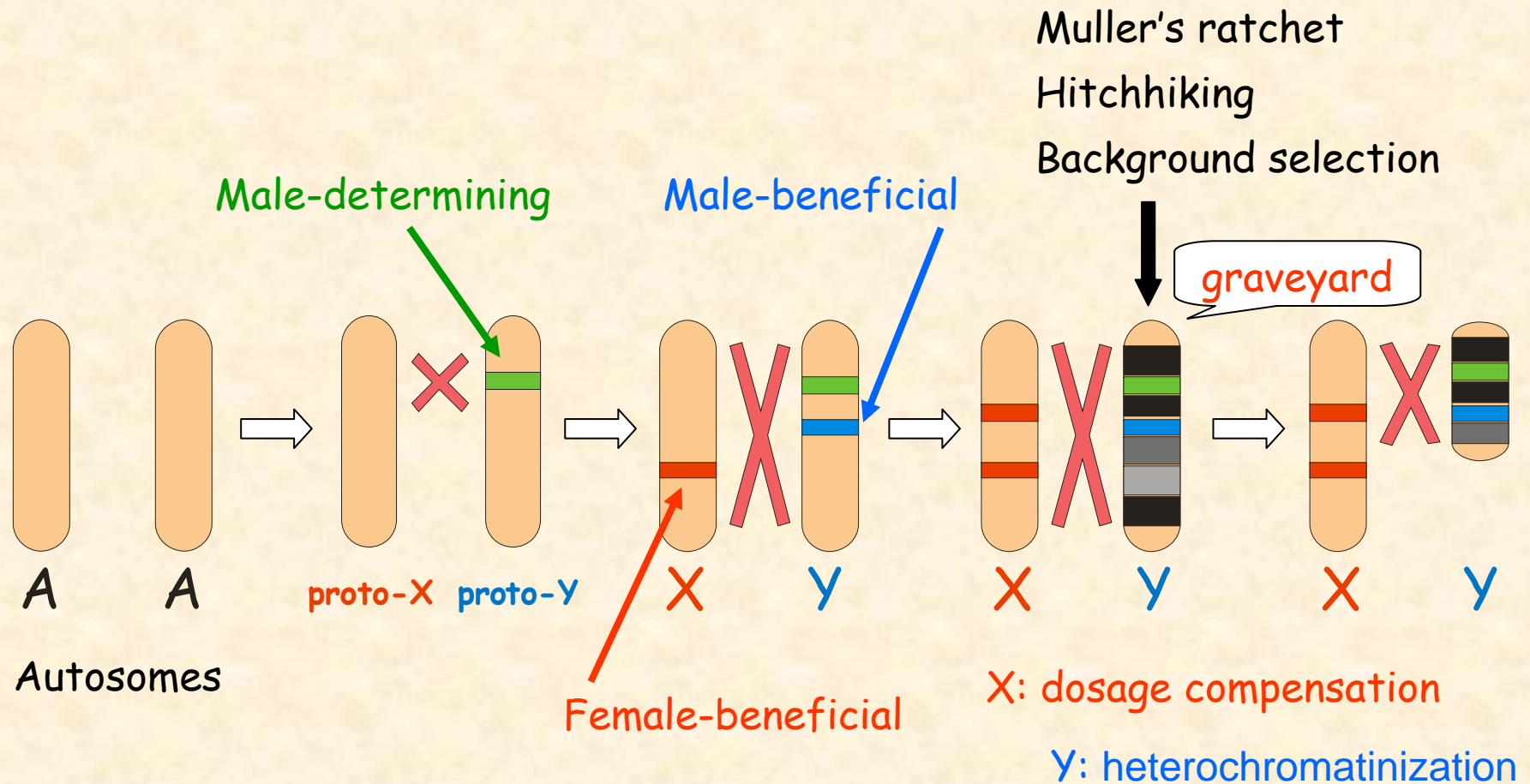
Frantisek Marec^{1,2}



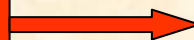
¹Biology Centre ASCR, Institute of Entomology, Ceske Budejovice, Czech Republic

²University of South Bohemia, Faculty of Science, Ceske Budejovice, Czech Rep.

Theory: evolution of sex chromosomes



Restriction of recombination

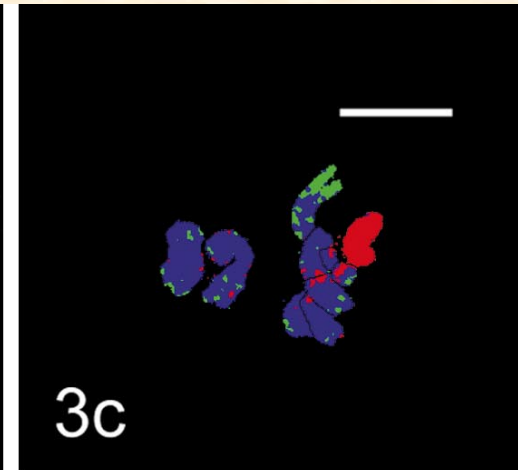
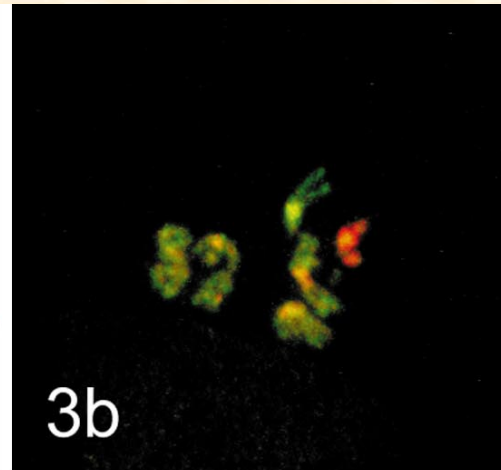
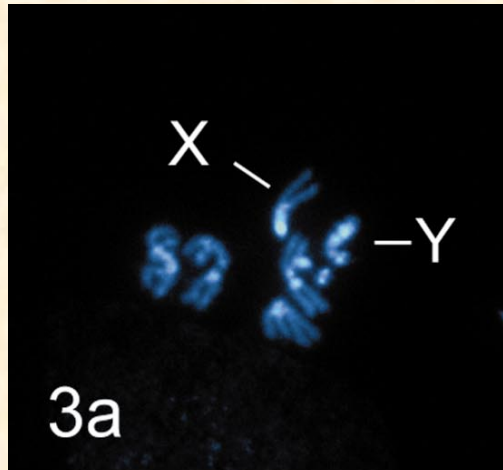


Molecular differentiation

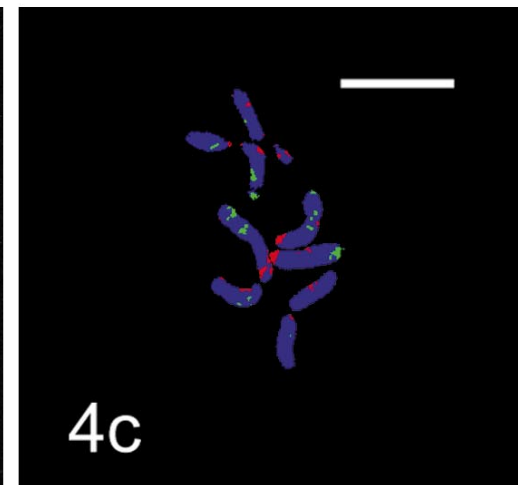
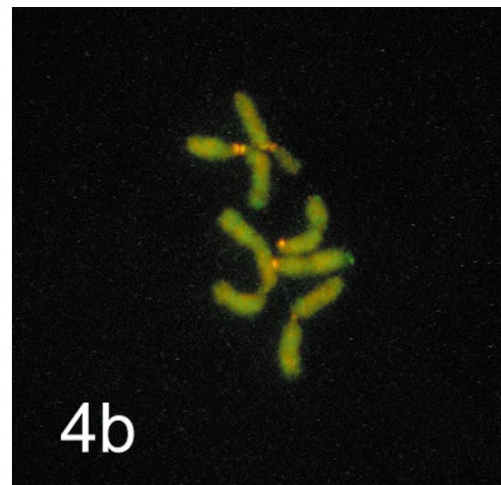
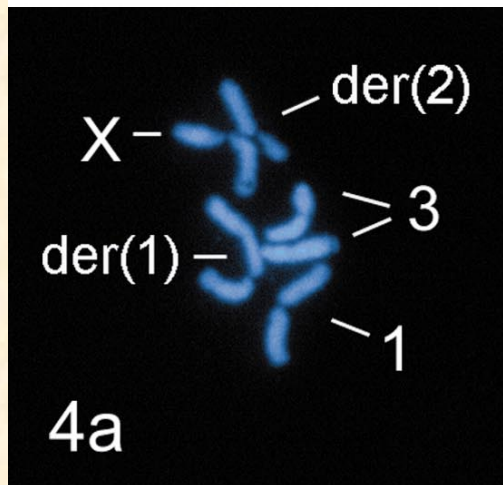
Morphological differentiation

Molecular differentiation of sex chromosomes by CGH

Drosophila melanogaster



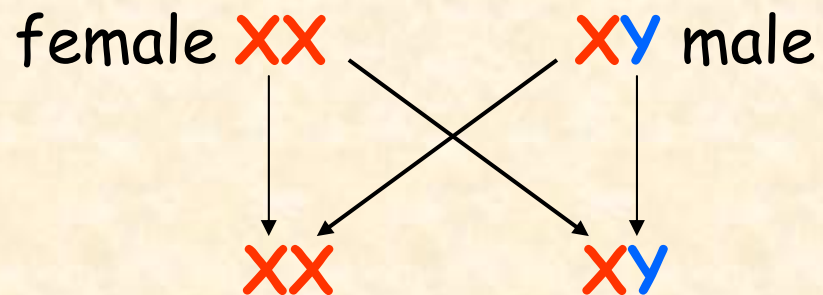
Megaselia scalaris



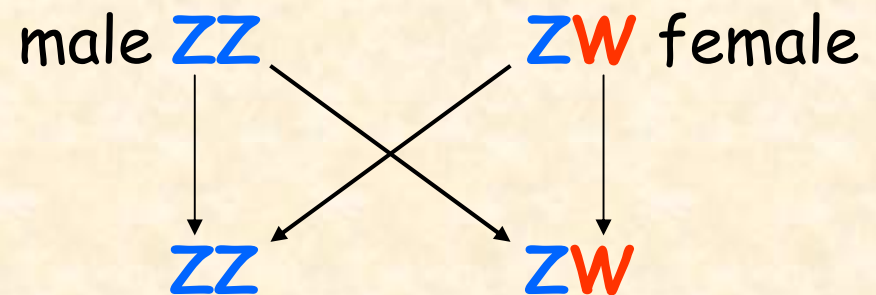
Traut et al. (1999) *Chromosoma* 108: 173-180.

Two basic sex-chromosome systems had evolved:

Male heterogamety
(mammals, *Drosophila* ...)



Female heterogamety
(birds, butterflies ...)



The conception of Y as a "graveyard" revised

- Y chromosomes of primates: palindrome-driven sister chromatid and intrachromatid recombination prevents degradation and loss of genes => continuing evolution

Lange et al. (2009) *Cell* 138: 855-69.

Hughes et al. (2010) *Nature* 463: 536-39.

- *Drosophila* Y chromosome: an important role in male fitness - it contributes to adaptive phenotypic variation through a regulatory role of Y-linked polymorphic elements in gene expression

Lemos et al. (2008) *Science* 319: 91-93.

WZ/ZZ system (female/male)

Heterogametic female sex inferred from sex-linked inheritance in *Abraxas* 100 ya
- sex chromosomes never seen.

- **Lepidoptera**
- Trichoptera (ZO)
- snakes (Reptilia: Serpentes)
- birds (Aves)



Abraxas glossulariata
(Geometridae)

- some fishes (*Xiphophorus maculatus*)
- number of amphibians (*Xenopus laevis*)
- some lizards (*Lacerta vivipara*)
- some parasitic worms (*Schistosoma*)
- several plants (*Fragaria elatior*) ???



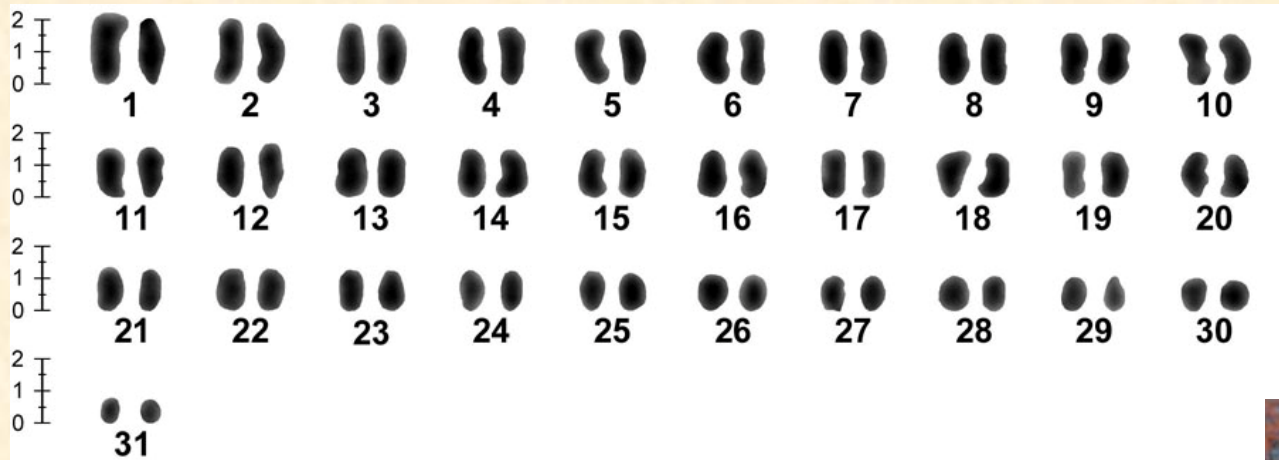
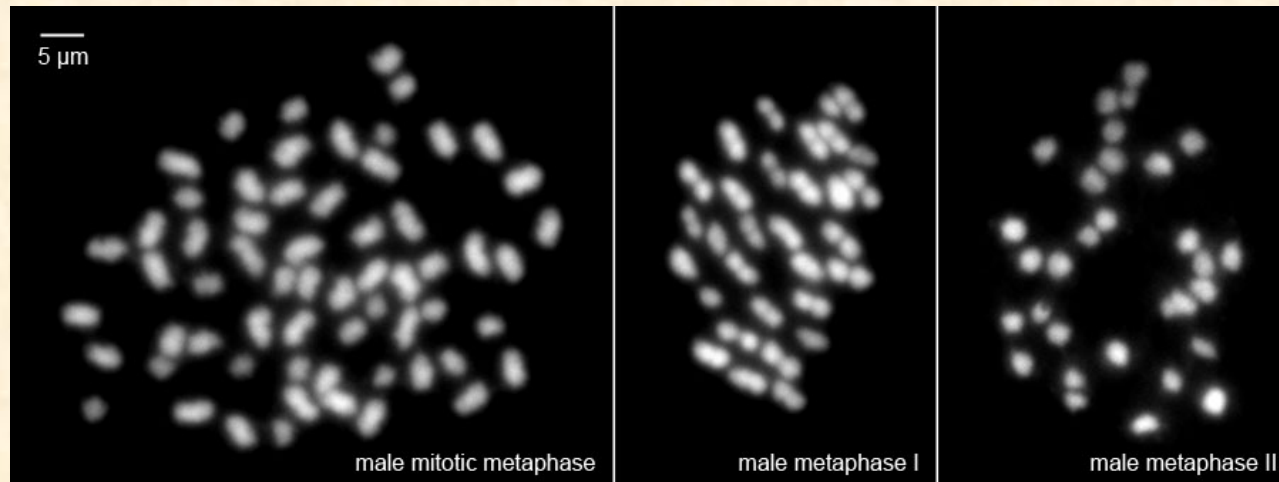
Lepidoptera

moths and butterflies



- Largest taxon with female heterogamety
 - more than 150.000 species described
- The only taxon with female heterogamety having holokinetic chromosomes
- Achiasmatic female meiosis => no cross-over
- Sex chromatin in somatic cells of females

Holokinetic chromosomes of Lepidoptera

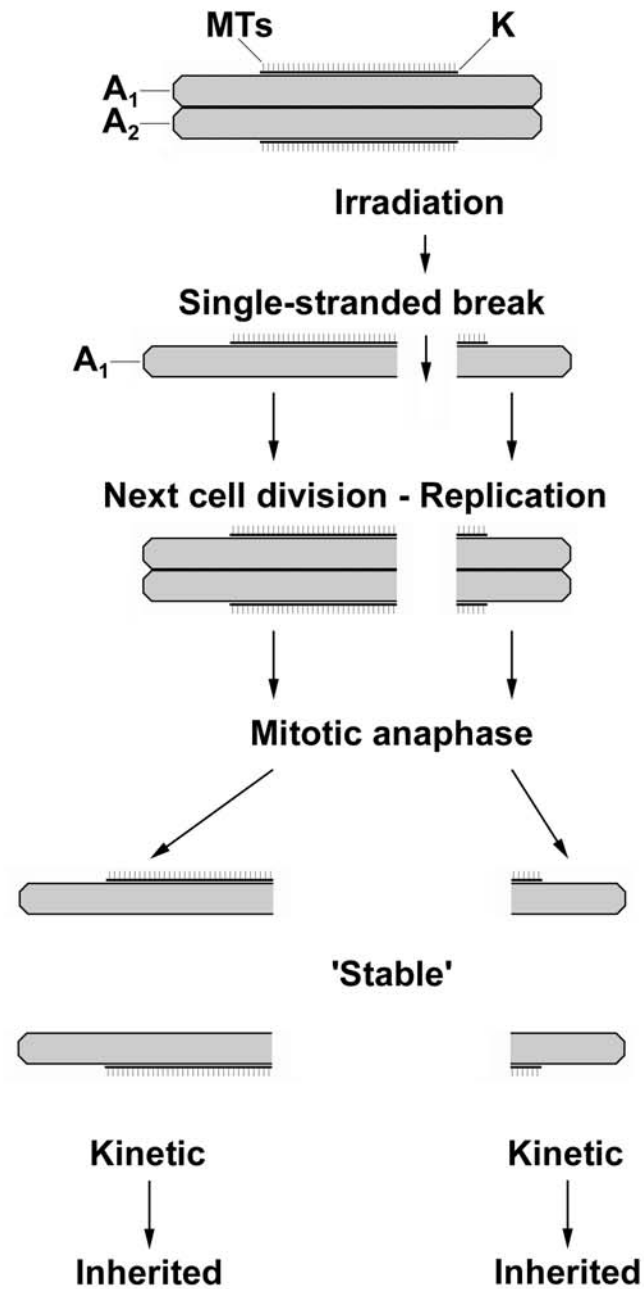


Carob moth, *Ectomyelois ceratoniae* (Pyralidae)

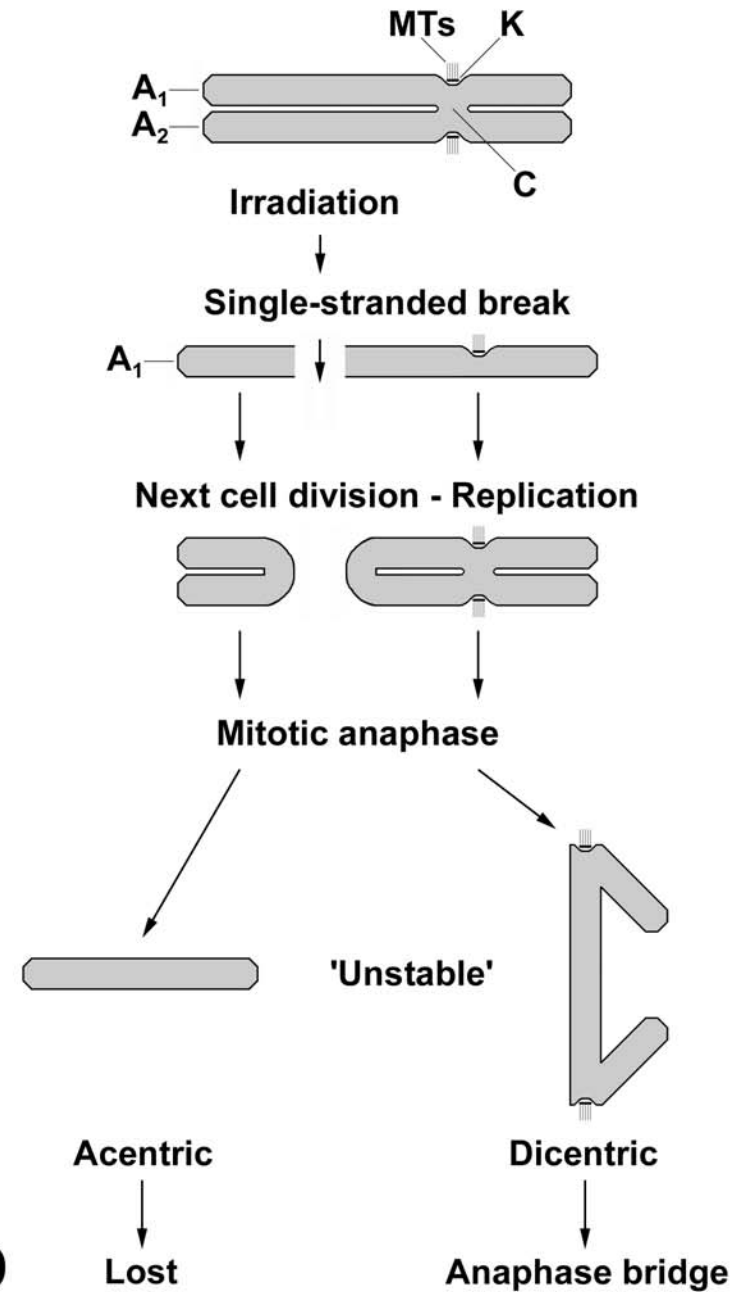
$n = 31$; $2n = 62$



Holokinetic chromosome of Lepidoptera



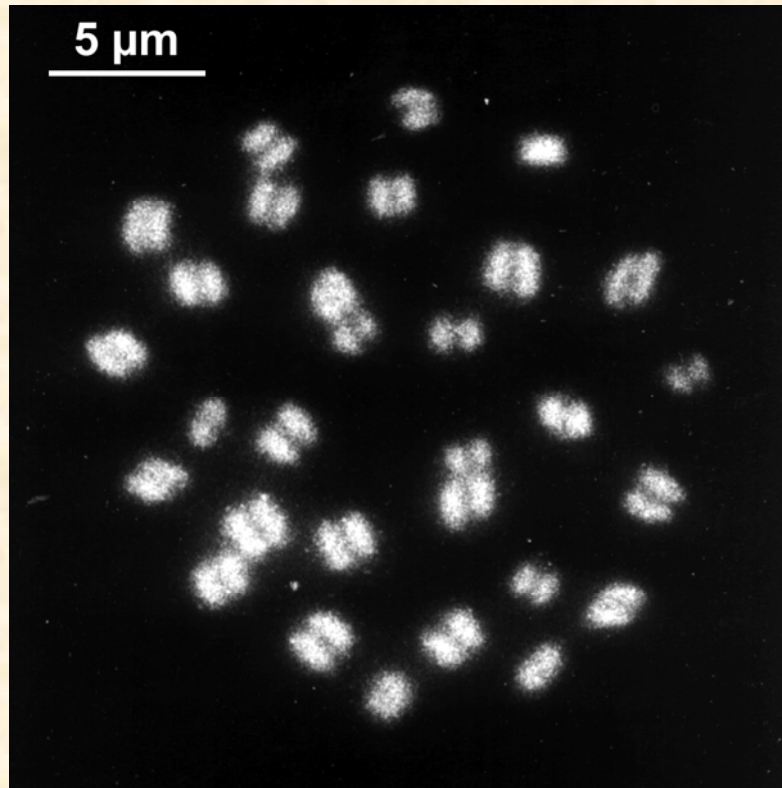
Typical monocentric chromosome



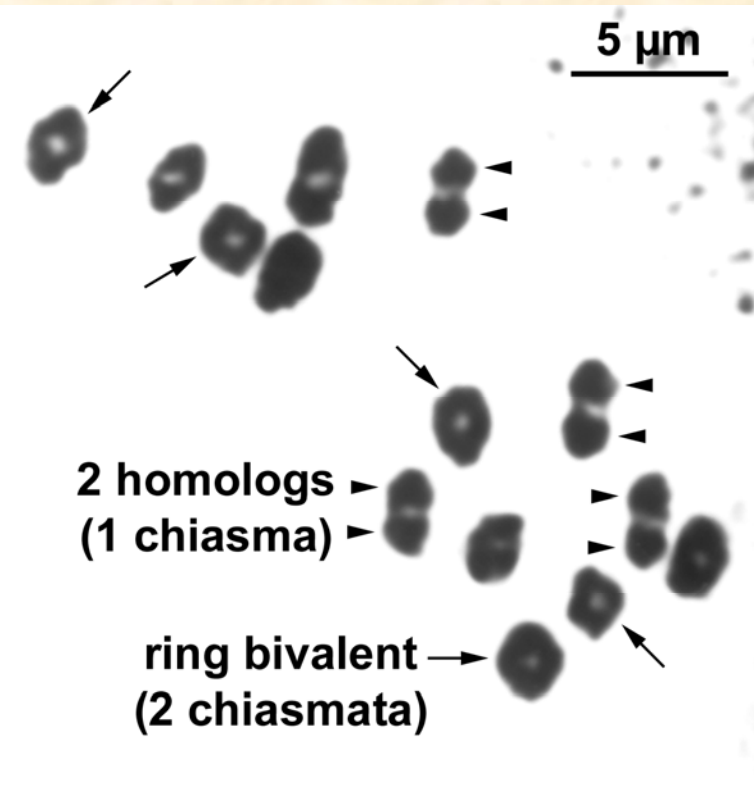
Holokinetic chromosomes of Lepidoptera

- contribute to resistance to clastogenic compounds and ionizing radiation
 - chromosome breaks do not lead to break-fusion-bridge cycle
 - chromosome fragments inherited
 - reduced risk of dicentric chromosomes
- facilitate karyotype evolution by rearrangements
 - fusion => reduced chromosome numbers
 - fission => increased chromosome numbers
- **BUT surprising stability of genomes/karyotypes**
 - common /ancestral chromosome number $n = 31$ (exceptions occur)
 - widely conserved synteny of genes between species

Achiasmatic versus chiasmatic meiosis in Lepidoptera



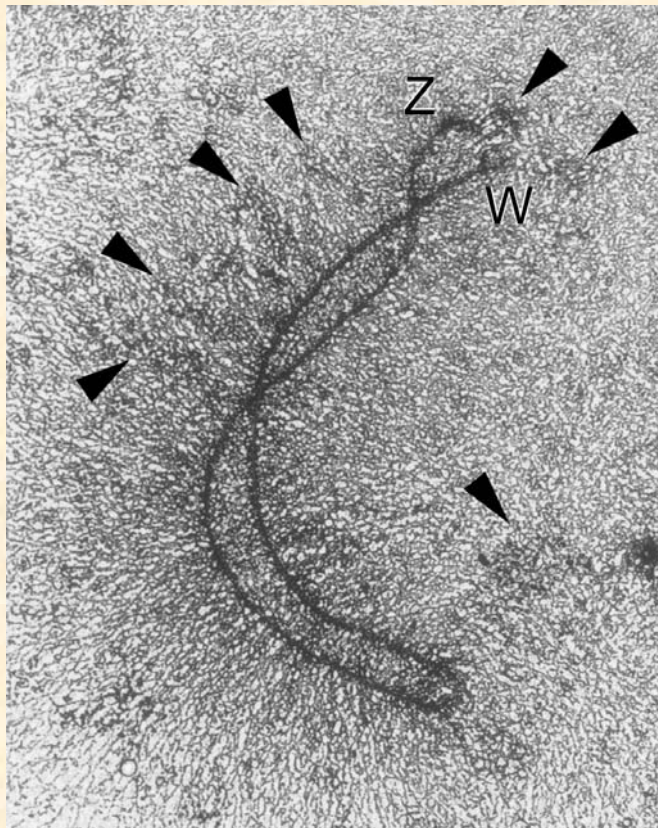
Metaphase I bivalents of the flour moth, *Ephestia kuehniella* (Pyralidae) - female (n=30)



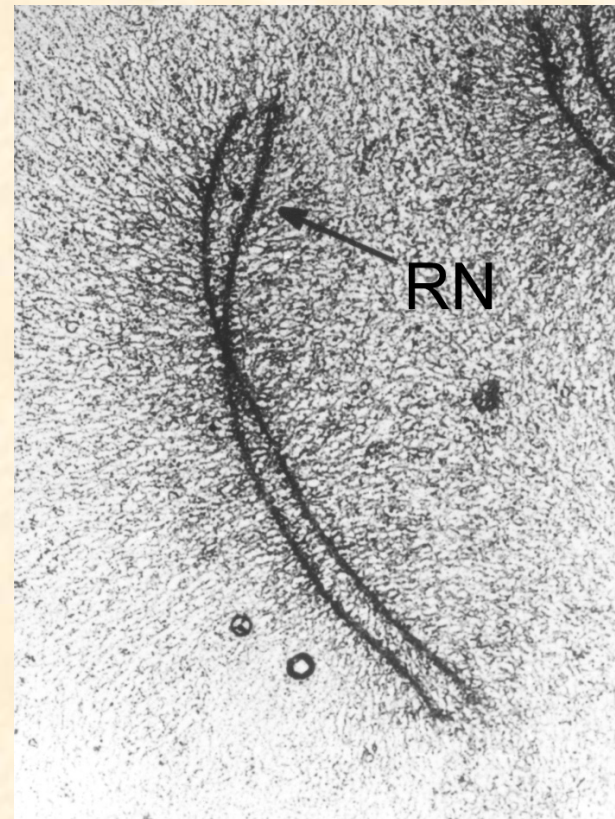
Metaphase I bivalents of the vapourer moth, *Orgyia antiqua* (Lymantriidae) - male (n=14)

⇒ no recombination between W-Z,
not even between autosomes

The flour moth, *Ephesia kuehniella* (Pyralidae)



female SC

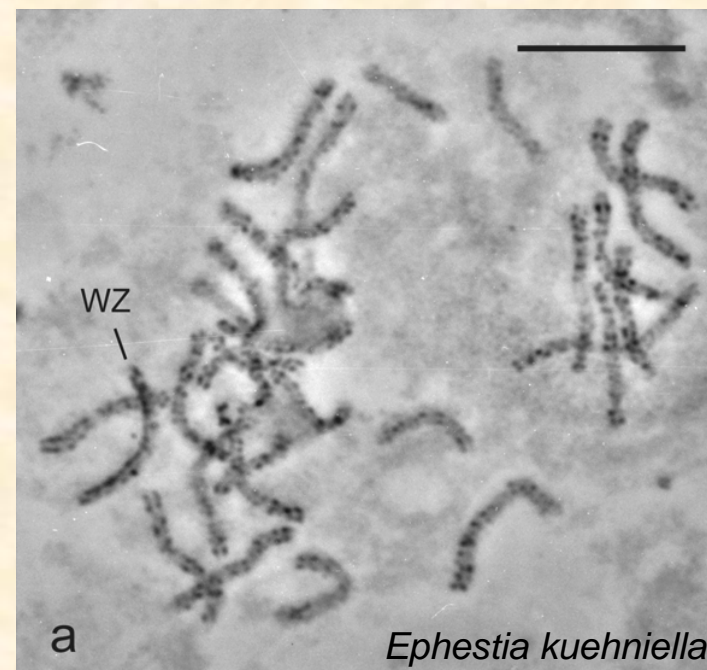
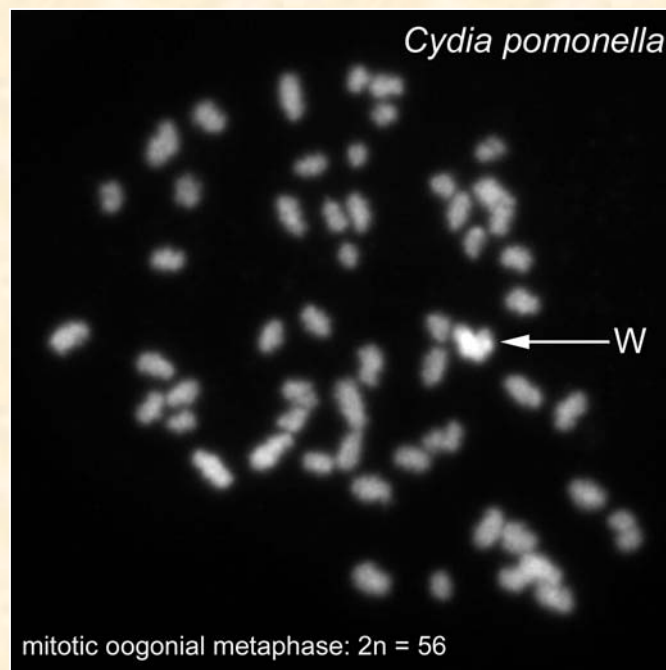


male SC

Marec and Traut (1993) *Heredity* 71: 394-404.

W chromosome in Lepidoptera

- consists of heterochromatin => mainly composed of repetitive DNA
- small holokinetic chromosomes => resist identification in metaphase
- distinguishable in pachytene; WZ bivalent
- in contrast to gene-rich Z, W carries only a few genes (*Fem* factor)

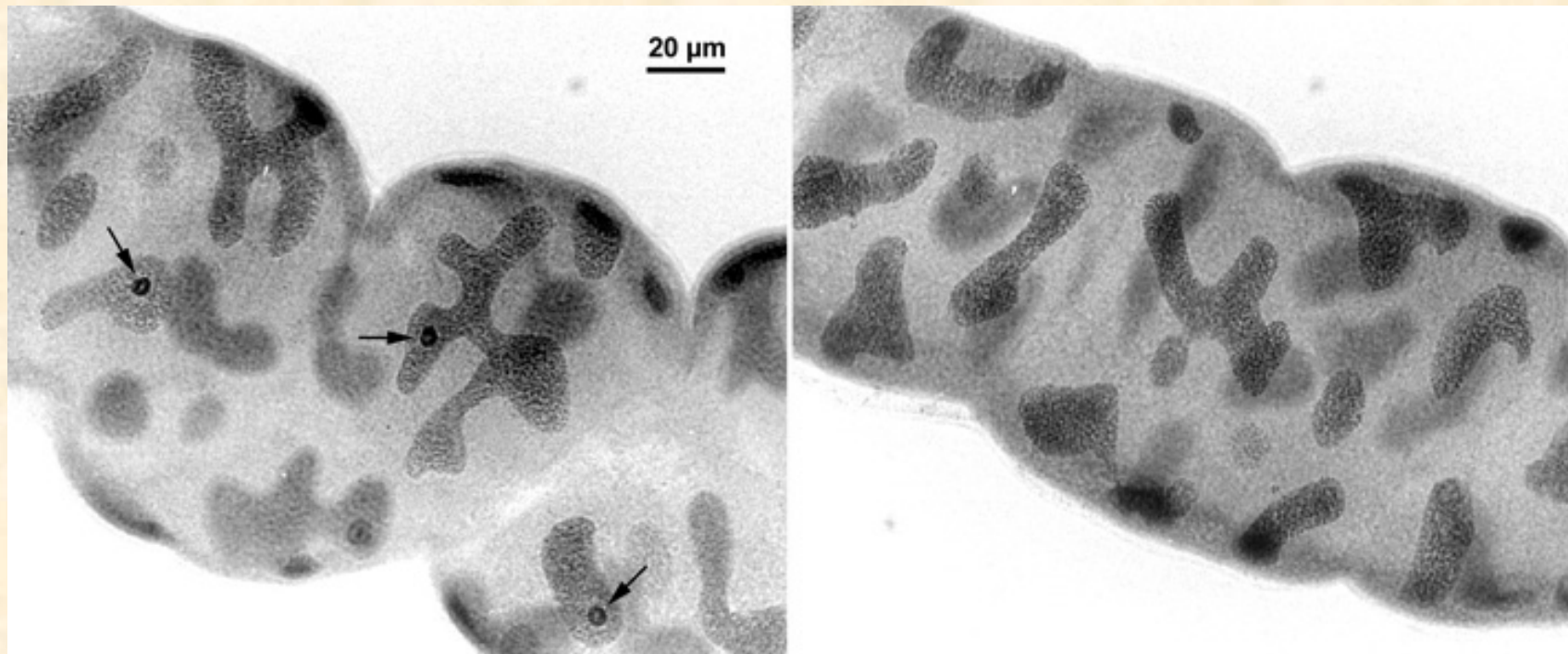


Sex chromatin in Lepidoptera

Ephestia kuehniella - Malpighian tubules

female

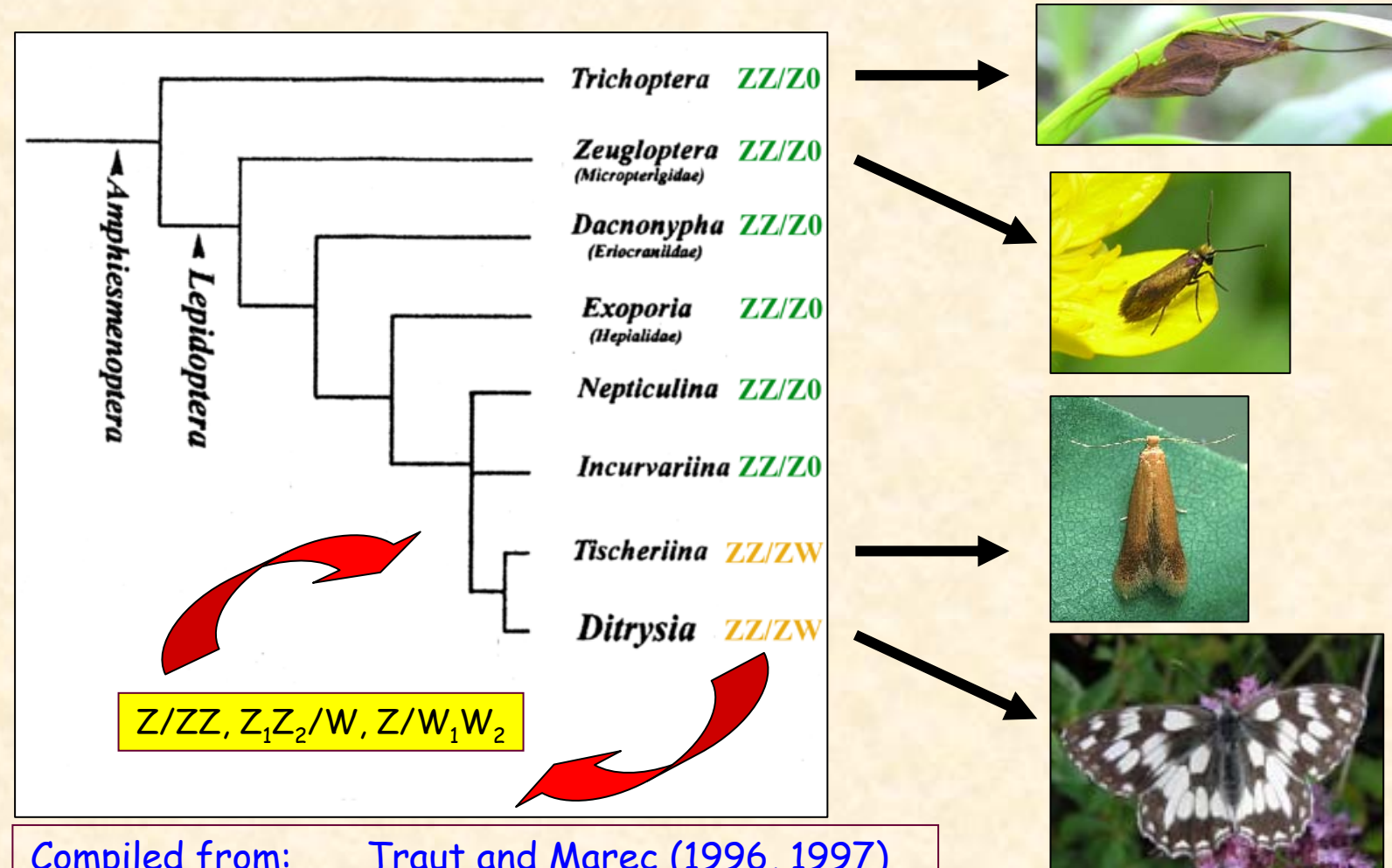
male



Characteristic trait of females in 80% of Ditrysia, derived from W

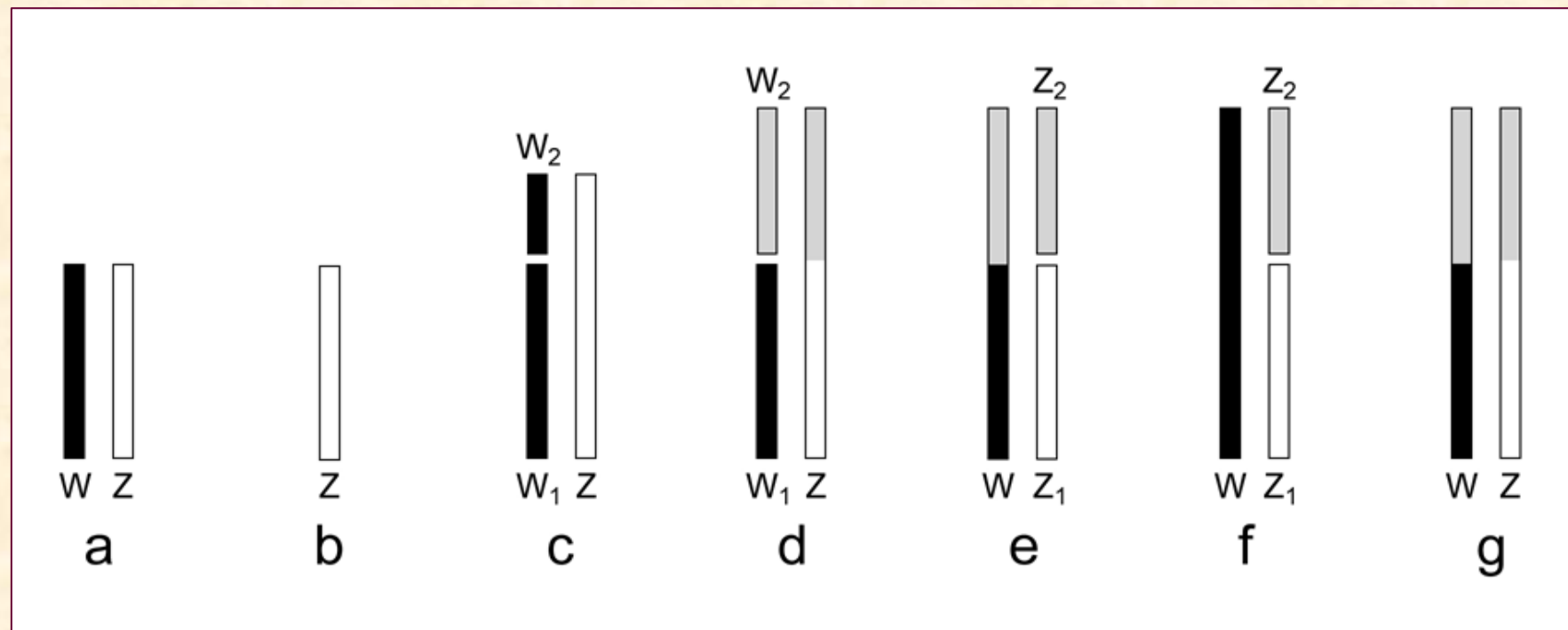
reviewed by Traut and Marec (1996) *Q. Rev. Biol.* 71: 239-256.

Phylogeny of sex chromosomes in caddis-flies, moths and butterflies



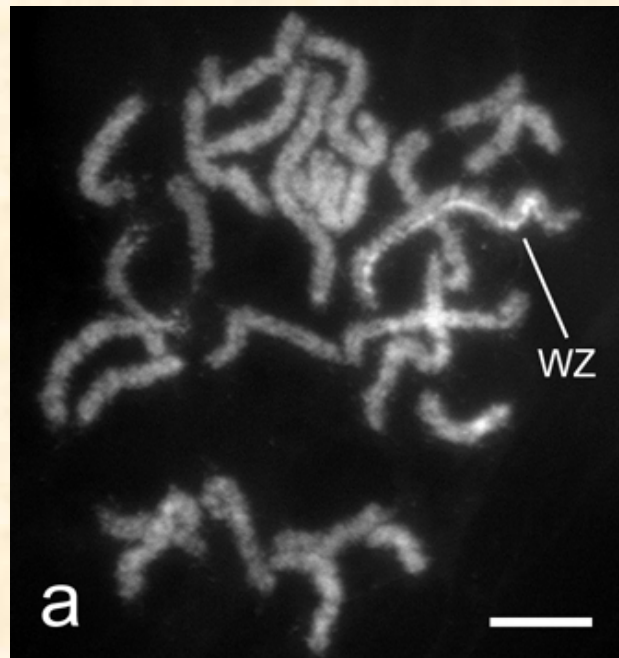
Compiled from: Traut and Marec (1996, 1997)
Marec and Novak (1998)
Lukhtanov (2000)

Variations of sex chromosomes in Lepidoptera

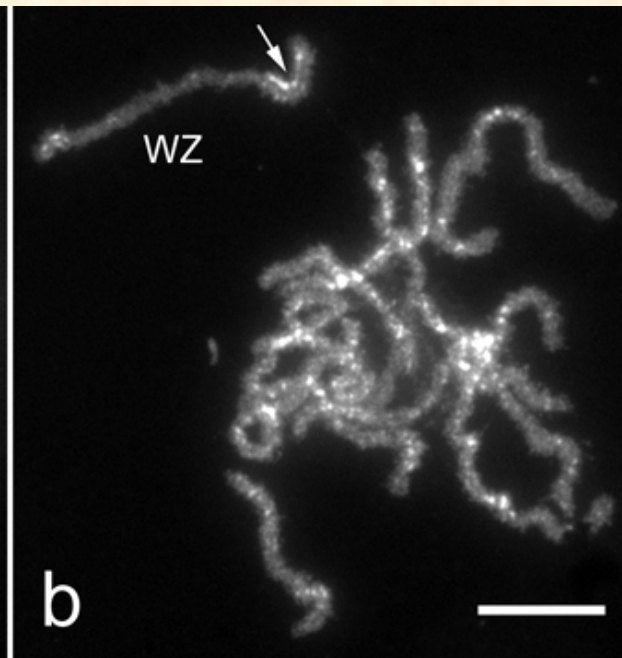


Marec et al. (2010) in: *Molecular Biology and Genetics of the Lepidoptera*. CRC Press.

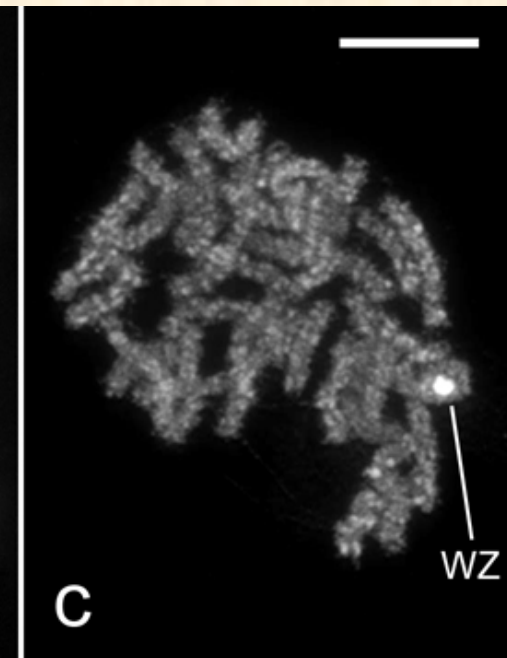
W-chromosome forms



Cydia pomonella



Orgyia antiqua



Bicyclus anynana

Marec et al. (2010) in: Molecular Biology and Genetics of the Lepidoptera. CRC Press.

Tools of molecular cytogenetics for W-chromosome identification



GISH

CGH

Our project:

Evolutionary history of
sex chromosomes in Lepidoptera

Origin of the W chromosome

W-chromosome origin Hypotheses

1) Translocation/Fusion of an autosome onto the Z chromosome:

- $T(Z;A) \Rightarrow$ neo-Z chromosome
- autosome homologue \Rightarrow neo-W chromosome

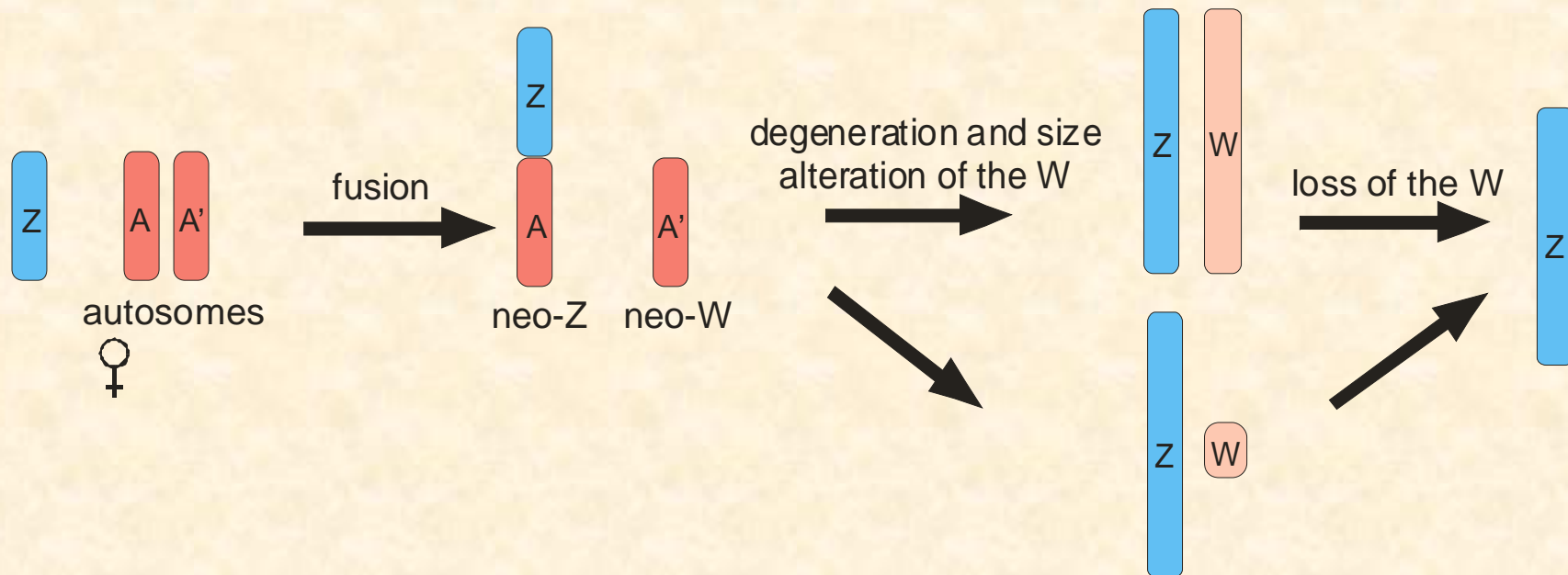
[Traut and Marec 1996]

2) B (supernumerary) chromosome \Rightarrow neo-W

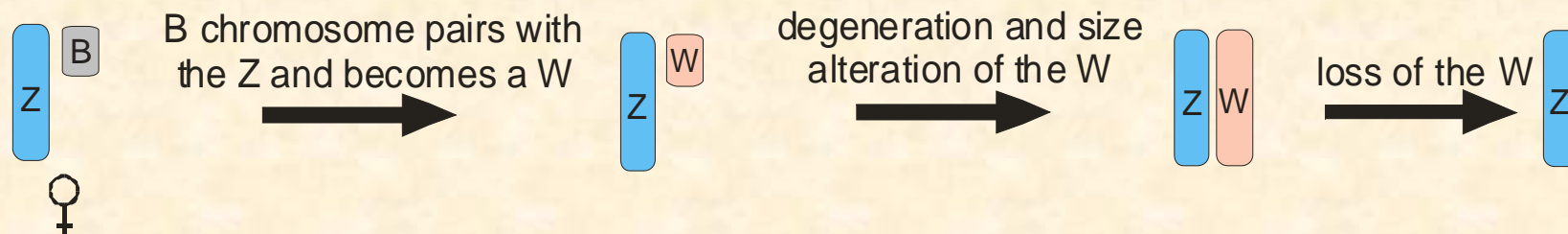
[Lukhtanov 2000]

Rise of the W chromosome

Hypothesis 1: fusion of the Z chromosome with an autosome



Hypothesis 2: rise of the W chromosome from a B chromosome

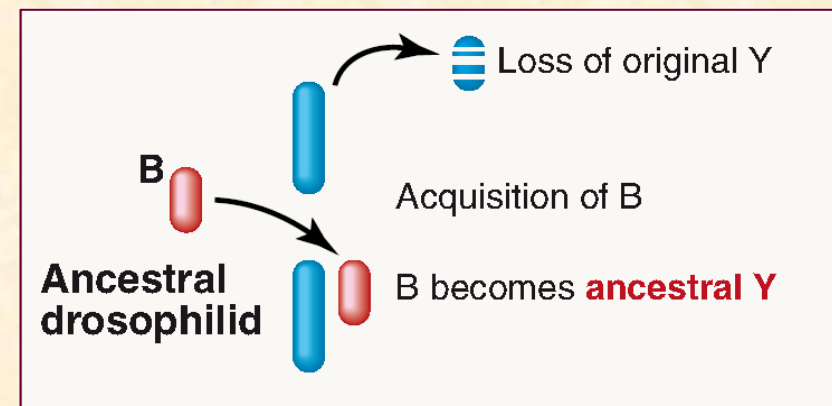
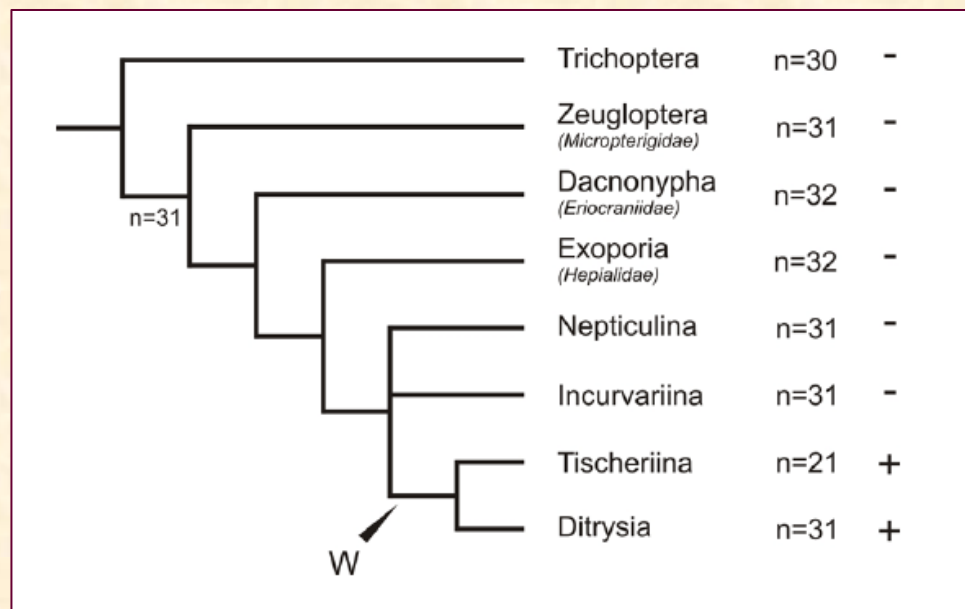


1) Autosomal origin of W – supported by:

- WZ pairs in a bivalent and form a regular SC
- neo-W repeatedly arose by T(W;A) in Ditrysia
- W and Z often belong to largest chromosomes

2) B-chromosome origin of W – preferred due to:

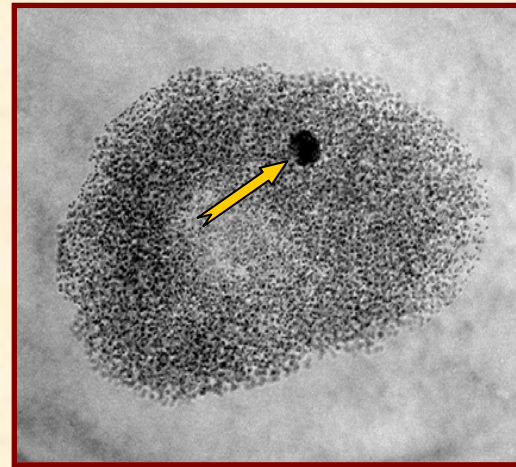
- autosomal origin lacks support in modal/ancestral chromosome number (n=31)
- similar origin predicted for Y in drosophilids



To answer the question of the W-chromosome origin

- 1) Does the W chromosome contain remnants of homology to an autosome or "autosomal part" of the Z chromosome? Or to a B chromosome???
- 2) Does the Z chromosome in Ditrysia contain an autosomal part or is it fully orthologous to the ancestral Z of primitive Lepidoptera and Trichoptera?

Probing the W chromosome for evolutionary studies in Lepidoptera



Our first approach:

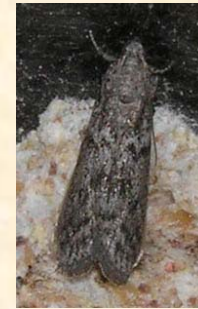
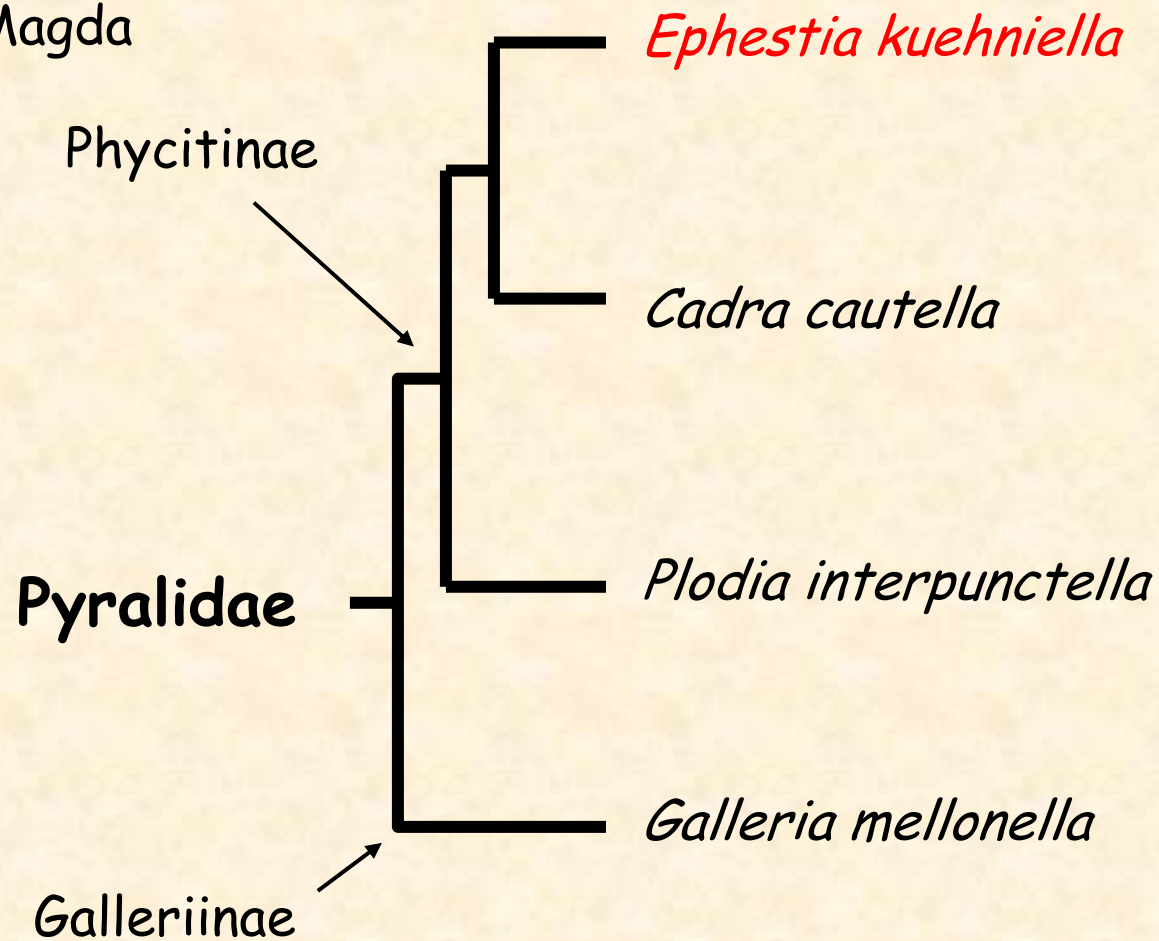
- preparation of W-chromosome painting probes by laser microdissection of W-chromatin bodies
- comparative chromosome painting (Zoo-FISH) of W chromosomes in related species





Magda

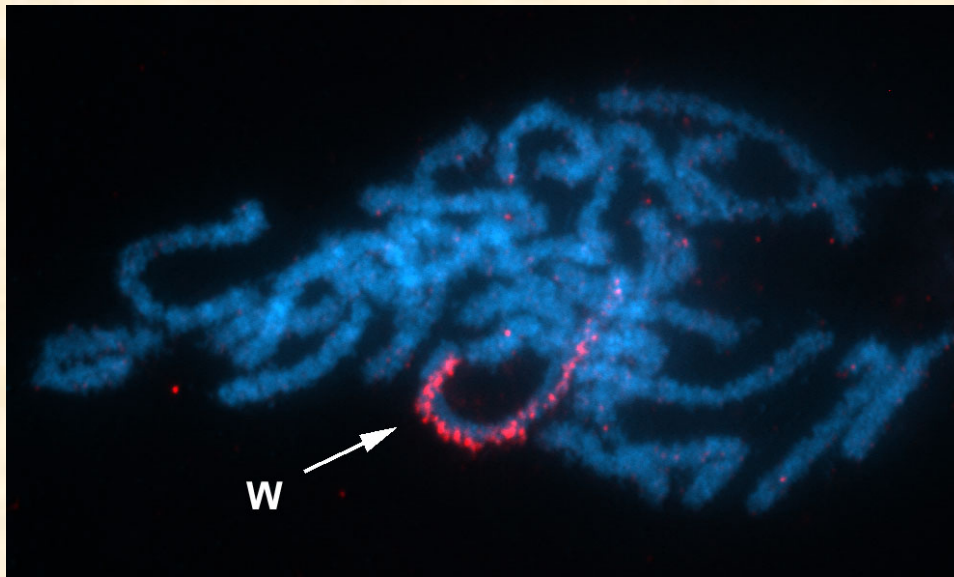
Zoo-FISH



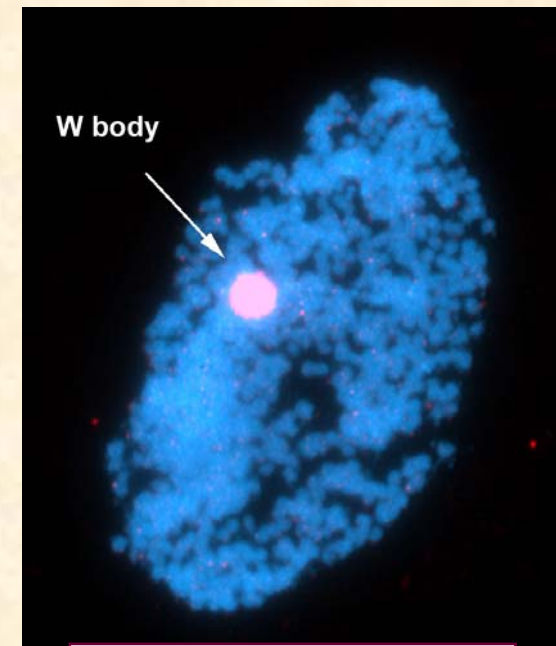
Vitkova et al. (2007) *Chromosome Res.* 15: 917-930.

Zoo-FISH with W-specific probe, derived from microdissected sex chromatin

- DOP-PCR amplification and labelling of sex chromatin sample
- Cross-hybridization of *Ephestia* W-probe to other pyralids



Ephestia kuehniella - pachytene oocyte



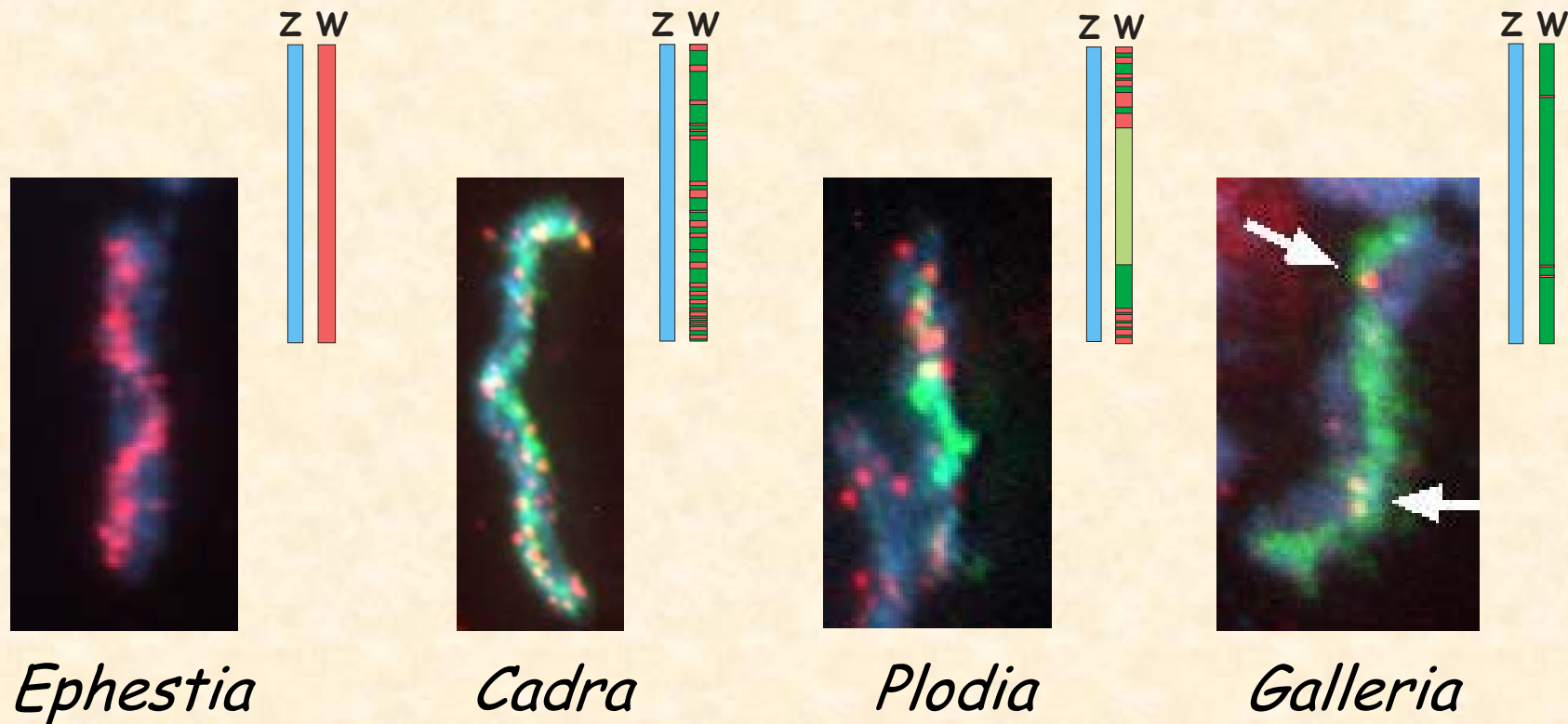
female interphase

Zoo-FISH with the W-specific probe from *Ephestia*

Red signal = W-probe from *Ephestia*

Green signal = W-probe from the same species as chromosomes

Blue = DAPI counterstaining



Conclusions

Results of Zoo-FISH in pyralids suggest rapid molecular evolution of the W chromosome.

W chromosome cannot be used for comparative evolutionary studies between higher taxa.

Z chromosomes for tracking sex-chromosome evolution in Lepidoptera

Two different approaches:

- (i) preparation of Z-chromosome painting probes by laser microdissection and their use for
 - creation of Z-DNA libraries and search for molecular markers of Z chromosomes
 - comparative chromosome painting (Zoo-FISH)
- (ii) conserved synteny mapping of orthologs of Z-linked genes known from *Bombyx* or other sp.
 - by BAC-FISH, if BAC library available
 - by direct gene mapping, if BAC library not available

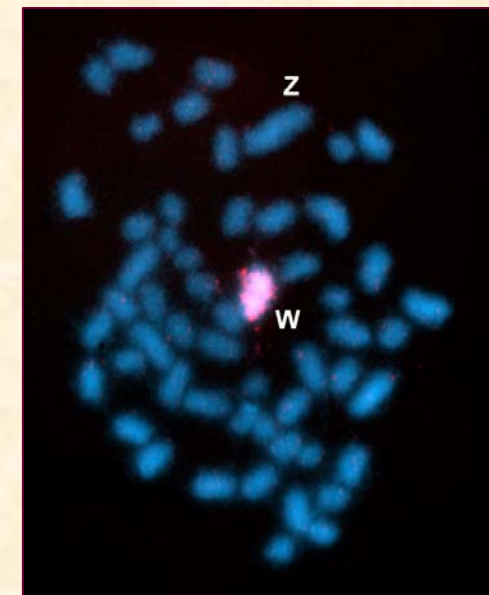
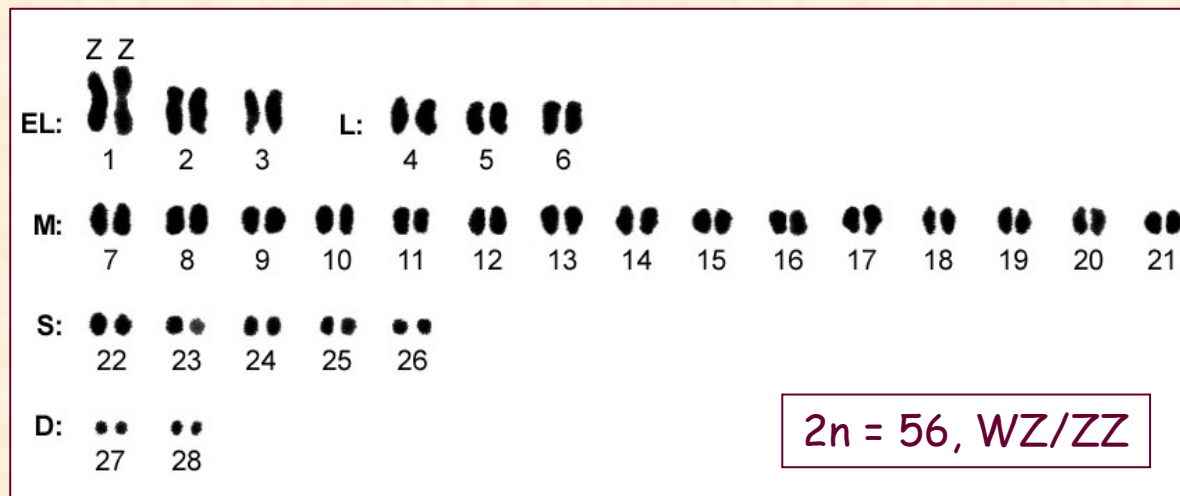
Codling moth (*Cydia pomonella*)

F: Tortricidae



Why codling moth?

- Karyotype and sex chromosome analysis done; Z largest element; tools of molecular cytogenetics established (Fukova et al. 2005, 2007), W/Z molecular markers developed (Fukova et al. 2009)
- Bacterial Artificial Chromosome (BAC) library available
- Most basal species of Lepidoptera phylogeny with some genetics
- Another motivation: development of genetic sexing strains for SIT control of this pest (Marec et al. 2005, 2007)

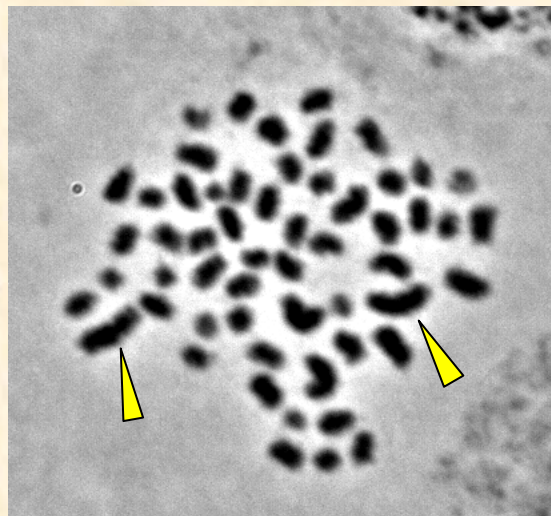


Creation of Z-chromosome library

- laser microdissection of Z chromosomes from spermatogonial metaphase sets
- DOP-PCR amplified samples tested as Z-probe by FISH
- samples cloned, sequenced and analysed

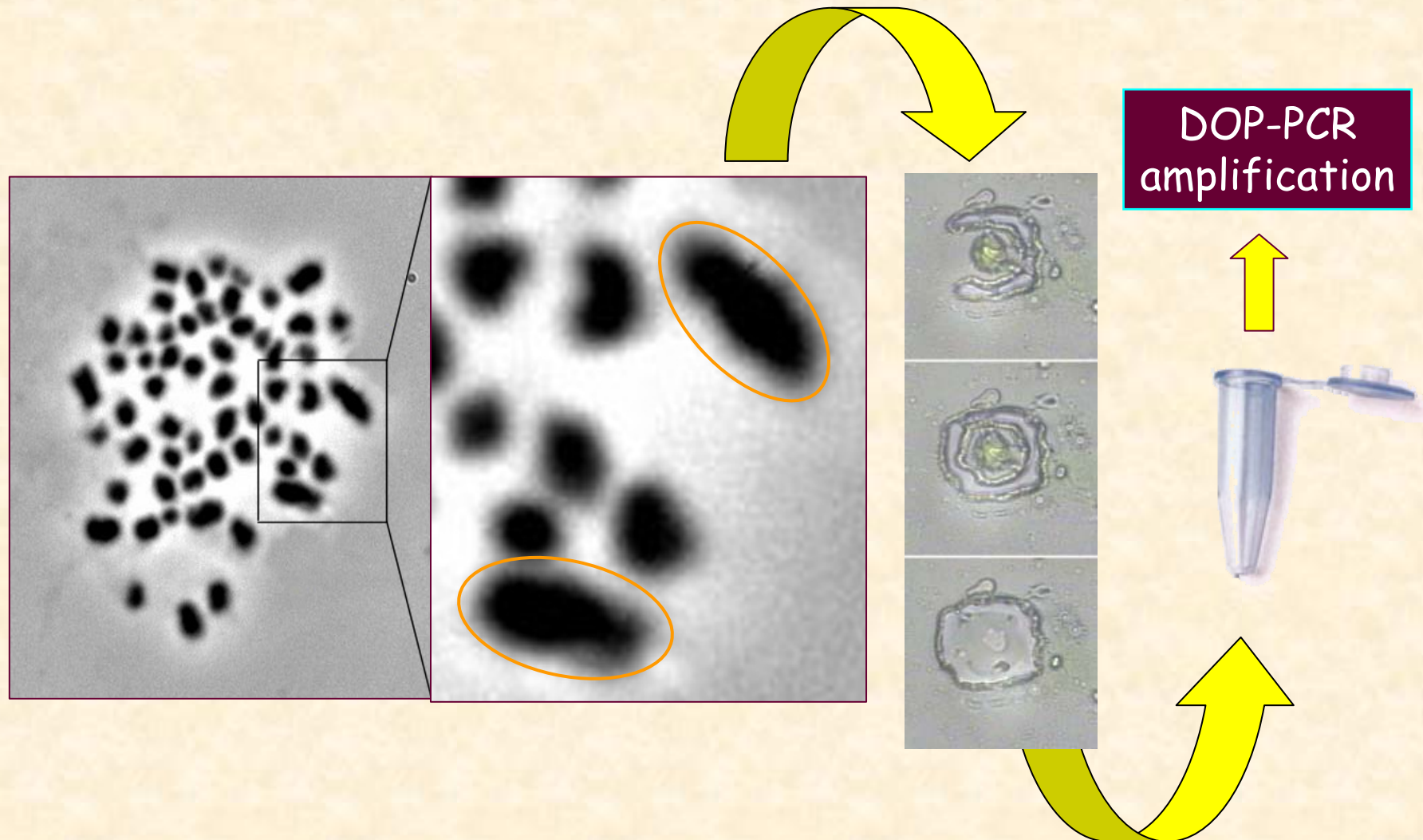


Jindra



Magda

Laser microdissection of Z chromosomes



A low specificity of FISH with
Z-chromosome-derived probes
in codling moth males

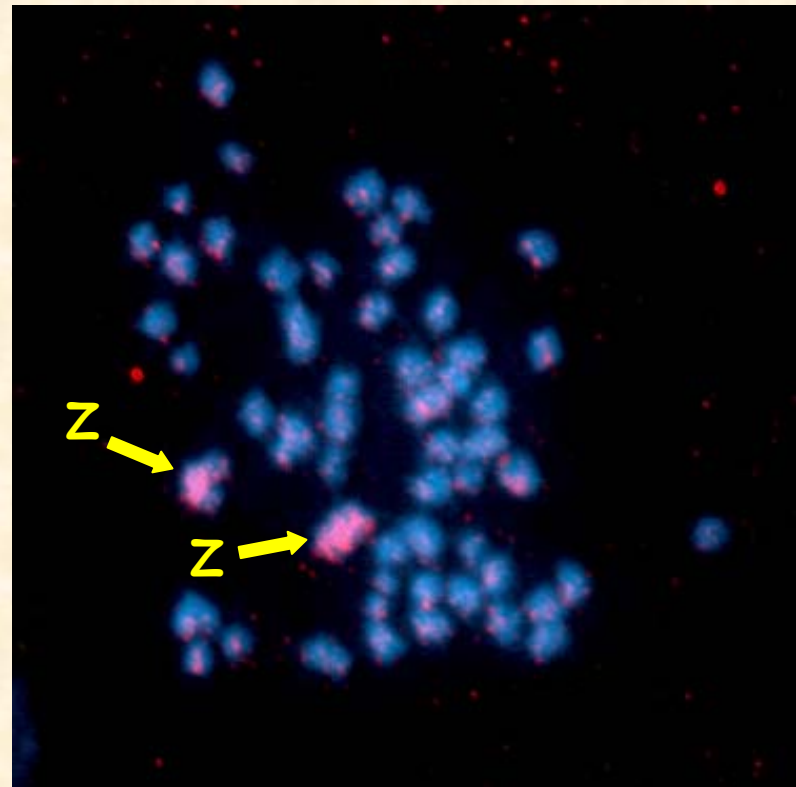


FISH with Z-chromosome-derived
probe highlighted W in females



Alternative system for balanced amplification of laser-microdissected Z-chromosome samples

- amplification using WGA4 kit (Sigma-Aldrich)
 - even a single chromosome can serve as a template
- re-amplification using WGA3 kit



Laser microdissection of Z chromosomes

Summary

- Low specificity of Z probes and W-painting: DOP-PCR amplified mainly repetitive DNA, enriched in W, as confirmed by cloning, sequencing, and Southern hybridization
- Balanced amplification using WGA4 / WGA3 system => sufficient specificity for Z chromosomes
- Z-painting probes will be used to determine similarity of Z chromosomes between related species of the family Tortricidae:
 - Plum fruit moth (*Cydia funebrana*)
 - Oriental fruit moth (*Grapholita molesta*)
 - European grapevine moths (*Lobesia botrana* and *Eupoecilia ambiguella*)

Synten mapping of *Bombyx mori*
Z-linked genes in codling moth by
BAC-FISH

Mapping codling moth orthologs of Z-linked genes by BAC-FISH

First step: search for orthologs of *Bombyx* Z-linked genes



Petr

Candidate genes:

- *period* => *Cpper* (Fukova et al. 2009)
- *Bmkettin*
- *apterous*
- *Triose-phosphate isomerase* (*Tpi*)
- *Lactate dehydrogenase* (*Ldh*)
- *6-Phosphogluconate dehydrogenase* (*6-pgd*)

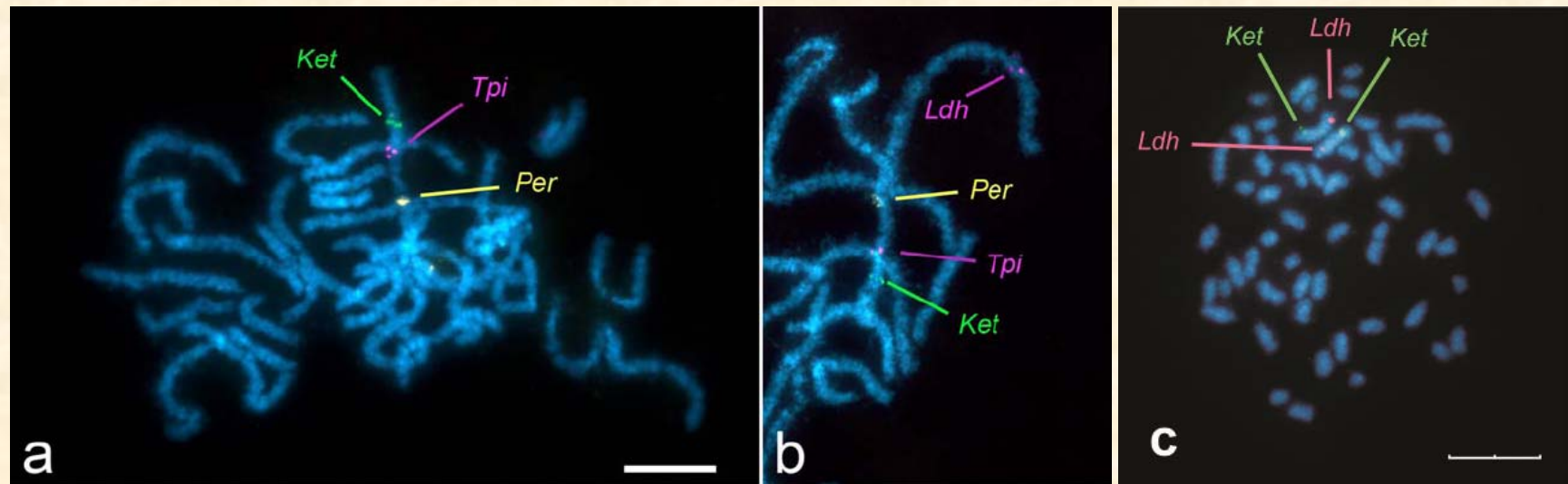


Martina

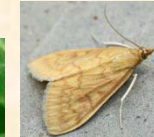
Second step: screening BAC library with isolated orthologs

Third step: BAC-FISH mapping

BAC-FISH mapping of codling moth orthologs of Z-linked genes



Comparison of Z-chromosome maps



Pap

Hel

Bm

Cp

On

Ap



ephrin, receptor
titin

EGF receptor

IDGF4

p-glykoprotein

*NADH ubichinon
oxidoreduktasa*

CyP303a1

kettin

apterous

Tpi

katalasa

period

6-pgd

Ldh

henna

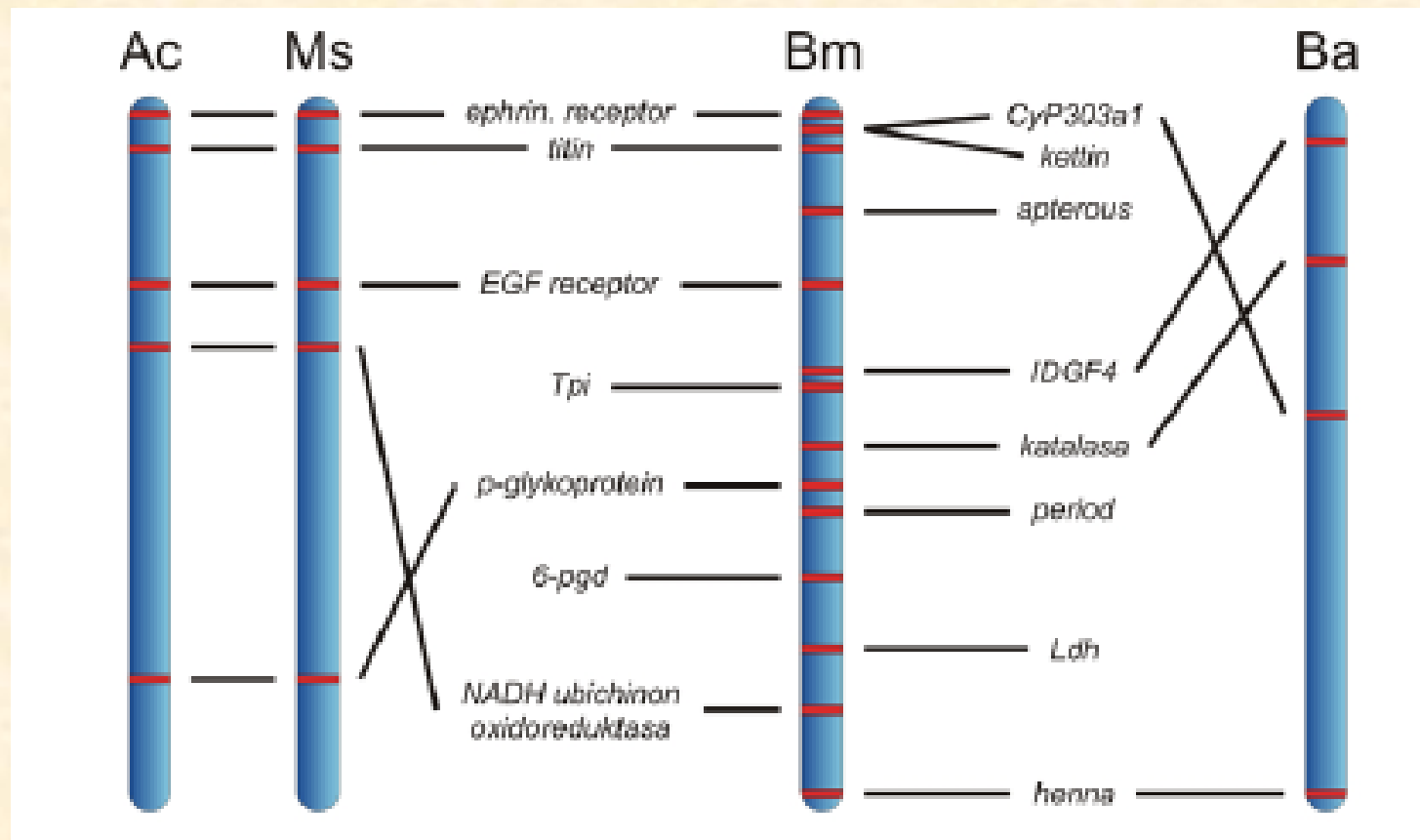
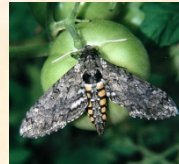
kettin, titin
Tpi, period
6-pgd, Ldh

apterous
Tpi

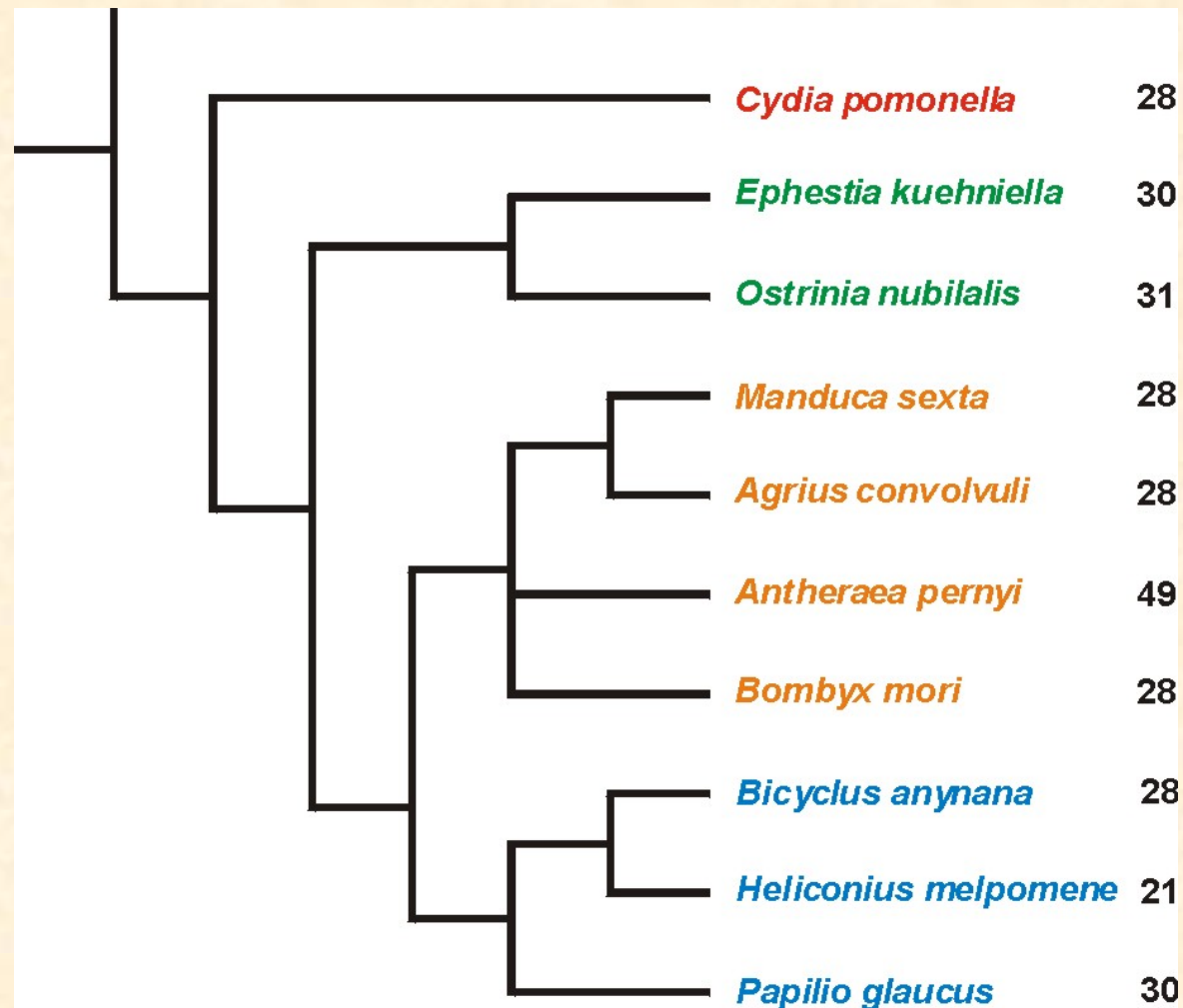
6-pgd
apterous

period

Comparison of Z-chromosome maps



Conserved synteny of Z-linked genes



BAC-FISH mapping of codling moth genes

Summary and Conclusions

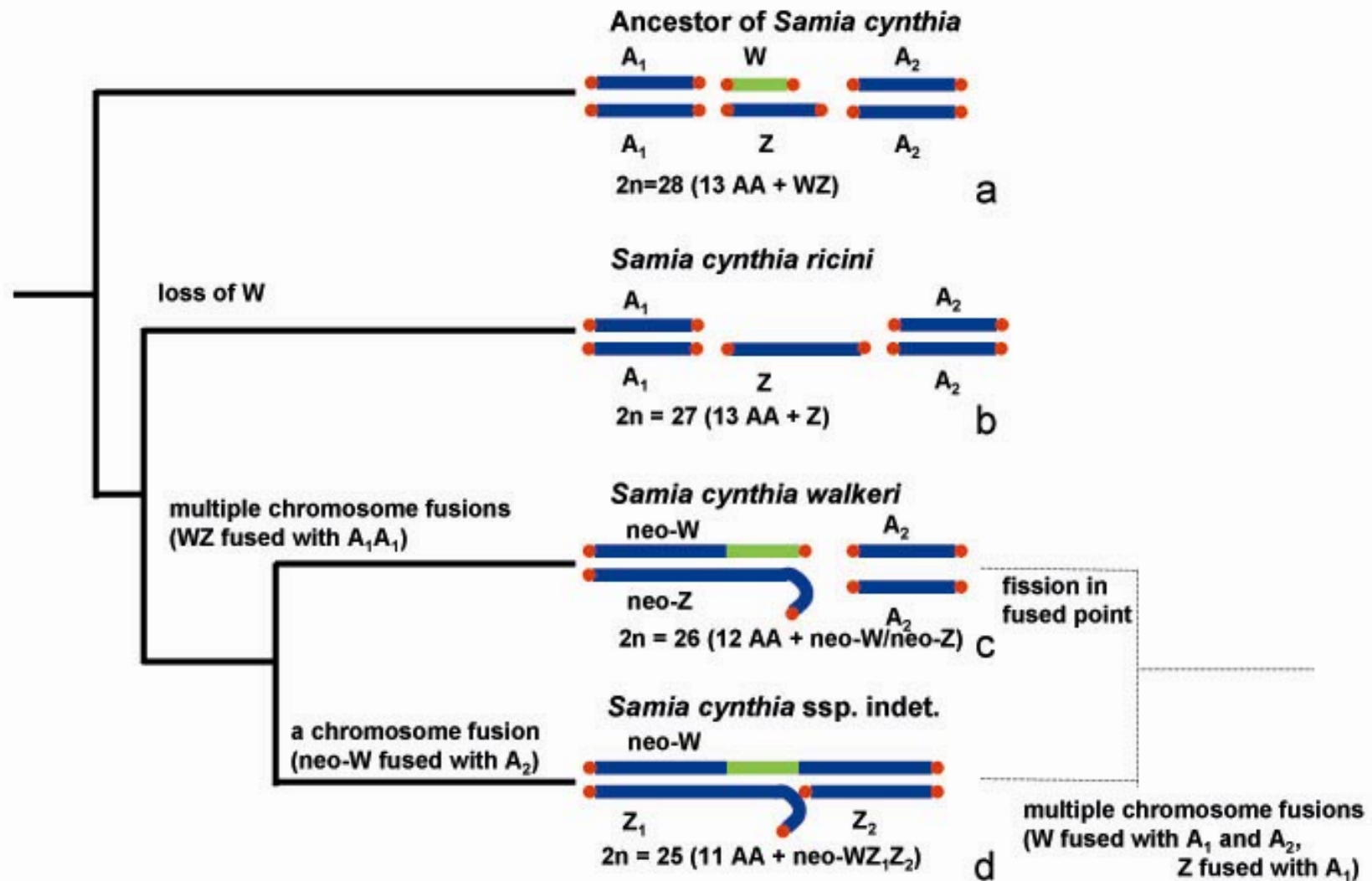
- Study confirmed conserved synteny of genes and conserved gene order between Z chromosomes of *B. mori* and *C. pomonella*
- Conserved synteny of Z-linked genes between all species studied including codling moth as representative of a basal ditrysian clade suggests the common origin of Z chromosomes in Ditrysia
- Results add further evidence to stability of lepidopteran genomes, in spite of holokinetic nature of their chromosomes

A case study

Wild silkworm (*Samia cynthia*) F: Saturniidae

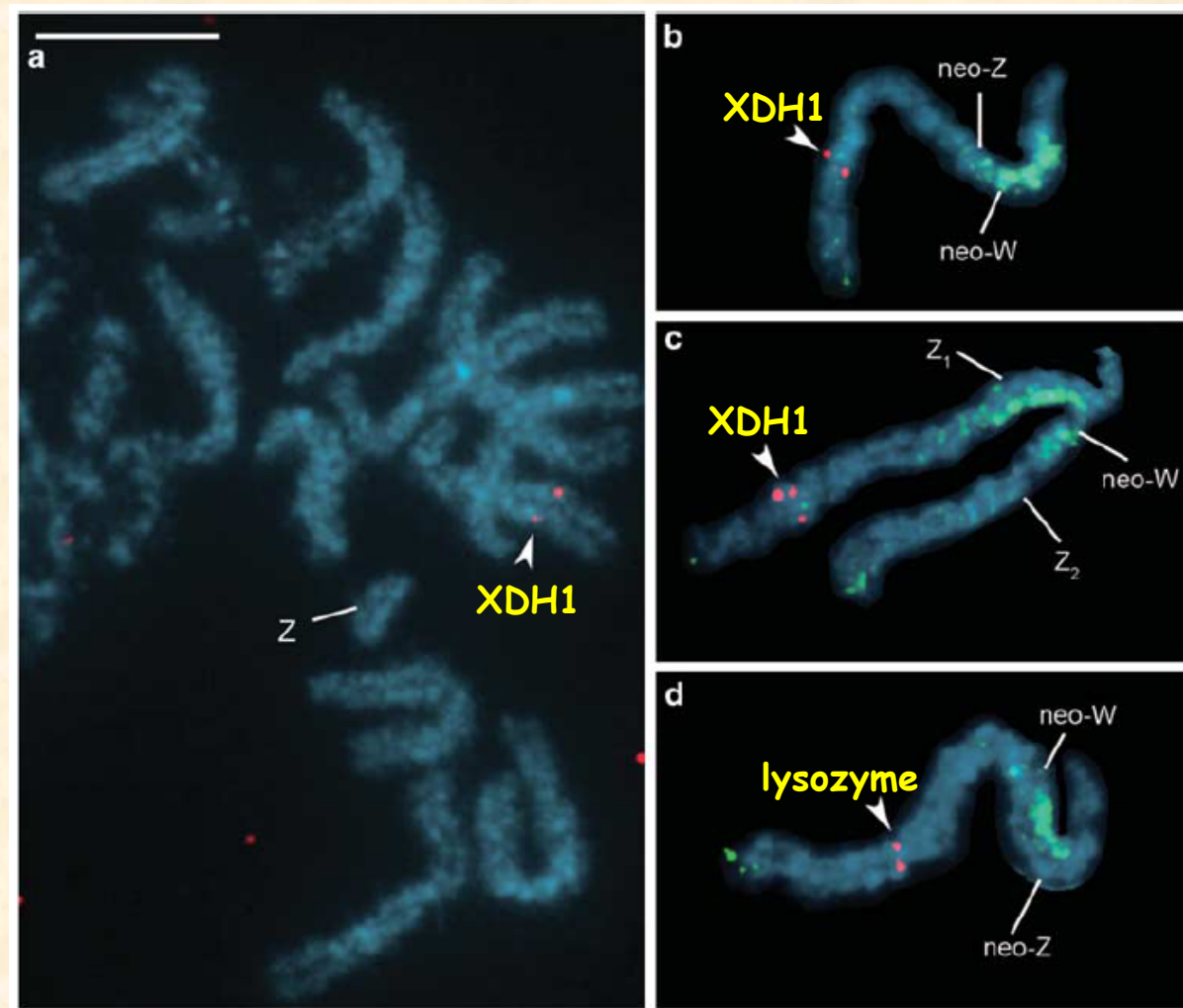
Sex-chromosome evolution in geographical subspecies

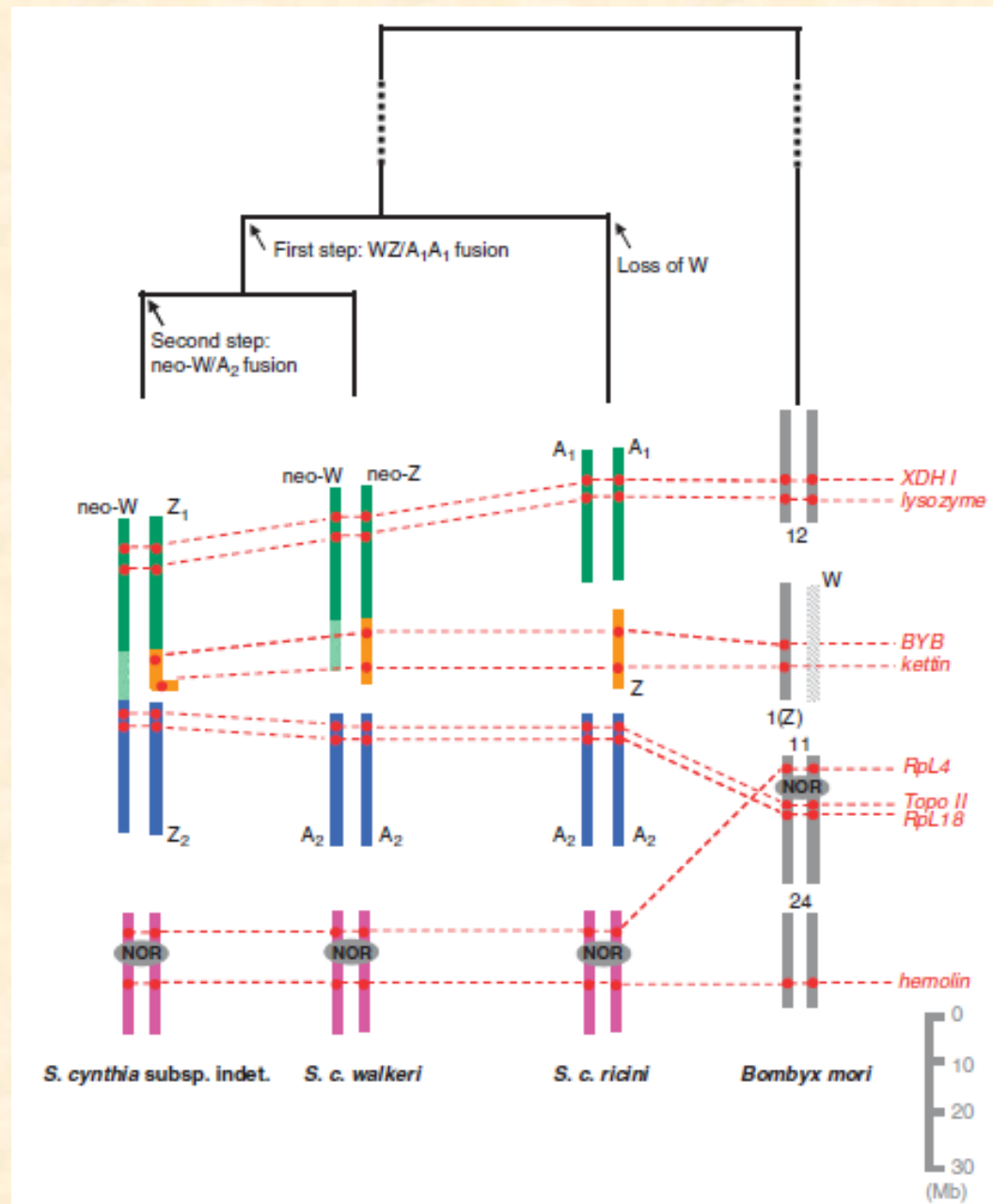




Yoshido, Marec, Sahara (2005) *Chromosoma* 114: 193-202.

Direct FISH-mapping of *Bombyx mori* orthologous genes on chromosomes of *Samia cynthia ssp.*





Yoshido, Sahara, Marec, Matsuda (2010) *Heredity*. Online 28 July 2010.

Acknowledgements



Walther Traut
Lübeck



Ken Sahara
Sapporo



Petr Nguyen



Magda Vitkova



Atsuo Yoshido
Sapporo



Iva Fukova



Jindra Sichova



Martina Dalikova

Svatava Kubickova
Brno

Lisa Neven
Yakima

Marie Korchova



*Budweiser
Budvár*



České Budějovice