B chromosomes

Jones, 1995; Bougard and Jones, 1997; Camacho et al., 2000; Pokorná & Reifová, 2021

Chromosome types:

- A = autosomal chromosomes
 - Contain all the genes
- B = supernumerary or accessory
- Discovered 1907. Wilson; Science 26:870-871 in bug, Metapodius
- Term 'B' derived from maize in 1928 (Randall; Anat. Record 41:102)

B chromosomes

- Are not duplicates of any A chromosome
- Exist only in some individuals of a population
 - Found in 1372 spp. of seed plants, of which 12 are conifers.
 - o Also in ferns & fungi
 - = 10-15% of flowering plants
 - o All are outbreeders
 - Found in 500 animal spp. (Insects, fish, birds)
 - Found in 1.94% of mammalian spp. (Vujosevic et al, 2018)
- Are dispensable
- Do not pair or recombine with A chromosomes
- Some pair with each other during meiosis
- About ¹/₂ are highly heterochromatic, esp. in animals
- Their number is very variable
 - 0 to 34 (in maize)
 - Accounts for up to 155% of the DNA
- Have genes that deteriorate in the absence of purifying selection
 - Also have genes for their maintenance



Figure 1. Niwa & Tsujimoto 1992, Plant Breeding109:78-81. Rye root-tip and diakinesis chromosomes. Arrows indicate B chromosomes (note pairing at diakinesis). Arrow heads are the satellite chromosomes. Bar = $10 \mu m$.

Figure 2. Top: A B-chromosome from maize. Bottom: B chromosomes from rye plants from different countries.



- Rhoades and Dempsey, 1973 [J. Hered. 64:12-18]
 - B chromosomes have the ability to eliminate knobs from A chromosomes

Rosato et al., 1998

As altitude increases, B number increases

Suggests function



Mitotic stability

Figure 3. Rosato et al., 1998

- In about ¹/₃ of cases, B chromosomes are mitotically stable
 esp. in grasses, but rare in Allium
- Can be lost in roots and other organs of some spp.

Mitotic drive

• In Crepis capillaris, Bs accumulate in reproductive organs

Meiotic drive

E.g., *Lilium callosum*: 1B × 0B ⇒ 80% 1B rec ⇒ 50% 0B, 50% 1B



Non-disjunction

- 60% of spp. exhibit B drive, most due to non-disjunction during the second pollen mitosis, as observed in maize
- The sperm cell with the 2 B's is produced 50-100% of the time, and will preferentially fertilize the egg 60- 70% of the time

Figure 4. bdlilies.com



Figure 5. Non-disjunction of B chromosomes during microgametogenesis of maize. Roman, 1947

- In rye, non-disjunction occurs at 1st mitosis (happens 85% of time for & and %)
- The phenomenon was first observed in rye by Hasegawa, 1934

Figure 6. From **Hasegawa 1934**, redrawn by **Houben 2017**. A- pollen mitosis; B- nondisjoined, lagging B chromosomes; C - disjoined B's; D - B's showing non-disjunction



Kamayo, 1957: B chromosomes in Lillium callosum

- In this species, it is the micropylar megaspore that gives rise to the gametophyte
- Bs tended to lie on the micropylar end of the Met I spindle, and again of the Met II spindle, and hence get included preferentially in the egg.

Stage:	% Bs @ micropylar pole
Metaphase I	75.4
Anaphase I	84.1
Met II / Ana II	80.0
Postmeiotic mitoses	73.1
% Eggs with B's	83.8

Table 1. Kamayo, 1957: B chromosomes in Lillium callosum

II - B Chromosomes • Spring 2025, PBGG 8890

Wu et al, 2019

Showing the 1st pollen mitosis. Note that 85% of the time, the B chromosomes will end up in the generative cell, which will divide to form the sperm.

Carlson and Roseman, 1992



Using translocations with chromosome 9 of maize, have identified 4 regions of the B chromosome necessary for non disjunction to occur:

- heterochromatic block = nonessential
- region 2 = responsible for preferential fertilization
- region 3 = sticking receptor
- region 4 = regulates rate

Regions 2, 3, and 4 also prevent meiotic loss during megasporogenesis. The unpaired B chromosomes go to the functional pole and get included in the egg (not due to nondisjunction).







Figure 7. Wu et al, 2019 (Houben lab)

Blavet et al, 2021



- B-repeat responsible for non-disjunction in cis
- Derived from knob sequence → delayed replication may cause sister chromatids to not separate in time for mitosis
- Preferential fertilization related to sequences linked to centromere
- Trans factor for non disjunction located at distal end, which contains 34 genes

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Jones and Puertas, 1993; Jones and Pasakinskiene, 2004

- In rye, a gene for non disjunction is located in the distal end of the long arm
- Causes sequences around the centromere to stick together during pollen mitosis
- If the end of the chromosome is missing, then centromeric sequences do not stick together
- However, it does not explain why there are different frequencies of B chromosomes in different populations



Banei-Moghaddam et al., 2012

• Non-coding RNA coded by the distal end involved in stickiness and asymmetrical spindle

Chen et al., 2024

- Candidate gene: DCR28 =coding a microtubule-associated protein
 - 15 copies in rye B chromosome
 - Highly expressed at 1st pollen mitosis

Balancing forces

Müntzing, 1963



As B number increases, plant growth decreases in Swedish varieties & increases in Korean ones, but, fertility decreases

Pereira et al, 2016

- B subtelomeric domain highly enriched in E3900-related sequences
 - B-specific repetitive sequences
- Full length E3900 is 3.9-kb long
- A truncated version is 2.7 kb
 - Contains heat-responsive cis-acting regulatory elements
- The short version is upregulated at pachytene under heat stress.
- Pachytene cells with non-truncated B's also exhibit fewer chromosomal abnormalities



0.0	410	AD 11	
0B	28	2B-del	
94 ± 9 (108)	$29\pm19~(90)$	55 ± 17 (30)	
t-test (P)	0.01	0.04	

How B's influence the A's



Figure 8. Differential expression of 18 known miRNAs



Genetic control

Romera, Jiménez, and Puertas, 1991

- Looked at different rye cultivars from Korea
 - o 'Paldang'-- 20% of its plants naturally have B chromosomes
 - 'Puyo' -- 60% of its plants naturally have B chromosomes
- Synthesized a population of Paldang with 60% B chromosomes, and one of Puyo with 20% B chromosomes

Population:	Year	Madrid		La Coruña		
		% B's	#B/plant	% B's	# B/plant	
Paldang 20	1987	17	0.33			
	1988	21	0.42	22	0.44	
	1989	20	0.40	30	0.63	
Paldang 60	1987	55 ↓	1.21			
	1988	42	0.89	54↓	1.19	
	1989	38	0.82	47	1.02	
Puyo 20	1987	22 [↑]	0.44			
	1988	47	0.98	19 ↑	0.39	
	1989	57	1.30	34	0.66	
Puyo 60	1987	57	1.36			
	1988	71	1.78	52	1.15	
	1989	74	1.74	50	1.04	

• These were grown in 2 locations in Spain for 2 seasons:

The trend is for the populations to return to their original frequency of B chromosomes, suggesting that genetic factors probably determine the equilibrium frequency of B chromosomes.

- selfish

- adaptive value

Interplay between these two forces determines the number of B's found in a given plant.

- Some Bs appear to survive as they benefit the host; others do so due to drive.
- This is balanced against detrimental effects their accumulation provides to the host.
- There must be some sort of genetic control

González-Sánchez et al., 2003



The mBt autosomal gene in corn controls B preferential fertilization in maize.

- Eggs with the mBt^h are preferentially fertilized by B-containing sperm
- B's in eggs controlled by the fBt gene in the A chromosomes

• The fBtl allele is dominant -> loss of B's at meiosis.

Origin & composition

Origin is unknown. In plants may be derived from fragment of centromere and surrounding heterochromatin

o Evolve rapidly to prevent pairing with A chromosomes

Martis et al 2012

3R

The rye B chromosome

Rye B's derived from parts of chromosomes 3 & 7

- Therefore, the first step appears to be a reciprocal translocation that results in a 'proto-B' chromosome
- Followed by loss of genes and other homologous sequences to A chromosomes to prevent pairing
- And accumulation of other DNA sequences

A chromosomes

Accumulation also happens in A chromosomes, but many of them get discarded by recombination and segregation

A chromosome rearrangements via reciprocal translocations



Marques et al, 2018



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Use in breeding

Birchler, 2015

Limitation to artificial chromosomes has been the inability to clone centromeres and transfer them due to their epigenetic nature. So, need an alternative

Transgenes

• Transgenes normally integrate into chromosomes- requires recombination on both sides

• But, if flank the transgene with telomeric sequences on one side, it will still integrate, but the side with telomeric sequences will not integrate

- o Therefore, end of chromosome is lost
- Integration into A chromosomes can be lethal, as lose necessary genes
- Integration into B chromosomes does not affect viability; therefore are preferentially recovered
- o Result is a minichromosome

