



Meiotic Recombination in Ruminant Livestock Species

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Goals

1. To quantify, characterize and compare the number of meiotic recombination/crossover (CO) between beefalo, cattle and sheep.
2. Test the relationship between the number of CO and chromosome length.
3. Examine the chromosomal structural integrity.

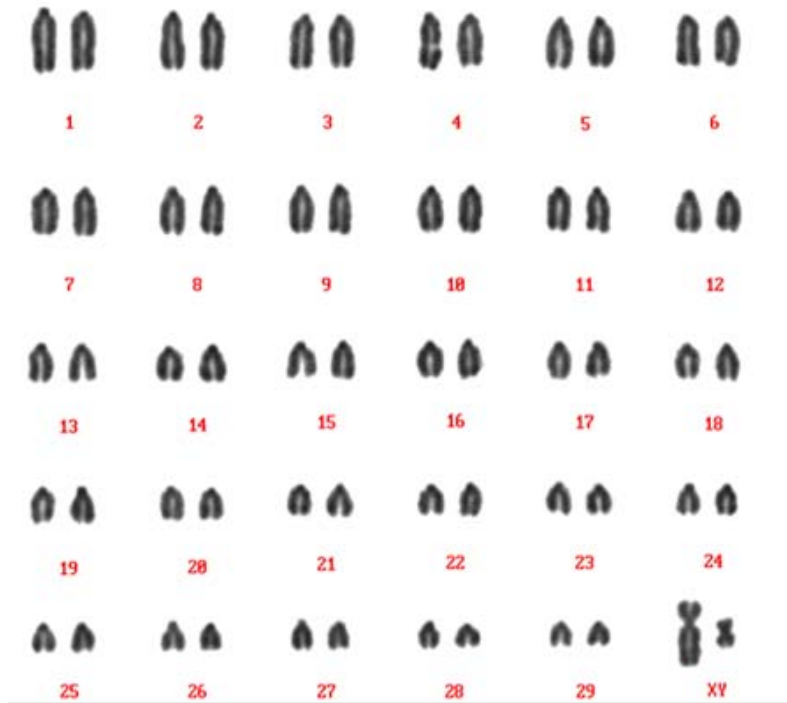


Meiotic Recombination

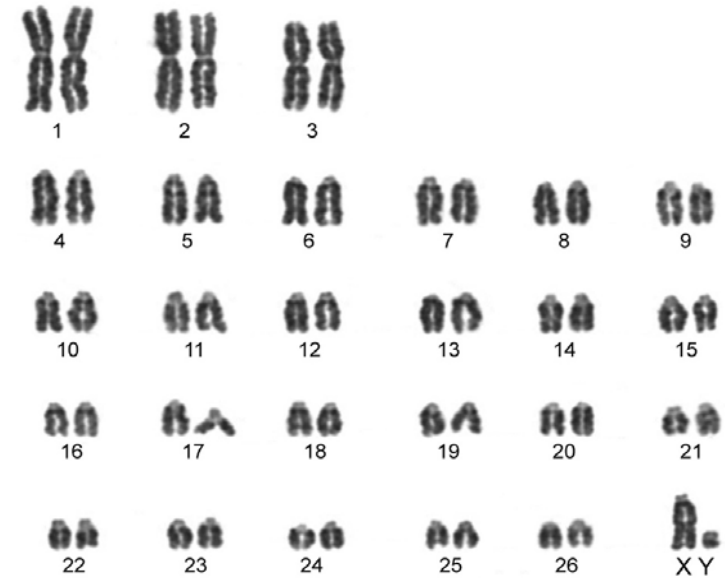
- Exchange of genetic material contributes to genetic diversity
 - NOT random
 - Associated with the number of chromosome arms
 - Interference
 - “Hotspots”
 - Sexual dimorphism
- Ensure faithful chromosome segregation

Karyotype: Chromosome Arms

Cattle

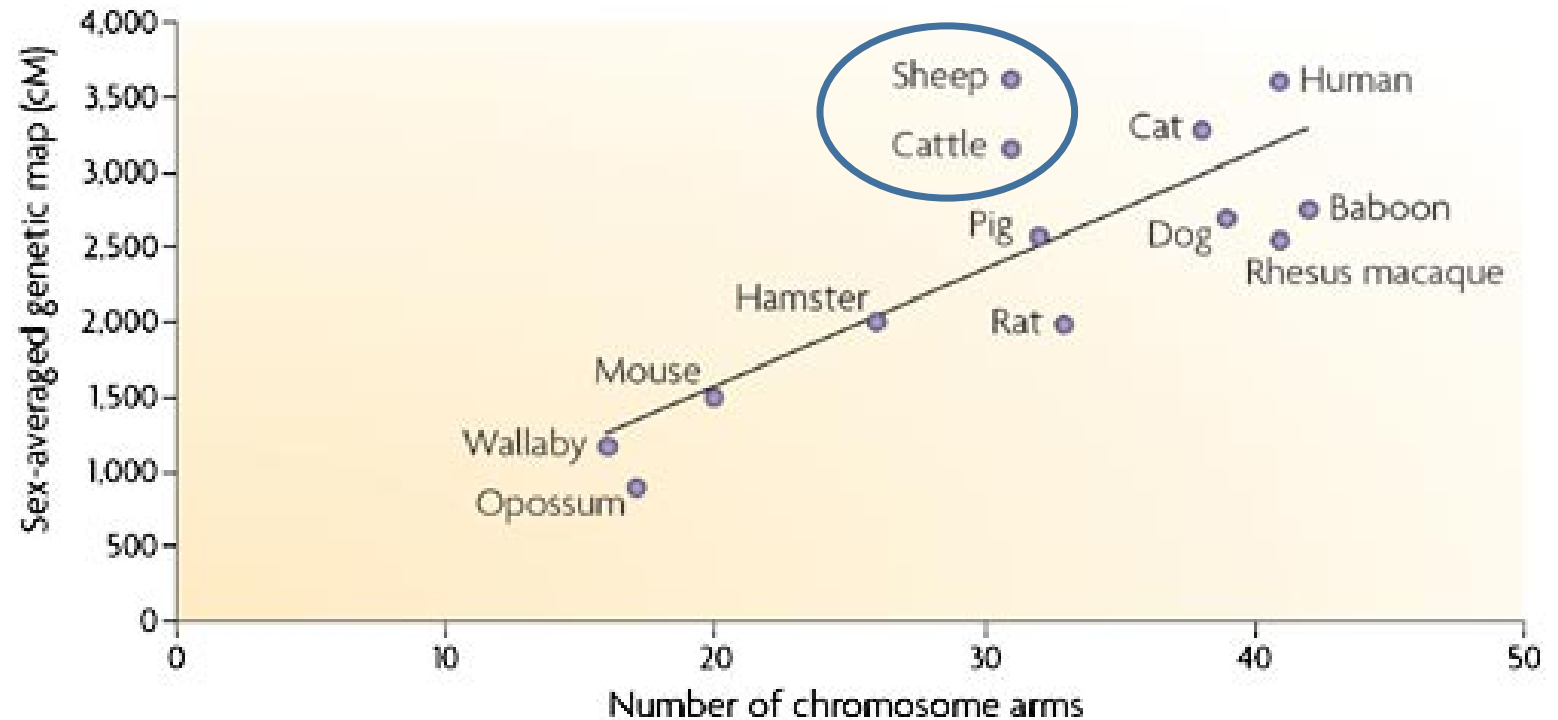


Sheep



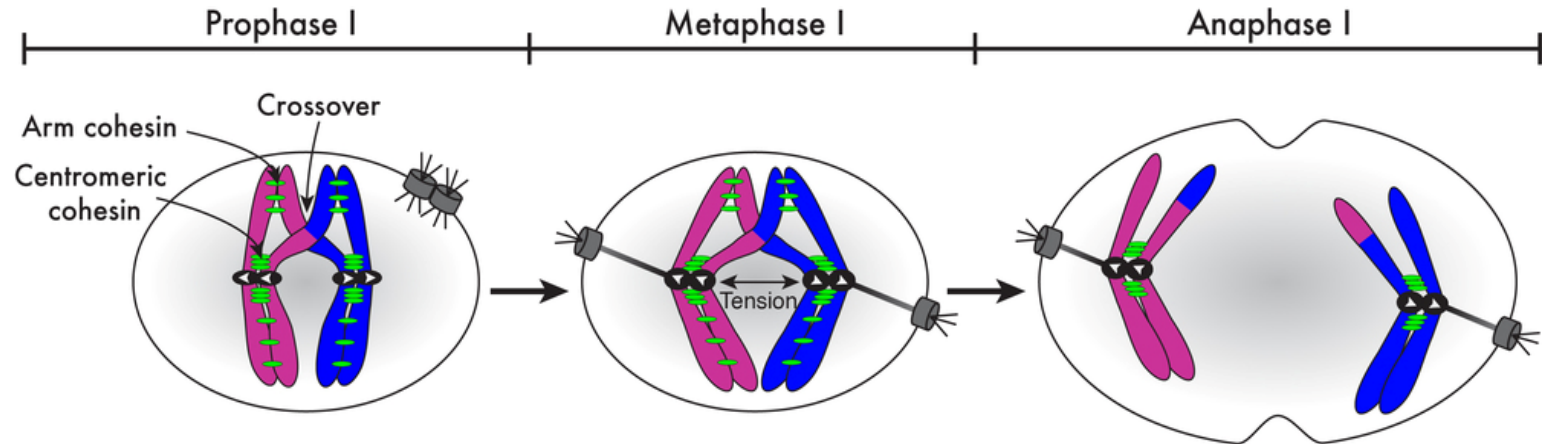
Mammalian Recombination

Sex averaged linkage map comparison to number of chromosome arms in different mammalian species.



Meiosis

- Ensure faithful chromosome segregation



- Mis-segregation or non disjunction leads apoptosis in males

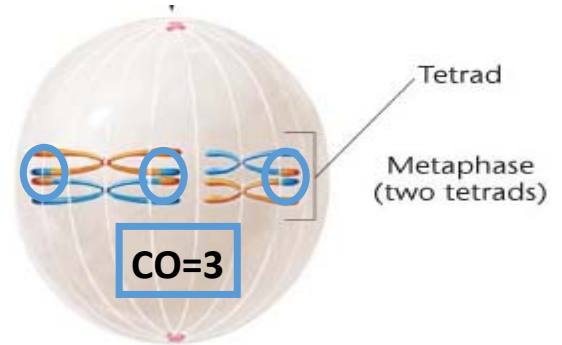


Method: Cytogenetics

- Allows us to characterize recombination directly
 - Has distinct advantages compared to linkage mapping
 - Reference genome
 - Large number of progeny
- Determine structural integrity



Prophase I
(synapsis)



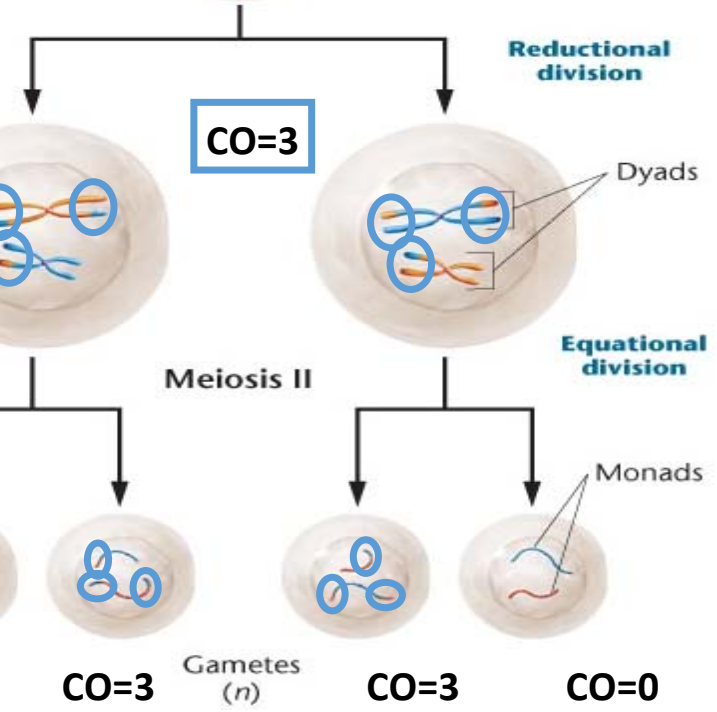
Tetrad
Metaphase
(two tetrads)

CO=3

Recombination in prophase (pachytene cell) observed by immunofluorescence.



Recombination observed in gametes are lower due to independent assortment of chromatids.



Reductional division

CO=3

Dyads

Equational division

Meiosis II

Monads

CO=0

CO=3

Gametes
(*n*)

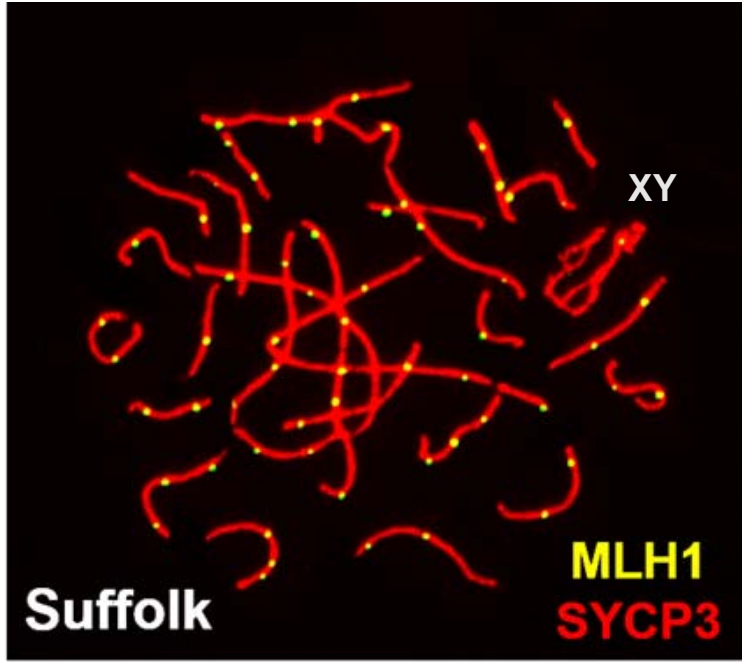
CO=3

CO=0

Average crossover counted in offspring = 1.5

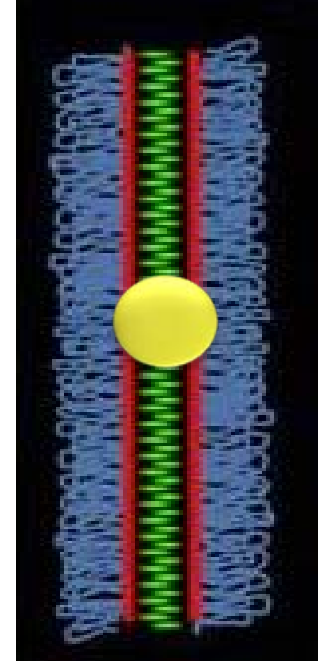


Immunofluorescence



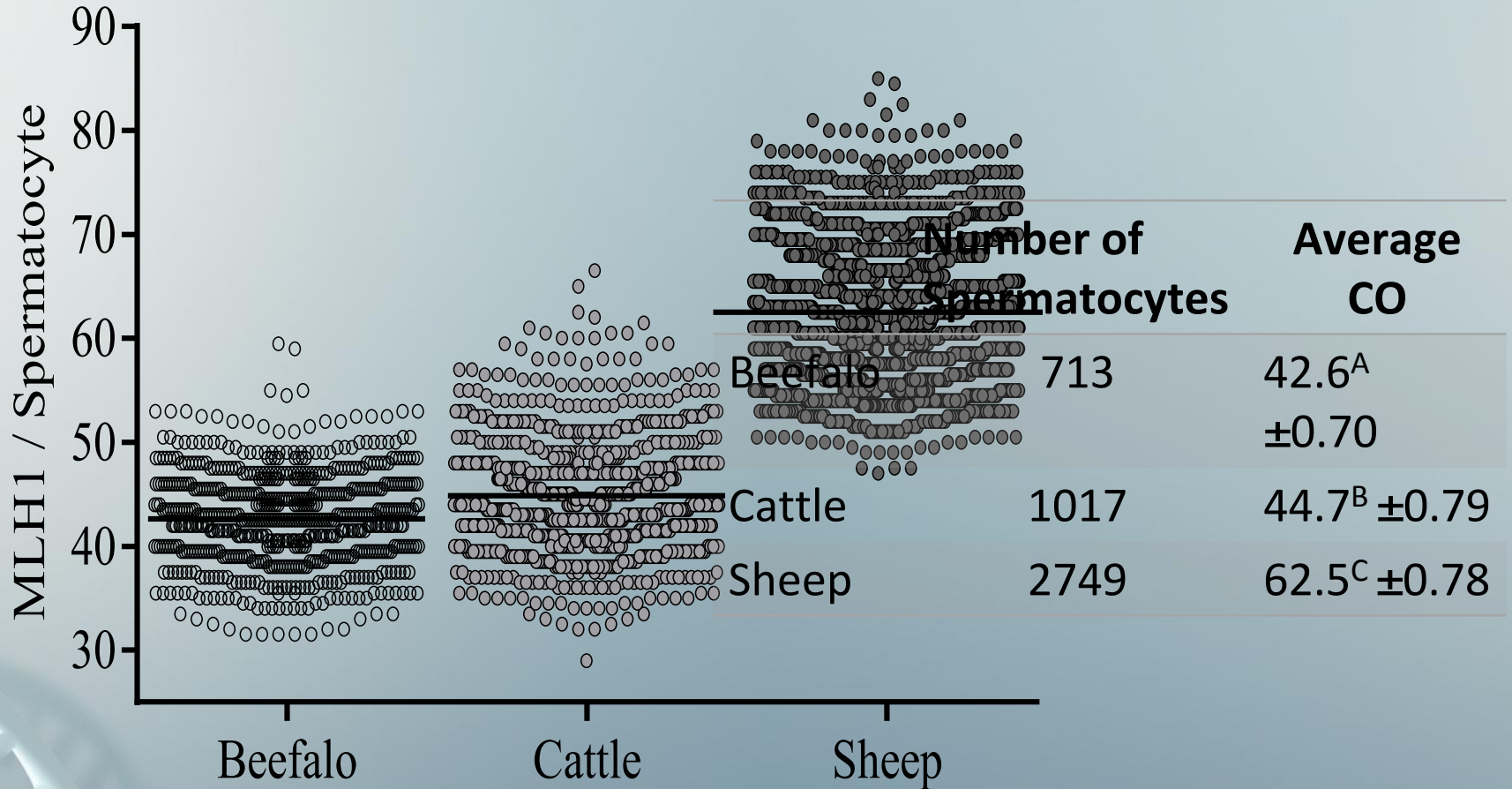
Synaptonemal
Complex (SC)

SYCP3



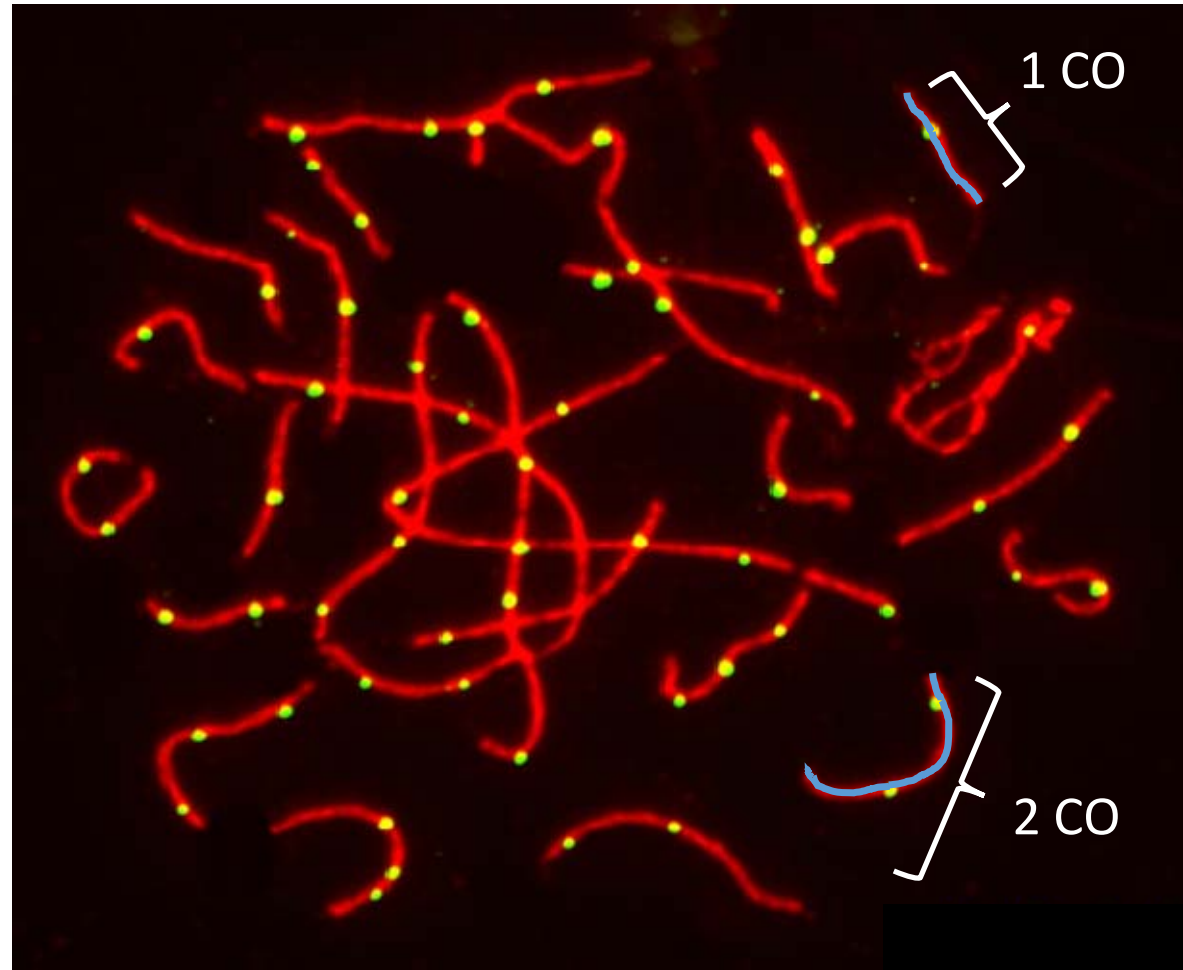
MLH1 ●
Crossover (CO)

Number of COs

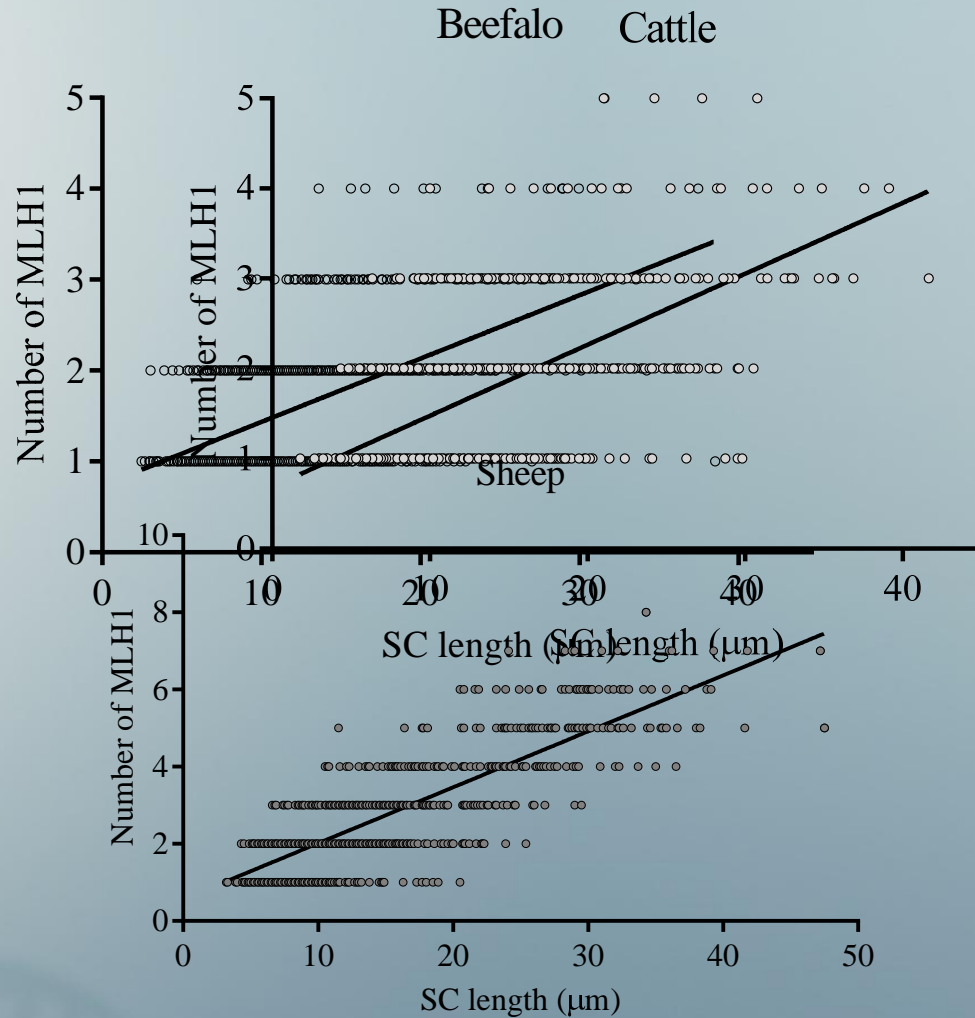


Each dot represents the number of COs from individual spermatoocyte, black bars represent breed mean, and the letters above denotes significant differences ($P < 0.01$).

Chromosome Length & CO Numbers



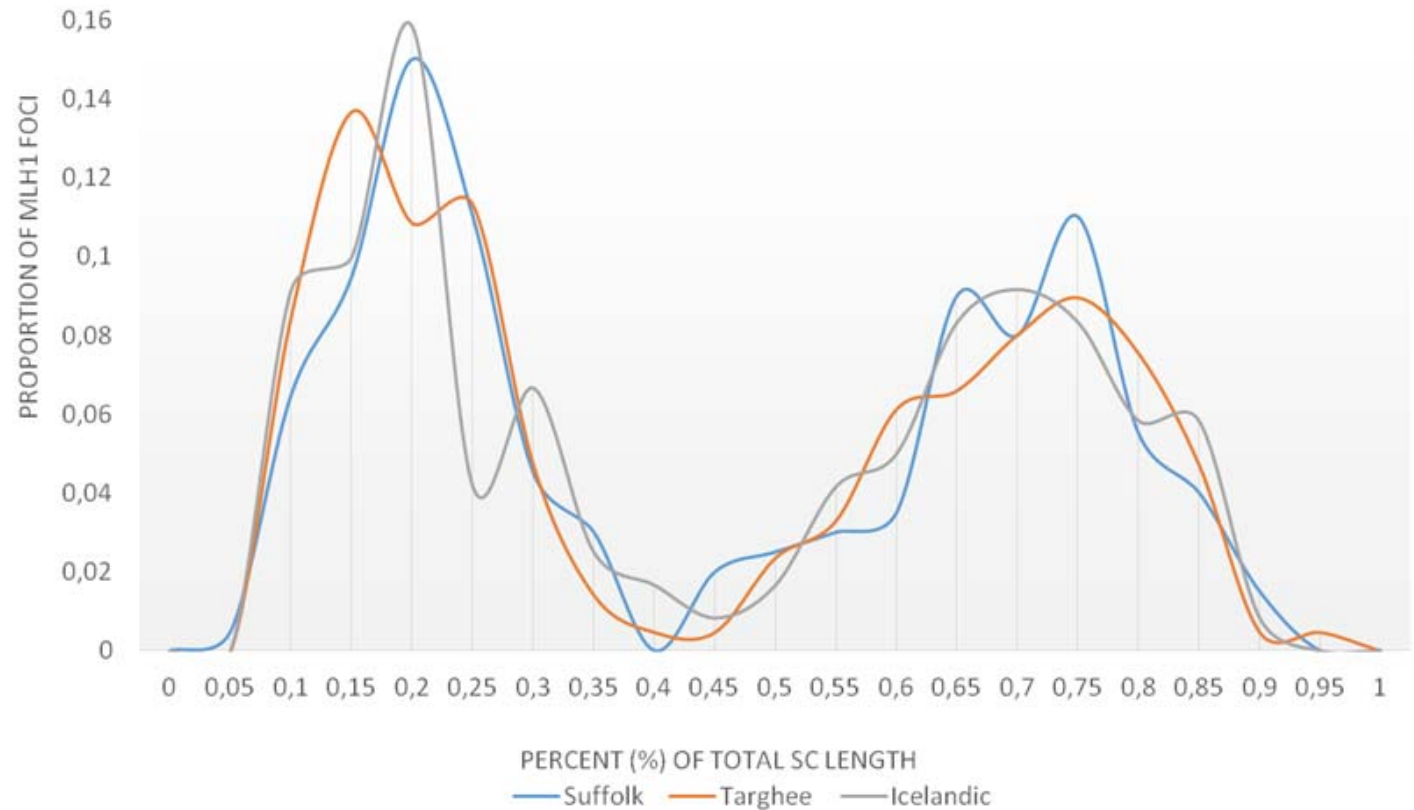
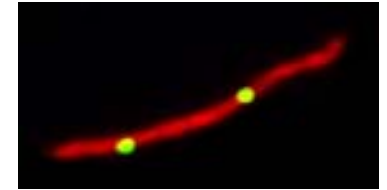
Length of Chromosome & CO Number



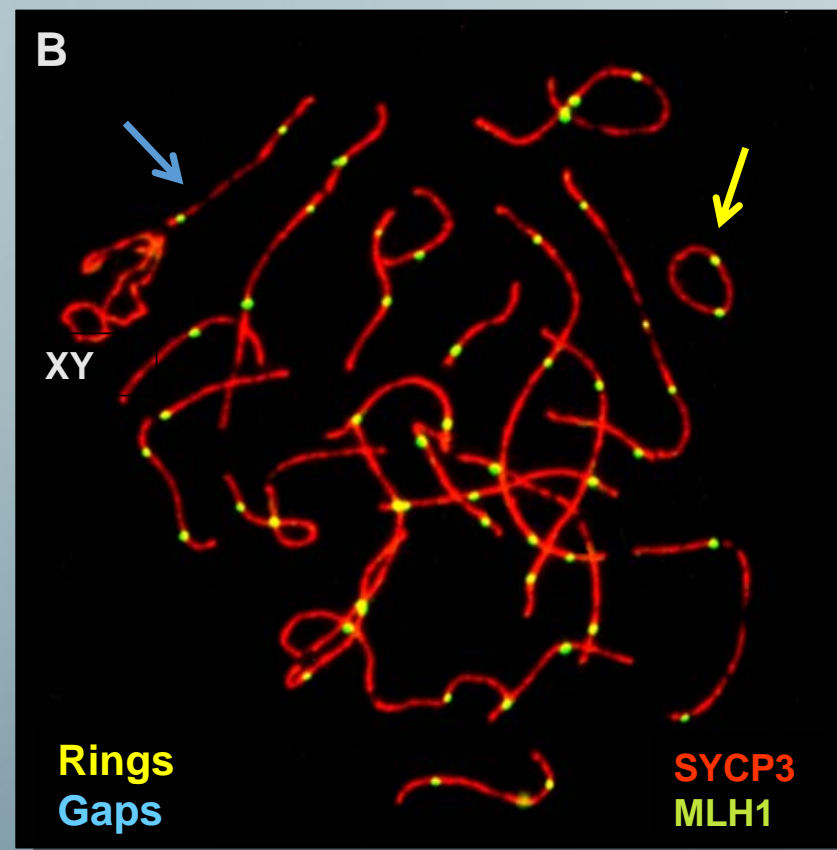
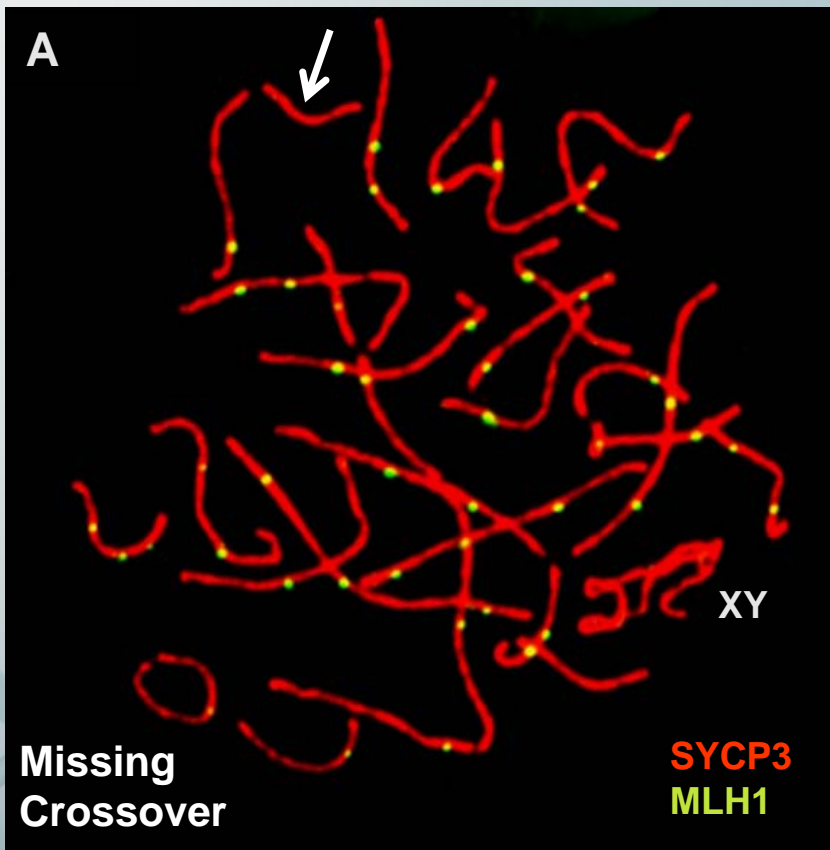
Beefalo (61 cells, $r=0.53$)
 Cattle (86 cells, $r=0.57$)
 Sheep (340 cells, $r=0.70$)

Chromosome Length & CO Position

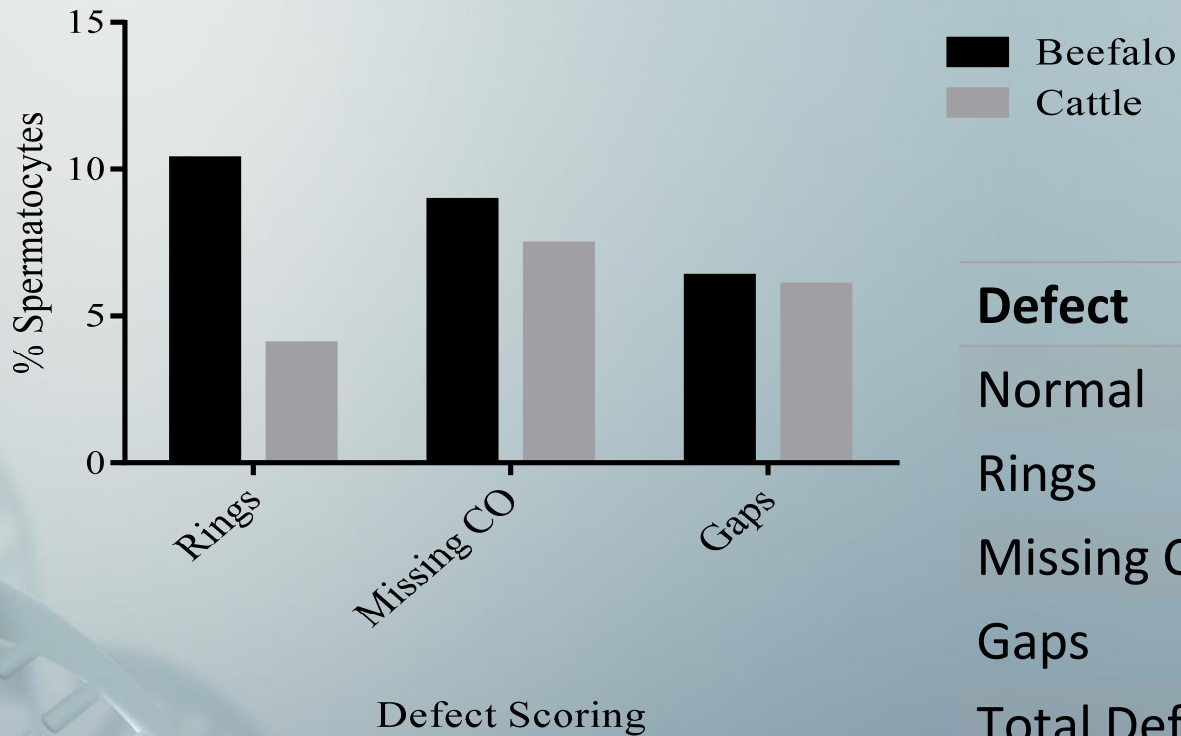
- Locations of CO on SCs with 2 CO



Chromosomal Defect Scoring



Chromosomal Defect Scoring



Defect	Beefalo	Cattle
Normal	74.6 ^A	83.6 ^B
Rings	10.3 ^A	3.8 ^B
Missing CO	8.9 ^A	6.2 ^A
Gaps	6.3 ^A	6.2 ^A
Total Defects	25.4 ^A	16.4 ^B

Letters denote significant differences (P<0.01)



Summary

- Sheep exhibit a significantly greater number of COs and cattle exhibit more COs in comparison beefalo.
- Larger chromosomes tend to have a greater number of COs.
- Beefalo spermatocytes have a greater number of structural defects.

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Lab

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- Hannah Jaeger
- Dominic De La Torre
- Eric Robinson

Testicular Samples

- Pure Country Harvest
- University of Idaho Vandal Meats
- University of Idaho Sheep Experiment Station
- C&L Meat Locker

Questions?





Number of COs in Cattle

Bulls	Spermatocytes	Average CO
Charolais	112	48.9 ^A ±0.54
Gelbvieh	97	47.6 ^A ±0.58
Jersey	100	48.6 ^A ±0.39
Angus 1	100	43.3 ^B ±0.33
Angus 2	100	43.4 ^B ±0.30
Angus 3	101	41.5 ^B ±0.36
Angus 4	100	42.4 ^B ±0.37
Angus 5	100	41.7 ^B ±0.41
Angus 6	97	43.2 ^B ±0.61
Angus 7	98	46.5 ^A ±0.52