

Minisatellites and microsatellites – similar names but different biology

Yuri E Dubrova

*Department of Genetics
University of Leicester*

Tandem repeat DNA loci

➤ Consist of repeats

$GAACA_n$

_GAACA_GAACA_GAACA_GAACA_GAACA

➤ Highly variable

up to 100+ alleles per locus

➤ High mutation rate

up to 0.15 per locus

➤ Because of this they are widely used in:

- forensics (paternity testing, identity identification)
- population genetics
- gene mapping
- mutagenesis (germline & somatic mutation induction)
- cancer studies (stability of cancer cells)

Tandem repeat DNA loci

➤ **Microsatellite loci**

PubMed 26,529 items

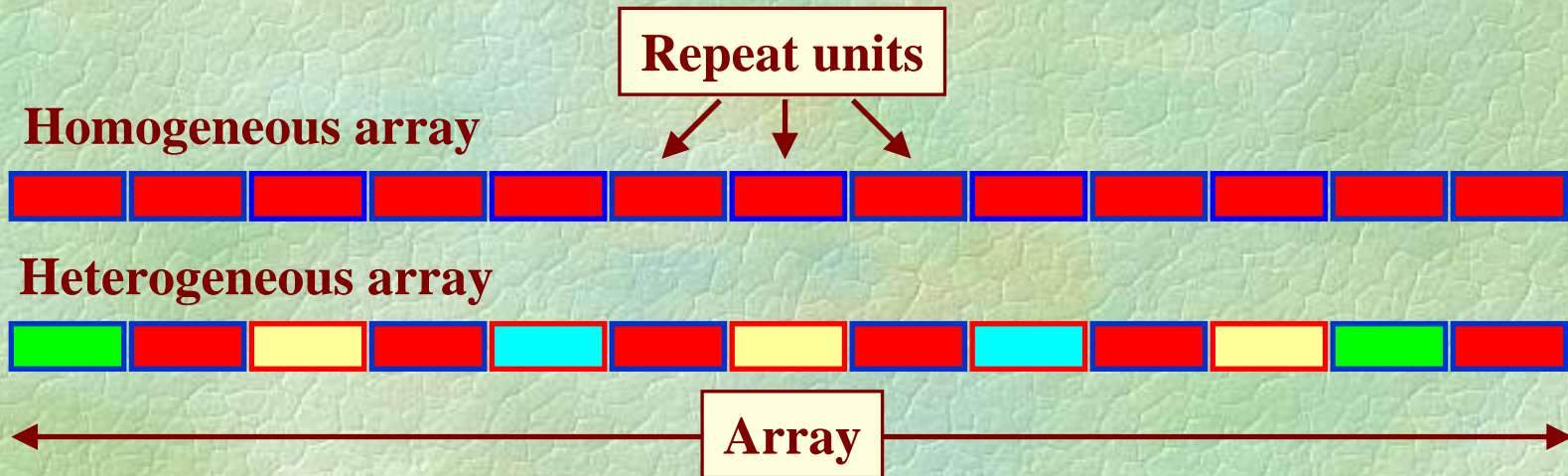
➤ **Minisatellite loci**

PubMed 3,120 items

➤ **Expanded Simple Tandem Repeat loci
(**ESTR**)**

PubMed 63 items

Tandem repeat DNA loci



Minisatellites

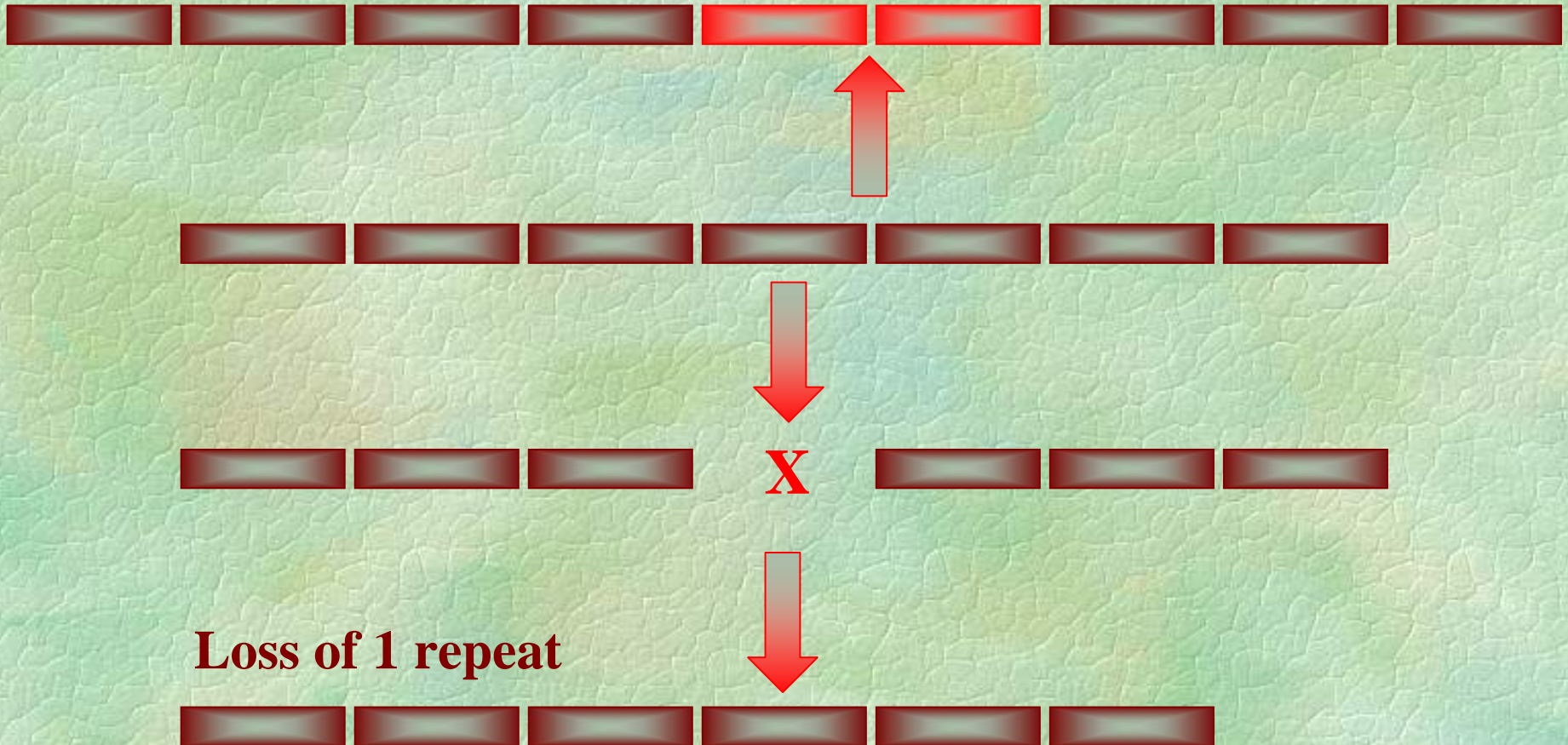
ESTRs

Microsatellites

| | | | |
|-----------------------|-----------------|--------------------|--------------------|
| ➤ Repeat unit | 10 – 60 bp | 4 – 10 bp | 2 – 6 bp |
| ➤ Size of array | 0.5 – 15 kb | 0.1 – 20 kb | 10 – 1000 bp |
| | 10 – 1,500 rpts | 10 – 2,000 rpts | 5 – 200 rpts |
| ➤ Complexity of array | heterogeneous | mostly homogeneous | mostly homogeneous |

Schematic examples of mutations at tandem repeat DNA loci

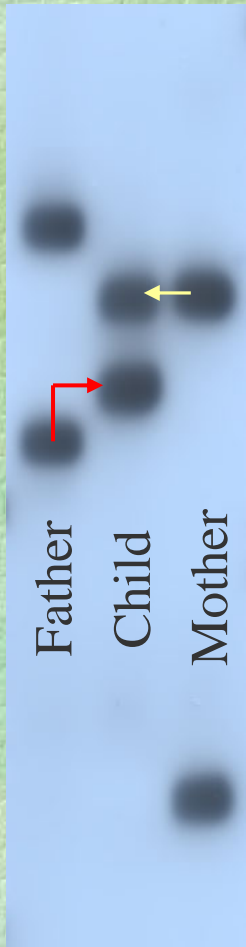
Gain of 2 repeats



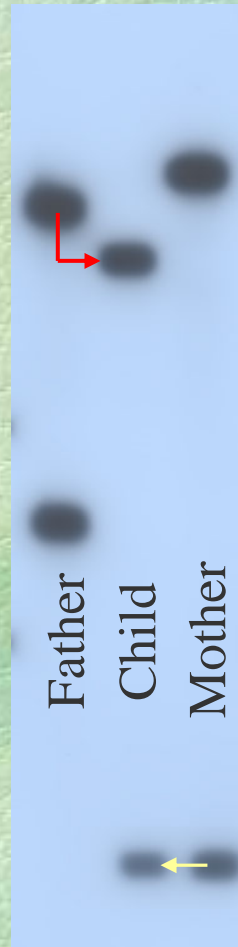
Loss of 1 repeat

Germline mutations at human minisatellite loci

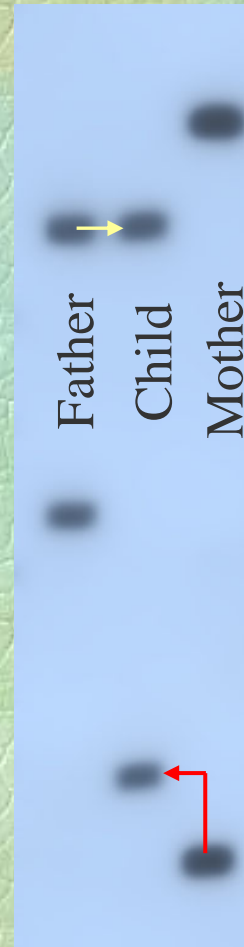
MS31
paternal/gain



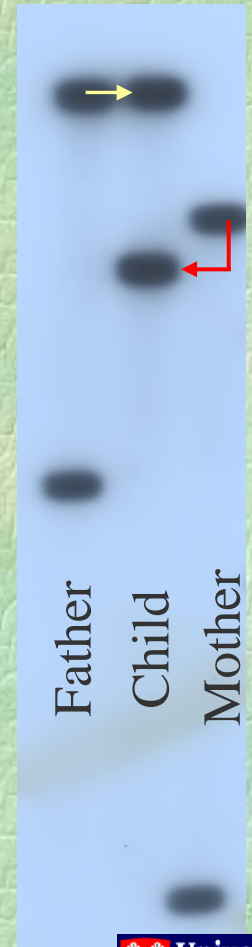
MS32
paternal/loss



MS1
maternal/gain



CEB25
maternal/loss



Spontaneous germline mutation rates at tandem repeat DNA loci and protein-coding genes

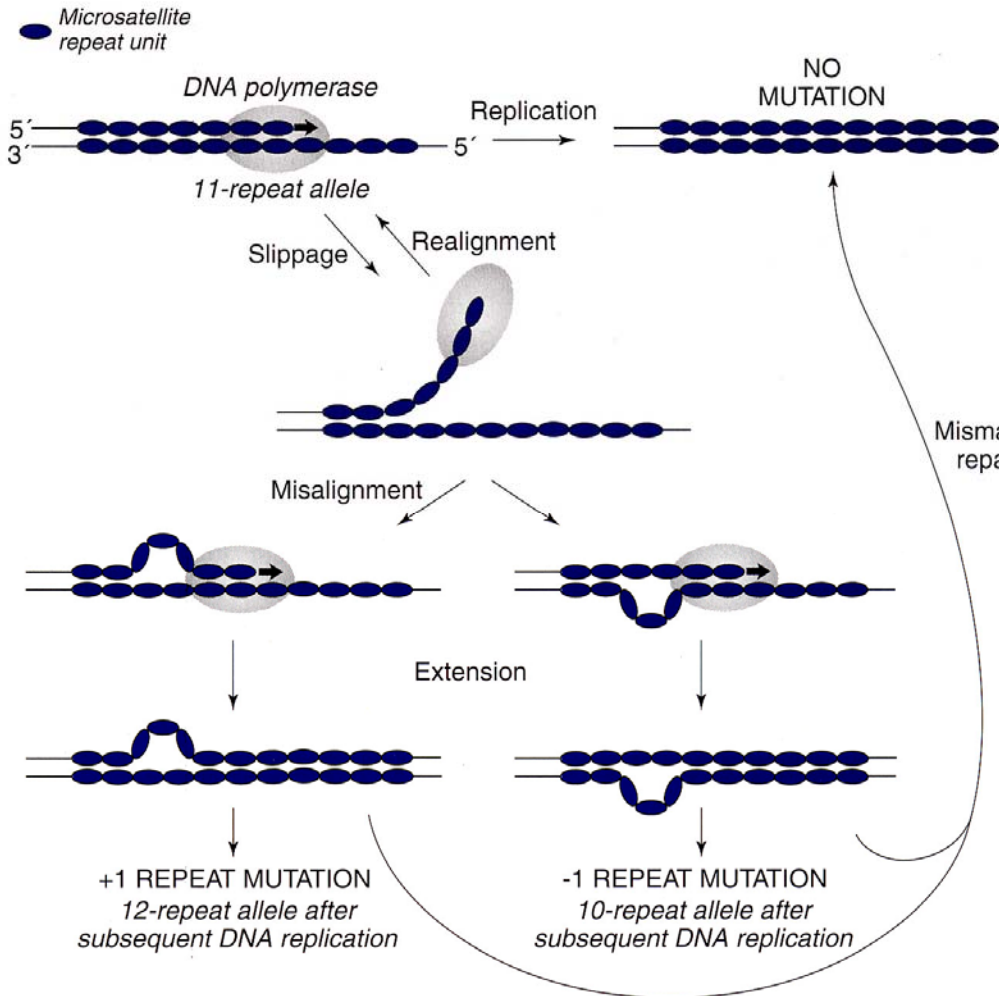
| <i>Probe (Locus)</i> | <u><i>Mutation rate per gamete</i></u> | | |
|------------------------------------|--|-----------------|--------------|
| | <i>Paternal</i> | <i>Maternal</i> | <i>Total</i> |
| ➤ Human minisatellite loci | | | |
| ✓ CEB1 (<i>D2S90</i>) | 0.161 | 0.003 | 0.082 |
| ✓ B6.7 (20q13) | 0.076 | 0.012 | 0.044 |
| ✓ MS1 (<i>D1S7</i>) | 0.055 | 0.049 | 0.052 |
| ✓ CEB25 (<i>D10S180</i>) | 0.035 | 0.019 | 0.027 |
| ✓ CEB36 (<i>D10S473</i>) | 0.018 | 0.018 | 0.018 |
| ✓ MS31 (<i>D7S21</i>) | 0.012 | 0.003 | 0.008 |
| ✓ MS32 (<i>D1S8</i>) | 0.009 | 0.006 | 0.007 |
| ➤ Mouse ESTR loci | | | |
| ✓ <i>Ms6-hm</i> | 0.10-0.18 | 0.06-0.09 | 0.09-0.13 |
| ✓ <i>Hm-2</i> | 0.02-0.07 | 0.04-0.14 | 0.04-0.11 |
| ➤ Human microsatellite loci | mean ~ 0.002 (0.0001 – 0.01) | | |
| ➤ Protein-coding genes | $10^{-6} - 10^{-5}$ (< 1 per 100,000) | | |

Data from numerous original publications

Mechanisms of mutation

Microsatellites

Replication slippage

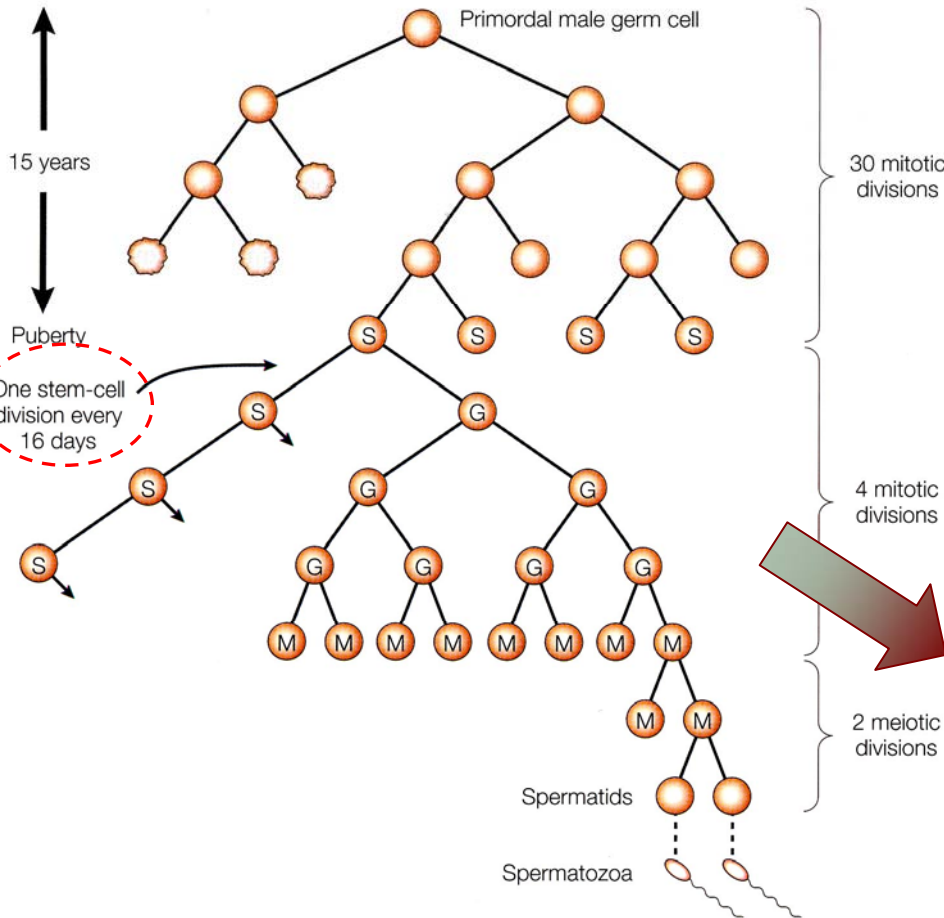


Predictions

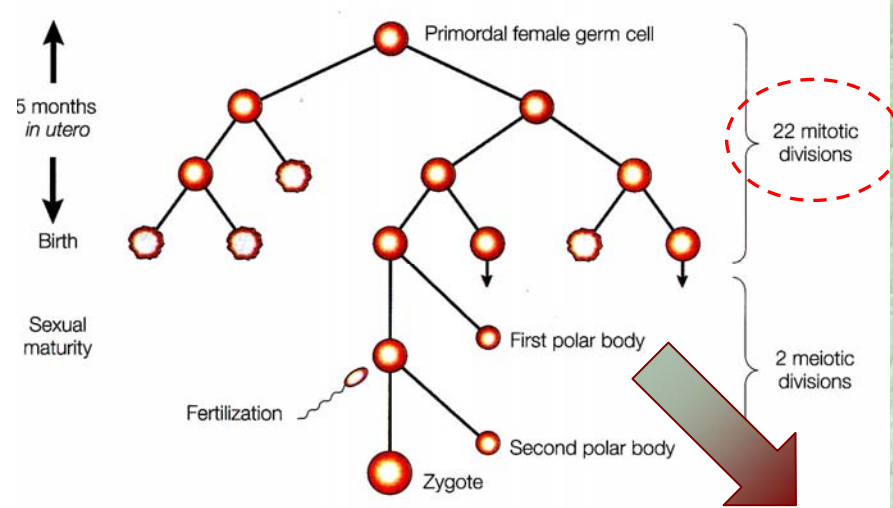
- Mutate in dividing cells only
- No germline specificity, *i.e.* mutations frequently occur in somatic tissues
- Germline mutation rate in males > females
- Age-related accumulation of mutations
- Simple mutation spectrum

Cell divisions during spermatogenesis & oogenesis in humans

Spermatogenesis



Oogenesis



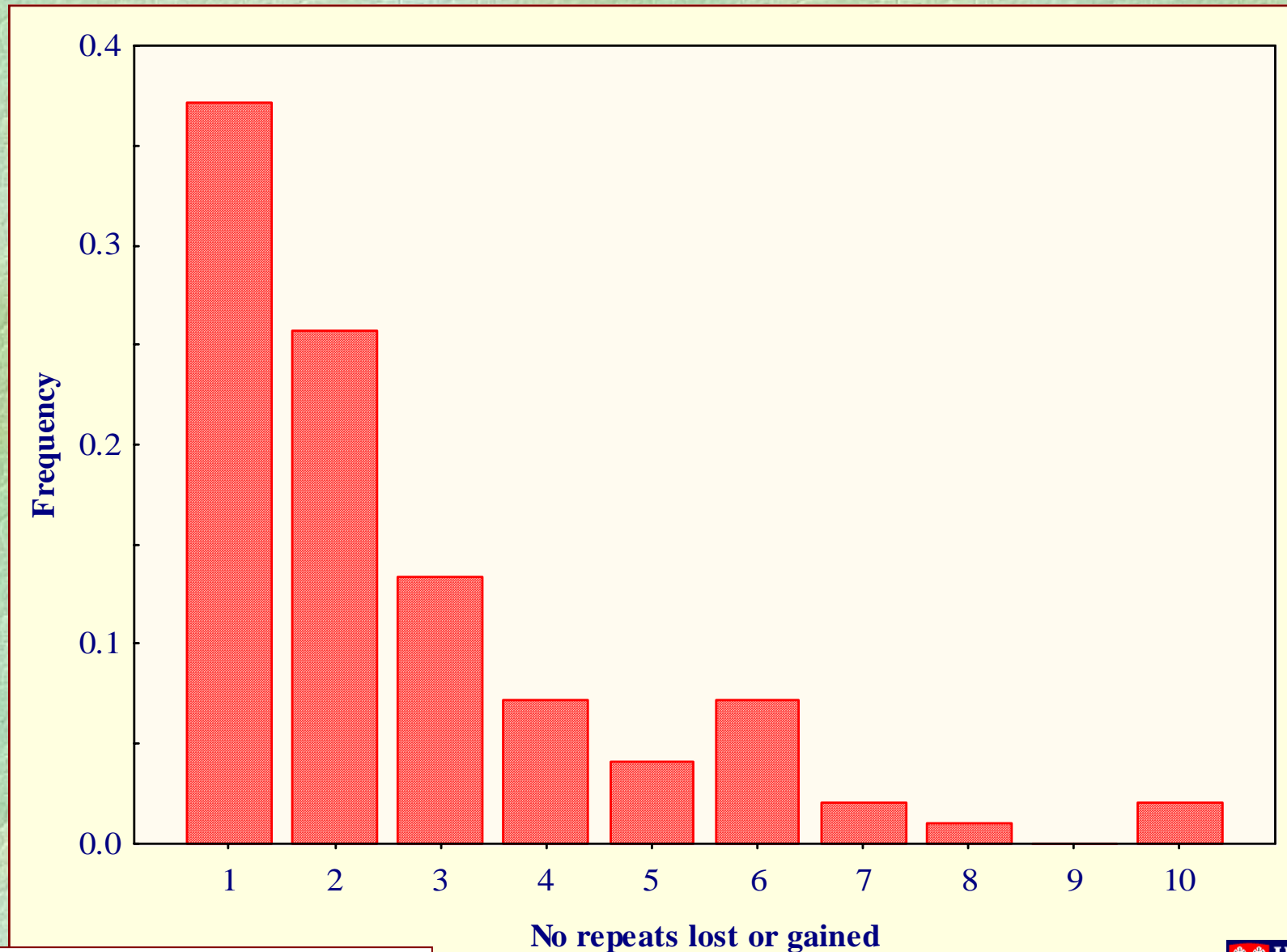
22 mitotic divisions for all ages

No male germ-cell divisions

| Age, years | No replications |
|------------|-----------------|
| 15 | 35 |
| 20 | 150 |
| 30 | 380 |
| 40 | 610 |
| 50 | 840 |

Microsatellite mutation rates
Male/Female ratio 5 : 1

Spectrum of spontaneous microsatellite mutation in humans (362 loci, 53 pedigrees, 630 subjects, 97 mutational events)

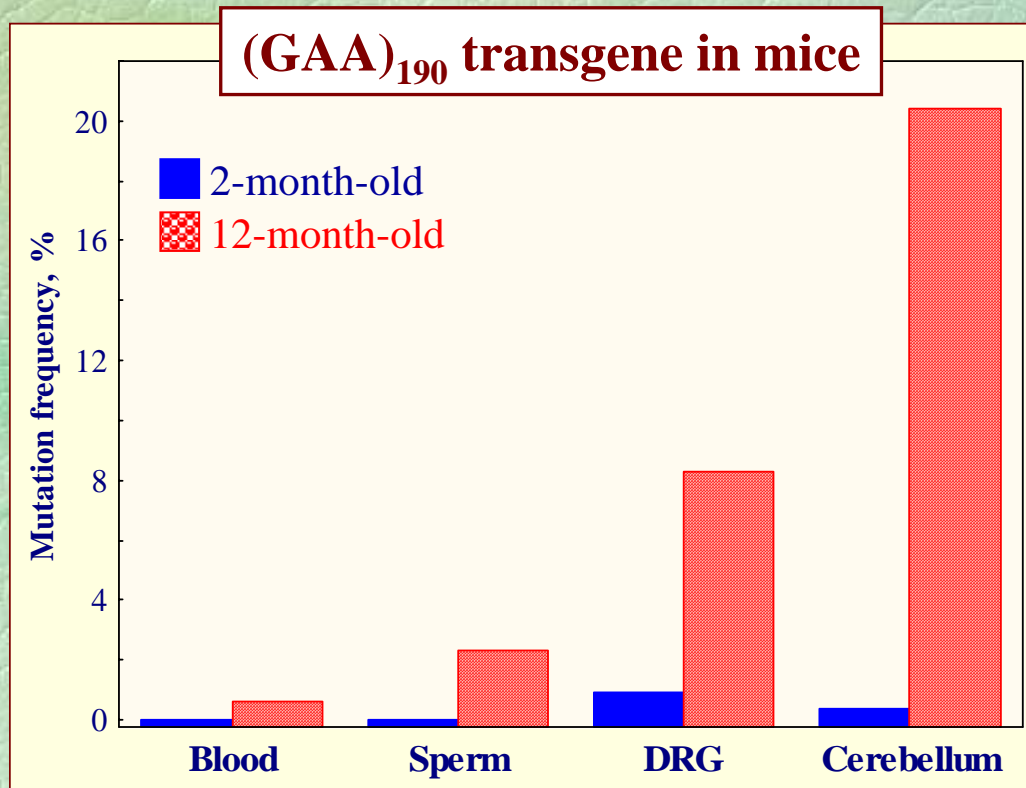


From: Huang *et al.*, 2002, *Am J Hum Genet* **70**, 625-34

The big, the bad, the ugly microsatellites

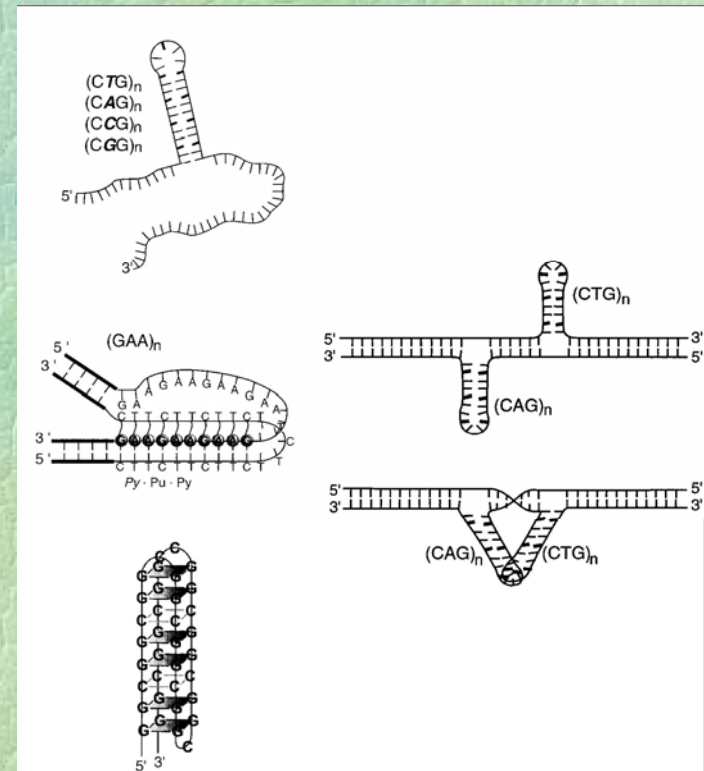
- Long arrays are highly unstable, '100%' mutation rate
- Associated with some human diseases

Mutate in non-dividing tissues



From: Clark *et al.*, 2007, *Hum Genet* **120**, 633

Secondary structures
DNA-repair is involved

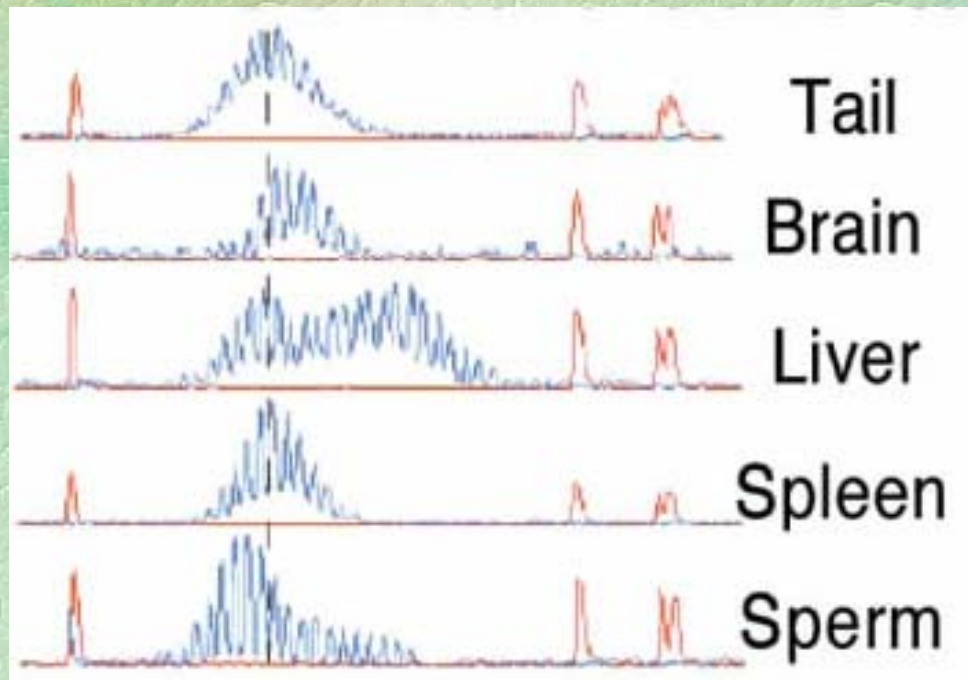
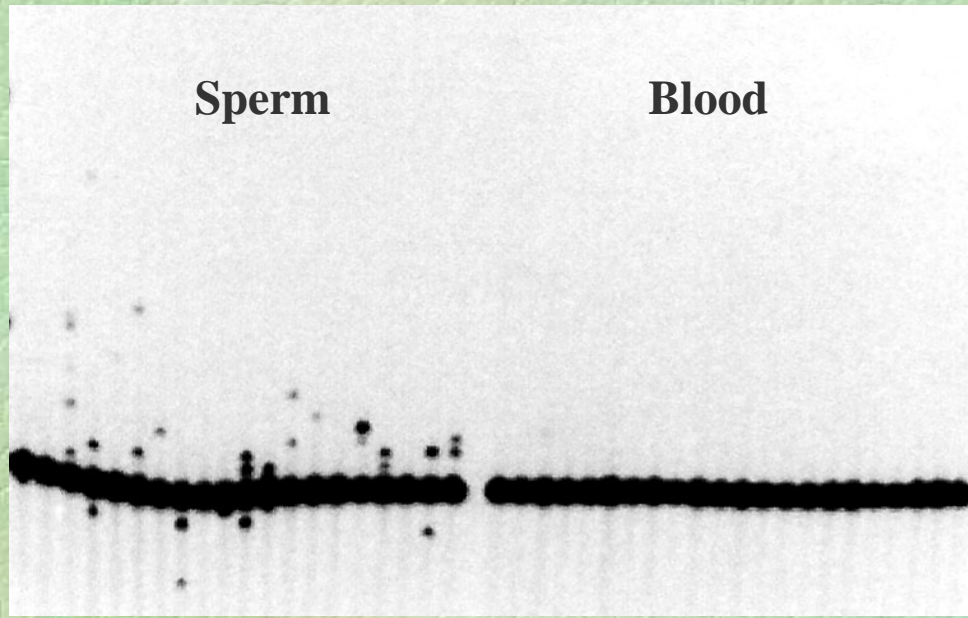


From: Sinden, 1999, *Am J Hum Genet* **64**, 346-53

Minisatellites

- **Very unstable in the germline
but ~ dead in somatic tissues**
- **Very complex mutations in the male germline
and very simple in somatic tissues**

Germline vs. somatic mutation at minisatellite loci

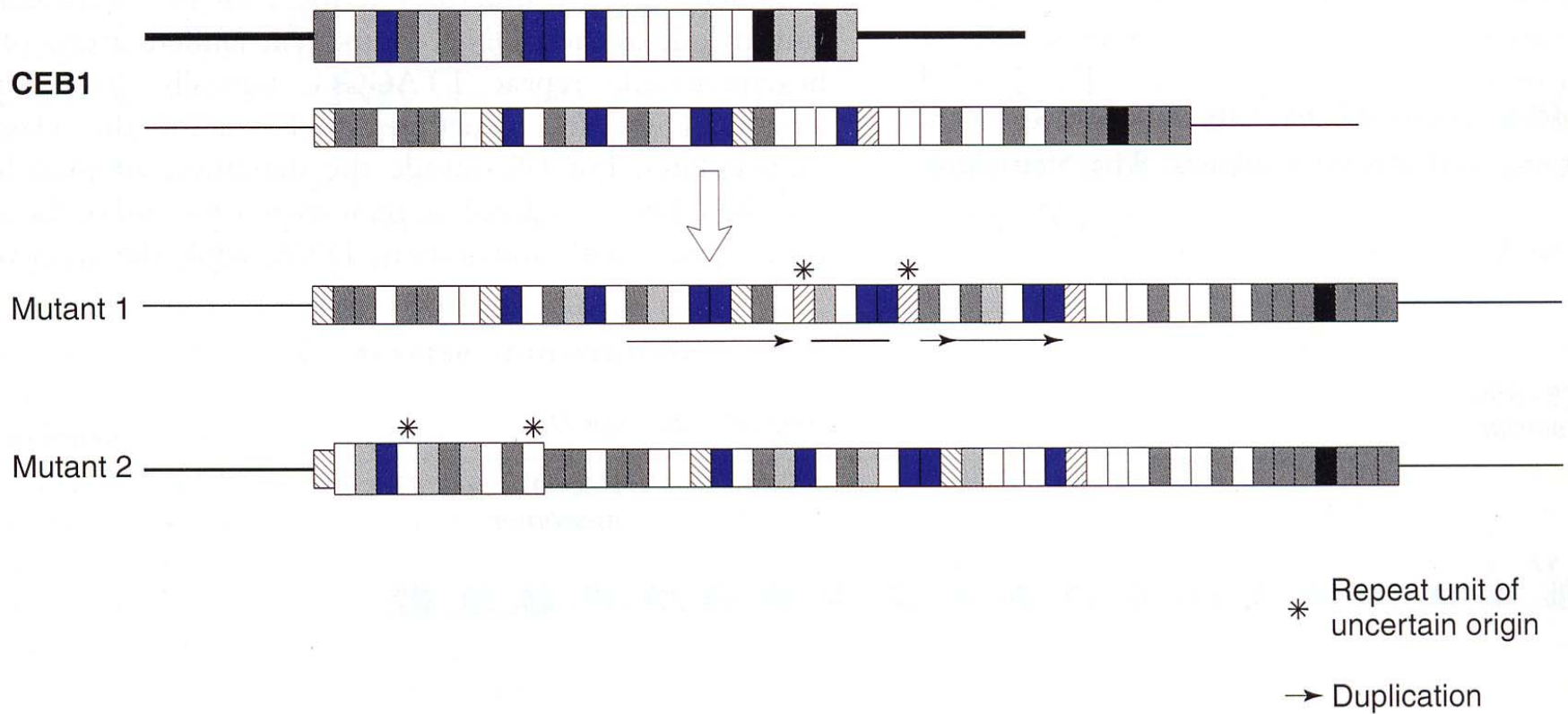


Human minisatellite B6.7

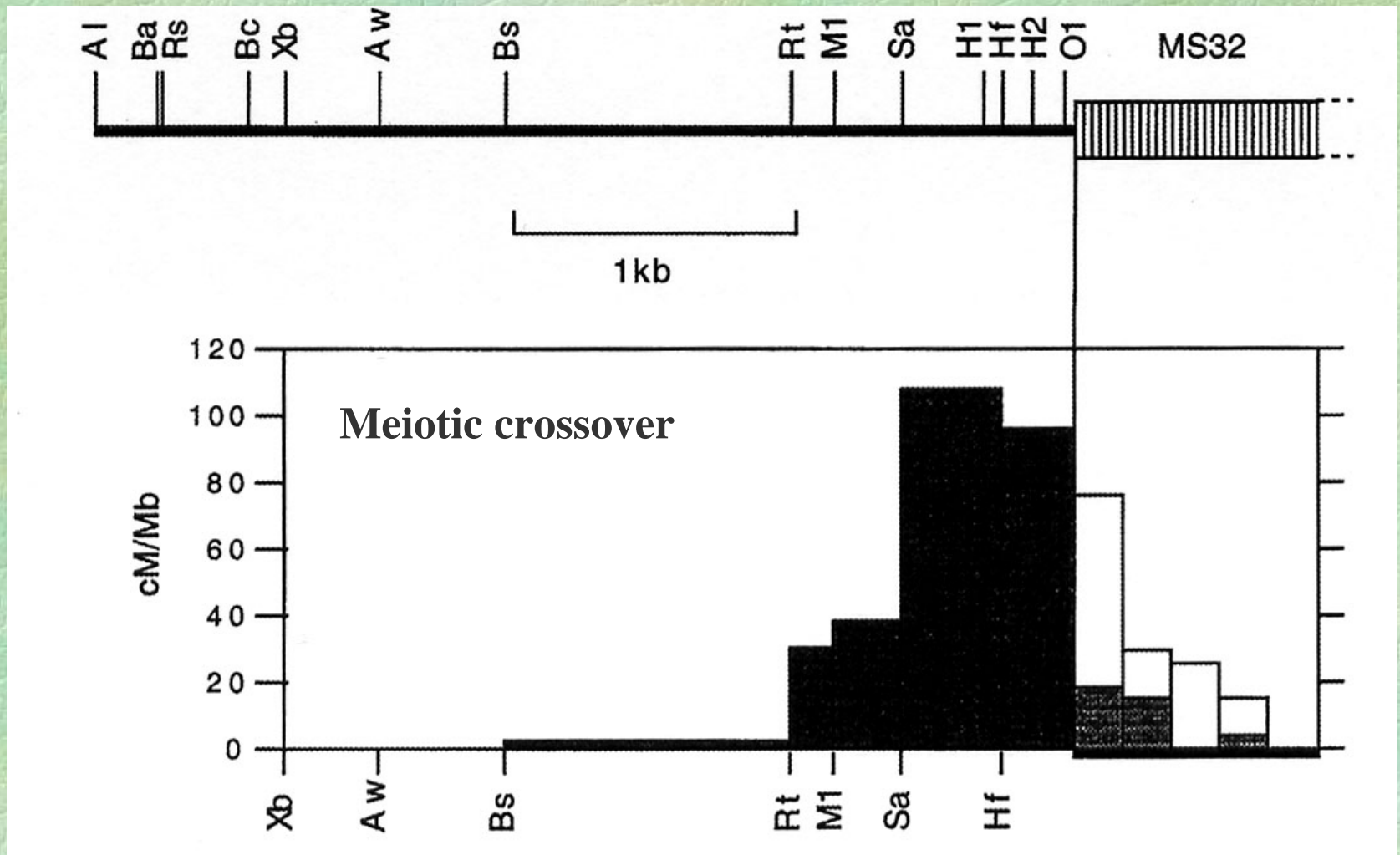
CAG₁₁₈ human microsatellite transgene in mice

From: Tamaki *et al.*, 1999, *Hum Mol Genet* **8**, 879-88
Kovtun & McMurray, 2001, *Nat Genet* **27**, 407-11

Spectrum of minisatellite mutants in sperm



Minisatellite mutation is somehow related to what is going on at meiotic crossover hot-spots



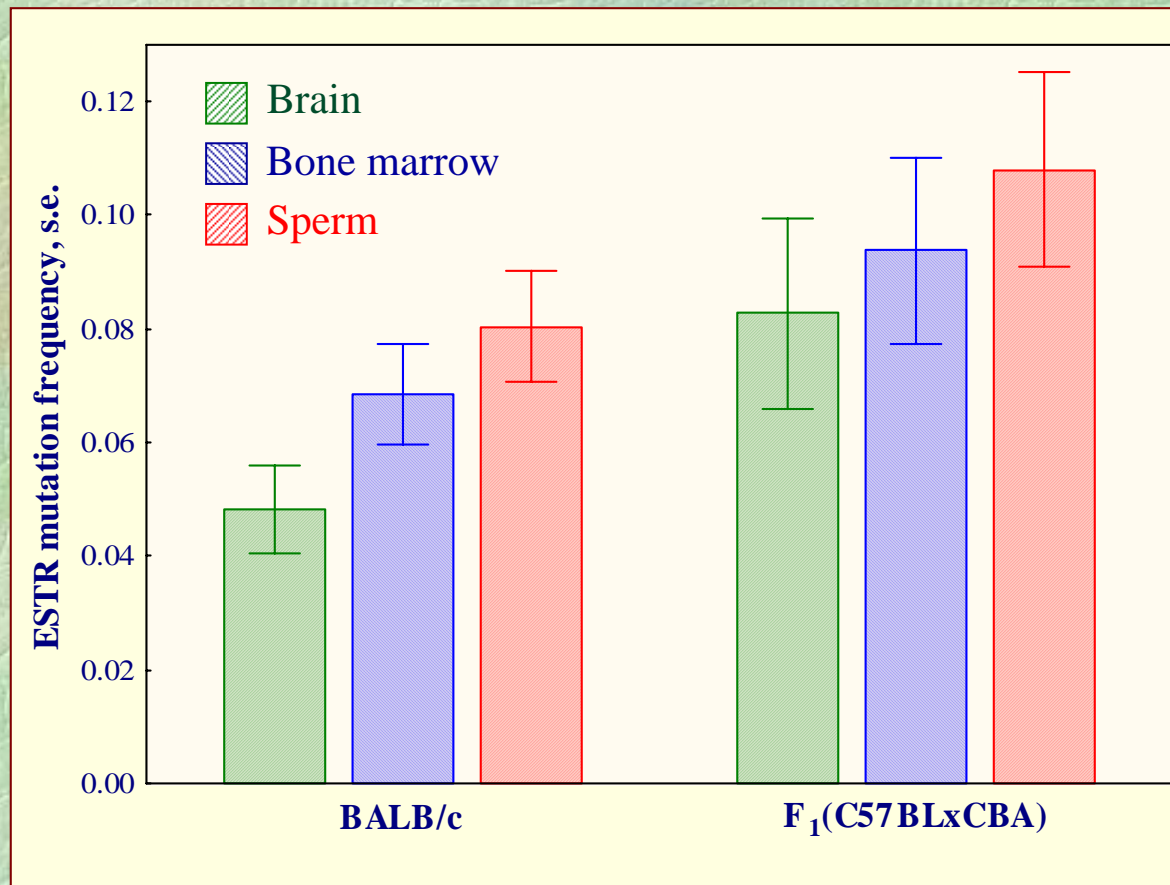
From: Jeffreys *et al.*, 1999, *Electrophoresis* **20**, 1665-75

**Expanded Simple
Tandem Repeat loci
or **ESTR****

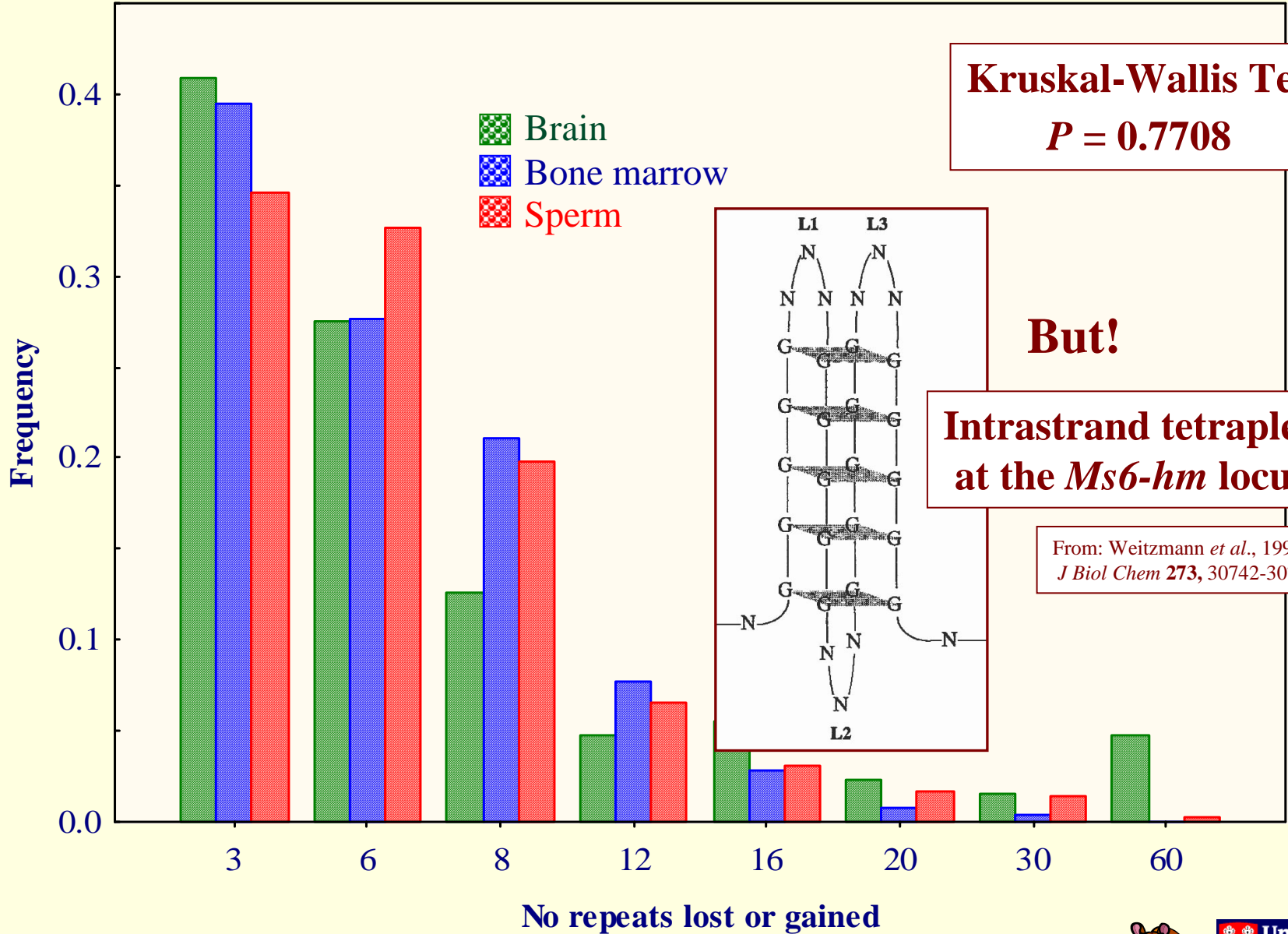


ESTR's

- Repeat size: 4-10 bp, *i.e.* between micro- (1-4bp) & mini-
- Array size: 10-2000 rpts, *i.e.* more mini-like
- Unstable in somatic tissues, *i.e.* micro-like



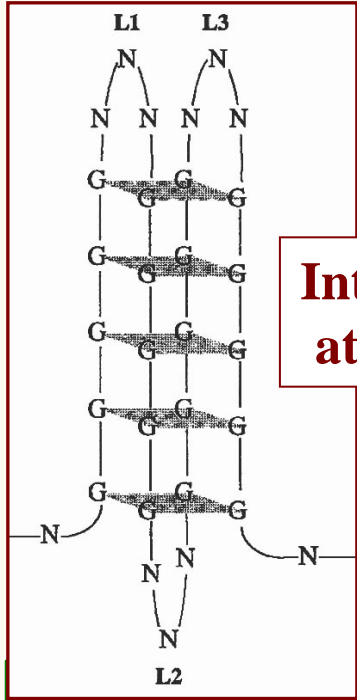
Similar spectra of spontaneous ESTR mutation in mouse tissues



Kruskal-Wallis Test
 $P = 0.7708$

But!

Intrastrand tetraplex at the *Ms6-hm* locus



From: Weitzmann *et al.*, 1998,
J Biol Chem **273**, 30742-30749

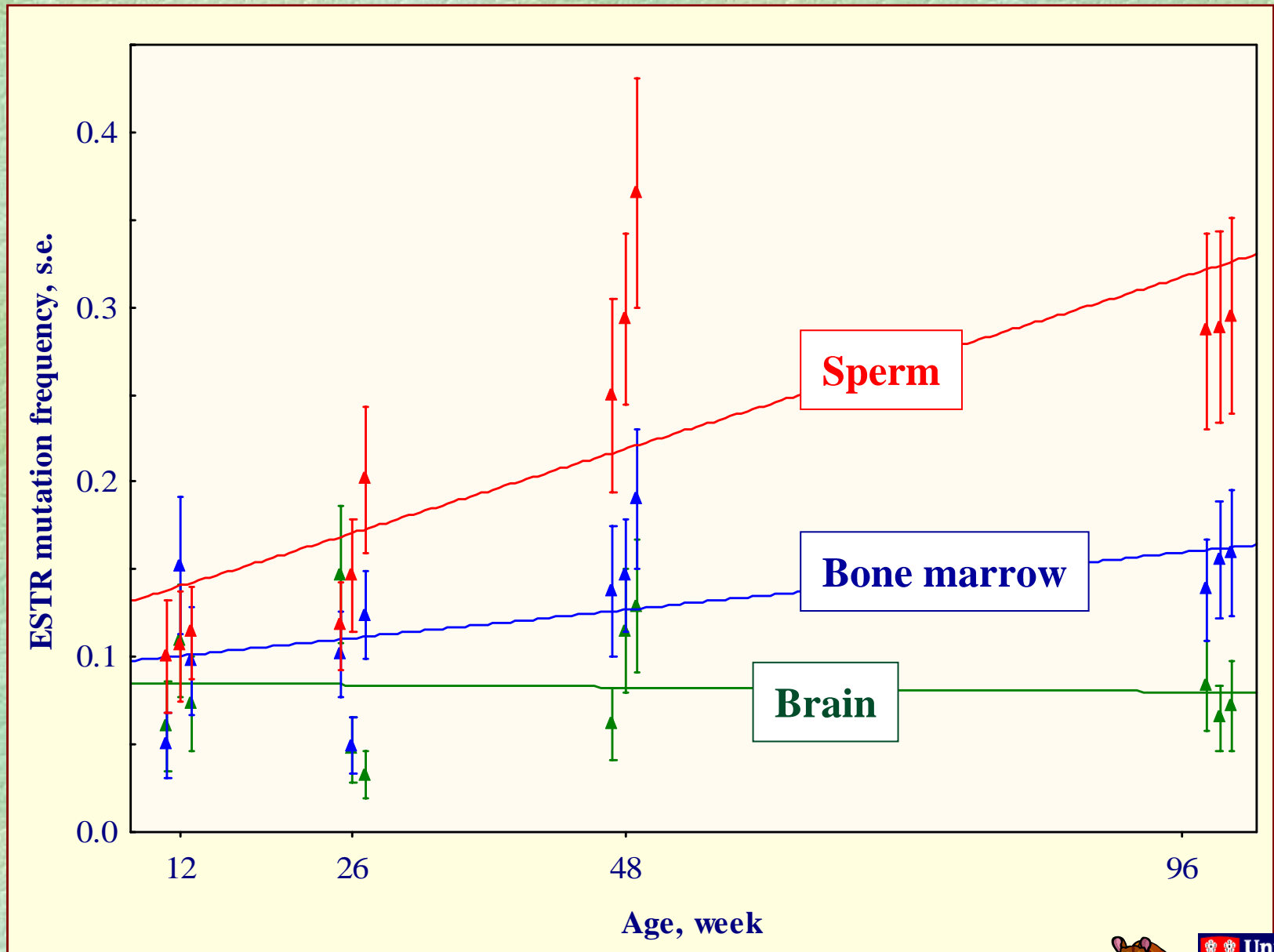


If ESTR loci ‘behave’ like true microsatellites then:

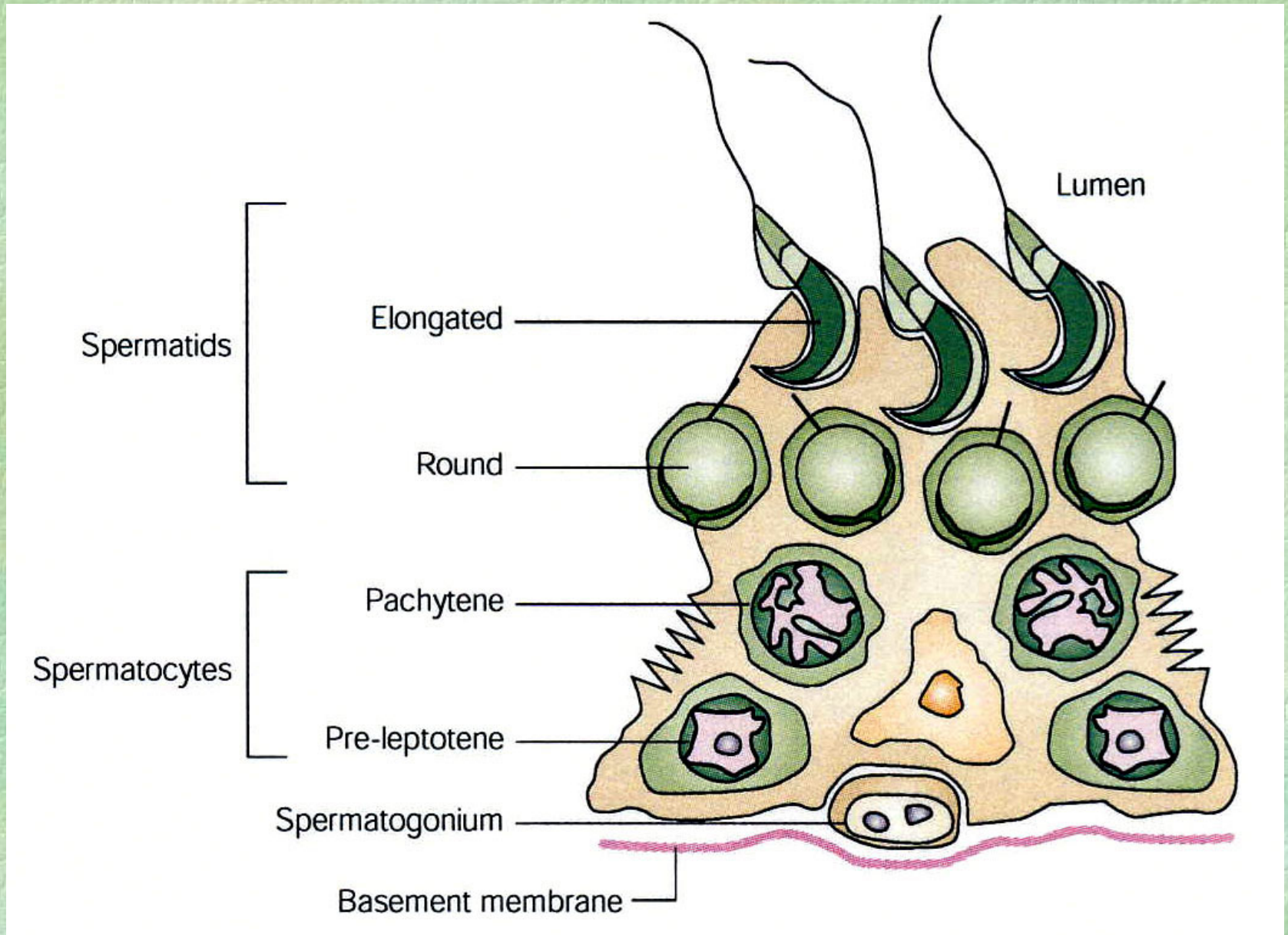
- **Age-related increases in replication-proficient tissues**
- **No age-related changes in non-dividing tissues**
- **Stage-specific pattern of spontaneous ESTR mutation in the male germline**



Age-related changes in ESTR mutation frequency in mice



A single Sertoli cell with its associated germ cells





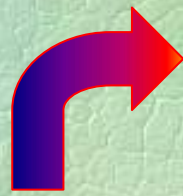
Meiotic spermatocytes I

Round spermatids

Elongated spermatids

Meiotic events

Pre-meiotic spermatogonia



Elongated spermatids

Late post-meiotic events

Pre-meiotic spermatogonia

Meiotic spermatocytes I

Round spermatids

Mitotic events

Pre-meiotic spermatogonia

Meiotic spermatocytes I

Round spermatids

Elongated spermatids



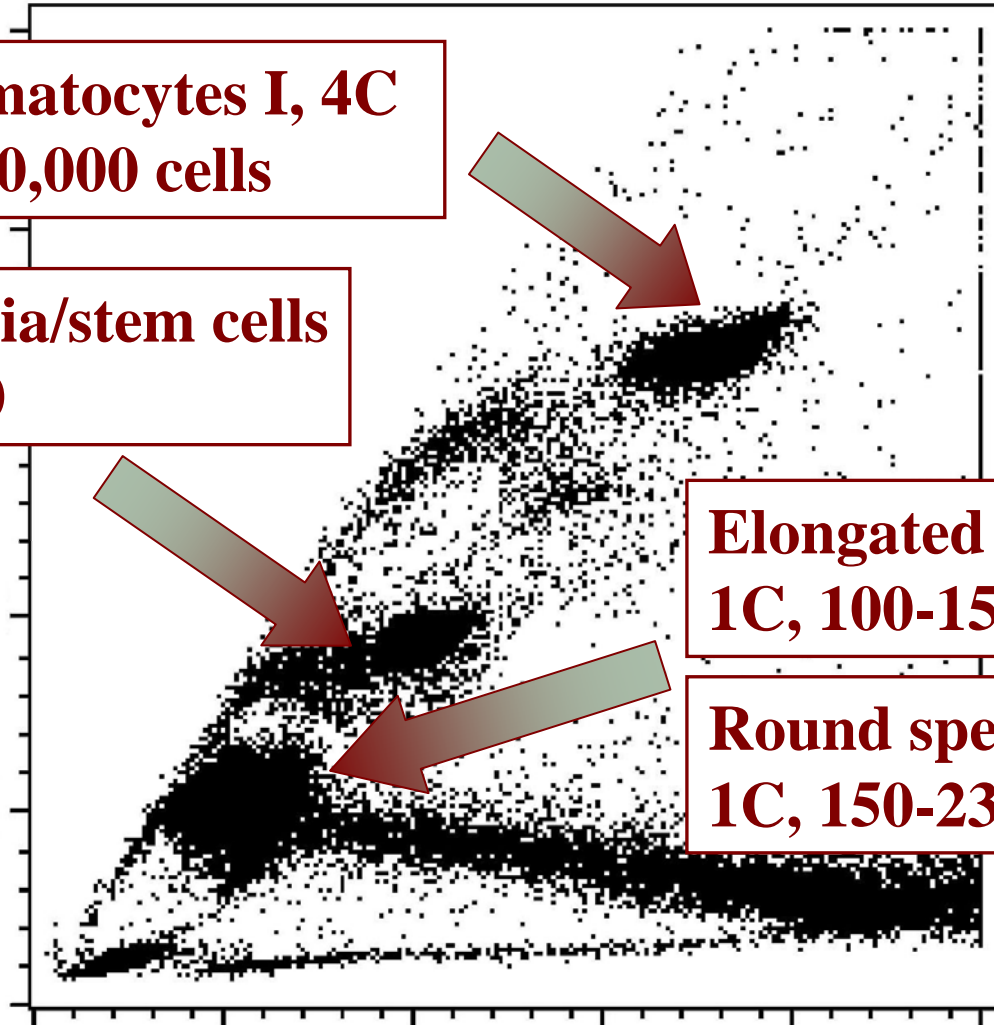
Separation of mouse germ cells by flow cytometry

Spermatocytes I, 4C
80-100,000 cells

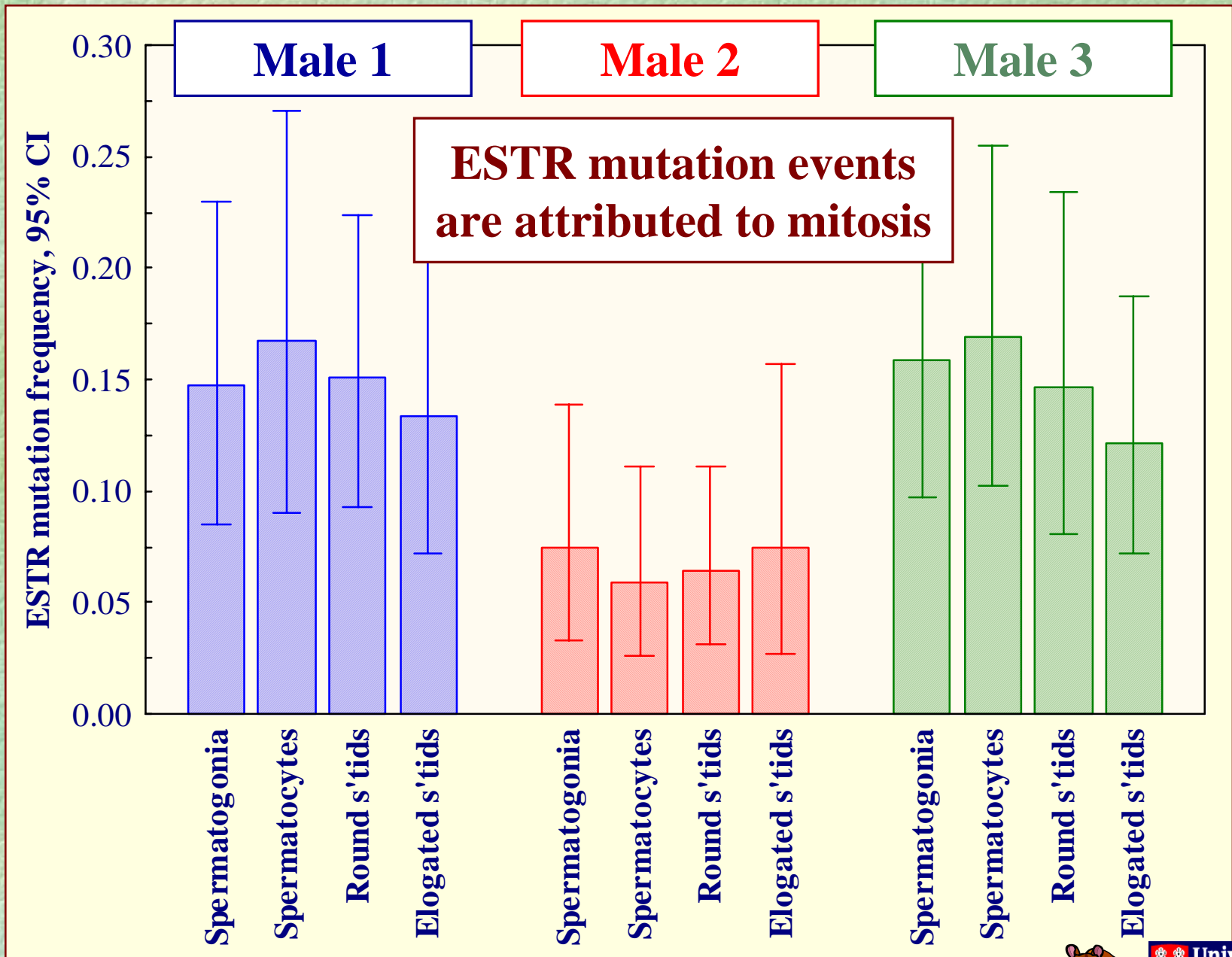
Spermatogonia/stem cells
2C, 50-80,000

Elongated spermatids
1C, 100-150,000 cells

Round spermatids
1C, 150-230,000 cells



ESTR mutation frequencies in mouse germ cells



Problems

We observe changes in mutation frequencies, p

$$p = f(u, N, s)$$

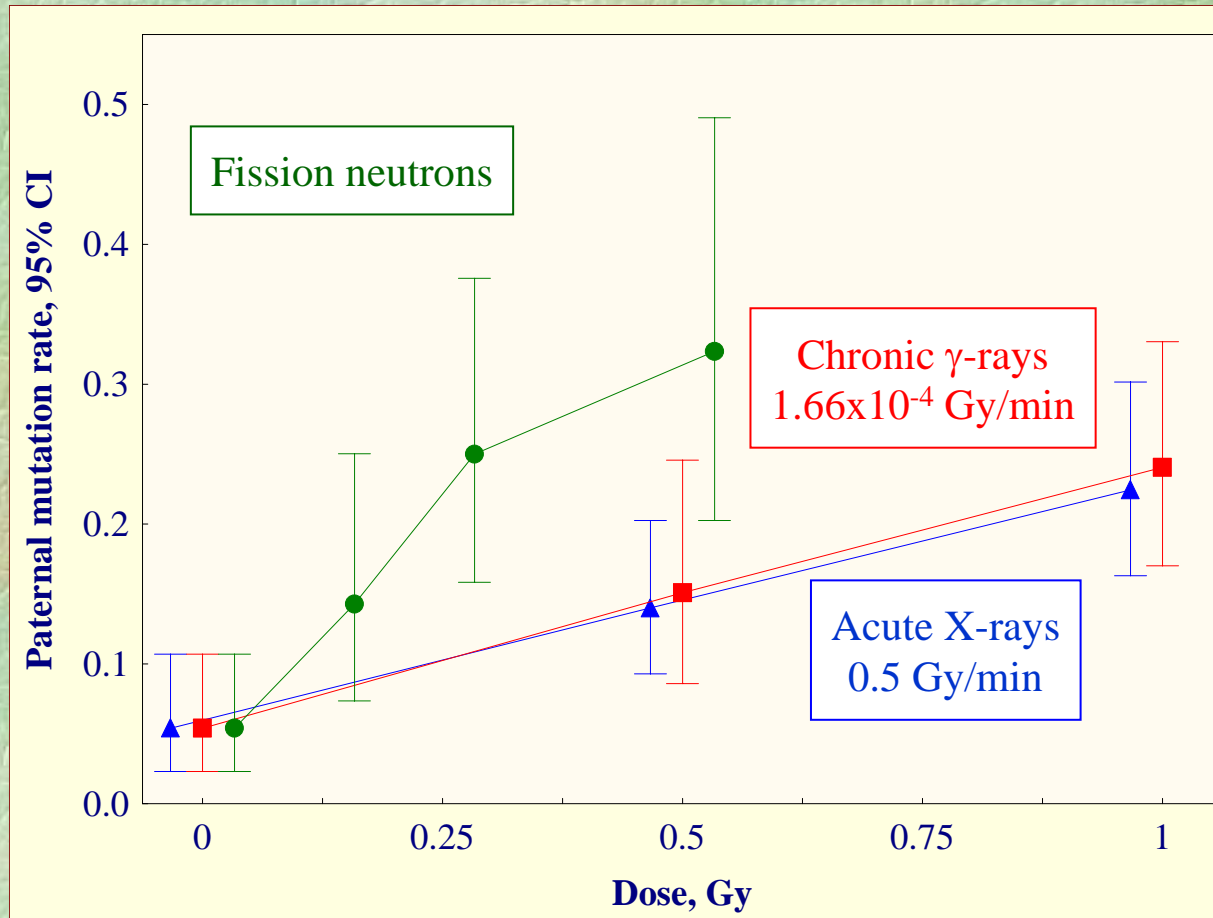
u – mutation rate per DNA replication/cell division

N – number of cell divisions
known for the male germ cells

s – selection against mutants, $s = 0$



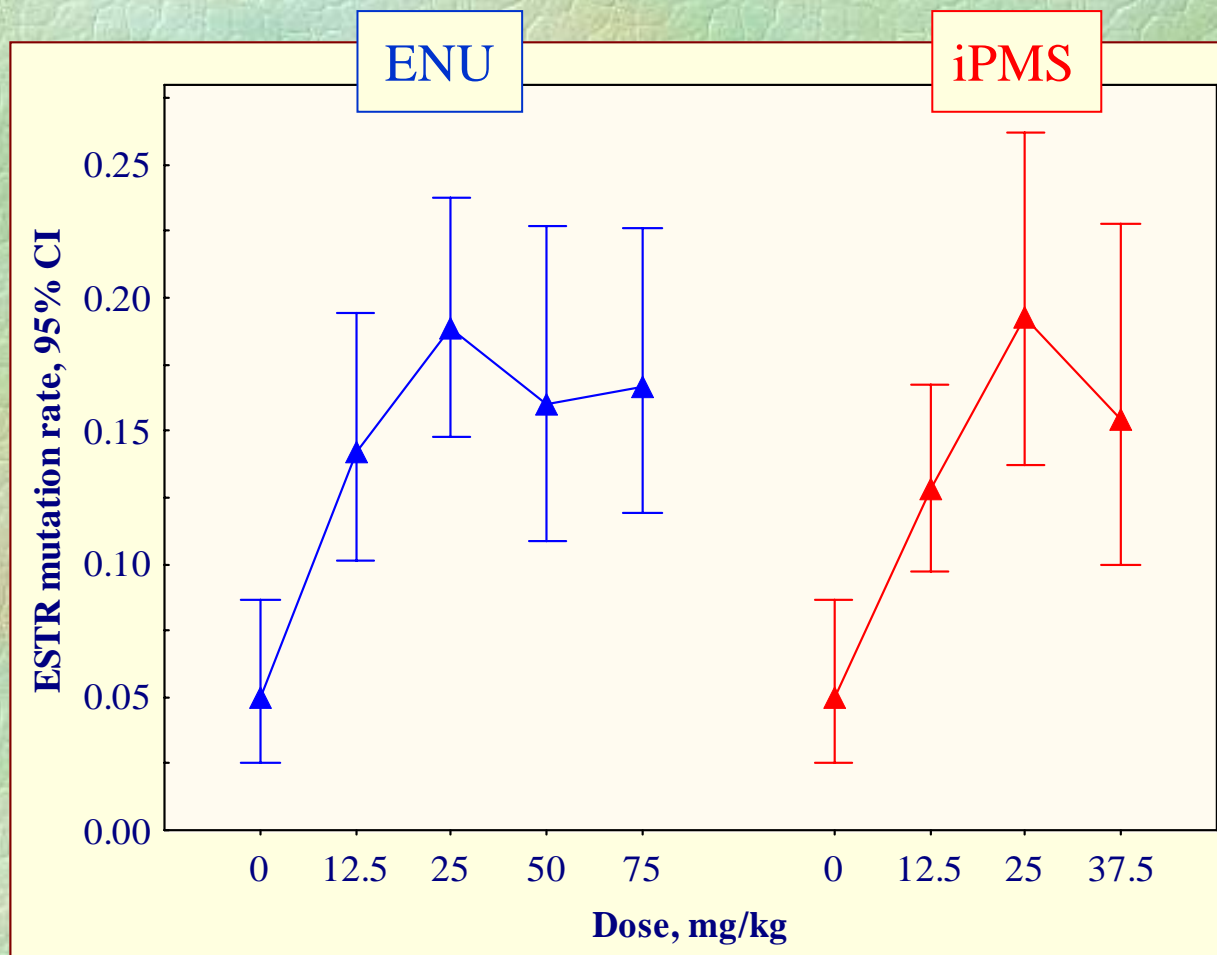
Mutation induction at mouse ESTR loci



From: Dubrova *et al.*, 1998, *PNAS* **95**, 6251-6255
2000, *Mutat Res* **453**, 17-24



Mutation induction at mouse ESTR loci



From: Vilariño-Güell *et al.*, 2003, *Mutat Res* **526**, 63-73



Non-targeted mutation induction at mouse ESTR loci

Expected genome damage from mouse ESTR data

- Spontaneous mutation rate (*Ms6-hm* + *Hm-2*) 0.055 per locus
- Induced paternal mutation rate (1 Gy) 0.225 per locus
- Radiation-induced increase in mutation rate 0.170 per locus
- Mean size for *Ms6-hm* & *Hm-2* (CBA/H mice) 5×10^3 bp
- Genome size 3×10^9 bp
- Damage to the whole genome $0.17 \times (3 \times 10^9) / (5 \times 10^3) = 100,000$

Radiation-Induced Damage in Eukaryotic Cells

- Base damage 2000 per 1Gy
- Single-strand breaks 1000 per 1Gy
- DNA-protein links 150 per 1Gy
- Double-strand breaks 40 per 1Gy
- Bulky lesions 40 per 1Gy
- Total 3300 per 1Gy $\ll 100,000$

From: Dubrova *et al.*, 1998, *PNAS* **95**, 6251-6255

Frankenberg-Schwager, 1990, *Radiat Environ Biophys* **29**, 273-292

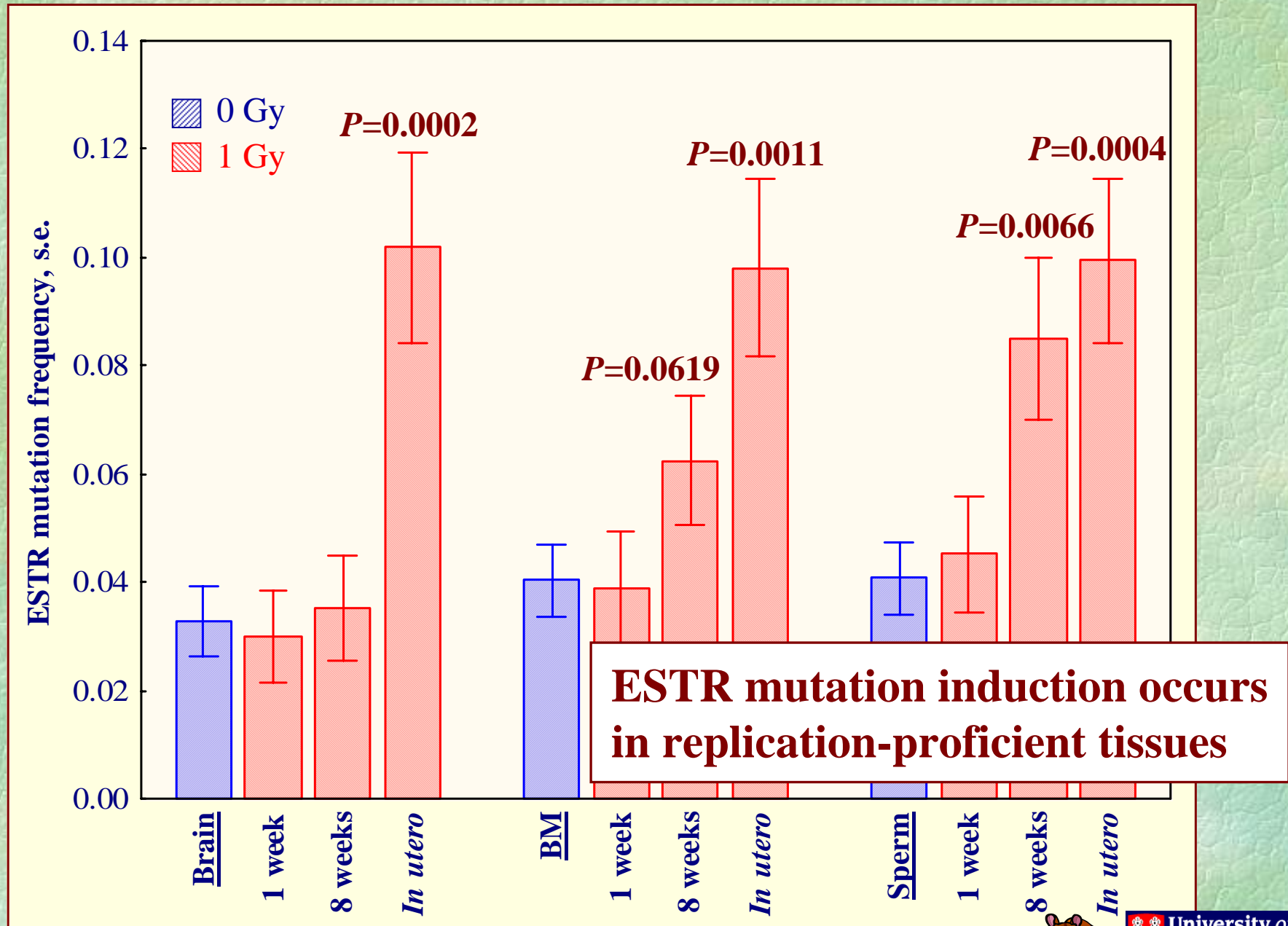


ESTR mutation induction and cell division

| Stage of irradiation | Tissue, proliferation capacity | | |
|--------------------------------|--------------------------------|-------------|-----------------|
| | Brain | Bone marrow | Male germ cells |
| Embryo males | | | |
| 12 days of gestation | + | + | + |
| Adult males | | | |
| 1 week after exposure | - | - | - |
| Adult males | | | |
| 10 weeks after exposure | - | +/- | + |



ESTR mutation induction in male mice exposed to 1 Gy X-rays



ESTR mutation induction occurs in replication-proficient tissues



Problems

- We observe changes in mutation frequency
- The magnitude of changes in mutation rate per cell division which corresponds to the observed increase

