

## Population genetics lab

(Please remind me to record if I forget)

14 October 2022

Taking attendance

BA-AZ-DO

- ▶ **Due: 28 OCT 2022 at 23:59 UK time**
- ▶ 25% of total grade
- ▶ Follow the [online lab](#)
- ▶ Turn in [on Canvas Quizzes](#)
- ▶ Emailed Tables 3 & 4

# The study system for the assignment

- ▶ Species: *Daphnia pulex*
- ▶ Small freshwater crustaceans
- ▶ Sensitive to environmental change<sup>1</sup>
- ▶ Can reproduce asexually & sexually<sup>2</sup>
- ▶ Sampled in 8 lakes near Chernobyl<sup>2</sup>



Figure 1: Close up picture of *Daphnia pulex*.

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<sup>1</sup>Flaherty, C M, & S I Dodson. 2005. *Chemosphere*. 61:200-207.

<sup>2</sup>Goodman, J, et al. 2019. *Ecol. Evol.* 9:2640-2650.

<sup>3</sup>Public Domain image by [Eric A. Lazo-Wasem](#)

## Microsatellites: repeated non-coding DNA sequences

- ▶ Repeated DNA sequences (e.g., 'AGTC')
- ▶ Do not code for proteins
- ▶ Not under selection (neutral)
- ▶ Mutations change repeat number

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### **Hypothetical microsatellite alleles:**

1. AGTCAGTCAGTCAGTC (4 repeats)
2. AGTCAGTC (2 repeats)
3. AGTCAGTCAGTC (3 repeats)

## Daphnia example with microsatellites

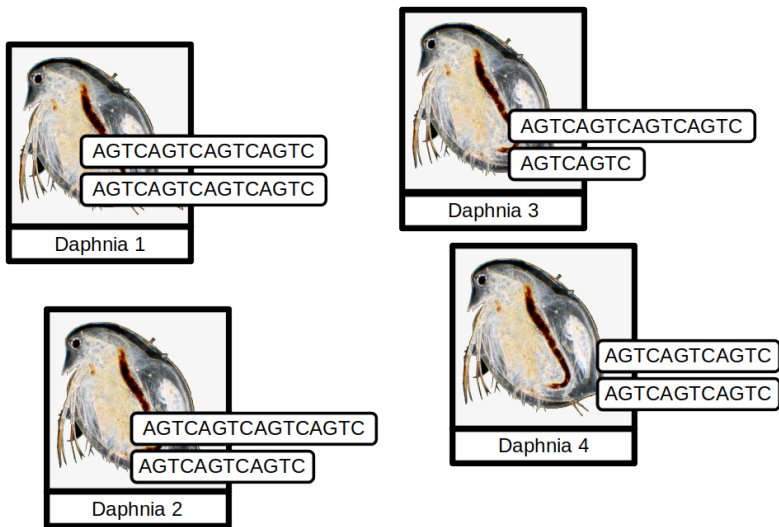


Figure 2: Example diploid population with microsatellite alleles.

## Can label microsatellites with a letter

1. AGTCAGTCAGTCAGTC (4 repeats)  $\rightarrow$  p
2. AGTCAGTC (2 repeats)  $\rightarrow$  q
3. AGTCAGTCAGTC (3 repeats)  $\rightarrow$  r



## Can label microsatellites with a letter

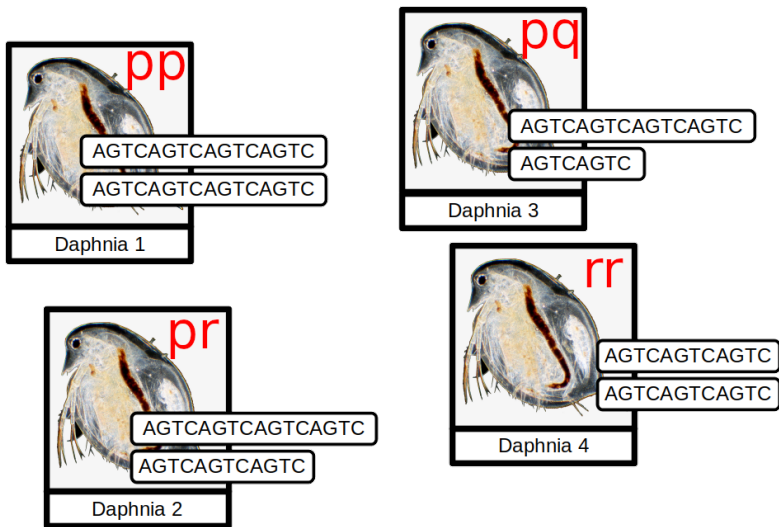


Figure 3: Example diploid population with microsatellite alleles labelled.

## Table 1 in the population genetics lab practical

Sample	Allele1	Allele2	Genotype
Ved1	152	152	pp
Ved2	152	152	pp
Ved3	152	152	pp
Ved4	152	152	pp
Ved5	152	152	pp
Ved6	152	152	pp
Ved7	152	152	pp
Ved8	152	152	pp
Ved9	144	152	pq
Ved10	148	152	pr
Ved11	152	152	pp
Ved12	152	152	pp
Ved13	152	152	pp
Ved14	152	152	pp
Ved15	152	152	pp
Ved16	144	152	pq
Ved17	152	152	pp
Ved18	152	152	pp
Ved19	152	152	pp
Ved20	152	152	pp
Ved21	152	152	pp
Ved22	144	152	pq
Ved23	152	152	pp
Ved24	152	152	pp
Ved25	152	152	pp
Ved26	148	152	pr
Ved27	152	152	pp

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<sup>1</sup>Auld, S. Radiation, genotypes and evolution in Chernobyl *Daphnia* – BIOU3GE. Evolution & Genetics. University of Stirling. 12 OCT 2022.

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Ved26	148	152	pr
Ved27	152	152	pp

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Ved27	152	152	pp

### ***Daphnia* samples from lake Vedilitsy**

- ▶ 22 pp genotypes
- ▶ 3 pq genotypes
- ▶ 2 pr genotypes

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### ***Daphnia* samples from lake Vedilitsy**

- ▶ 22 pp genotypes
- ▶ 3 pq genotypes
- ▶ 2 pr genotypes
- ▶ 49 p alleles
- ▶ 3 q alleles
- ▶ 2 r alleles

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### ***Daphnia* samples from lake Vedilitsy**

- ▶ 22 pp ( $22/27 = 0.814$ )
- ▶ 3 pq ( $3/27 = 0.111$ )
- ▶ 2 pr ( $2/27 = 0.074$ )
- ▶ 49 p ( $49/54 = 0.90$ )
- ▶ 3 q ( $3/54 = 0.06$ )
- ▶ 2 r ( $2/54 = 0.04$ )

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- ▶ 49 p ( $49/54 = 0.90$ )
- ▶ 3 q ( $3/54 = 0.06$ )
- ▶ 2 r ( $2/54 = 0.04$ )

(Note, there are zero observed qq, rr, or qr genotypes)

## Table 2 in the population genetics lab practical

Allele	Number	Freq
p (152)	59	0.90
q (144)	3	0.06
r (150)	2	0.04

Genotype		Exp. Freq.	Obs. Freq.
pp	$p^2$	?	0.81481481
qq	$q^2$	?	0
rr	$r^2$	?	0
pq	$2pq$	?	0.11111111
pr	$2pr$	?	0.07407407
qr	$2qr$	?	0



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Allele	Number	Freq
p (152)	59	0.90
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r (150)	2	0.04

### Exp. Freq.

- **pp**: Probability of sampling  $p$  for the first allele, then  $p$  again for the second allele is  $p \times p = p^2$  ( $0.90^2 = 0.81$ ).

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### Exp. Freq.

- ▶ **pp**: Probability of sampling  $p$  for the first allele, then  $p$  again for the second allele is  $p \times p = p^2$  ( $0.90^2 = 0.81$ ).
- ▶ **pq**: Probability of sampling  $p$  for the first allele, then  $q$  for the second allele, **or** sampling  $q$  for the first allele, then  $p$  for the second allele is  $(p \times q) + (q \times p) = 2pq$  ( $2 \times 0.90 \times 0.06 = 0.108$ ).

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Allele	Number	Freq
p (152)	59	0.90
q (144)	3	0.06
r (150)	2	0.04

Genotype		Exp. Freq.	Obs. Freq.
pp	$p^2$	0.81	0.81481481
qq	$q^2$	0.0036	0
rr	$r^2$	0.0016	0
pq	$2pq$	0.108	0.11111111
pr	$2pr$	0.072	0.07407407
qr	$2qr$	0.0048	0

<sup>1</sup>Auld, S. Radiation, genotypes and evolution in Chernobyl Daphnia – BIOU3GE. Evolution & Genetics. University of Stirling. 12 OCT 2022.

## Table 2 in the population genetics lab practical

$$H_O = Obsf_{pq} + Obsf_{pr} + Obsf_{qr}$$

$$H_O = 0.111 + 0.074 + 0 = 0.185$$

$$H_E = Expf_{pq} + Expf_{pr} + Expf_{qr}$$

$$H_E = 0.108 + 0.072 + 0.005 = 0.185$$

Genotype		Exp. Freq.	Obs. Freq.
pp	$p^2$	0.81	0.81481481
qq	$q^2$	0.0036	0
rr	$r^2$	0.0016	0
pq	$2pq$	0.108	0.11111111
pr	$2pr$	0.072	0.07407407
qr	$2qr$	0.0048	0

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## Question 1 of the practical

- ▶ Use Table 3 to complete Question 1 A-F
- ▶ Procedure similar to what we just did
- ▶ New data set is from a different lake
- ▶ Note that there will be one more allele

## F statistics: What is identity-by-descent?

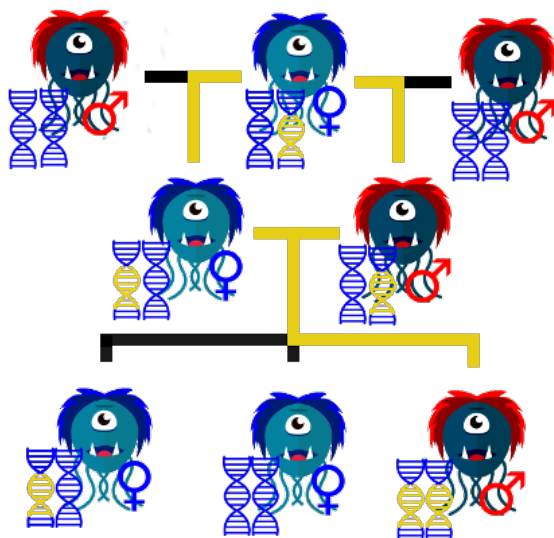


Figure 4: Cartoon figure illustrating identity-by-descent

## Probability of identity-by-descent

The probability two alleles are identical-by-descent,

$$F_{IS} = \frac{H_E - H_O}{H_E}.$$

This is also called the *inbreeding coefficient*.

Proportion of total genetic variance contained in the subpopulation

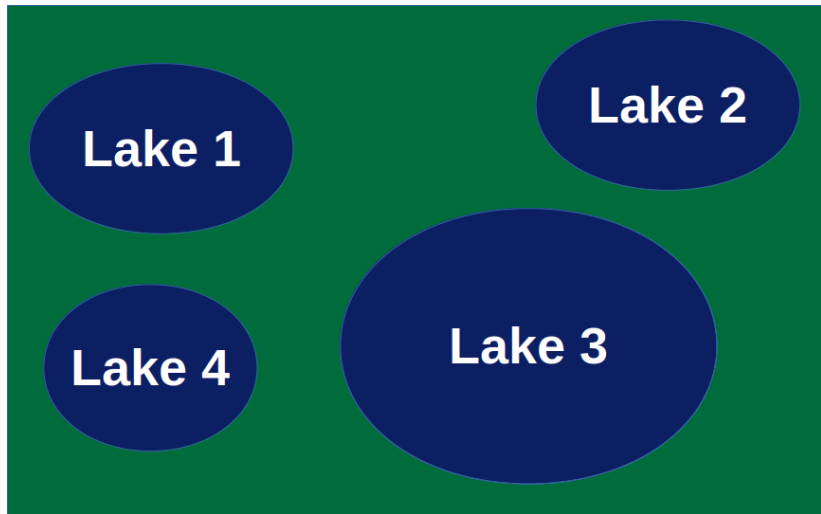


Figure 5: Four lakes (subpopulations) on a landscape.



## Proportion of total genetic variance contained in the subpopulation

$$F_{ST} = \frac{H_T - H_S}{H_T}$$

The above  $F_{ST}$  measures genetic drift.

- ▶  $H_T$  is total heterozygosity (if it were all one big lake)
- ▶  $H_S$  is average heterozygosity of the subpopulations
- ▶ If  $H_T = H_S$ , then  $F_{ST} = 0$ .
- ▶ If  $H_S = 0$ , then  $F_{ST} = 1$ .

## Population genetics lab practical Table 4

Population	Dose	Log10(Dose)	N	MLG	HE	HO	PA	TAR
Vediltsy	0.1	-1.00	27	27	0.447	0.564	1	29.202
Smolin	0.12	-0.92	28	28	0.517	0.452	2	34.687
Yampol	0.2	-0.70	28	28	0.416	0.408	2	29.424
Glinka	1.17	0.07	28	28	0.427	0.282	1	28.771
Buryakovka	1.77	0.25	28	28	0.448	0.325	0	31.427
Krasnyansky	55.79	1.75	38	38	0.611	0.491	3	40.704
Gluboke	181.15	2.26	28	27	0.599	0.650	4	40.429