

Students 1 2 3 4 5 6 7 8 ... 22

DNA

extraction
(n=22)

✓ ✓ ✓ ✓ ✓ ✓ ✓ ✓ ... ✓

Groups
(n=6)

N=3 N=3 N=3 N=3 N=3 N=3 N=4
A B C D E F G

DNA Frag

Vortex / Needle / Temp / Control

Barcode

✓ ✓ ✓ ✓

Library prep

✓

Sequencing

A B C D E F

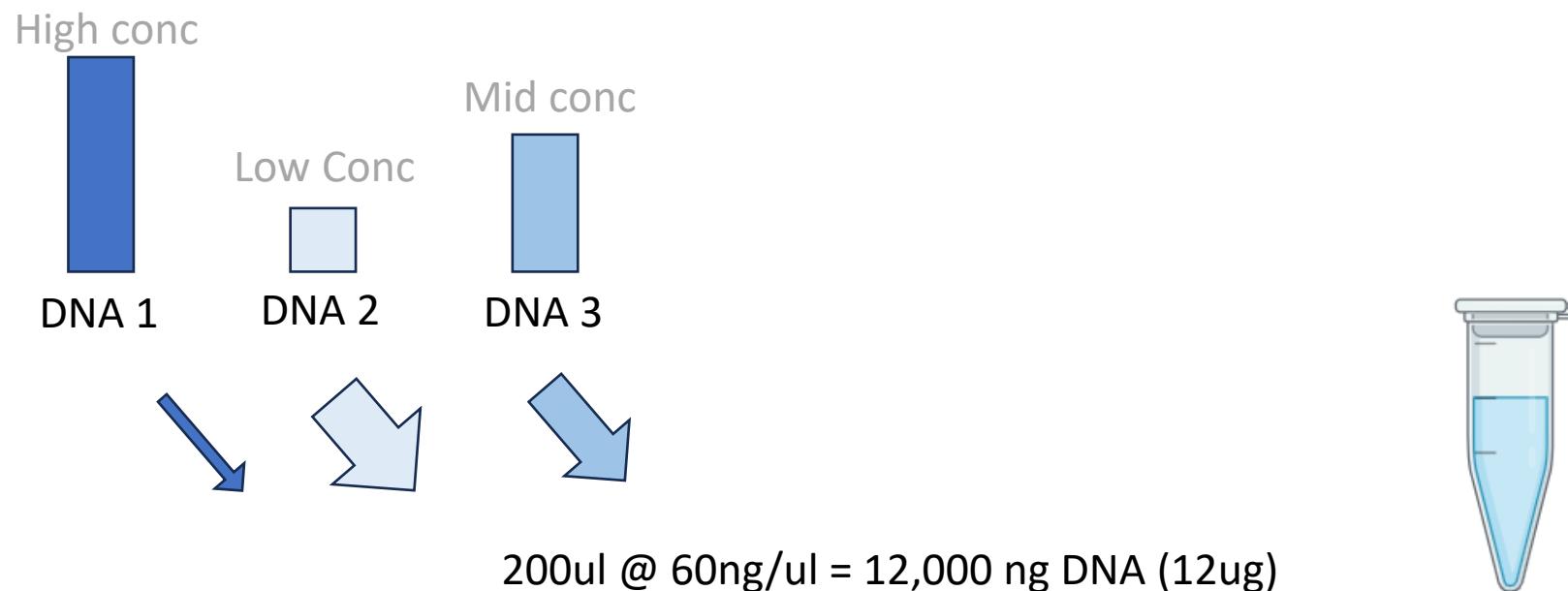
YES

Key points, Day 1

- Day 1 : extract DNA from blood, measure concentration with qubit
 - We will **re-quantify** after 3 days, run on gel and perform nanodrop.

Key points, Day 2

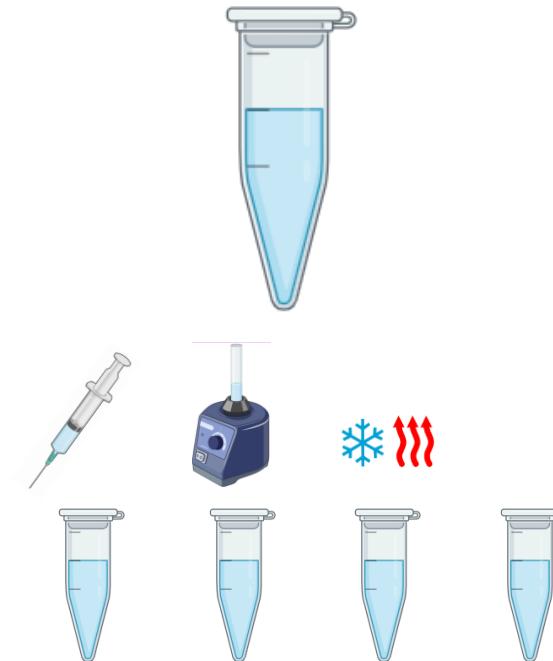
- Day 2 : using the re-quantify qubit values, within each group, combine an equal mass of DNA from each extraction ($n=3$) to make a final solution in water with a volume of 200ul and concentration of 60ng/ul.



Key points, Day 2

- Divide this into four microfuge tubes as follows:

- Needle shearing = 70ul
- Vortexing = 70ul
- Temperature extremes = 20ul
- Control = 20ul



DNA Treatments

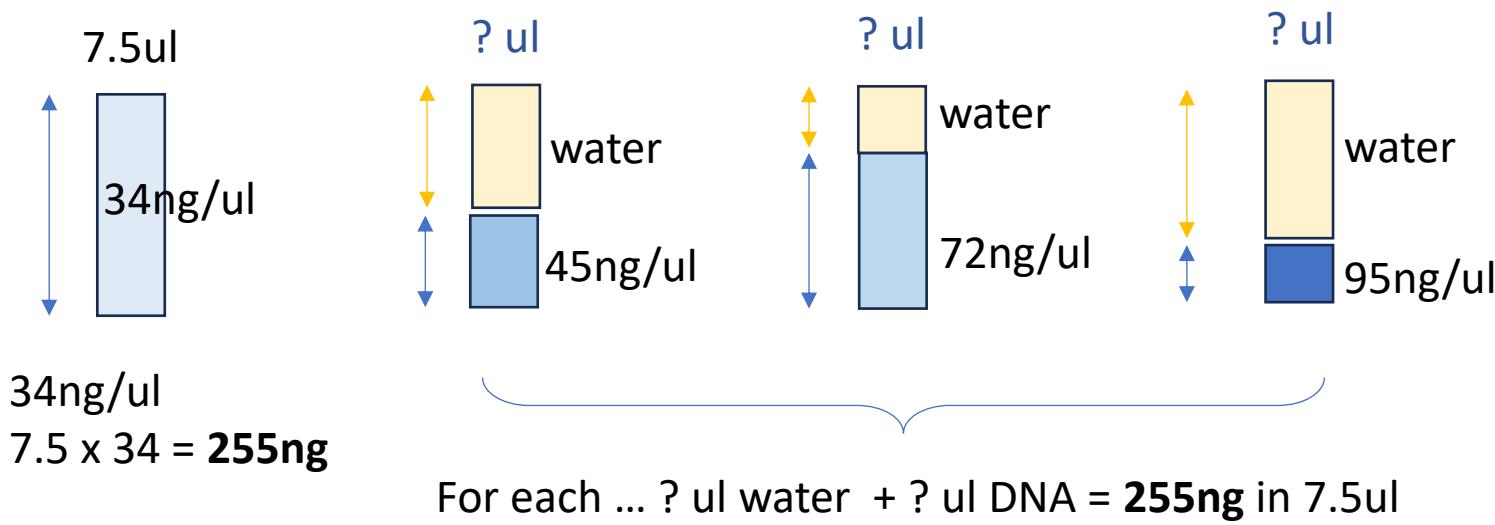
- **70ul** Needle shearing : pass DNA in solution up and down quickly through a blunt needle 30X.
- **70ul** Vortex : vortex DNA and metal bead in solution for 60 seconds at full speed.
- **20ul** Temp x-trem : Heat for 45s, freeze for 45s, repeat for total 4X.
- **20ul** Control : no treatment

Key points, Day 2 cont'

- Treated DNA will now be nick repaired and end-prepped, and then cleaned with beads and quantified (again!).

Key points, Day 3

- Calculate the total mass (ng) of DNA in 7.5ul of the treatment with lowest conc.
- For the three other treatments, calculate how much DNA and H₂O to combine to give a final mixture of 7.5ul containing the same mass of DNA as the lowest conc treatment.



Key points, Day 3

- Each sample will receive a different barcode.
- Samples within group will be combined (pooled) to one.
- This will be cleaned, quantified and used as input to adapter ligation.
- Adapter ligation is done on one sample per group. Finish with quantification.
- Prior to loading, Mariann will combine equal mass from each group.
- Video on loading here...
 - <https://www.youtube.com/watch?v=pfftqlldRSbk>

Dry lab intro

