

# Thursday, 26 July

- Convert (remapped), sorted, filtered, (phased) .vcf to OneMap input file (.raw format)
- Using R
- Linkage mapping with OneMap

# Cygwin: a LINUX-like environment for Windows

<https://www.cygwin.com/>

<https://cygwin.com/faq.html>

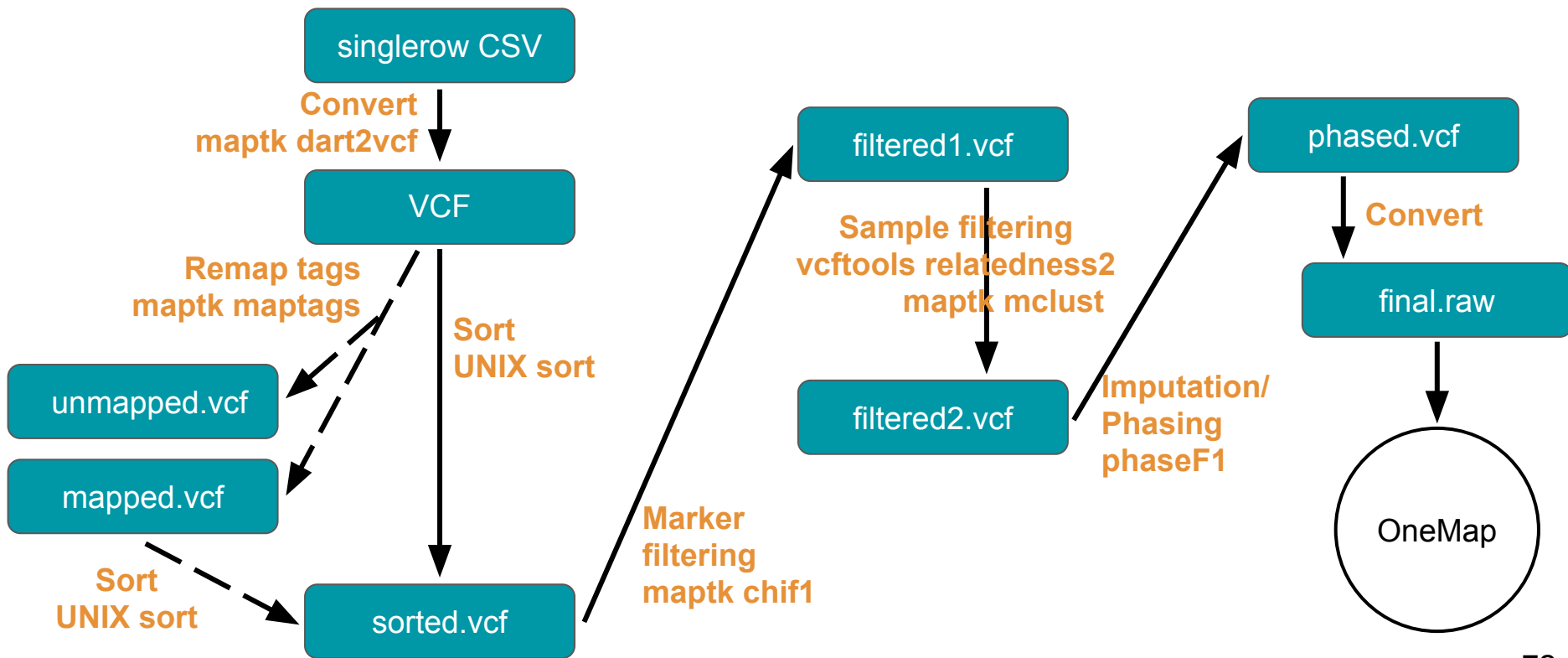
<https://cygwin.com/cygwin-ug-net/cygwin-ug-net.html>

for e.g. vcftools, you will have to compile it on your computer

any arguments that reference the filesystem must be in Windows (or DOS) format or translated

→ `cygpath` utility (<https://cygwin.com/cygwin-ug-net/using-effectively.html>)

# DArTseq to mapping: analysis flow chart



# maptk vcf2raw

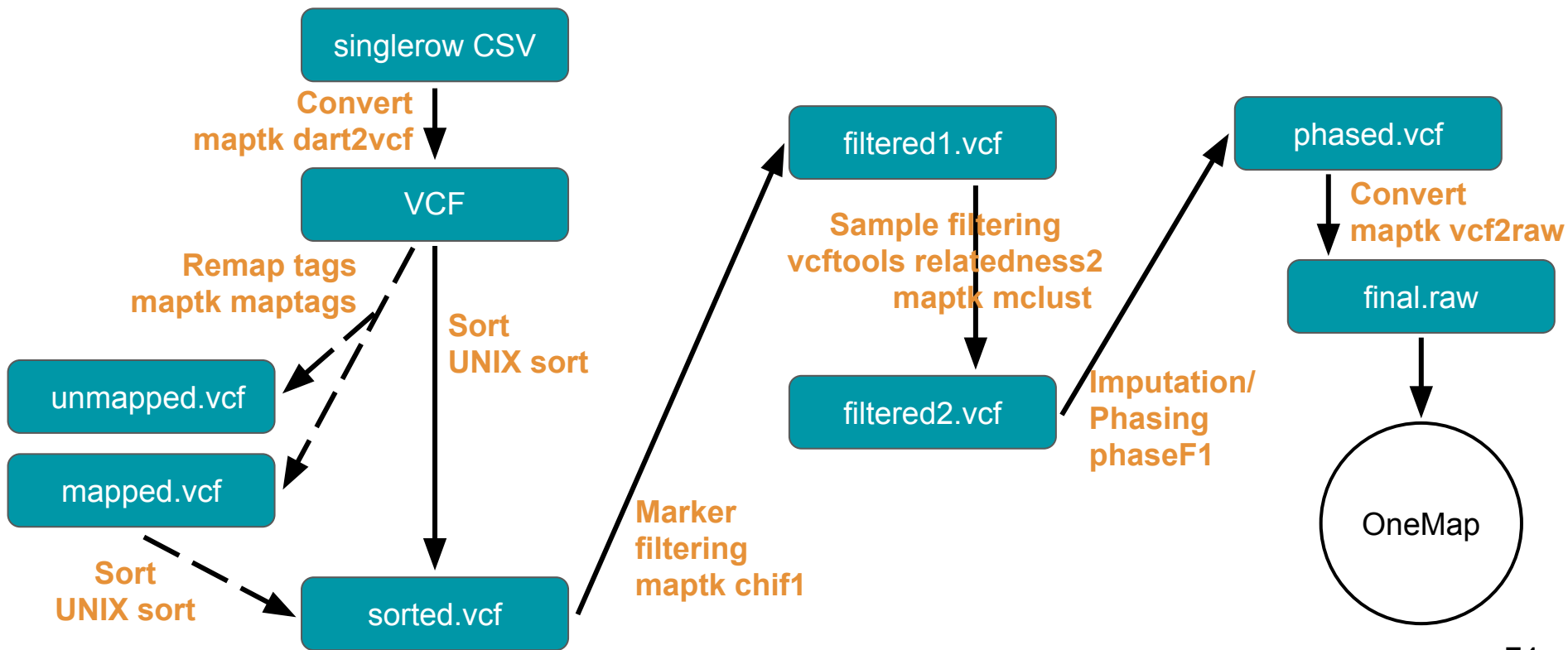
jess — lyons@ip-172-31-31-27:~ — ssh -i ~/Dropbox/DARTseq\_to\_mapping\_workshop/lyons.pem lyons@ec2-13-57-194-80.us-west-1.compute.amazonaws...

```
data type outcross
```

```
146 348 0 0 0
```

```
F1-0A01 F1-01A02      F1-01A03      F1-01A04      F1-01A05      F1-01A06      F1-01A07      F1-01A08      F1-01A09      F1-01A
*000020132|F|0-37 D2.15 ab a ab a a ab a ab a a ab a ab ab ab a ab a ab a a a a a ab a ab a ab a a a a ab
*000020135|F|0-37 D1.10 ab ab ab a a ab a a ab ab ab a a ab a a a ab a ab ab ab a a ab a ab a ab ab
*000020528|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a ab a a ab ab a a a ab ab ab a
*000020529|F|0-37 B3.7 b ab b a a b a ab ab b ab a ab ab ab ab ab ab ab a b ab ab a ab ab b a b b a a ab b ab ab a ab ab
*000020286|F|0-37 B3.7 a ab a b b a b ab ab a ab b ab ab ab ab ab ab ab b a ab ab b ab ab a ab ab b ab ab
*000020287|F|0-37 B3.7 a ab a b b a b ab ab a ab b ab ab ab ab ab ab ab b a ab ab b ab ab a ab ab b a b a b b ab a ab ab b ab ab
*000020266|F|0-37 D1.10 ab ab ab a a ab a a ab ab ab a a ab a a a ab a a ab ab ab a ab a a ab a ab ab a a a ab a ab a ab ab
*000020337|F|0-37 D1.10 a a a ab ab a ab ab a a a ab ab a ab ab ab a ab ab a a a ab a a ab a a ab ab a ab a a a a
*000020338|F|0-37 D1.10 a a a ab ab a ab ab a a a ab ab a ab ab ab a ab ab a a a ab a a a ab a ab ab a ab a ab a a a a
*000020339|F|0-37 D1.10 a a a ab ab a ab ab a a a ab ab a ab ab ab a ab ab a a a ab a a a ab a ab ab a ab a ab a a a a
*000020341|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a ab ab ab ab a a ab a ab a a a ab ab ab a
*000020343|F|0-37 D2.15 ab a ab a a ab a ab a ab a a ab a ab ab ab a ab a ab a a a a a ab ab a ab a ab ab ab a a a a ab
*000020345|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a ab ab ab ab a a ab a ab a a a ab ab ab a
*000020346|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a ab ab ab ab a a ab a ab a a a ab ab ab a
*000020347|F|0-37 D2.15 ab a ab a a ab a ab a ab a a ab a ab ab ab a ab a ab a a a a a ab ab a ab a ab ab ab a a a a ab
*000020348|F|0-37 D2.15 ab a ab a a ab a ab a ab a a ab a ab ab ab a ab a ab a a a a a ab ab a a ab ab ab a a a a ab
*000020349|F|0-37 D1.10 a ab a ab ab a ab ab a a a ab ab a ab ab ab a ab ab a a a ab a a a ab a ab a a ab ab a ab a a a
*000020478|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a ab ab ab ab a a ab a ab a ab a a a ab ab ab a
*000020535|F|0-37 D2.15 ab a ab a a ab a ab a ab a a ab a ab ab ab a ab a a a a ab ab a a a ab ab ab a a a a ab
*000020536|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a ab ab ab ab a a ab a ab ab a a a ab ab a
*000020537|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a ab ab ab ab a a ab a ab ab ab a a a ab ab ab
*000020538|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a ab ab ab ab a a ab a ab ab a a a ab ab ab
*000020539|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a ab ab ab ab a a ab ab ab a a a ab ab ab
*000020540|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a ab ab ab ab a a ab ab ab a a a ab ab ab
*000020541|F|0-37 D2.15 ab a ab a a ab a ab a ab a a ab a ab ab ab a ab a ab a a a a a ab ab a a ab ab ab a a a a ab
*000020543|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a ab ab ab ab a a ab ab a a a ab ab ab
*000020544|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a a ab ab ab a a ab ab a a a ab ab ab a
*000020545|F|0-37 D2.15 a ab a ab ab a ab a ab a ab ab a ab a a a ab a ab a a ab ab ab a a ab ab a a a ab ab ab a
*000020444|F|0-37 B3.7 b a b a b a ab ab b ab a ab ab ab ab ab ab a b b b a ab ab b ab ab ab a ab a b b a a ab b ab ab a ab ab b
*000020443|F|0-37 B3.7 b a b a b a ab ab b ab a ab ab ab ab ab ab a b b b a ab ab b ab ab ab a ab a b b a a ab b ab ab a ab ab b
*000020442|F|0-37 D2.15 ab a ab a a ab a ab a ab a a ab a ab ab ab a ab a ab ab ab a a a ab ab a ab a a a ab ab ab a a a a a
```

# DArTseq to mapping: analysis flow chart



# RRRRR matey

## **R statistical programming**

Very good linear algebraic support (vectors, matrices)

Publication-quality figures

Interactive interpreter and Rscript

Portable on all operating systems

Goals: be able to use R/OneMap and understand what you are doing



# Teaching yourself R

Comments:

# Write yourself lots of notes

Learn more about a function/package:

?thing

Rstudio, a GUI for R:

<https://www.rstudio.com>



# R: Variables, Math, Logic

Valid variable names may contain A-Z, a-z, 0-9 characters, as well as “\_” and “.”  
(Cannot begin with a number)

## Assignment:

`x = 5`

`x <- 5`

## Comparison:

Equivalent to: `x == y`

Not equivalent: `x != y`

Less-than: `x < y`

Less-than-or-equal-to: `x <= y`

Greater-than: `x > y`

Greater-than-or-equal-to: `x >= y`

## Logic:

And: `x & y`

Or: `x | y`

Negate: `!x`

## Math operations:

Addition: `x + y`

Subtraction: `x - y`

Multiplication: `x * y`

Division: `x / y`

Power: `x^y` or `x**y`

Modulo: `x %% y`

Log: `log(x)`

Square root: `sqrt(x)`



# R: Objects

Variables: are “objects” (containers for data):

`object = data`

Belong to a class/have “type” (character, numeric, vector, factor, etc.):

`class(object)`

May contain attributes:

`names(object)` # what attributes are there?

`object$attribute` # use the attribute as a variable



# variable types and coercion

**NULL:** NULL

**numeric:** -1, 0, 1, -1.0, 0.05, 1e-6

**character:** "string", "1", "FALSE"

**vector:** 1D sequence

```
V = c(1, 5, 10);  
value = V[index];  
V[index] = value
```

**matrix:** 2D table (typically numeric)

```
M = matrix(1:9, nrow=3, ncol=3)  
value = M[i,j]  
M[i,j] = value
```

**data.frame:** Excel-like table

```
D = data.frame(field1=1:9, field2=11:19)  
value = D[i,j]; D[i,j] = value  
value = D$field1; D$field1 = value
```

**logical:** TRUE, FALSE, NA

**list:** Multi-D table

```
L = list(field1=1:9, field2=11:19)  
value = L[i,j]; L[i,j] = value  
value = L$field1; L$field1 = value  
L = list(1:9, 11:19)  
value = L[[i]][j]; L[[i]][j] = value
```

**Testing for type:**

`is.type(object)`

**Coerce to type:**

`new.type = as.type(old.type)`

# Conditional statements

Template:

```
if (test1) {  
    execute this line  
  
} else if (test2) {  
    execute this line  
  
} else { # defaults to this  
    execute this line  
}
```

Template:

```
if (test1) {  
    if (test2) {  
        execute this line  
    }  
    execute this line  
  
} else { # defaults to this  
    execute this line  
}
```

# Conditional statements

**Example:**

```
i = 0
if (i < 10) {
    print("< 10")
} else { # defaults to this
    print(">= 10")
}
```

**Example:**

```
i = 5
if ((0 <= i) & (i <= 5)) {
    print("in range [0, 5]")
} else if (i < 0){
    print("negative")
} else {
    print("greater than 5")
}
```

# for and while

## Template:

```
for (variable in sequence) {  
    execute this code  
}
```

```
while (condition-is-true) {  
    execute this code  
}
```

## Example:

```
for (i in 1:10) {  
    print(i)  
}
```

```
j = 0  
while (j < 15) {  
    print(j)  
    j = j + 1  
}
```

# functions

## Template:

```
f = function(x=default) {  
  compute something  
  return(result)  
}  
  
y = f(x)
```

## Example:

```
f = function(x, add=FALSE) {  
  if (add) {  
    return(x + 1)  
  } else {  
    return(x - 1)  
  }  
}  
  
y = f(x)
```

# Other useful functions

<code>plot</code>	Plot X, Y data	<code>read.csv</code>	Read CSV file
<code>hist</code>	Plot a histogram	<code>write.csv</code>	Write CSV file
<code>seq</code>	Create a sequence	<code>length</code>	Calc. length of object
<code>pdf</code>	Open a PDF device for writing	<code>names</code>	Get attribute names
<code>dev.off</code>	Close device to save	<code>append</code>	Append item to vector
<code>paste</code>	Paste together data to make a string	<code>setwd</code>	Set working dir
<code>sort</code>	Sort a vector, list, etc.	<code>getwd</code>	Get working dir
<code>read.table</code>	Read in a space-/tab-delimited file	<code>ls</code>	Show defined variables
<code>write.table</code>	Write a table to file	<code>summary</code>	Summarize object data
<code>install.packages</code>	Install R packages from CRAN	<code>lapply</code>	Iterate and create list

# R exercises

1. Use R to calculate the sum of 5 and 8 and assign the output to a variable x.
2. Use R to determine if x is divisible by 2 (hint: see modulo operator).
3. Use `is.integer` type test to determine whether x is an integer. If not, what type is it?
4. Use a conditional statement to test whether x is equivalent to 12 AND 13
5. Use a conditional statement to test whether x is equivalent to 12 OR 13
6. Use the `seq` function to output even numbers between 0 and 50
7. Combine your solution to question 6 with a `for` loop to print odd numbers.
8. Create a function to calculate the difference between two numbers.
9. Use `data(cars)` to load some example data. The `cars` variable becomes defined. What columns does it contain? Plot a histogram of the first column. What is the mean? Use `sd` to calculate the standard deviation.



# R exercises

10. To save a plot to PDF, you must first call `pdf("name_of.pdf")`, then the plotting function of interest, then `dev.off()` to close the file. Save a PDF of your histogram from question 9.

# Genetic linkage map estimation

# Linkage mapping with OneMap

## Why OneMap?

- Free!
- Supports many different cross types (F1, F2, RIL, BC)
- Parallelizable
- Very flexible (not only one analysis)
- Portable to any system

# Loading the OneMap package and reading data

To load the OneMap package, use `require()`  
`require(onemap)`

When you load mapping data, cross type is read from the file  
`cross = read_onemap("/path/to/dir", "F1.raw")`

# Checking your data quality

Plots color genotype matrix

```
plot(cross)
```

Was our chi-squared test stringent enough?

```
plot(test_segregation(cross))
```

OneMap can give you a recommended LOD score

```
lod.suggested = suggest_lod(cross)
```

$$LOD = Z = \log_{10} \frac{\text{probability of birth sequence with a given linkage value}}{\text{probability of birth sequence with no linkage}} = \log_{10} \frac{(1 - \theta)^{NR} \times \theta^R}{0.5^{(NR+R)}}$$