https://jolars.shinyapps.io/eulerr/

#domesticated is blue 1C, wild is orange EE, sensitivity is black 05

myColors <- c("#1C86EE", "#050505", "#EE7600")

opacity = 0.6

Order for figure: D, S, SW, W, DS, DW, DWS

Color order: #1C86EE, #050505, #EE7600

Solid borders. Opacity 0.6

Save as each file format.

Gimp

Editing: default font, size 18

Compression level: 5

B07\_GEMMA\_genes.R

Domestication SNP level Venn

ALL D DS DW S W WS

1 2199 75 1369 1195 1020 6

Domestication Gene level Venn

This is only 1 gene / SNP (arbitrary, first record)

And then calculated overlap of traits per gene:

ALL D DS DW S W WS

237 1147 201 1180 625 502 79