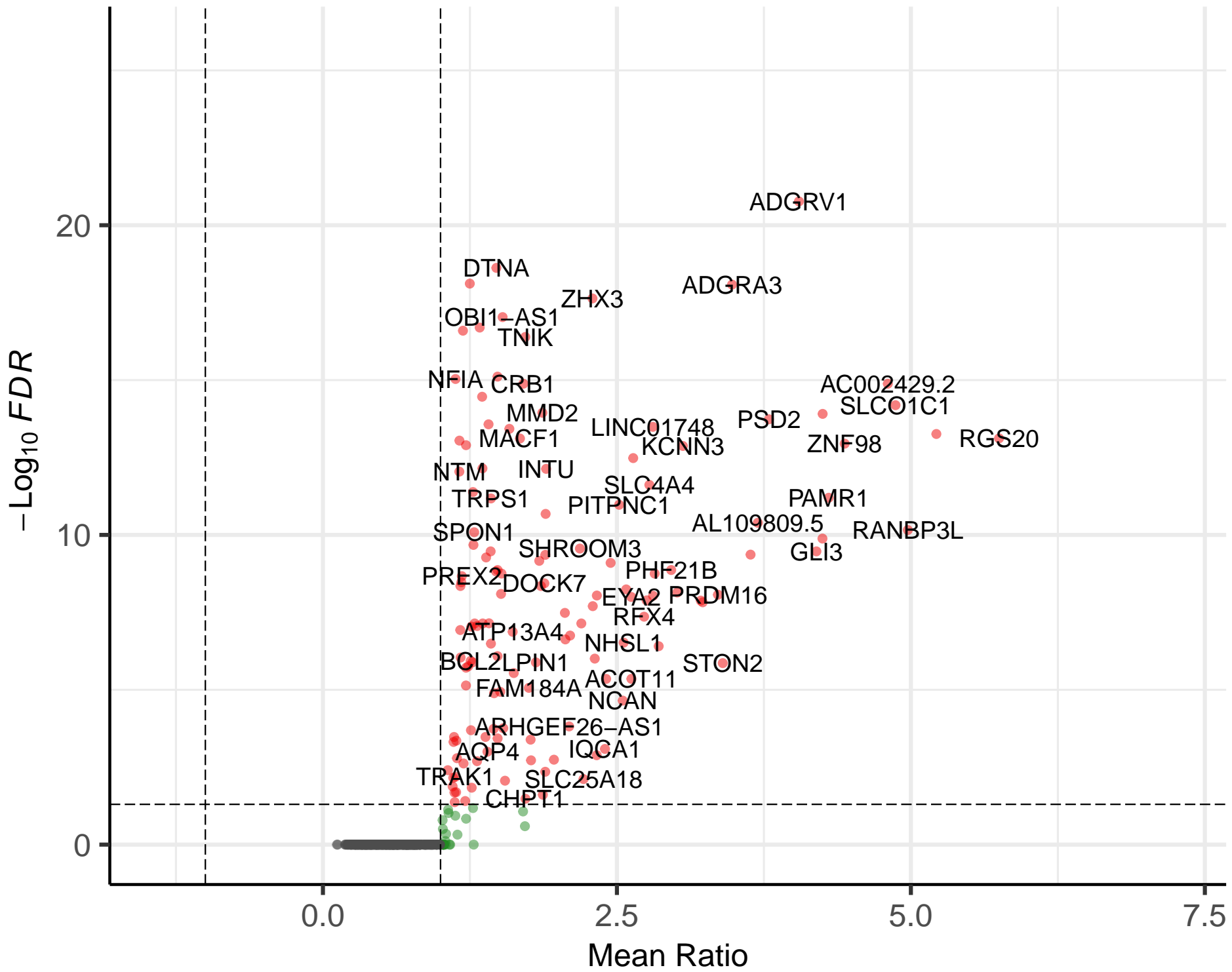


Astro

n nuc=3979, n genes FDR<0.05=136

● NS ● \log_2 FC ● p-value ● p-value and \log_2 FC

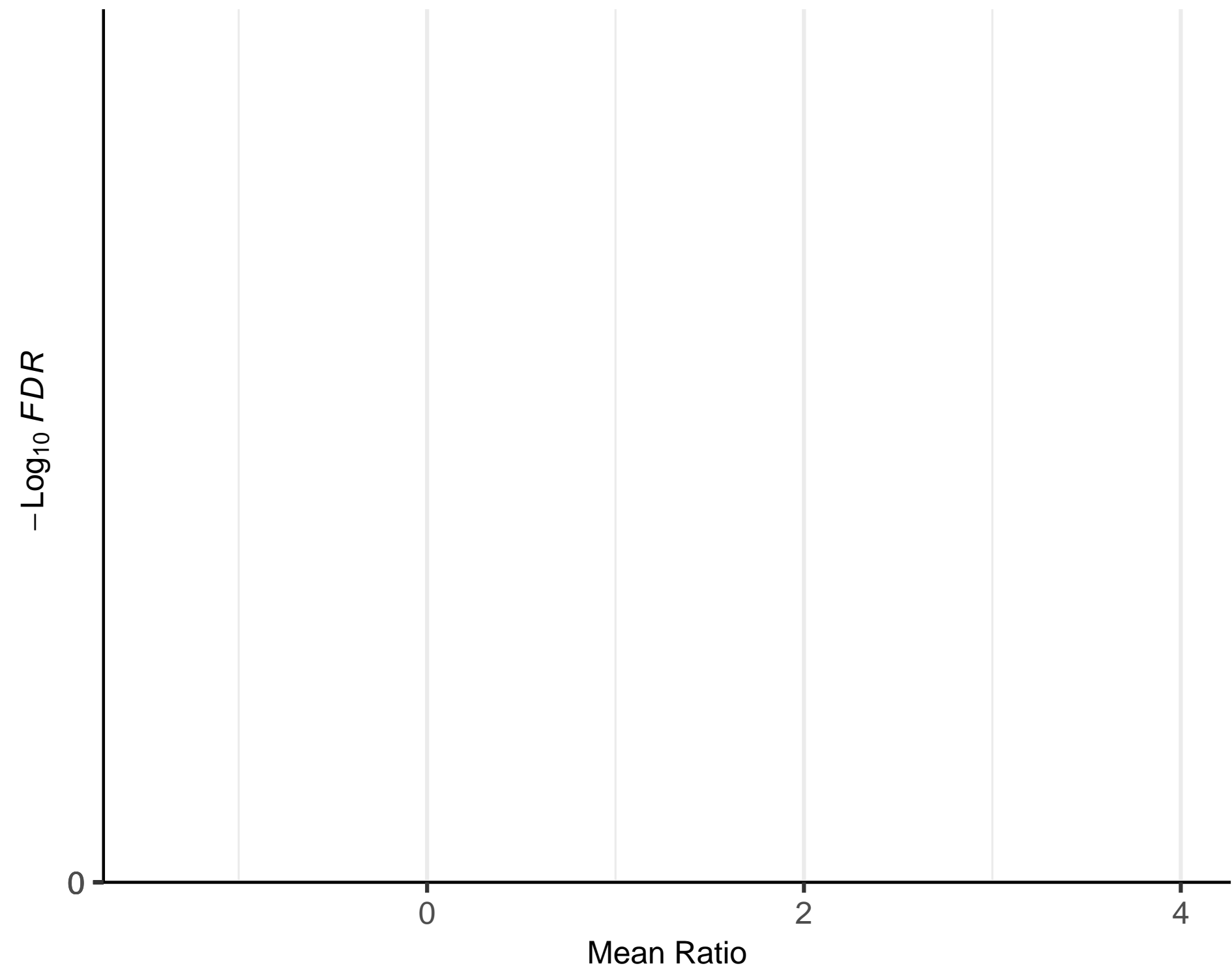


total = 595 variables

Endo.Mural_01

n nuc=446, n genes FDR<0.05=NA

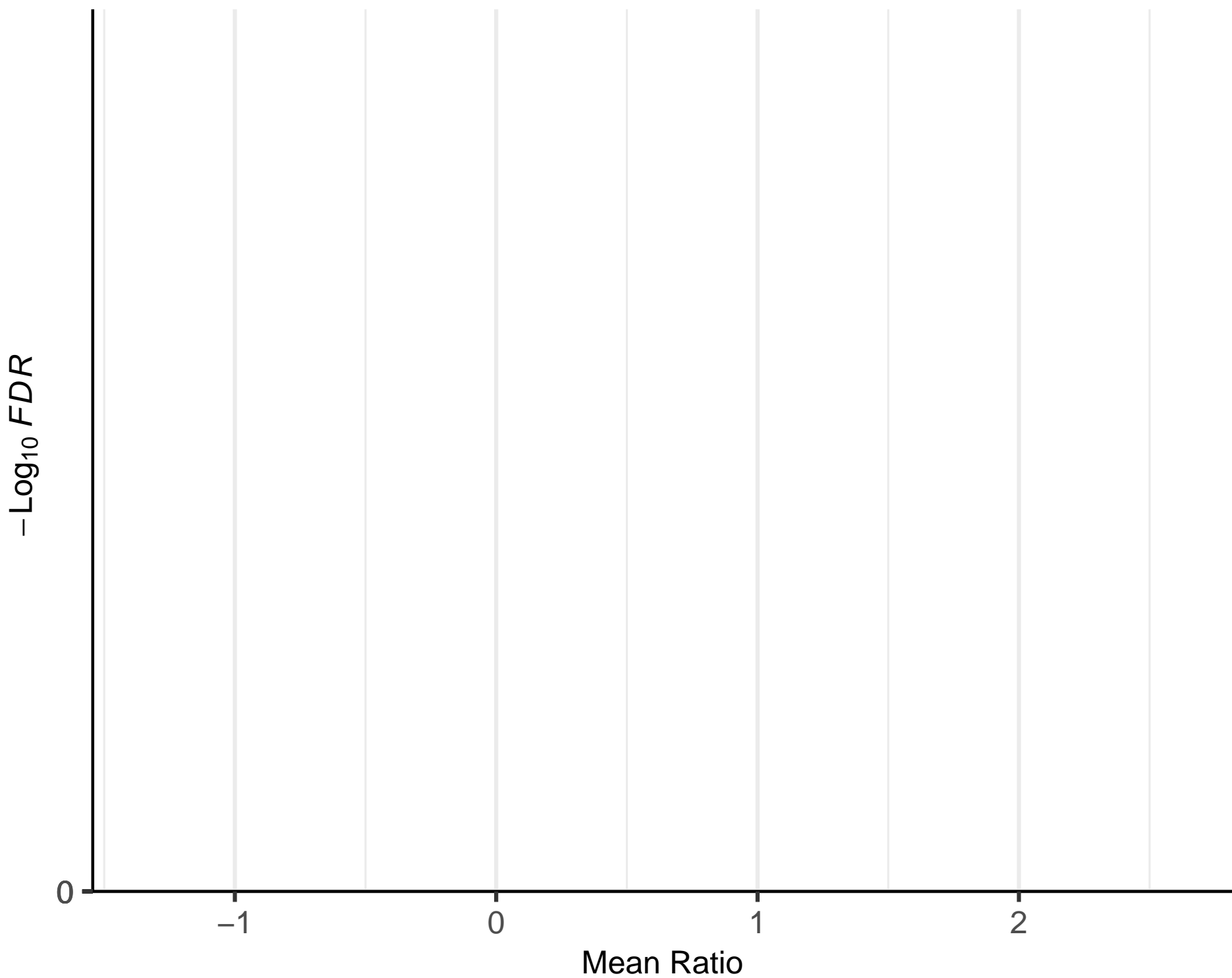
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Endo.Mural_02

n nuc=1711, n genes FDR<0.05=NA

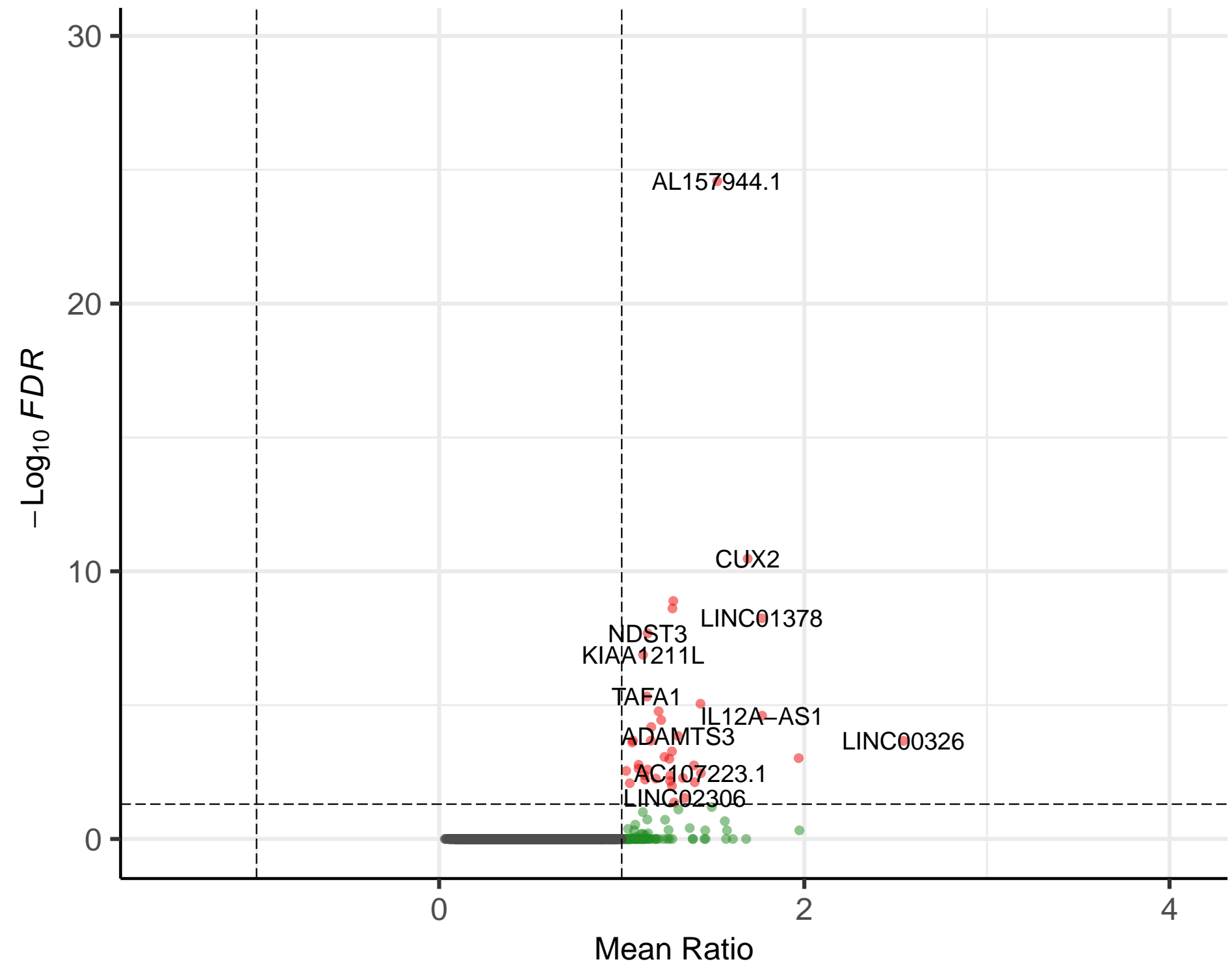
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Excit_01

n nuc=7927, n genes FDR<0.05=39

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

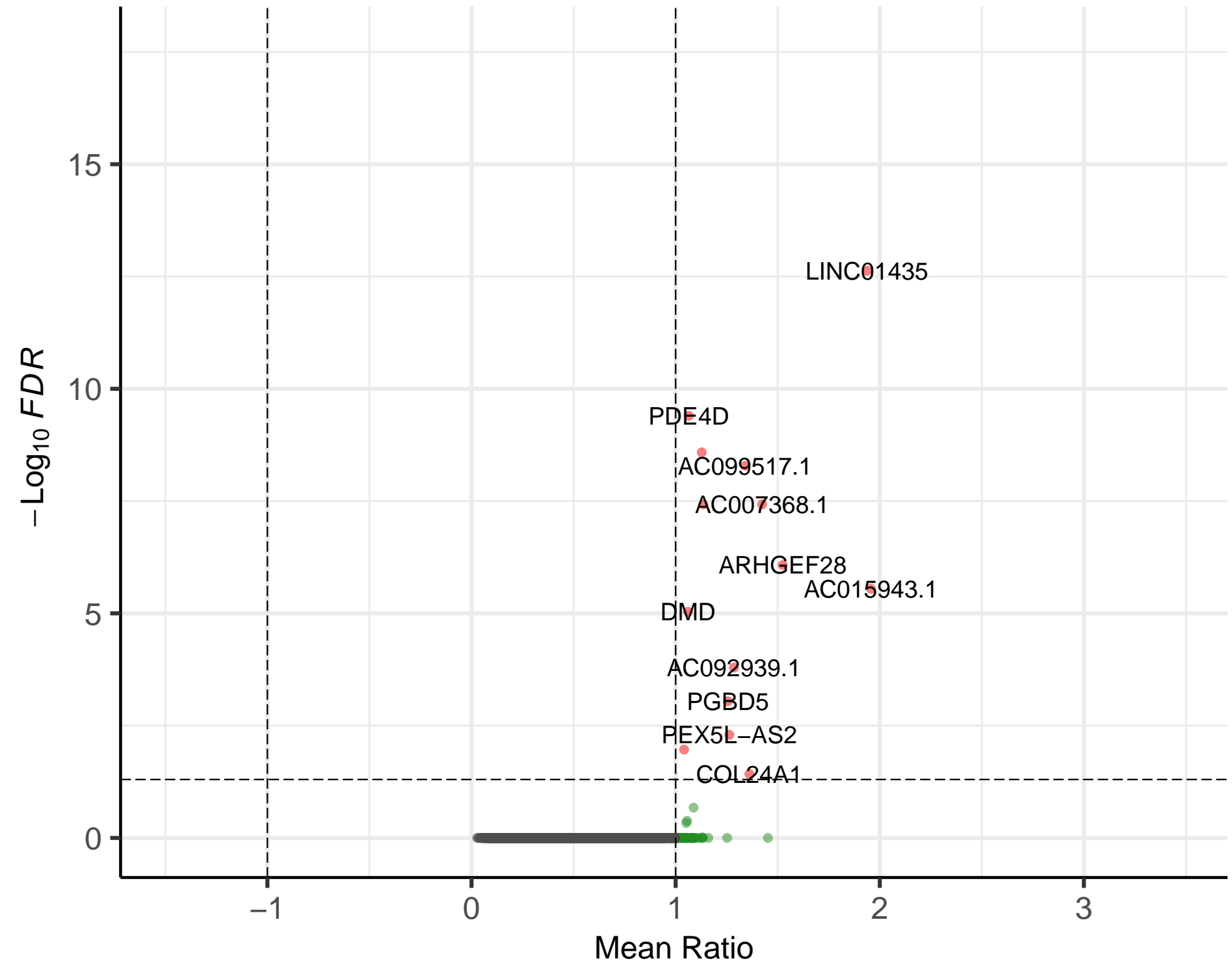


total = 6923 variables

Excit_02

n nuc=2487, n genes FDR<0.05=14

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

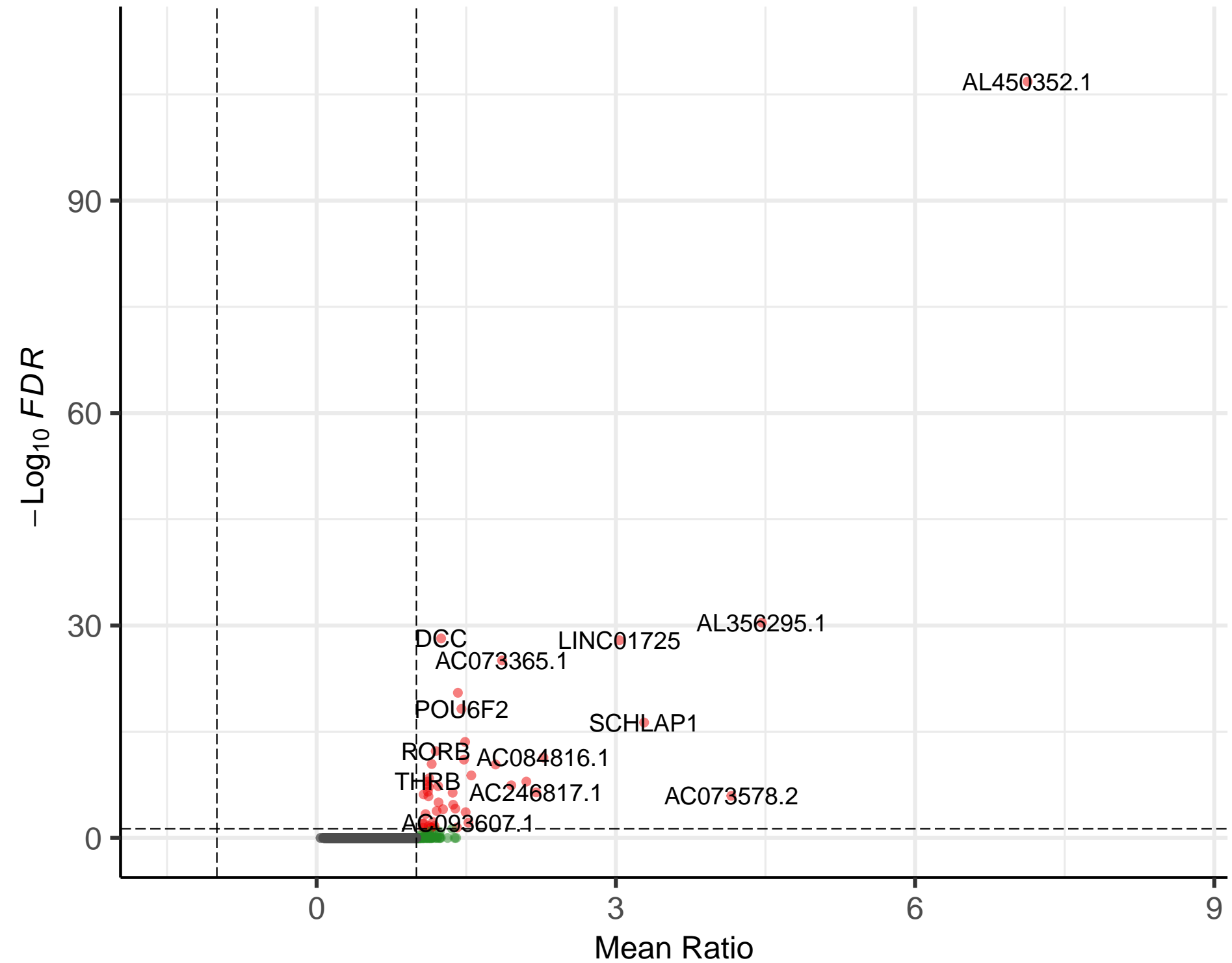


total = 7290 variables

Excit_03

n nuc=1309, n genes FDR<0.05=45

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

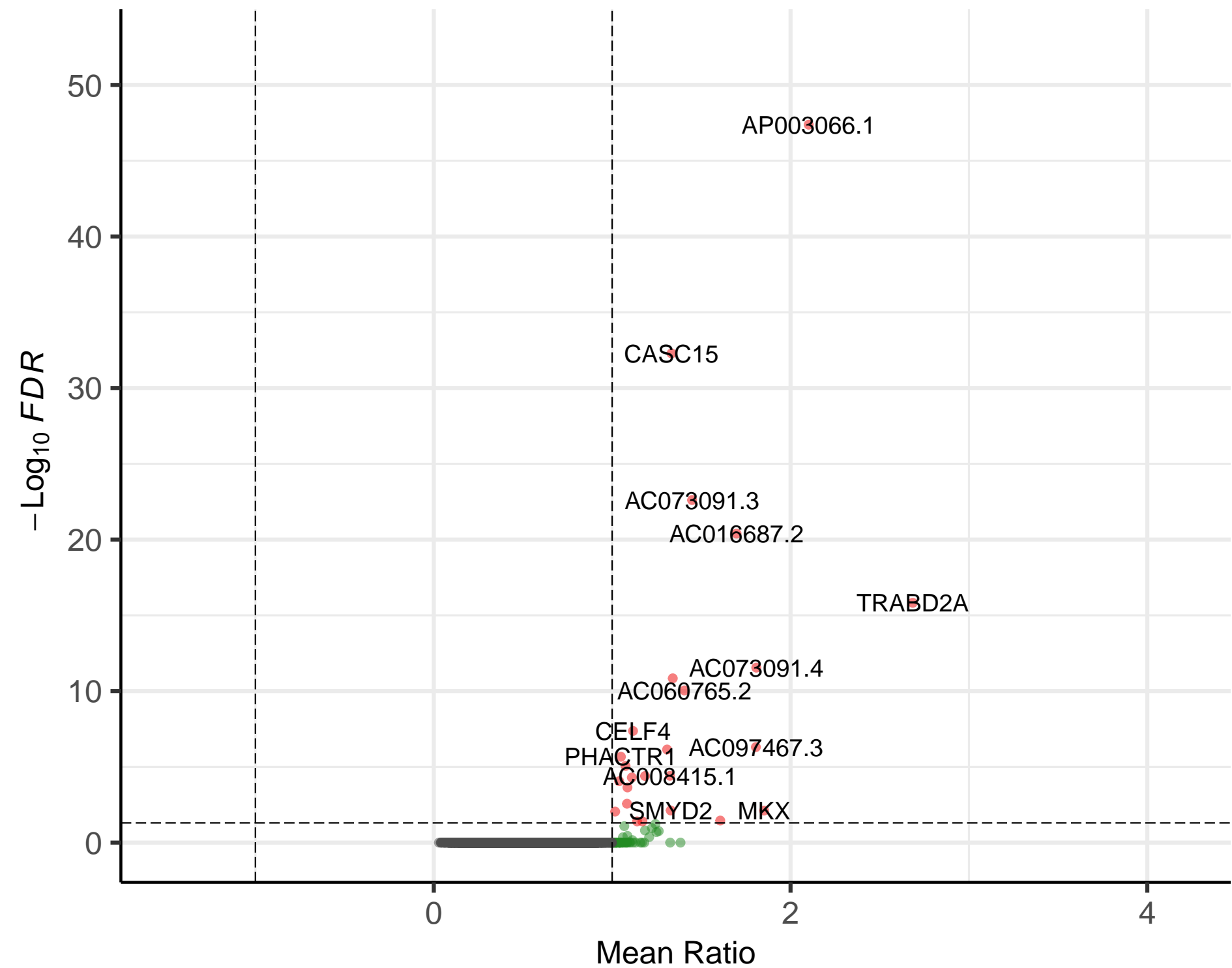


total = 5107 variables

Excit_04

n nuc=2171, n genes FDR<0.05=25

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

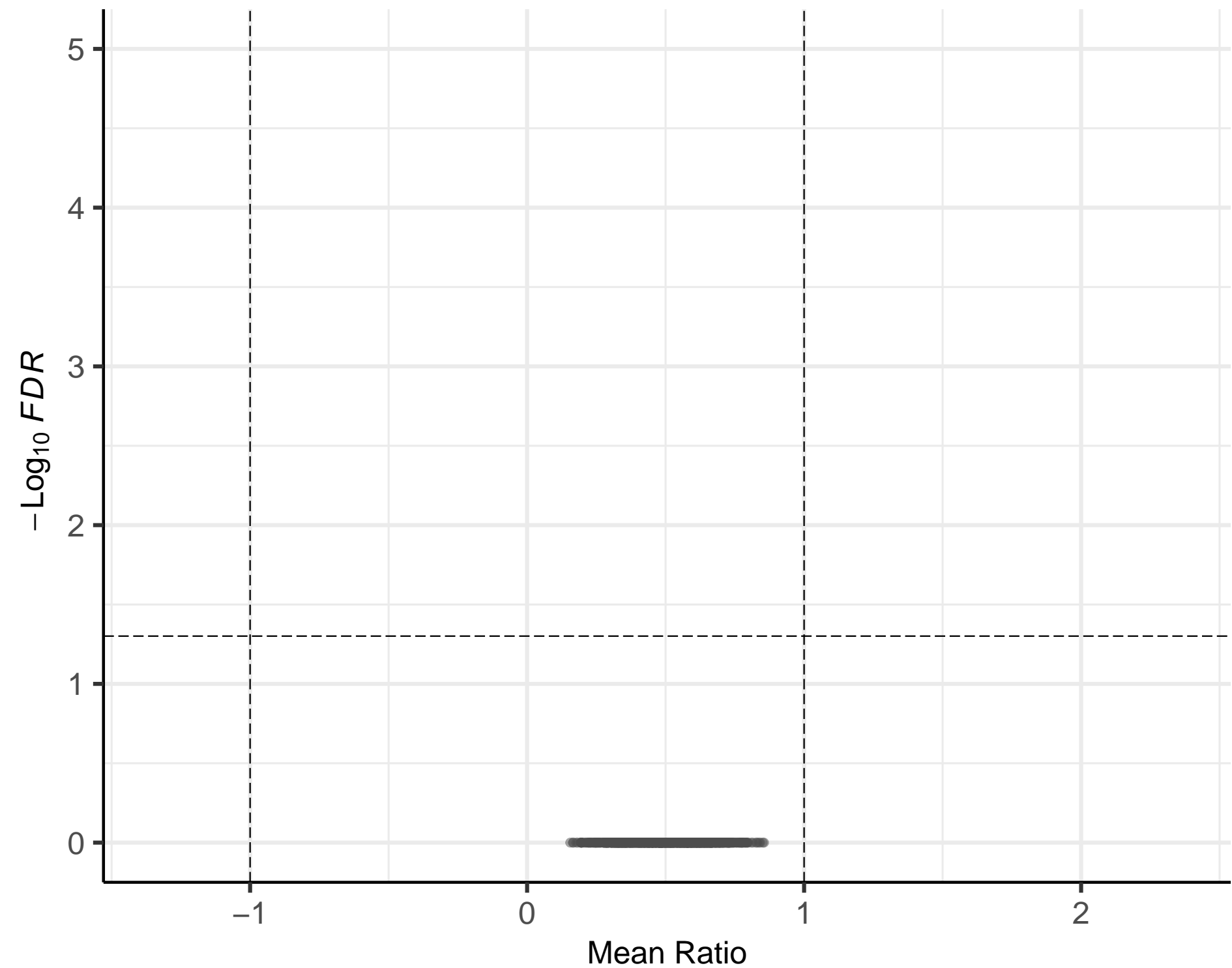


total = 7004 variables

Excit_05

n nuc=2532, n genes FDR<0.05=0

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

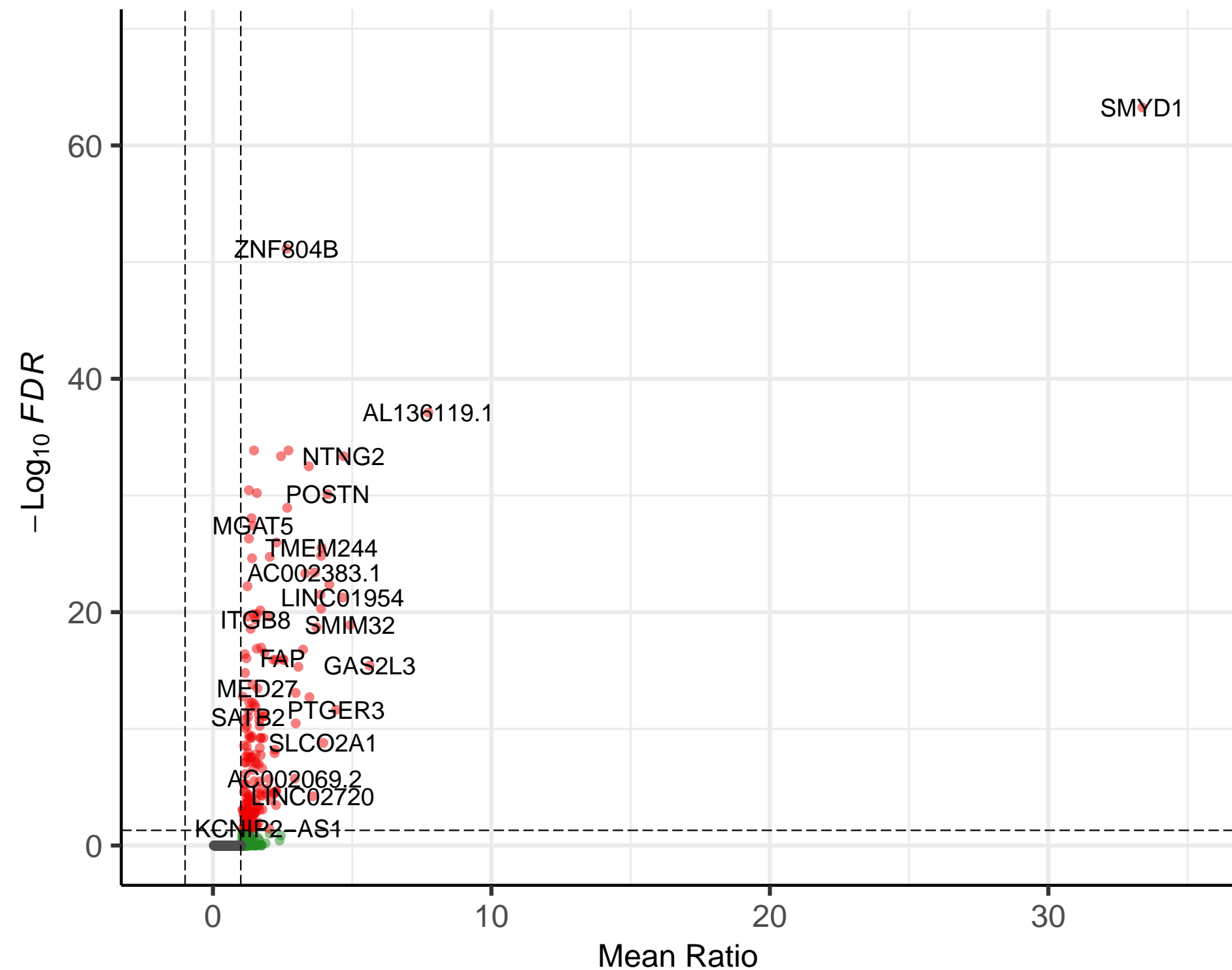


total = 1086 variables

Excit_06

n nuc=329, n genes FDR<0.05=182

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

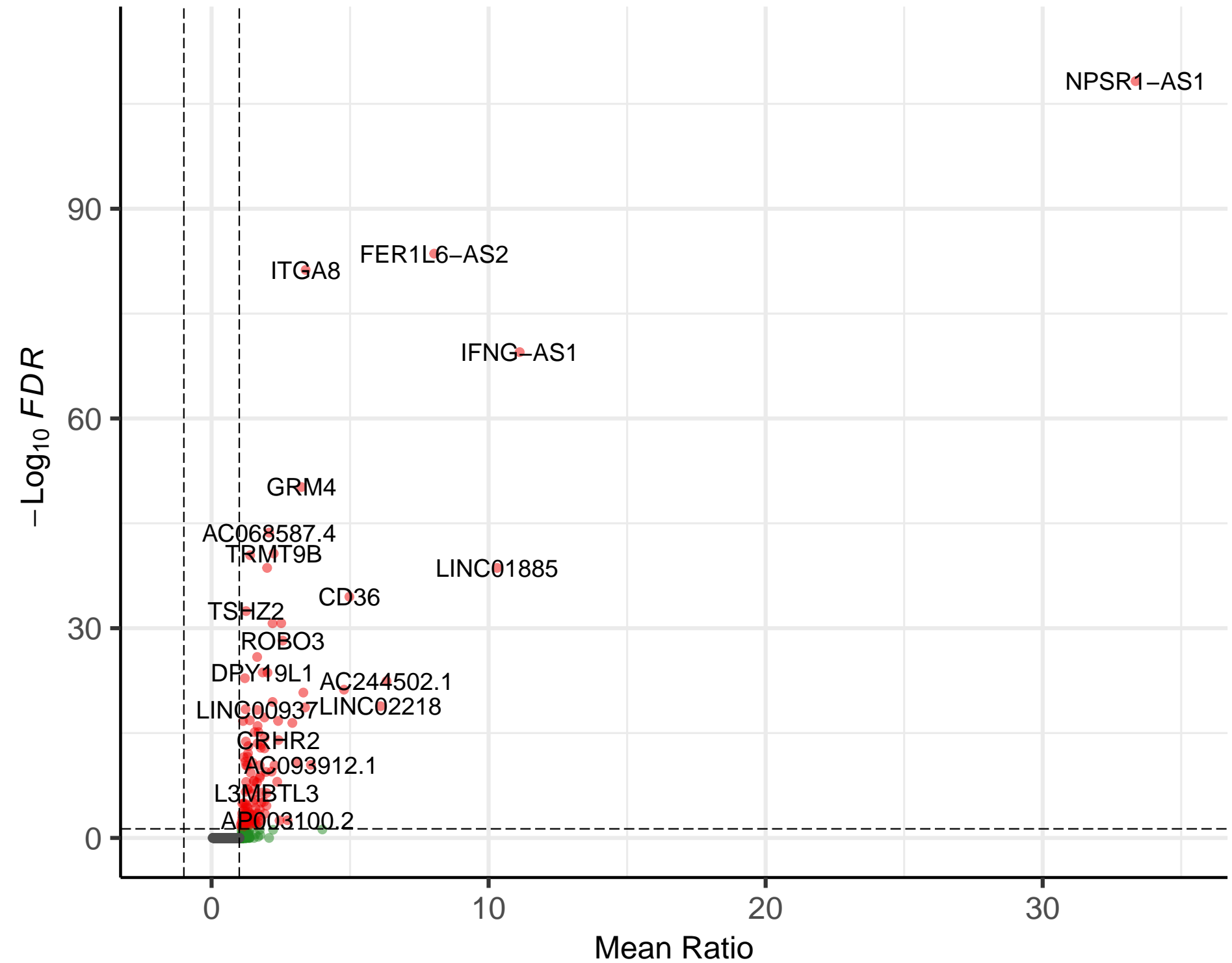


total = 6718 variables

Excit_07

n nuc=334, n genes FDR<0.05=129

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

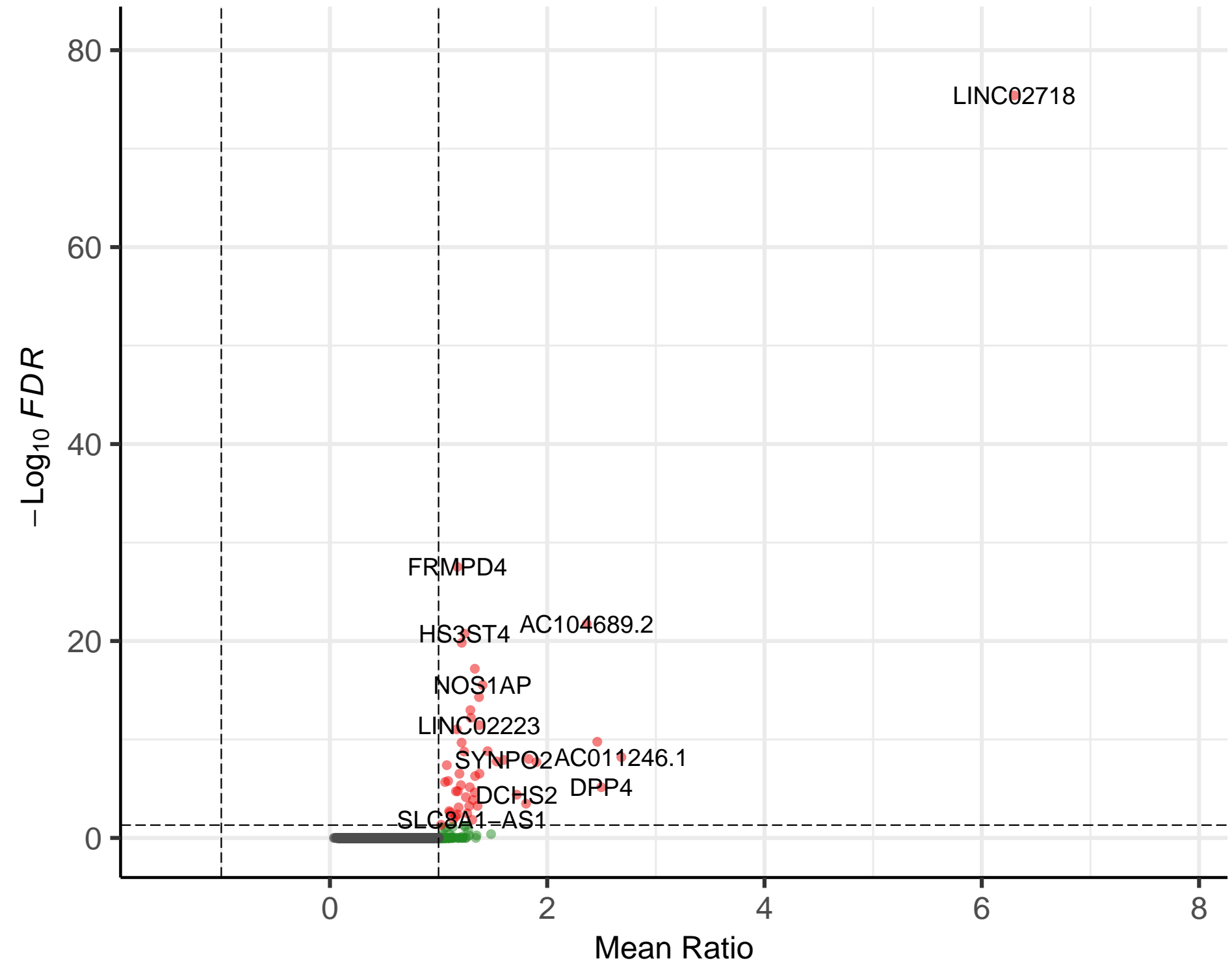


total = 5942 variables

Excit_08

n nuc=1463, n genes FDR<0.05=48

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

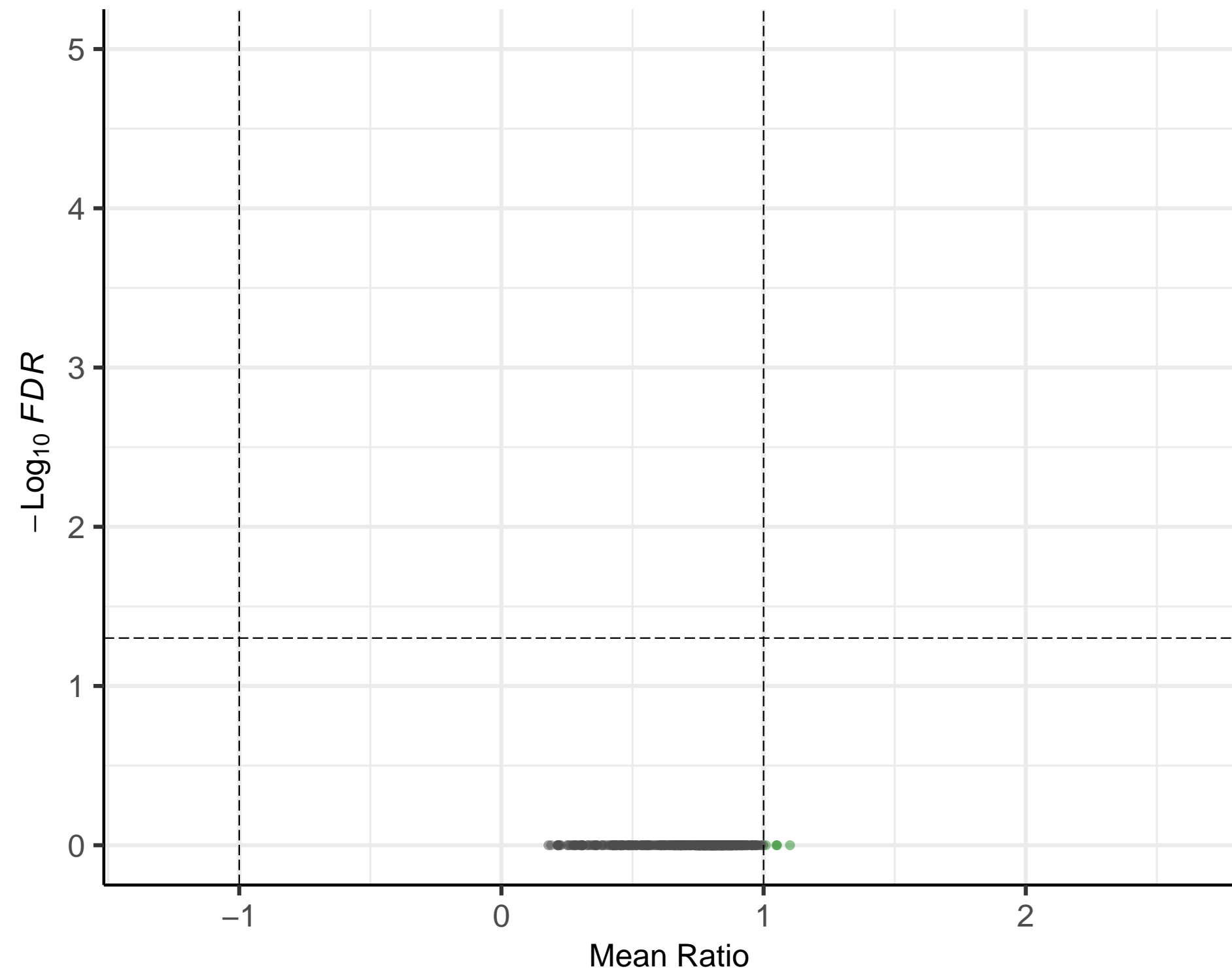


total = 6311 variables

Excit_09

n nuc=2561, n genes FDR<0.05=0

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

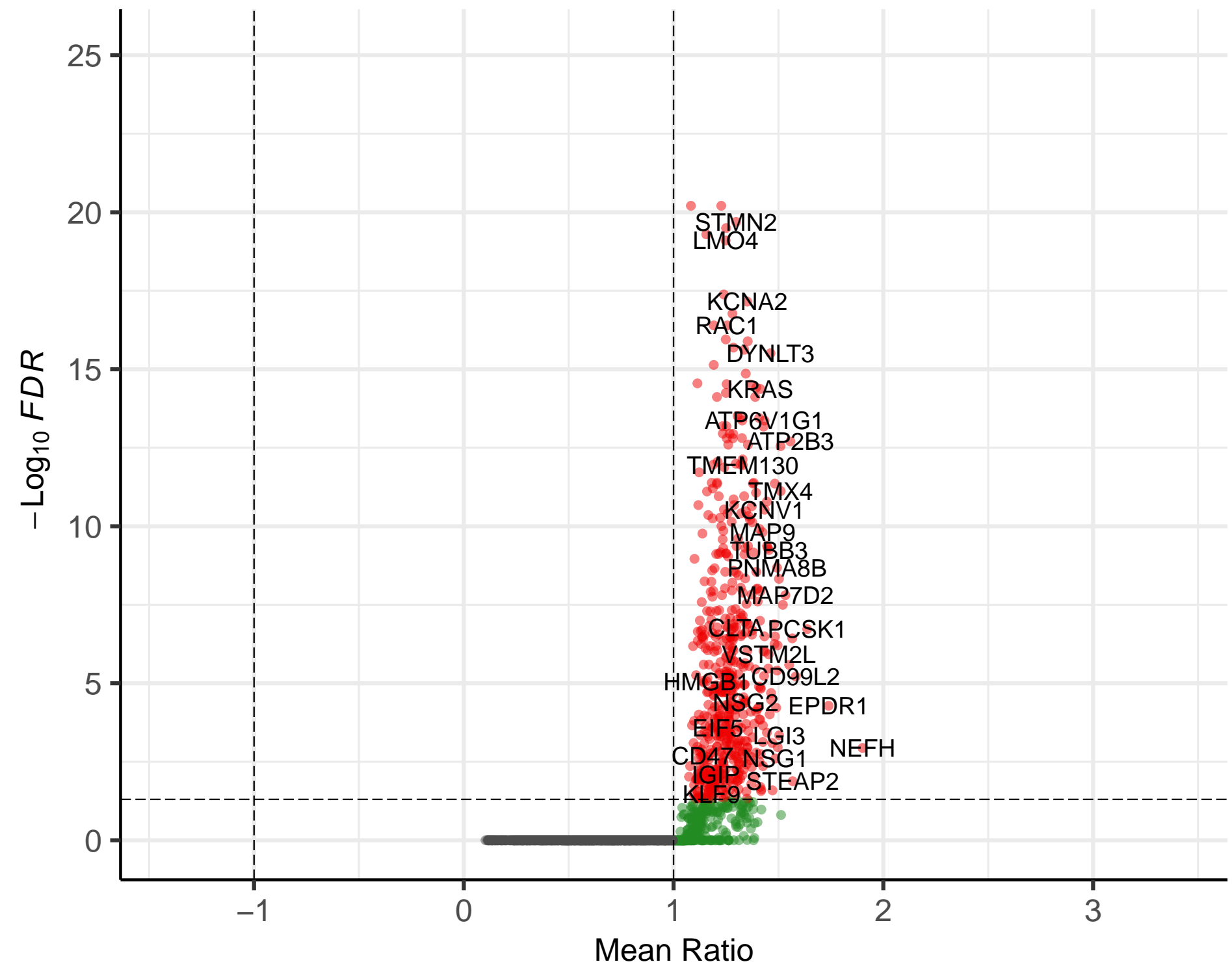


total = 453 variables

Excit_10

n nuc=1079, n genes FDR<0.05=550

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

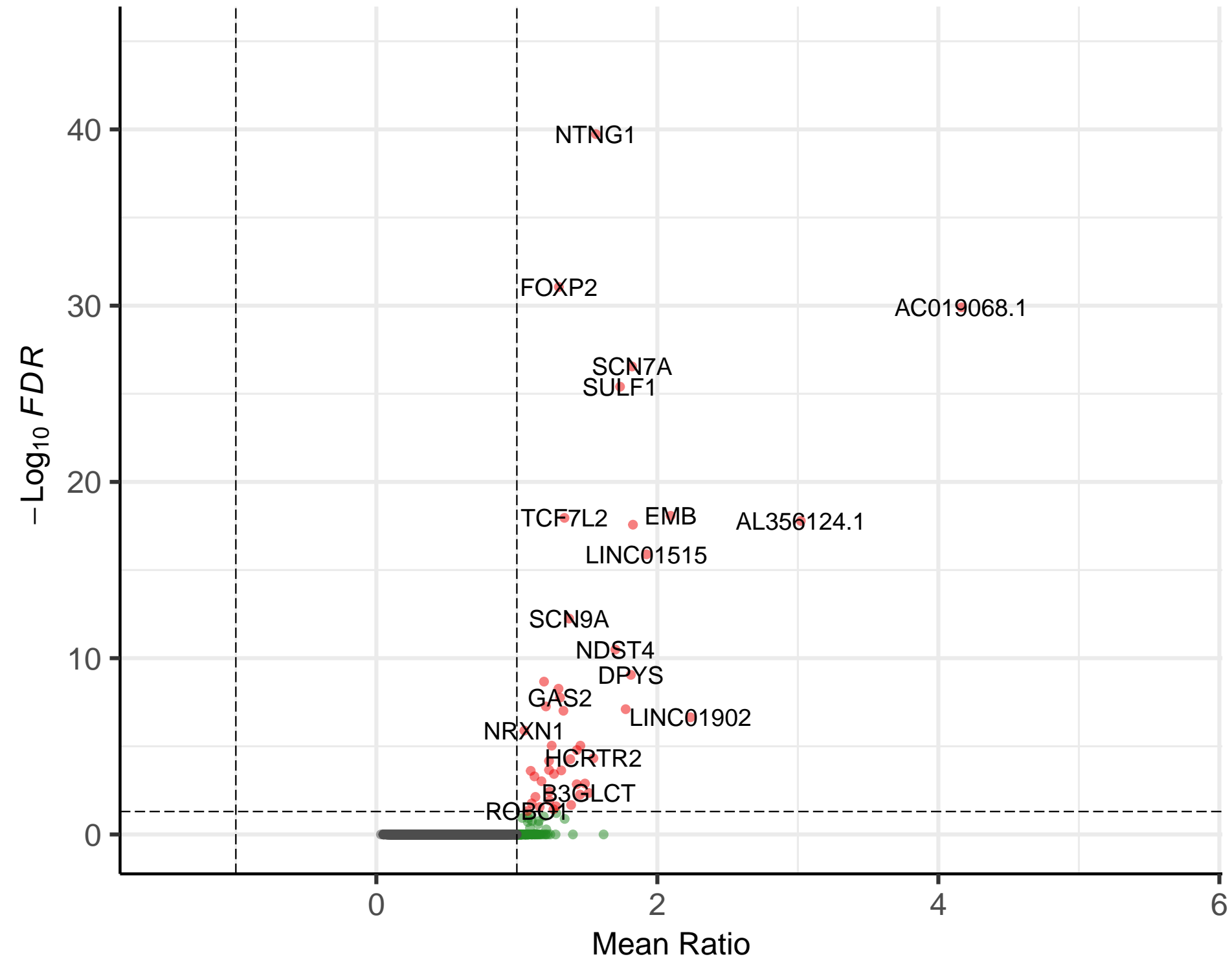


total = 1618 variables

Excit_11

n nuc=482, n genes FDR<0.05=46

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

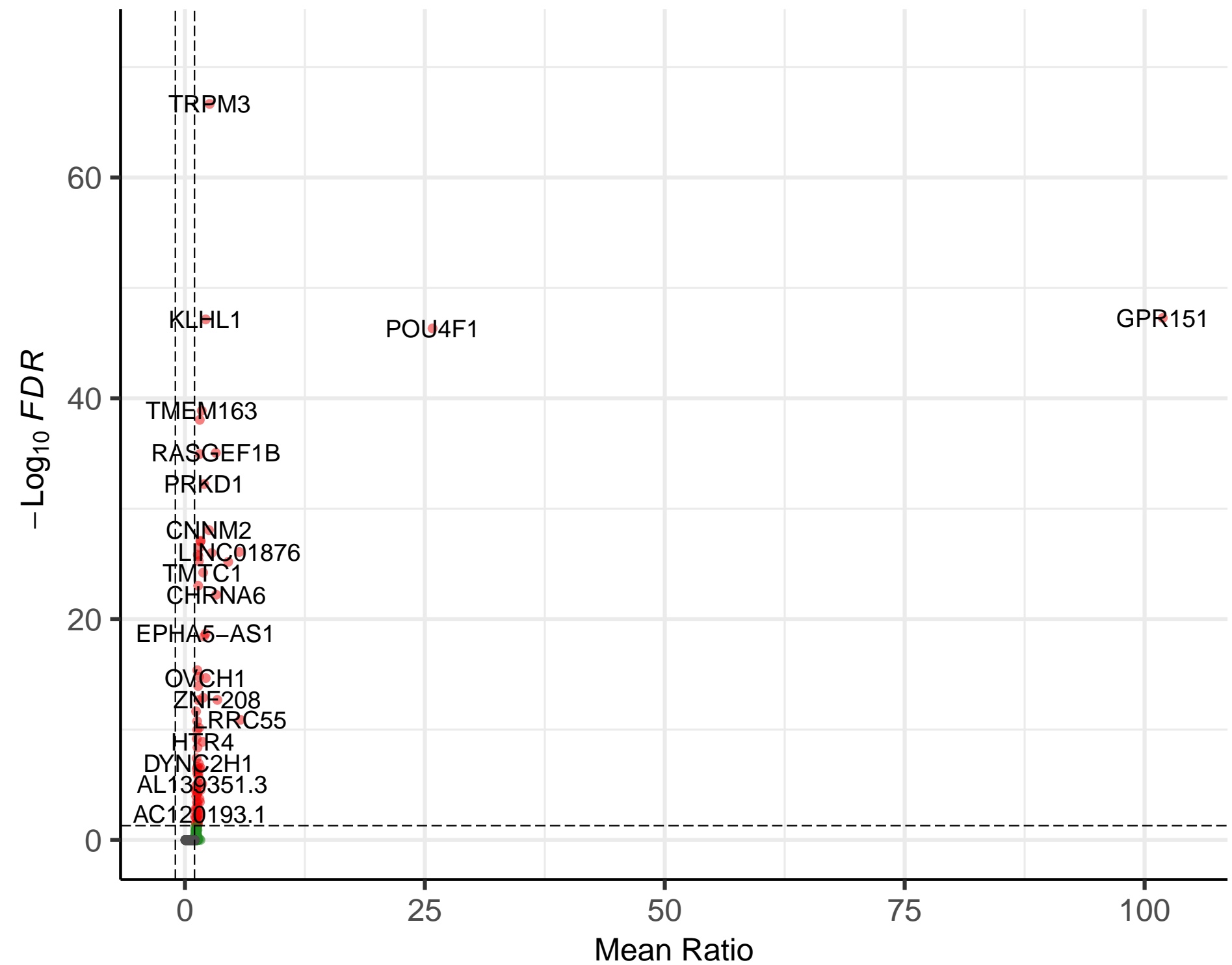


total = 4577 variables

Excit_12

n nuc=420, n genes FDR<0.05=77

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

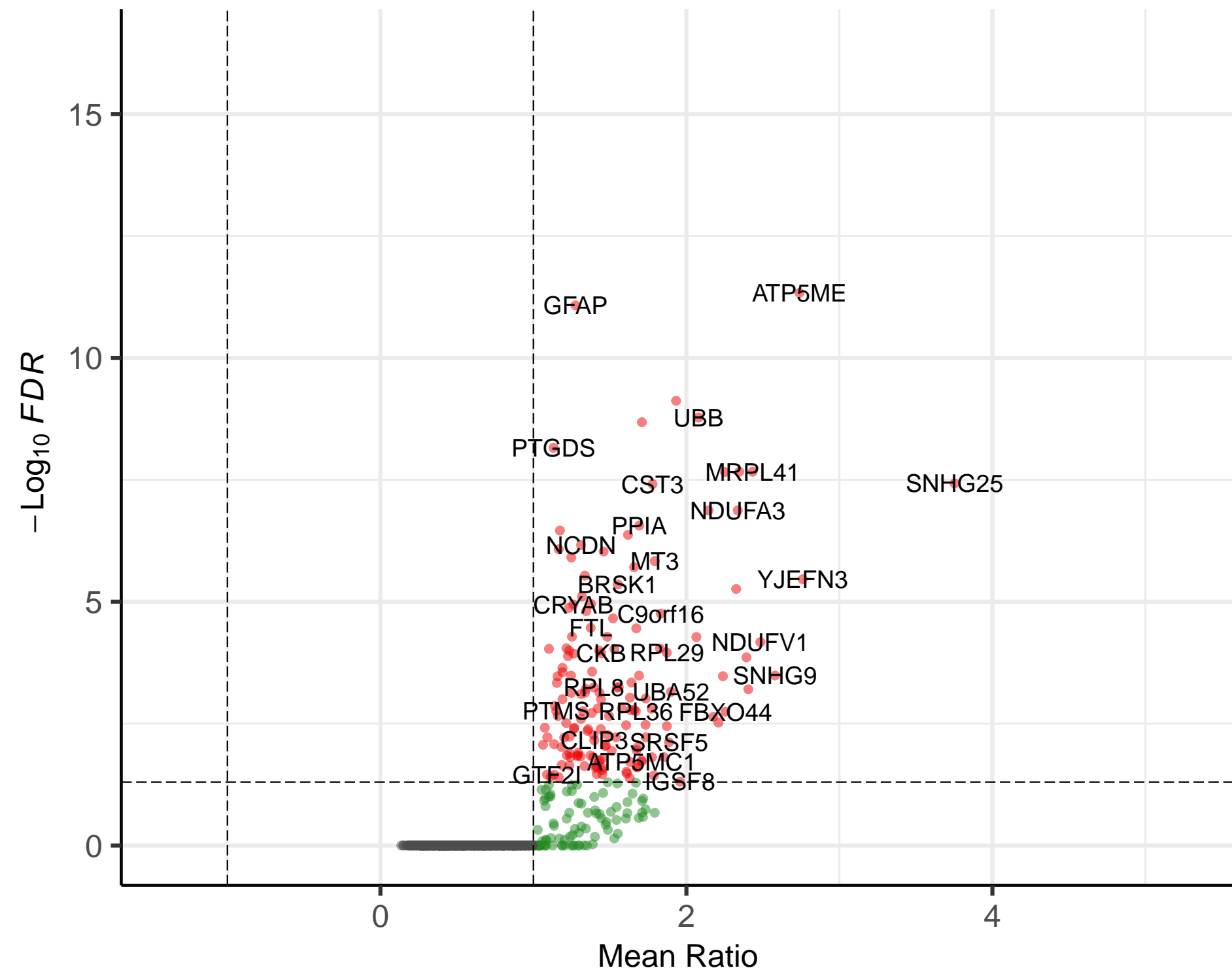


total = 4732 variables

Excit_13

n nuc=1567, n genes FDR<0.05=155

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

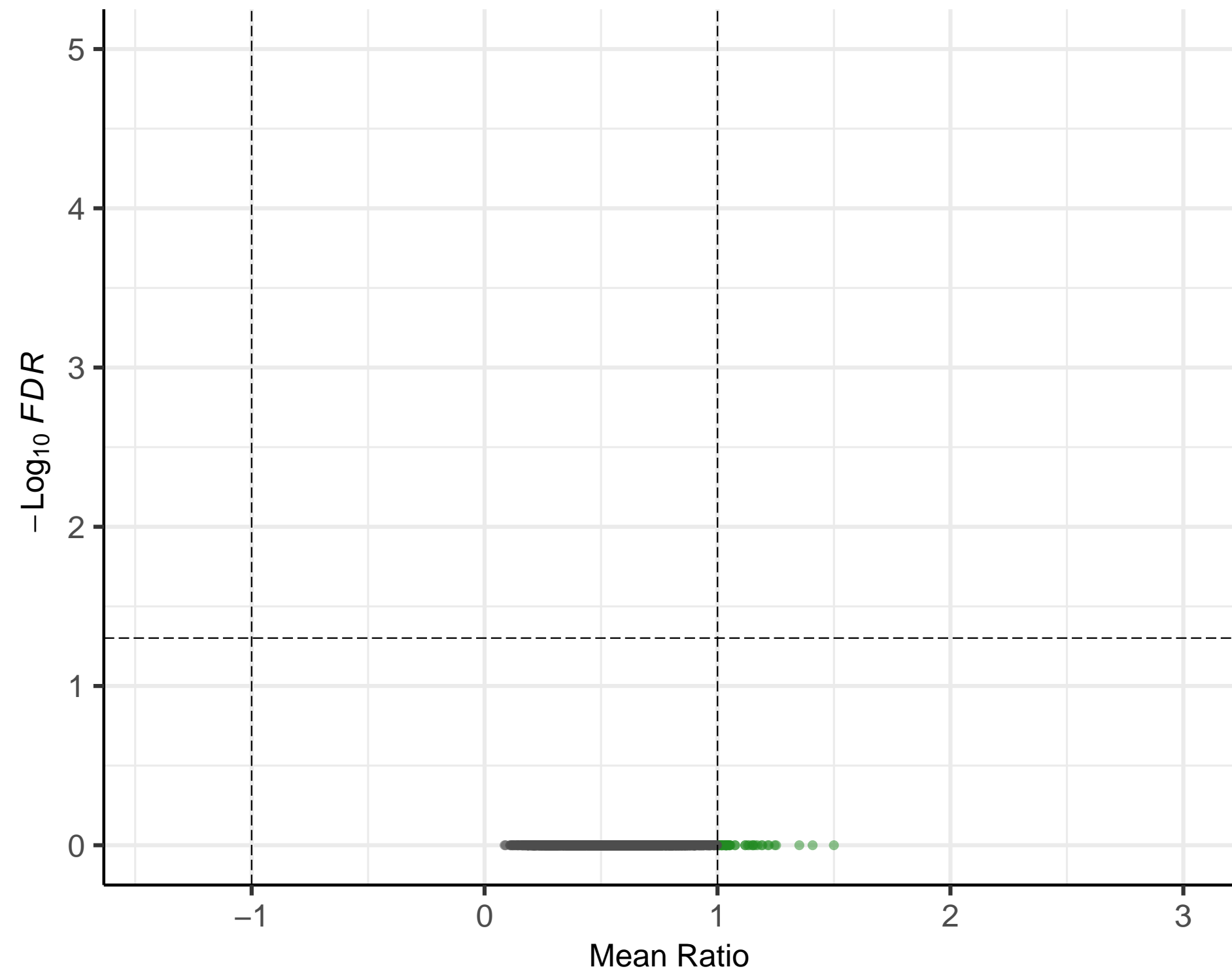


total = 839 variables

Excit_14

n nuc=82, n genes FDR<0.05=0

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

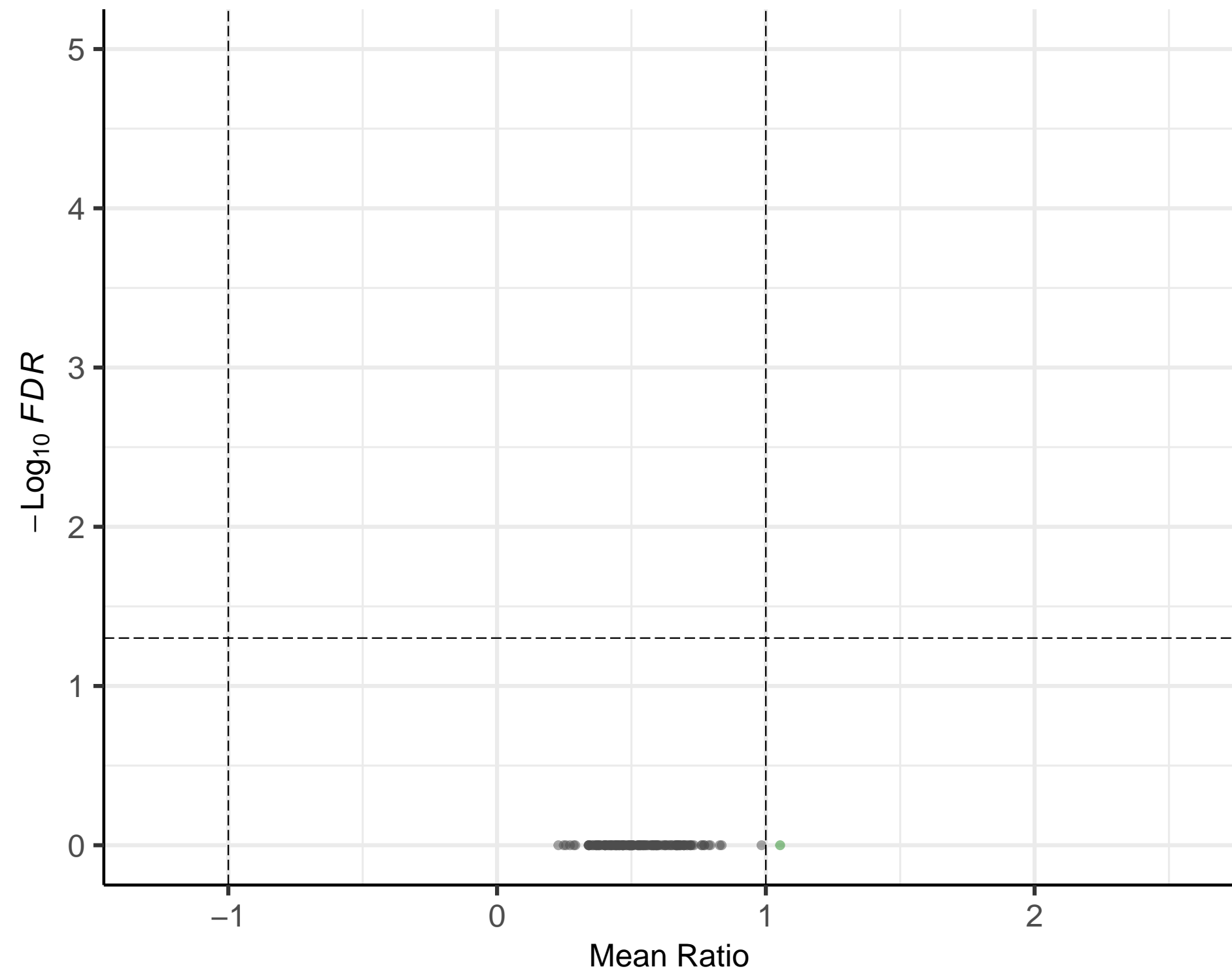


total = 2566 variables

Excit_15

n nuc=66, n genes FDR<0.05=0

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

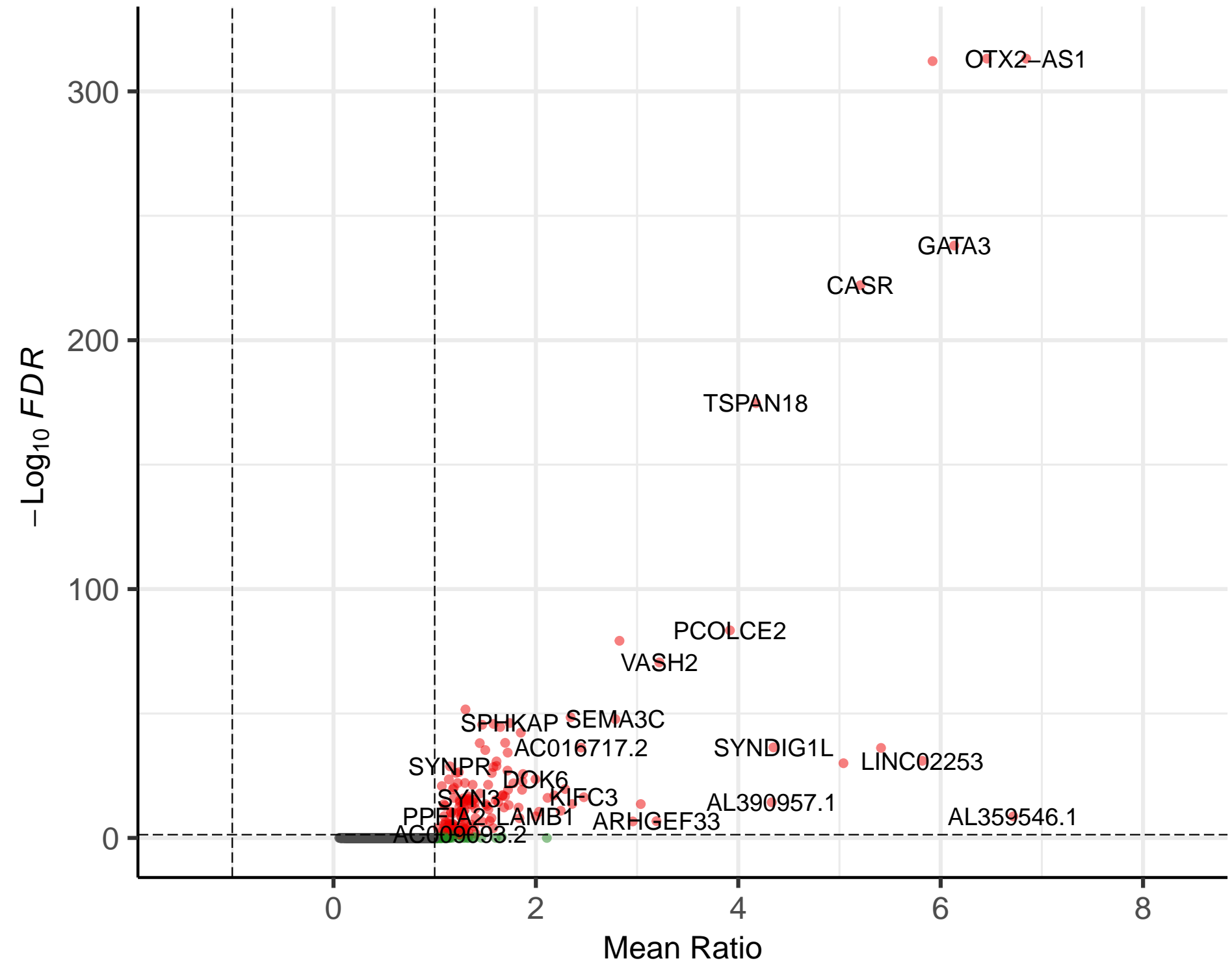


total = 148 variables

Inhib_01

n nuc=5366, n genes FDR<0.05=153

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

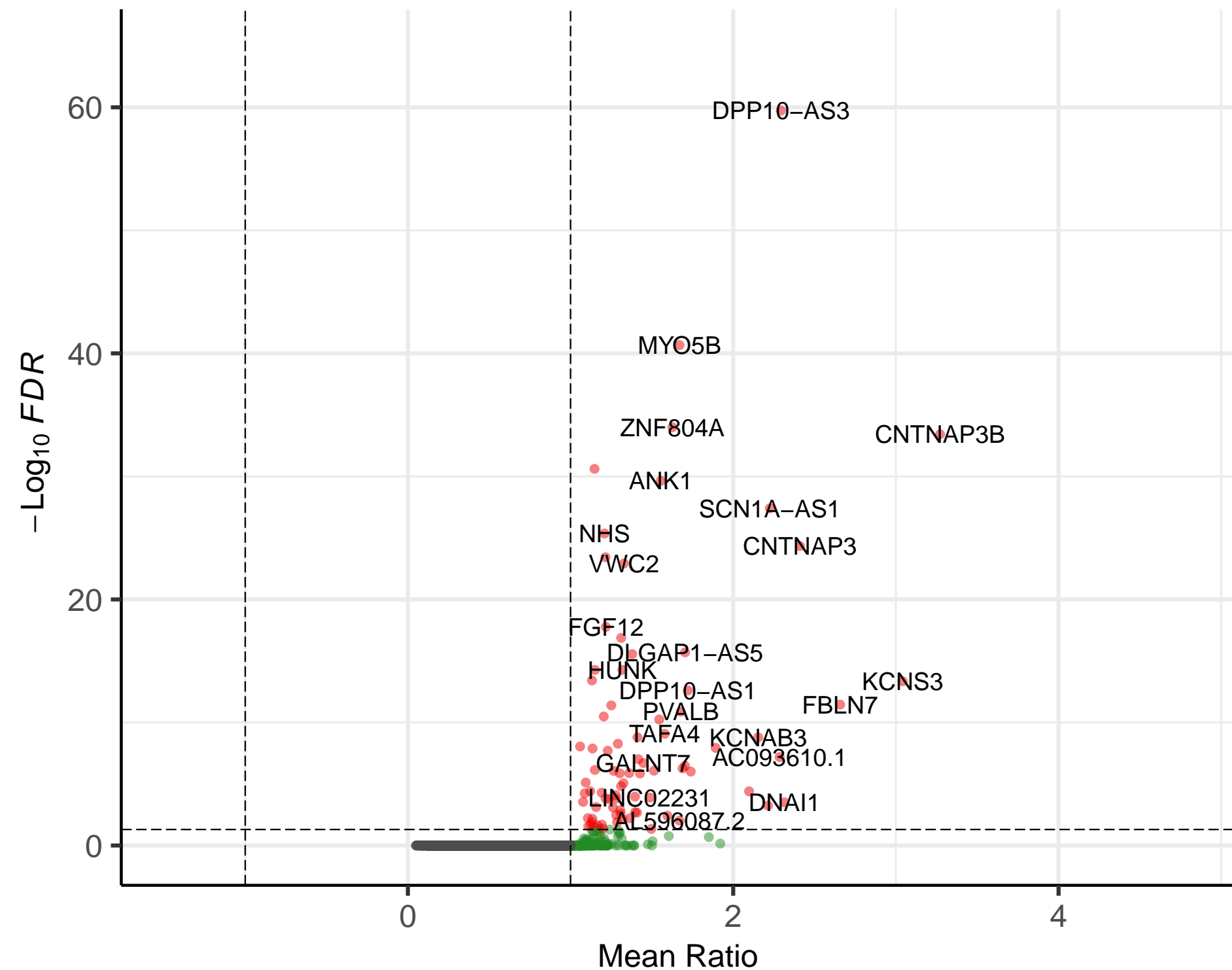


total = 3569 variables

Inhib_02

n nuc=1267, n genes FDR<0.05=84

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

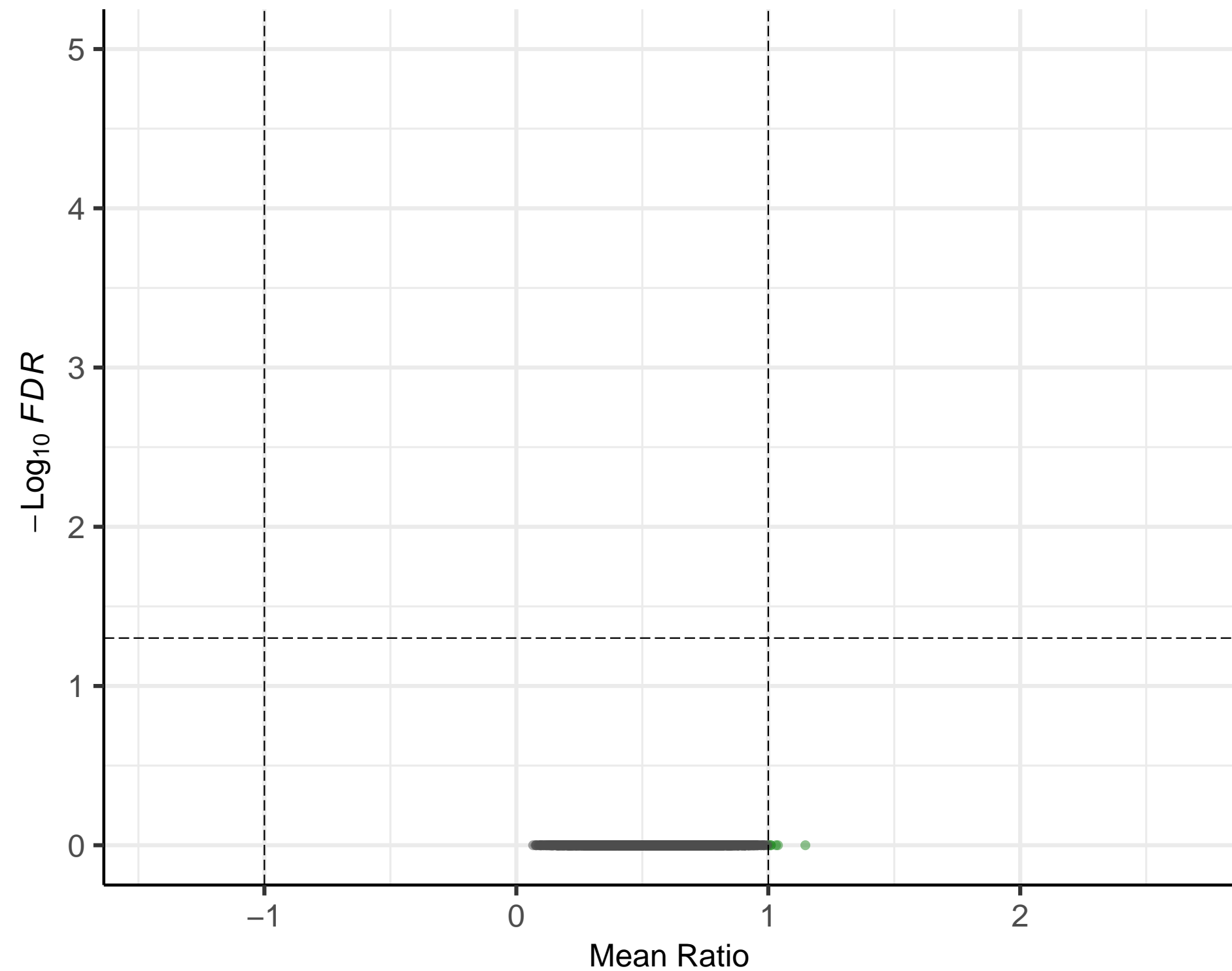


total = 5717 variables

Inhib_03

n nuc=1310, n genes FDR<0.05=0

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

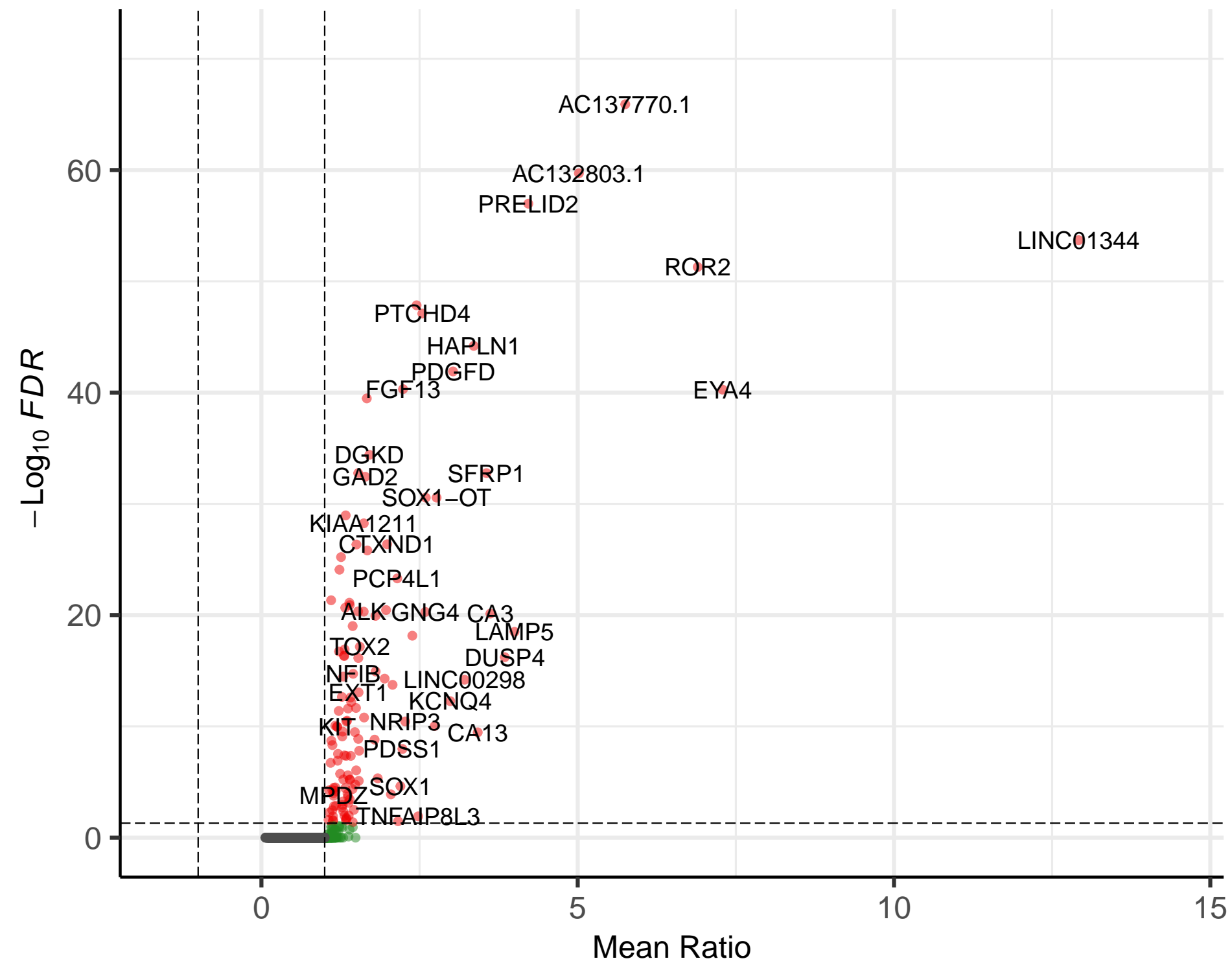


total = 2969 variables

Inhib_04

n nuc=565, n genes FDR<0.05=129

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

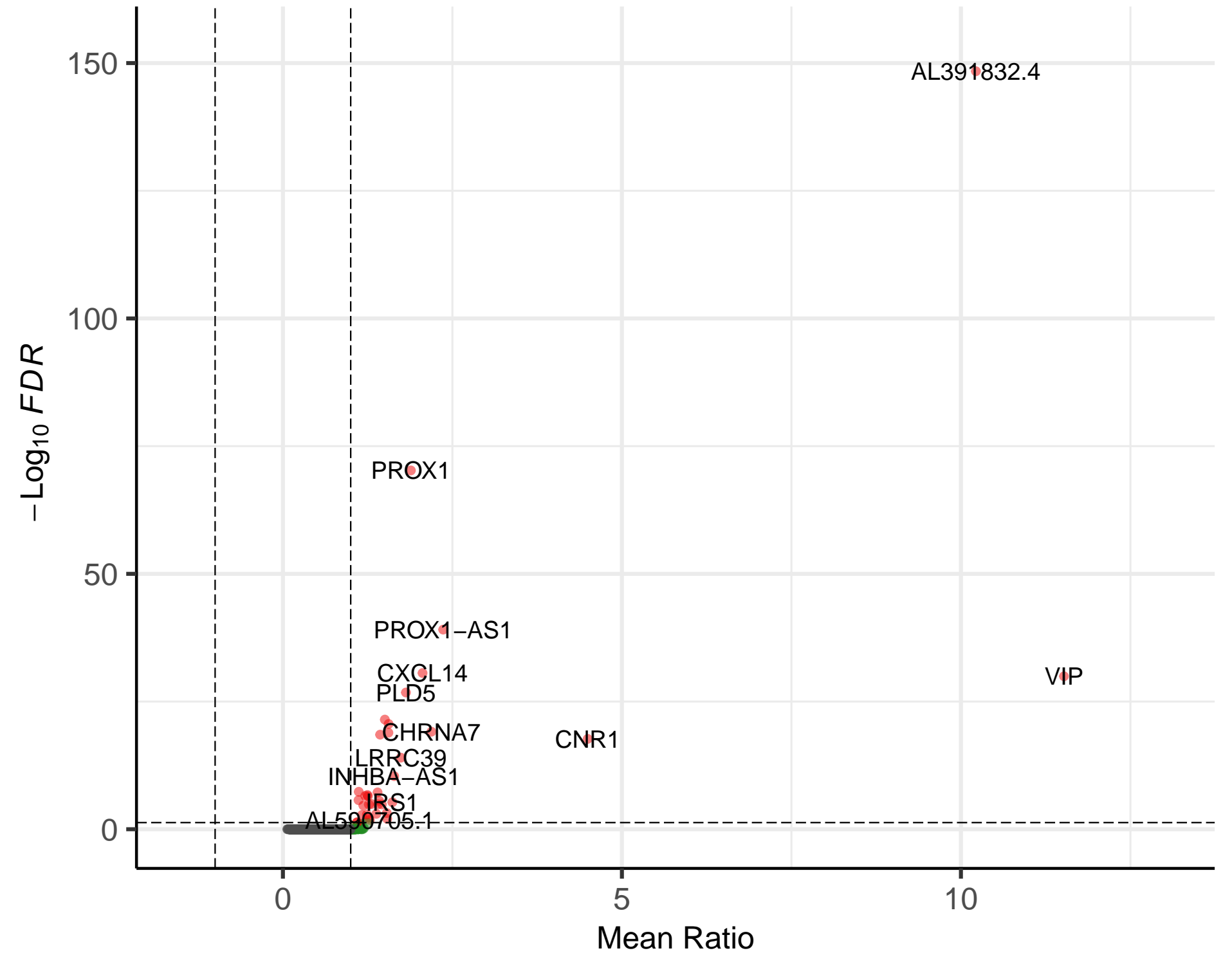


total = 4652 variables

Inhib_05

n nuc=1192, n genes FDR<0.05=36

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

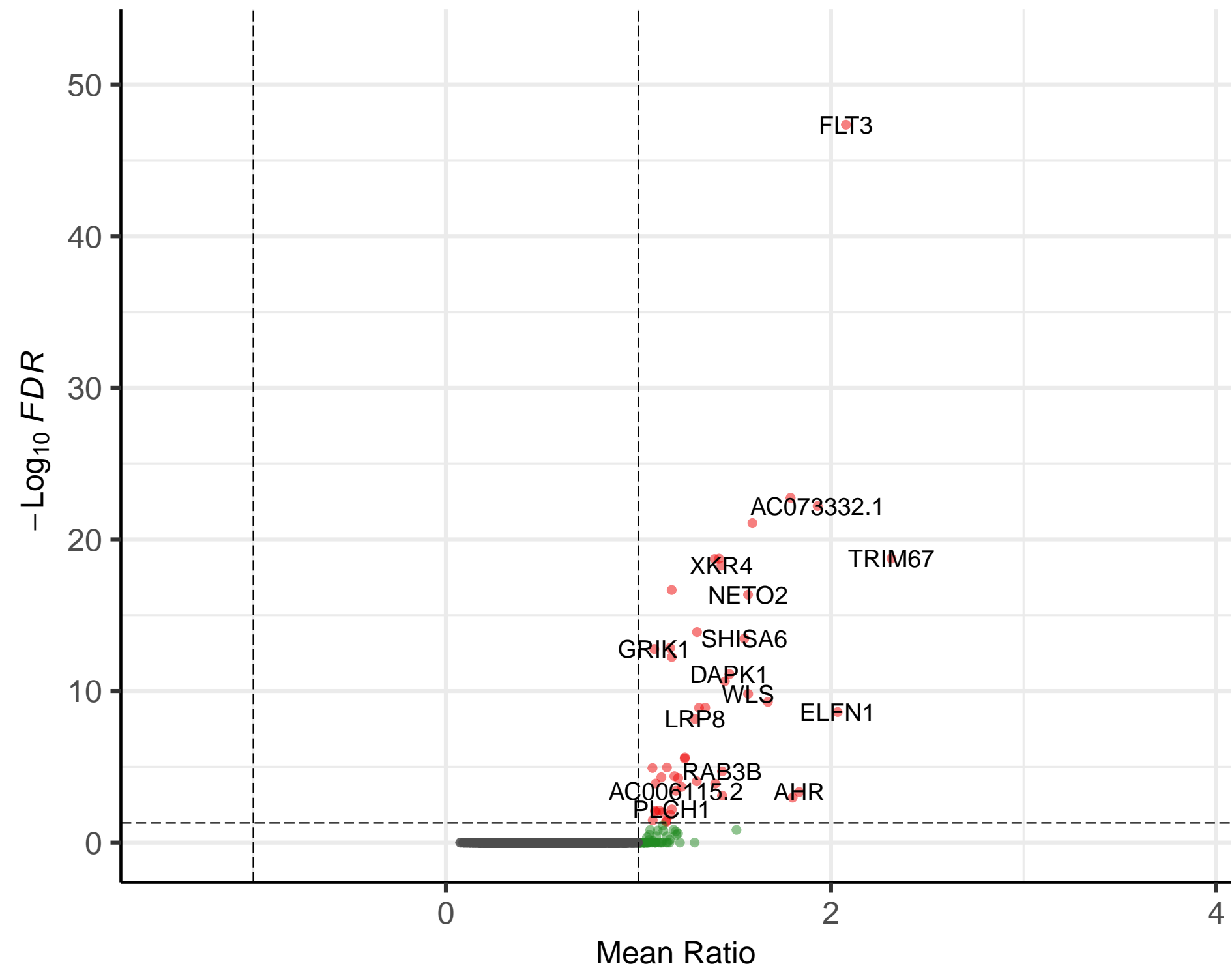


total = 3906 variables

Inhib_06

n nuc=1367, n genes FDR<0.05=48

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

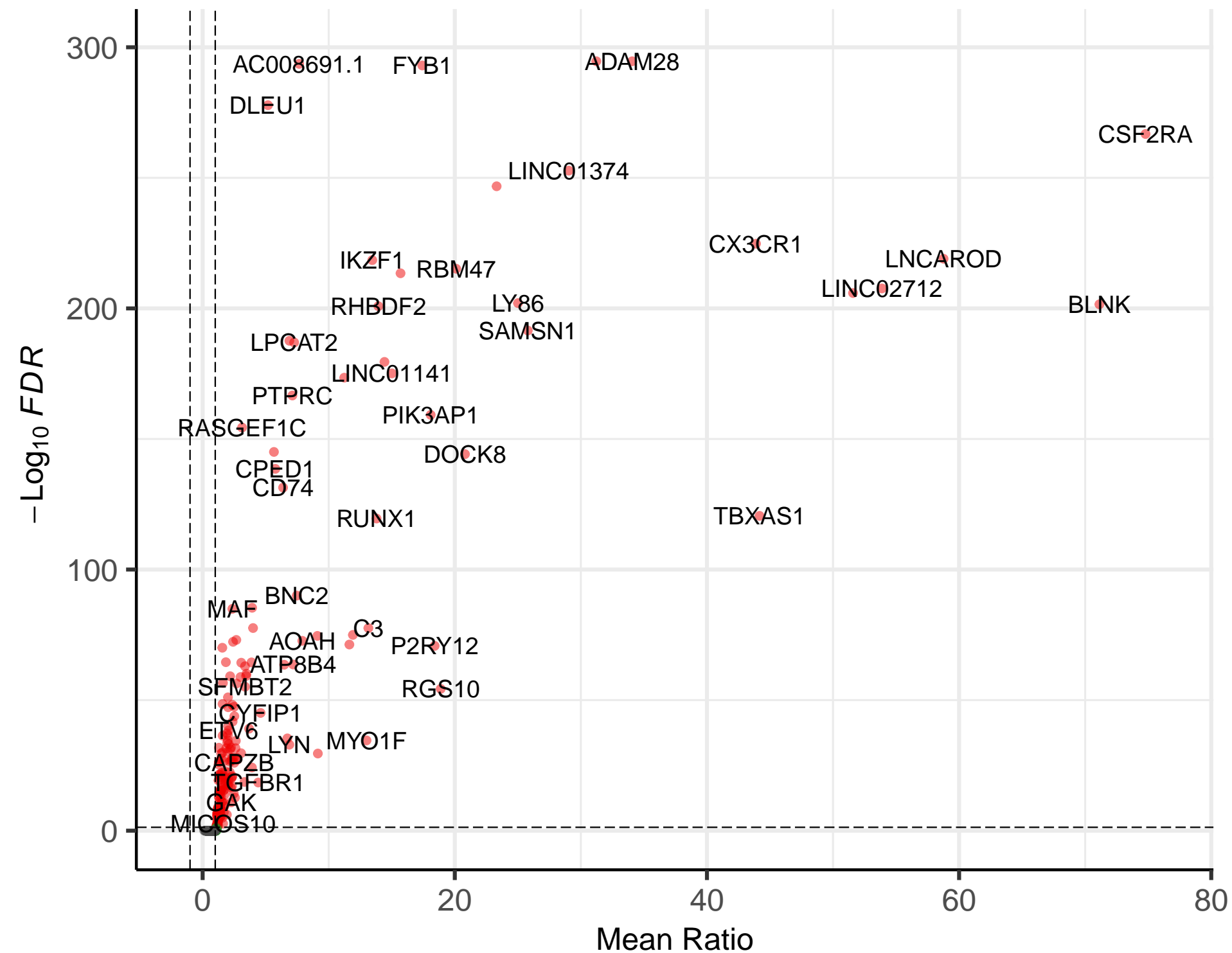


total = 3518 variables

Micro

n nuc=1601, n genes FDR<0.05=196

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

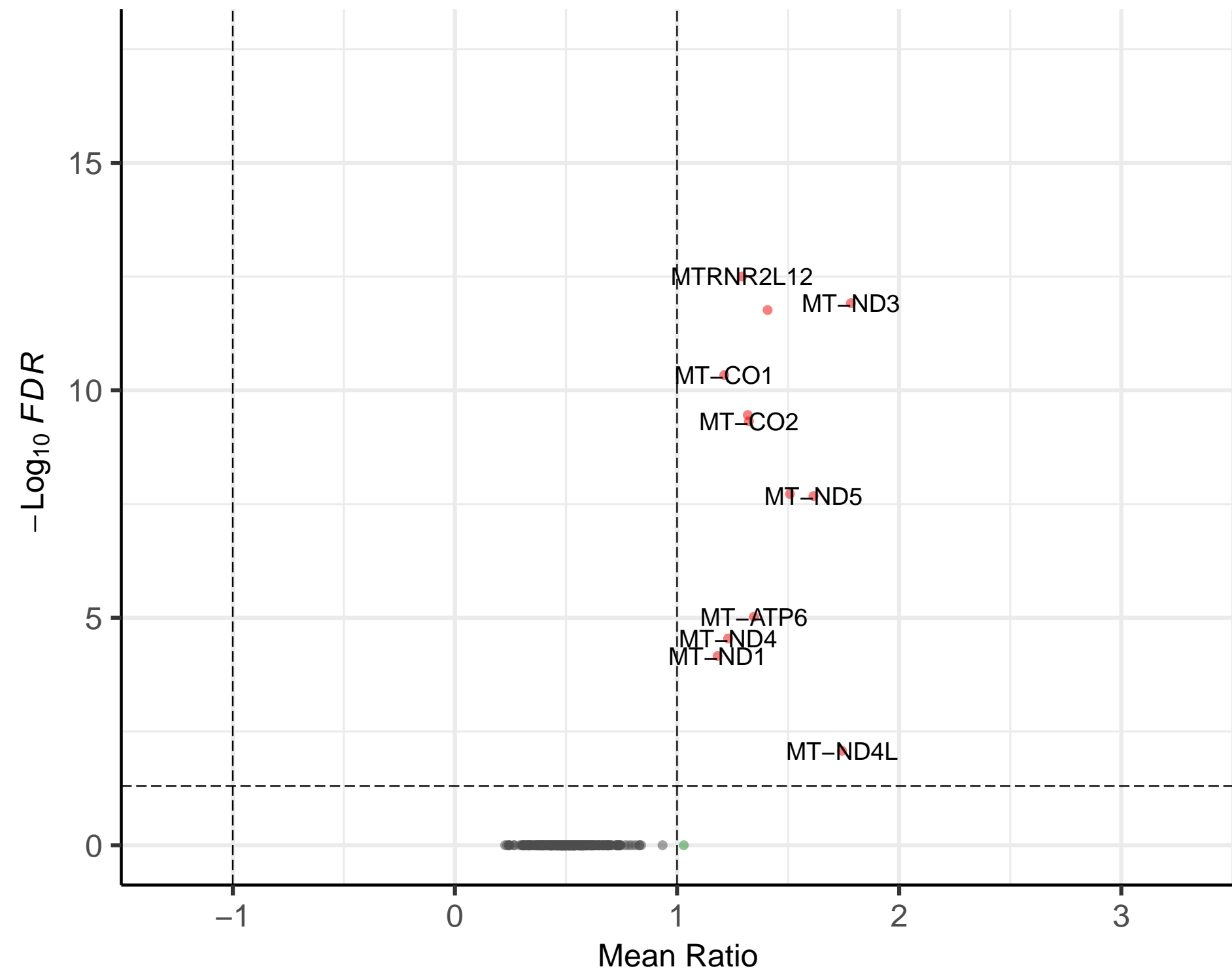


total = 473 variables

Oligo_01

n nuc=23025, n genes FDR<0.05=12

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

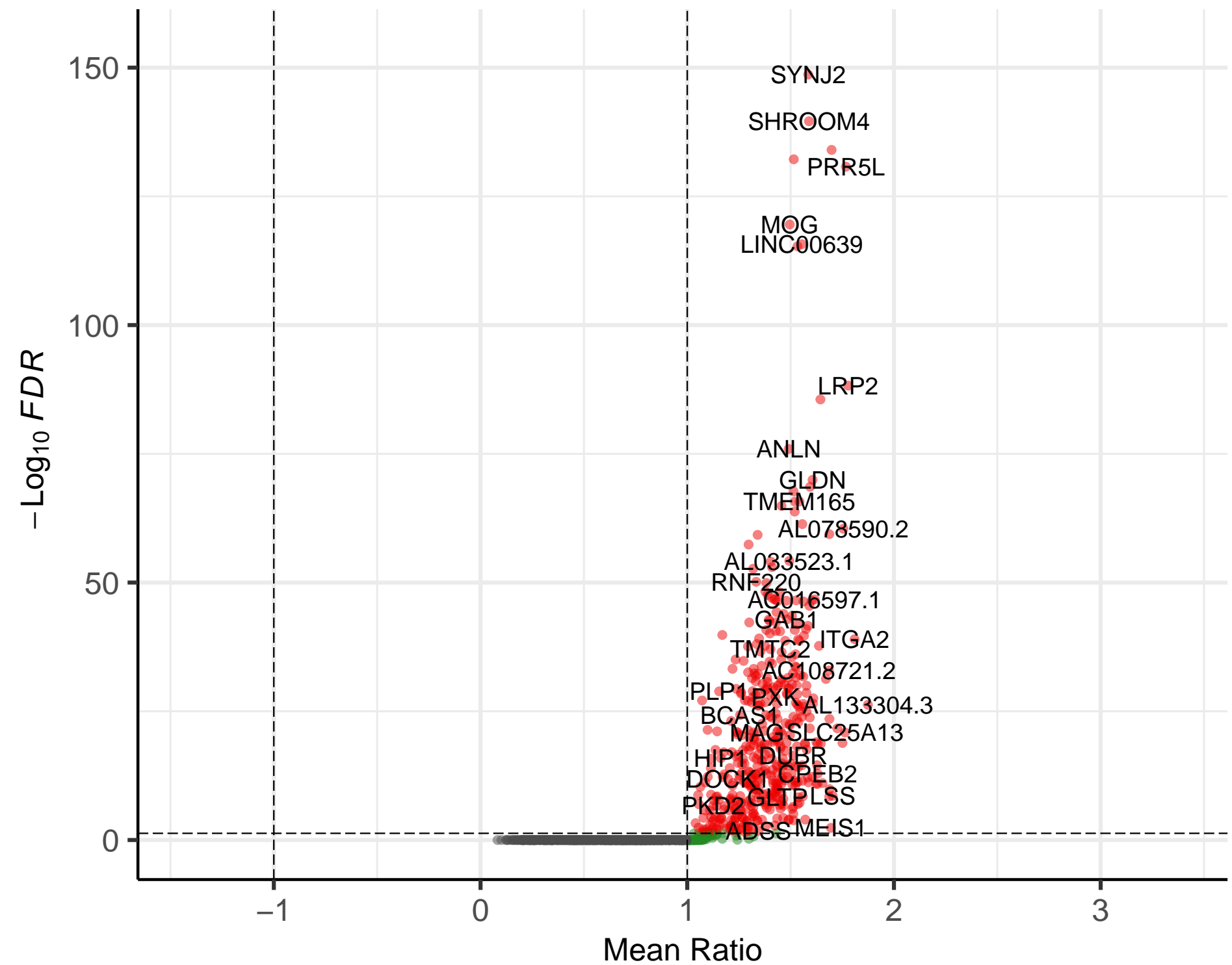


total = 284 variables

Oligo_02

n nuc=4732, n genes FDR<0.05=448

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

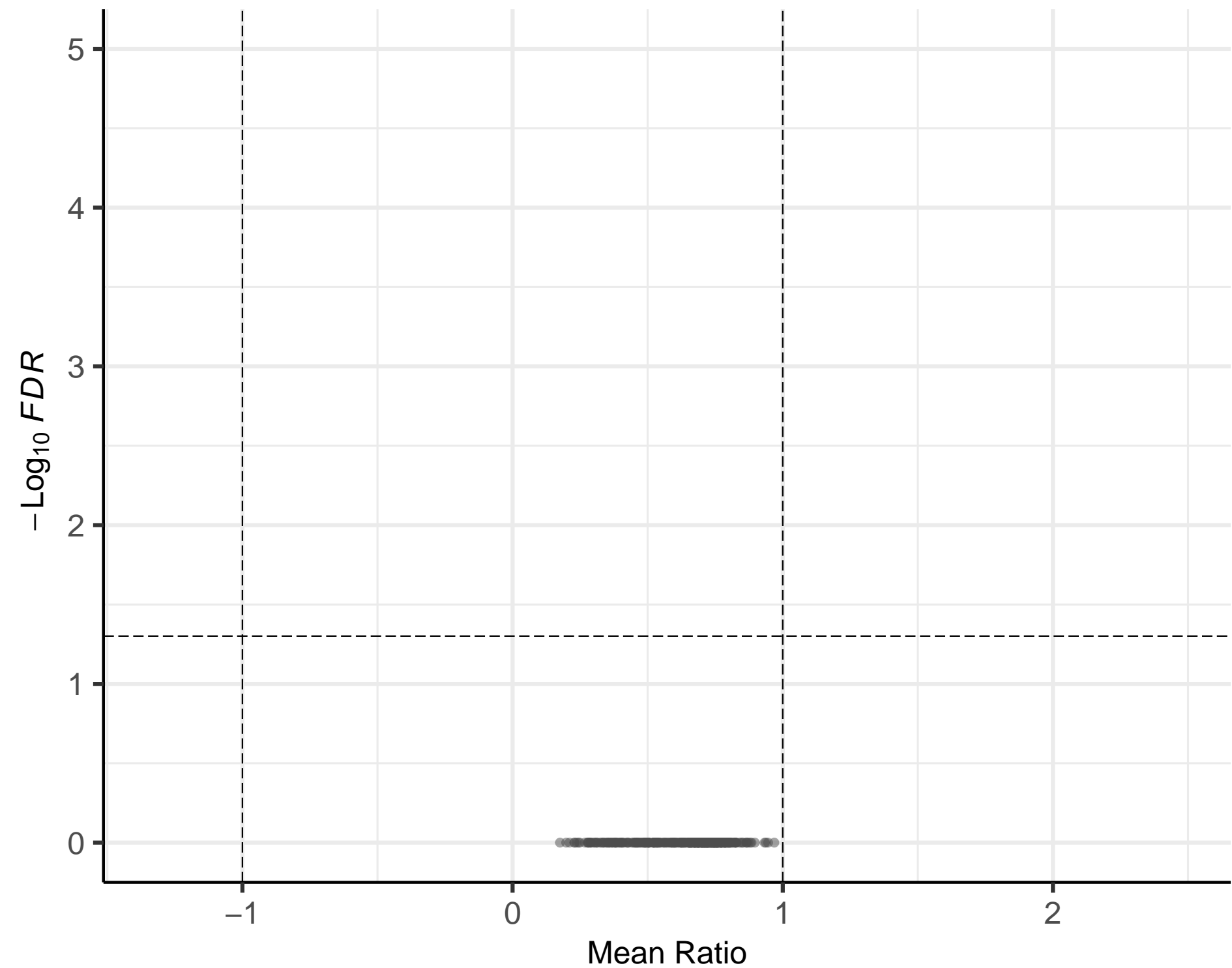


total = 1213 variables

Oligo_03

n nuc=4294, n genes FDR<0.05=0

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC

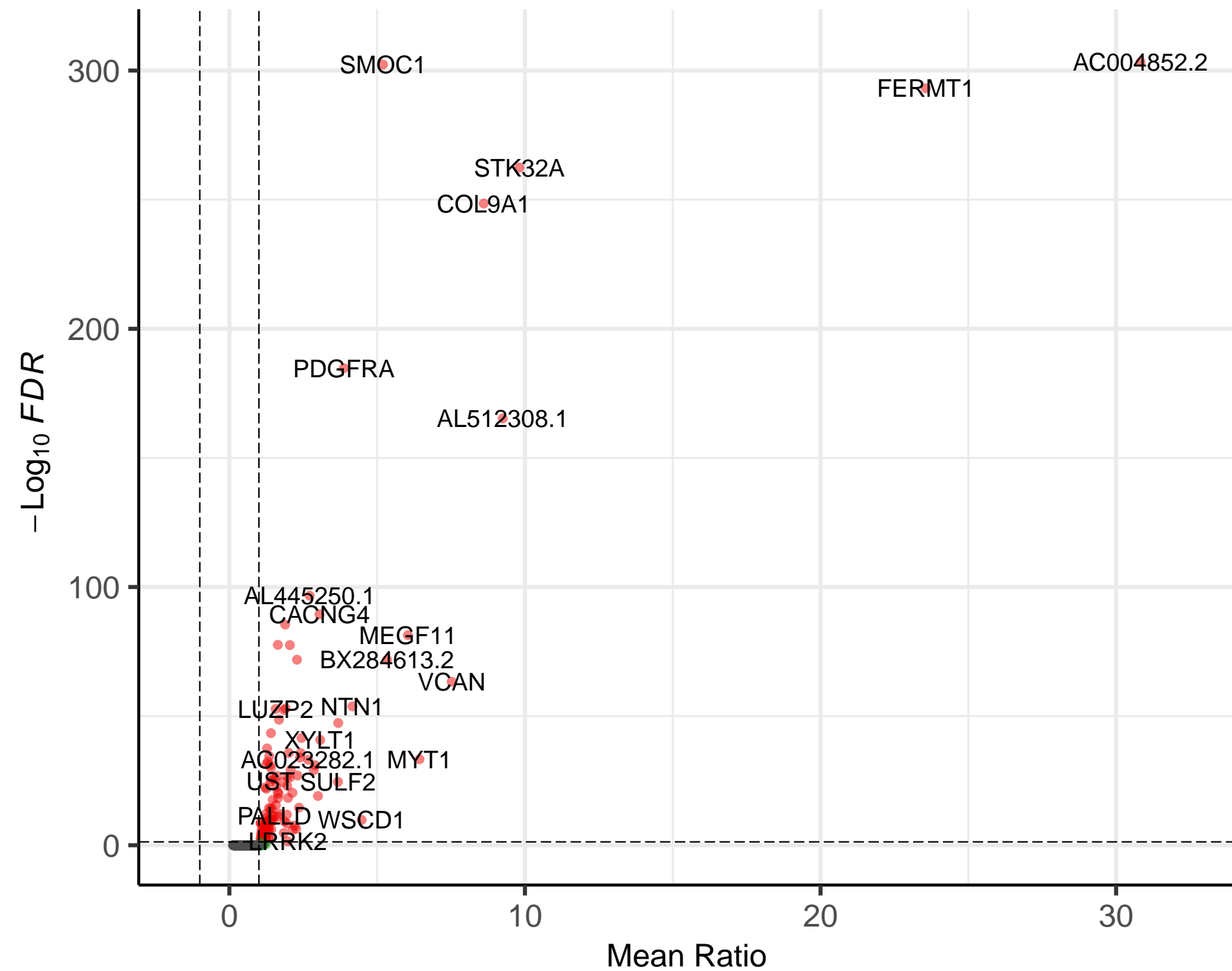


total = 363 variables

OPC

n nuc=1940, n genes FDR<0.05=125

● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



total = 1333 variables