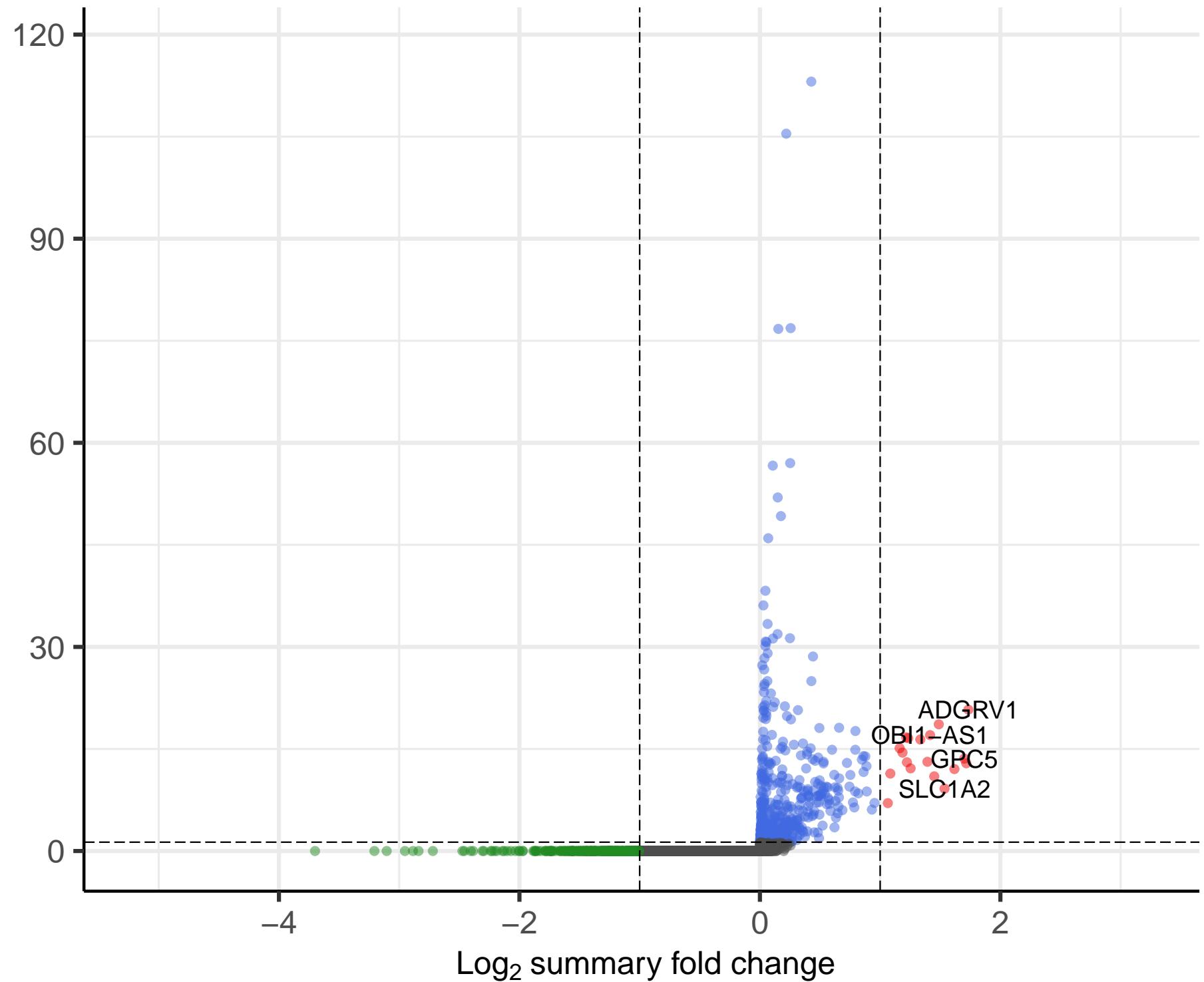


Astro

n nuc=3979, n genes FDR<0.05=482

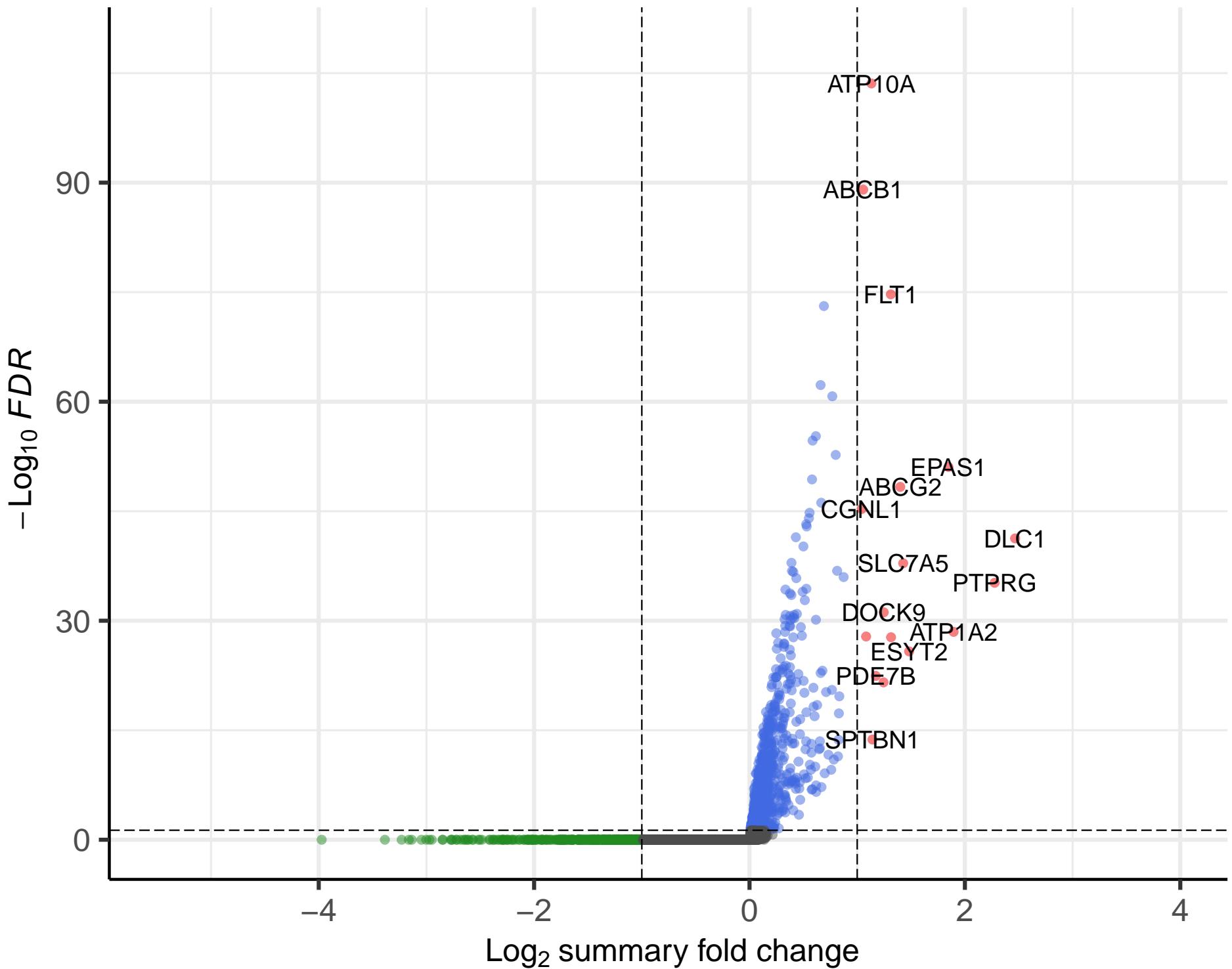
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Endo.Mural_01

n nuc=0, n genes FDR<0.05=914

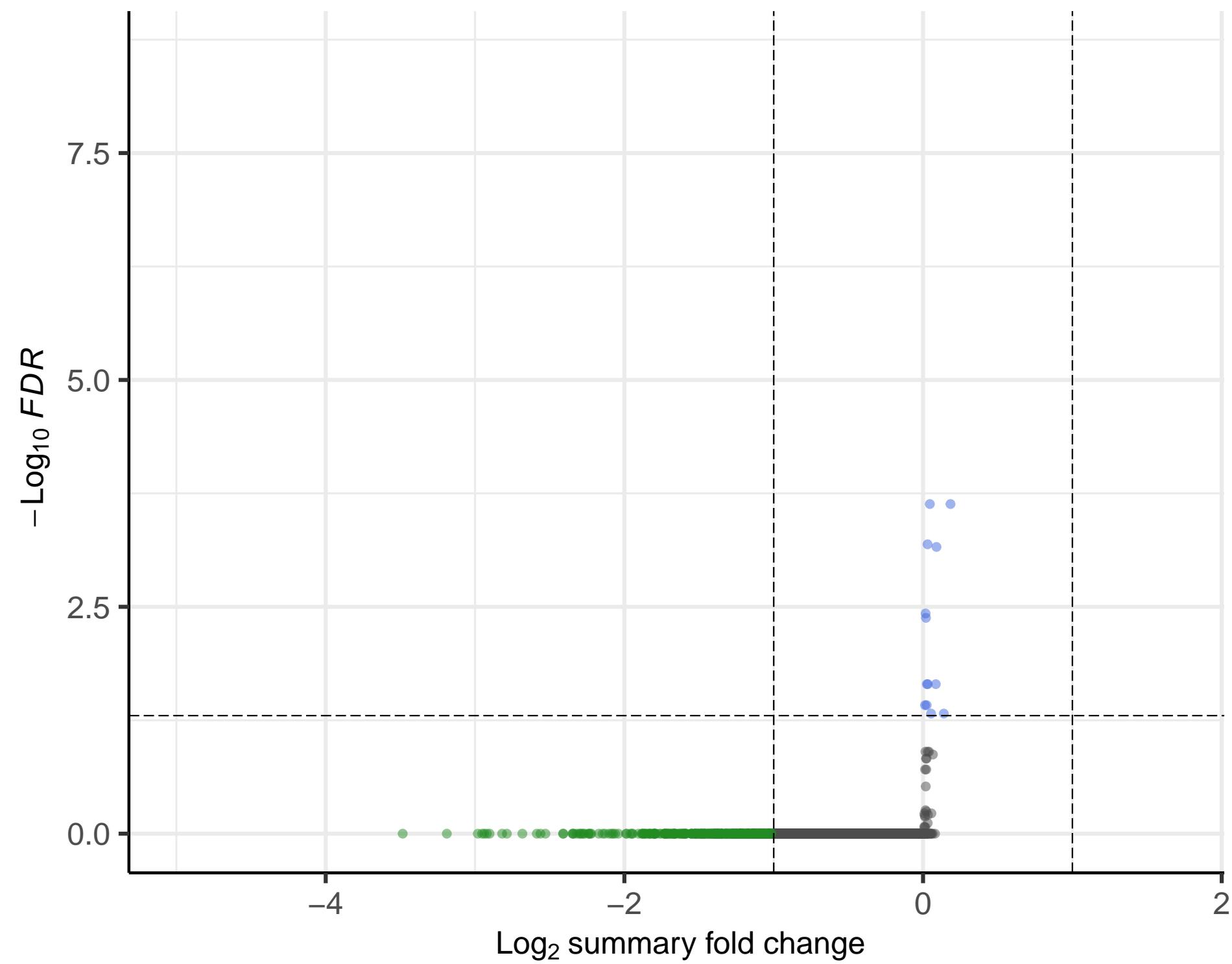
● NS ● Log₂ FC ● p-value ● p – value and log₂ FC



Endo.Mural_02

n nuc=0, n genes FDR<0.05=13

● NS ● Log₂ FC ● p-value

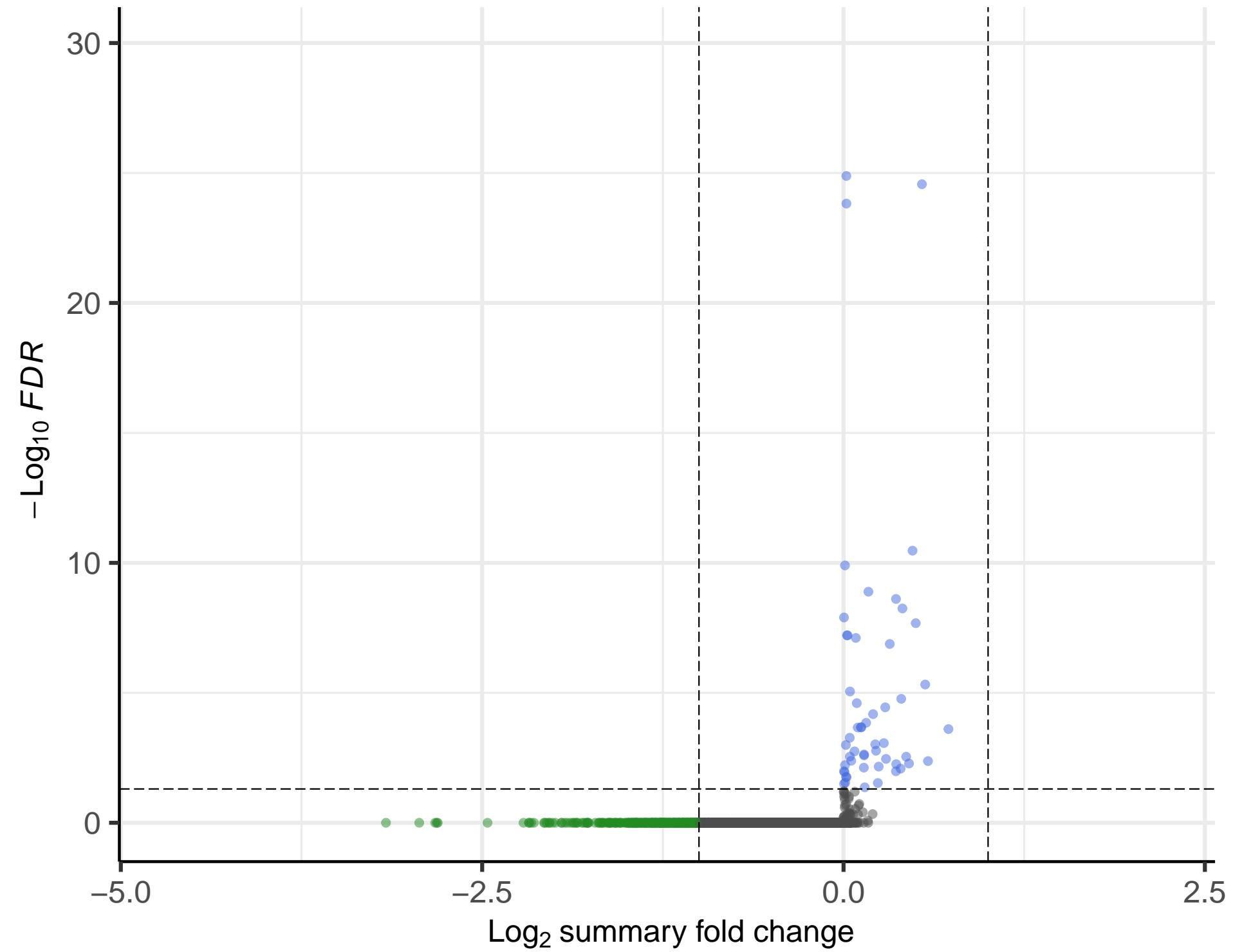


total = 36601 variables

Excit_01

n nuc=7927, n genes FDR<0.05=53

● NS ● Log₂ FC ● p-value

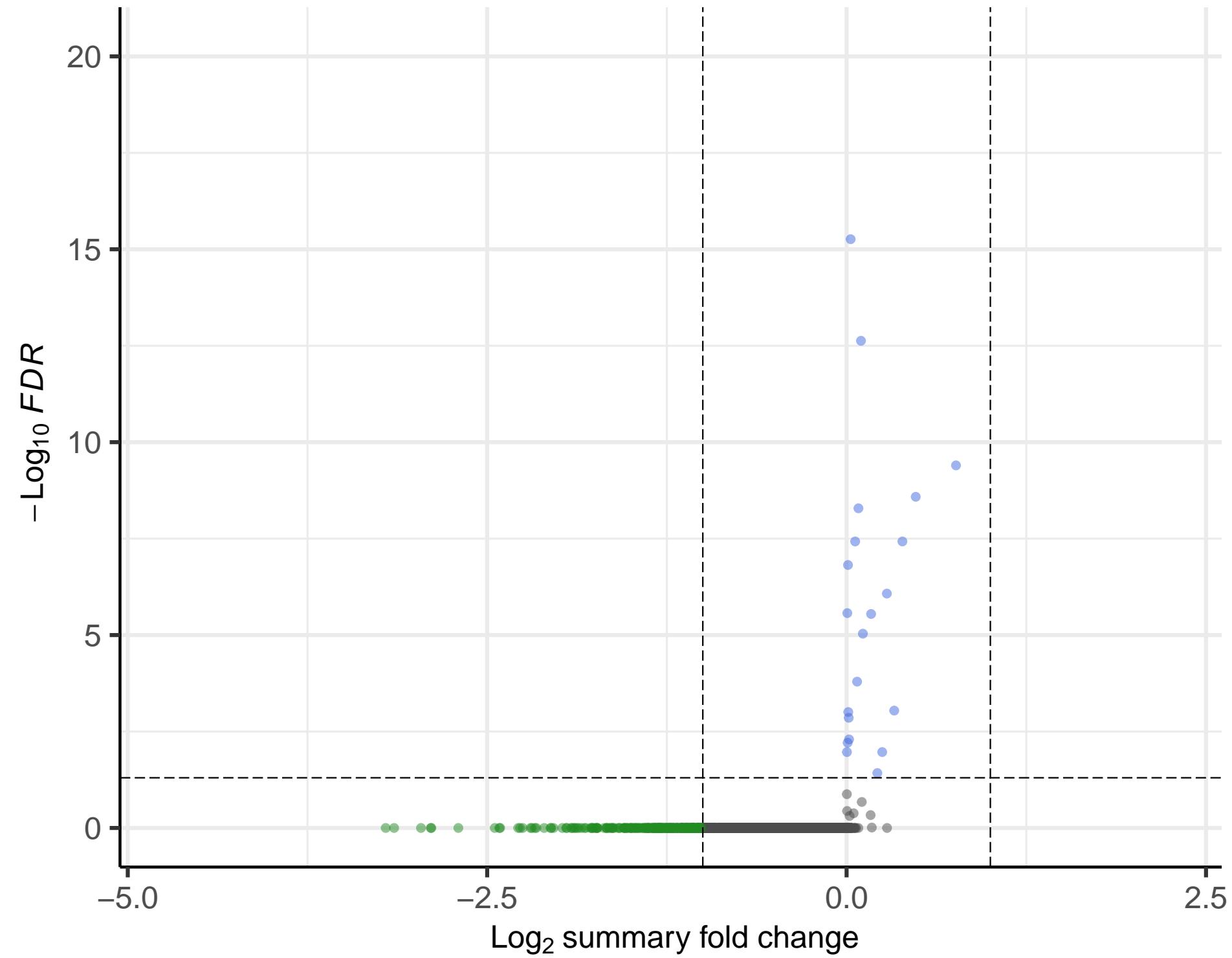


total = 36601 variables

Excit_02

n nuc=2487, n genes FDR<0.05=21

● NS ● Log₂ FC ● p-value

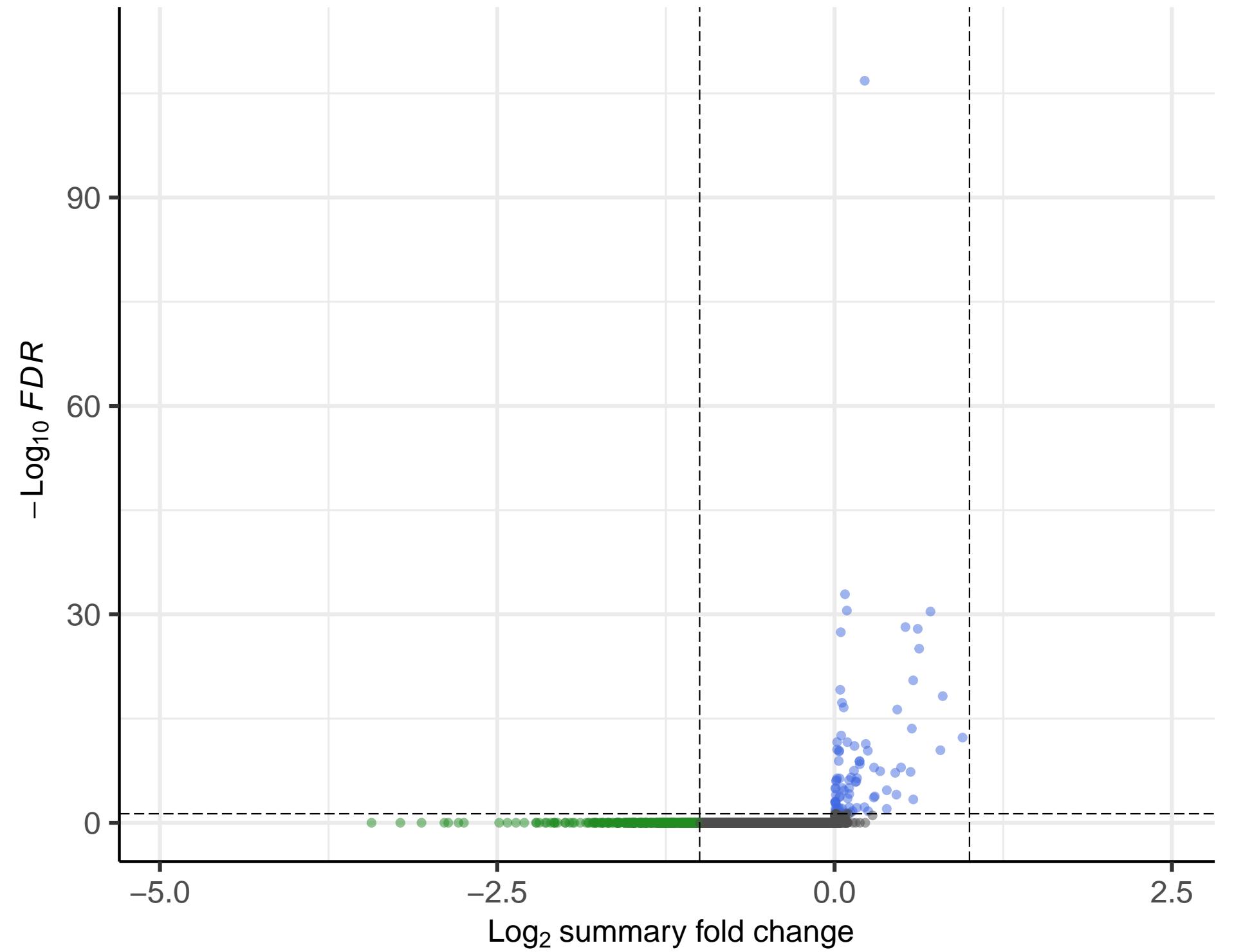


total = 36601 variables

Excit_03

n nuc=1309, n genes FDR<0.05=81

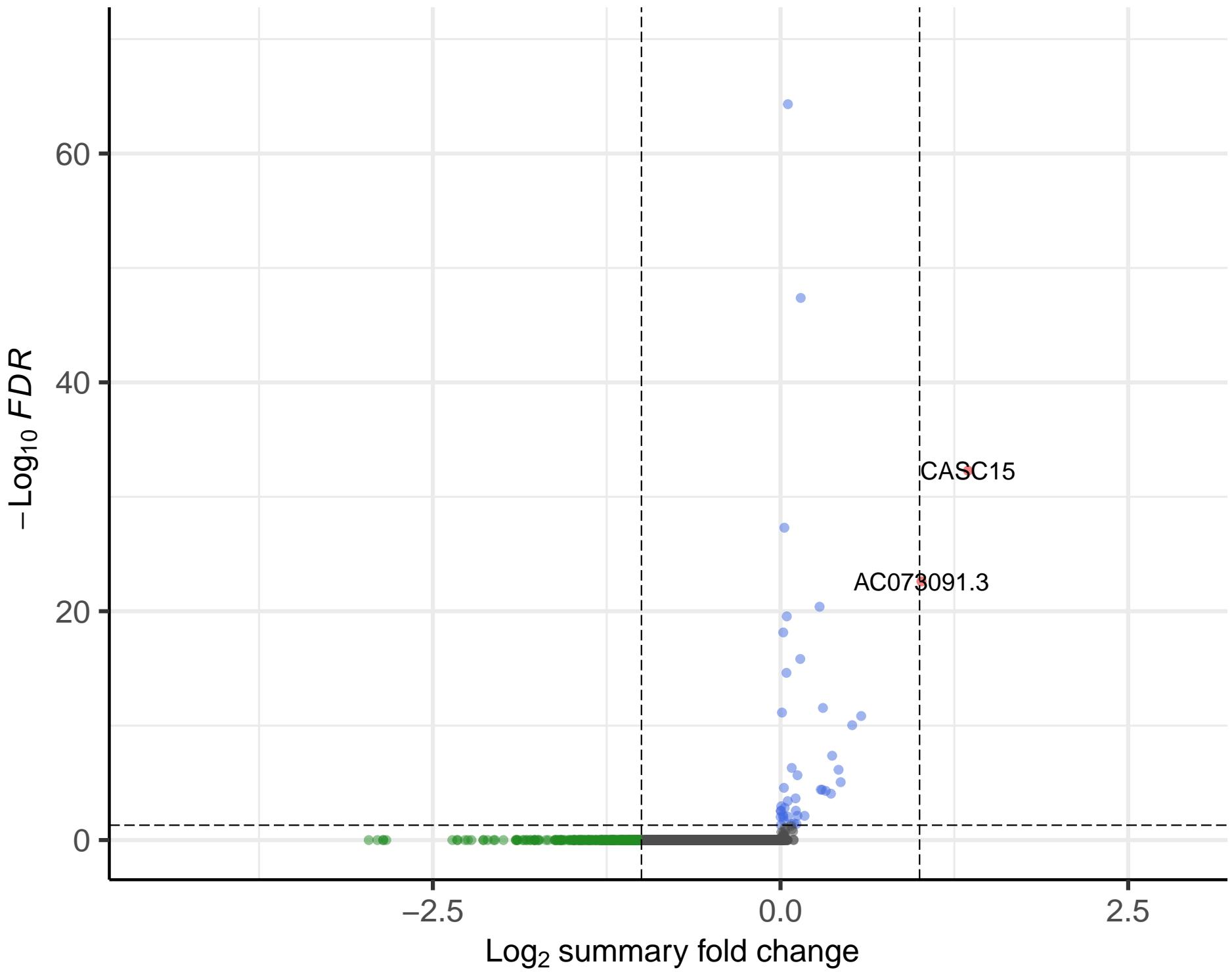
● NS ● Log₂ FC ● p-value



total = 36601 variables

Exit_04

n nuc=2171, n genes FDR<0.05=42

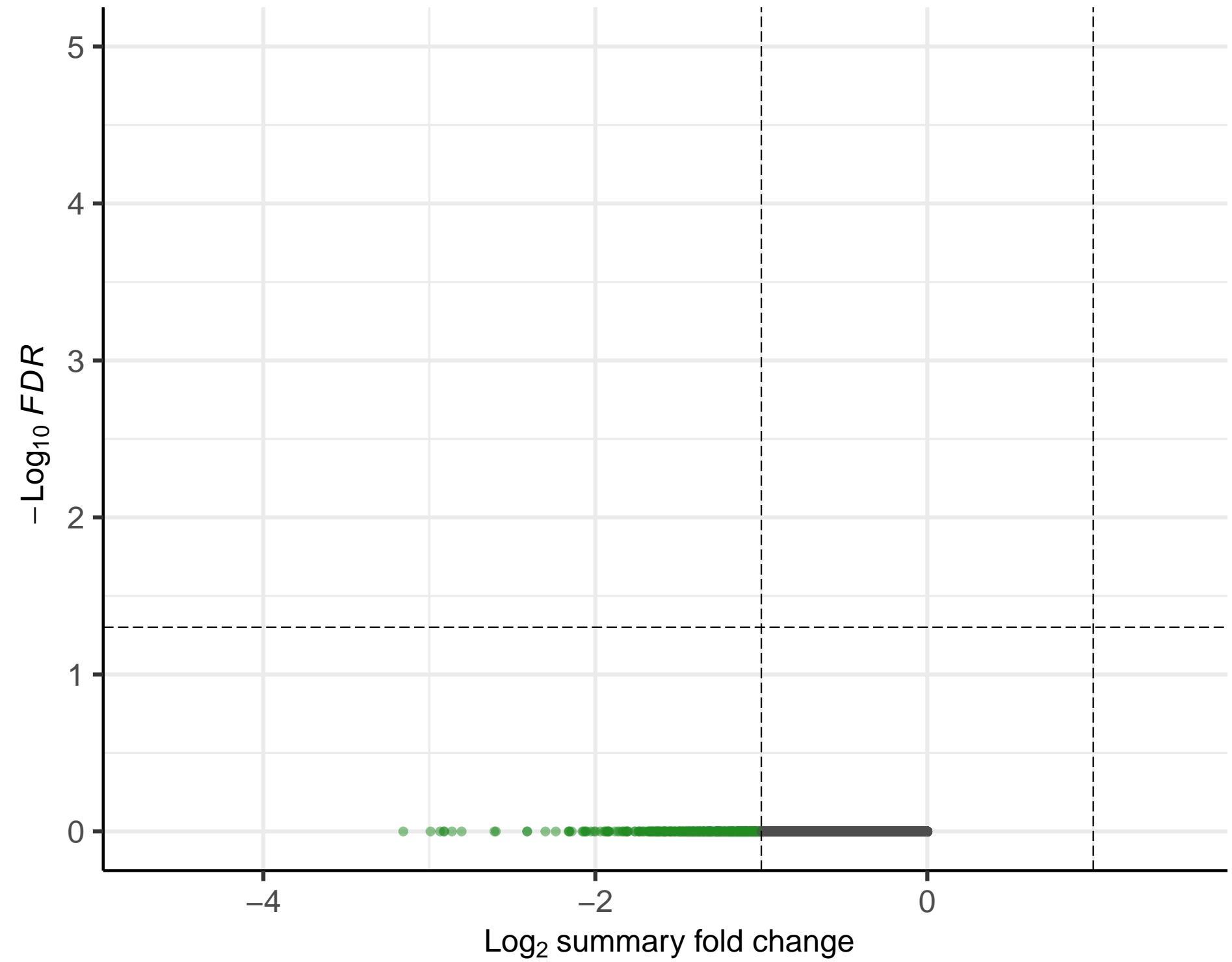


total = 36601 variables

Excit_05

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

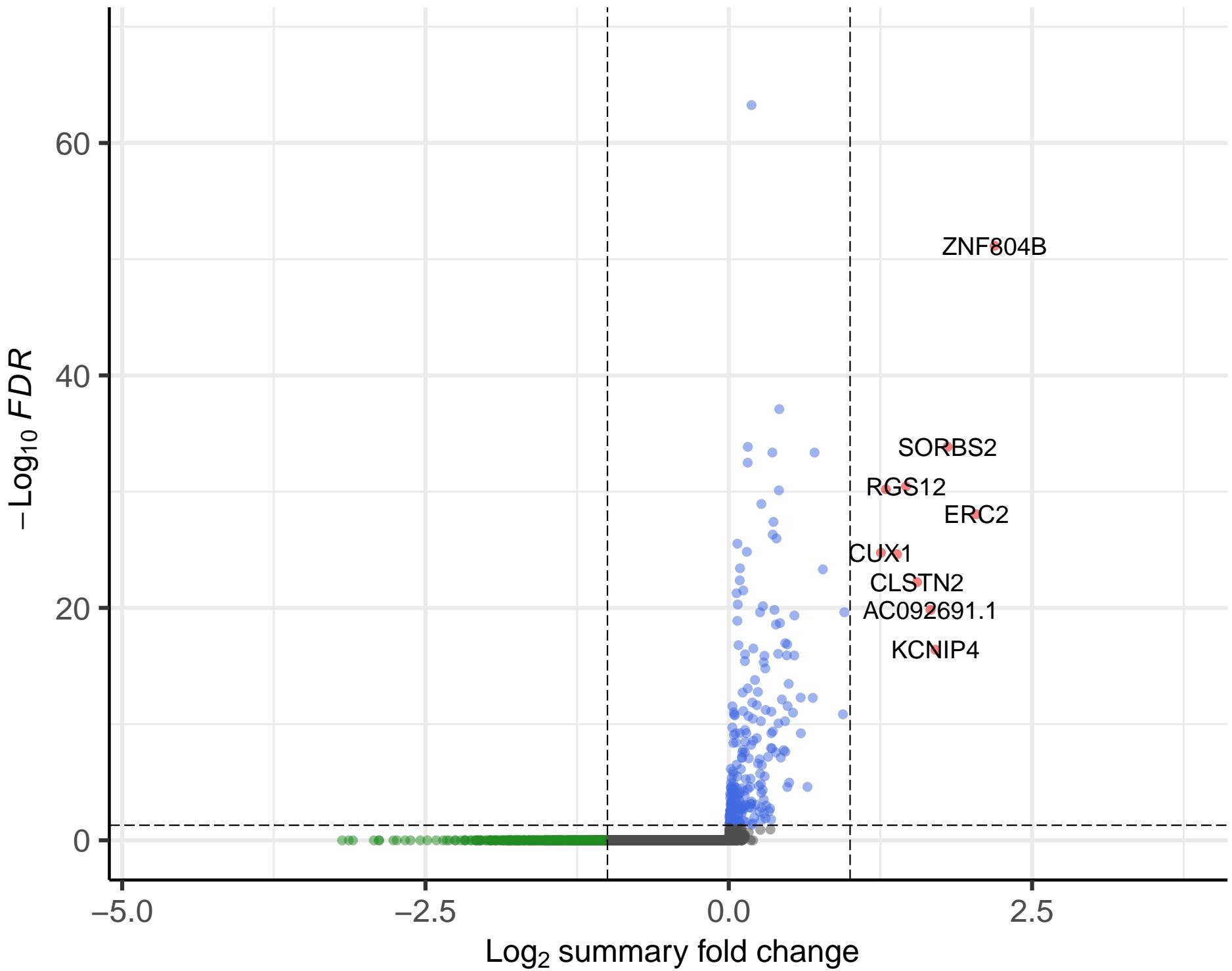
● NS ● Log₂ FC



Excit_06

n nuc=329, n genes FDR<0.05=237

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

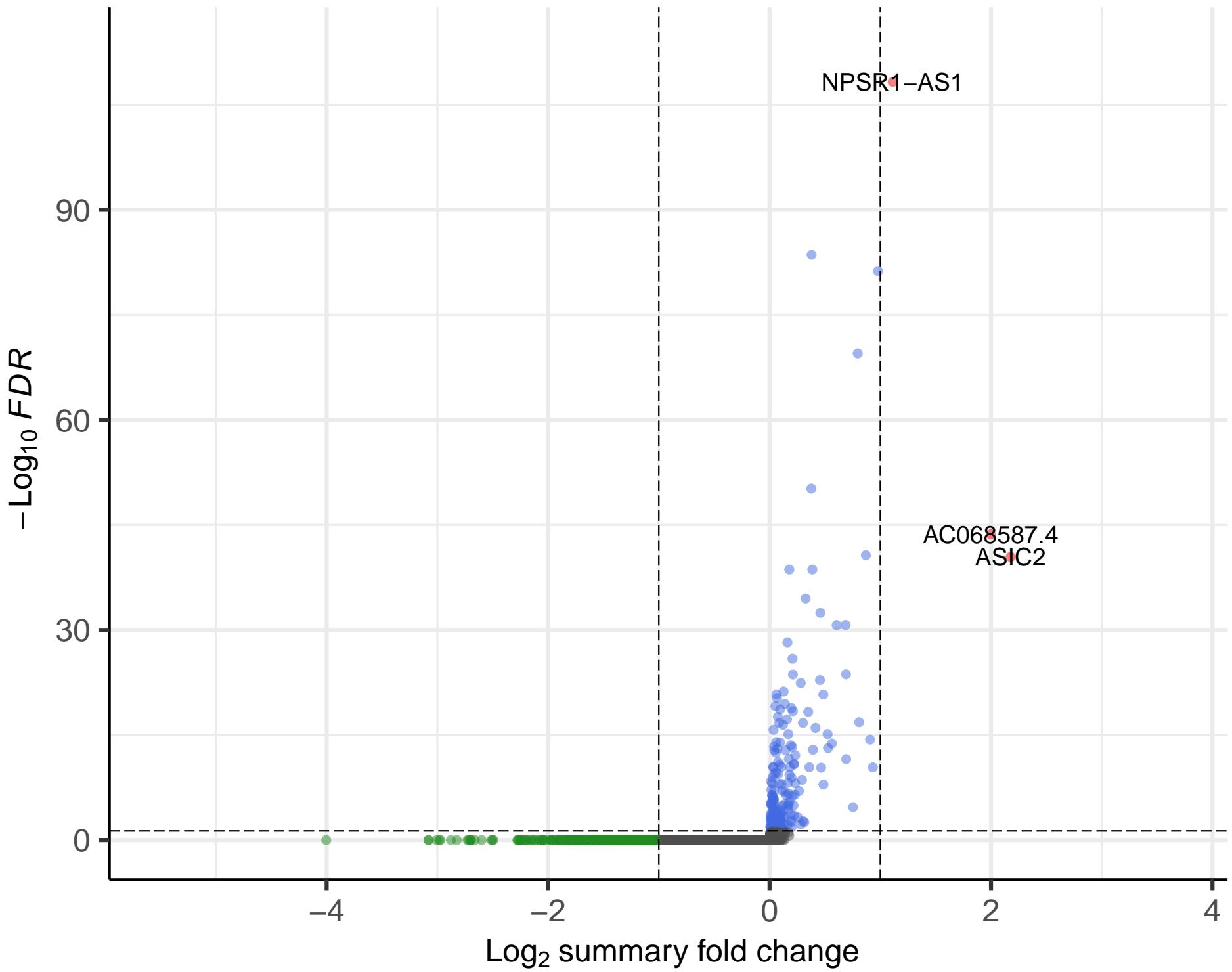


total = 36601 variables

Excit_07

n nuc=334, n genes FDR<0.05=185

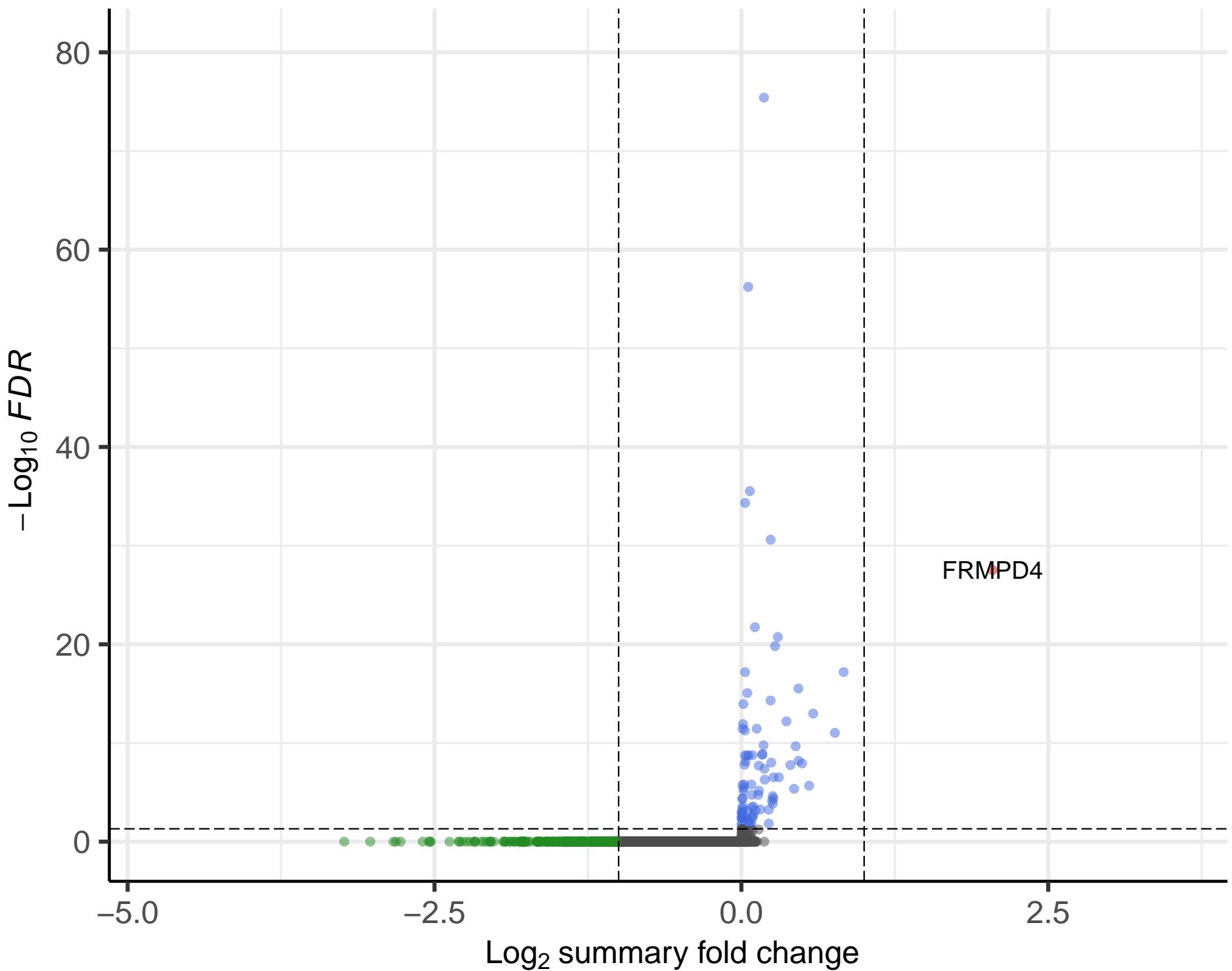
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Excit_08

n nuc=1463, n genes FDR<0.05=86

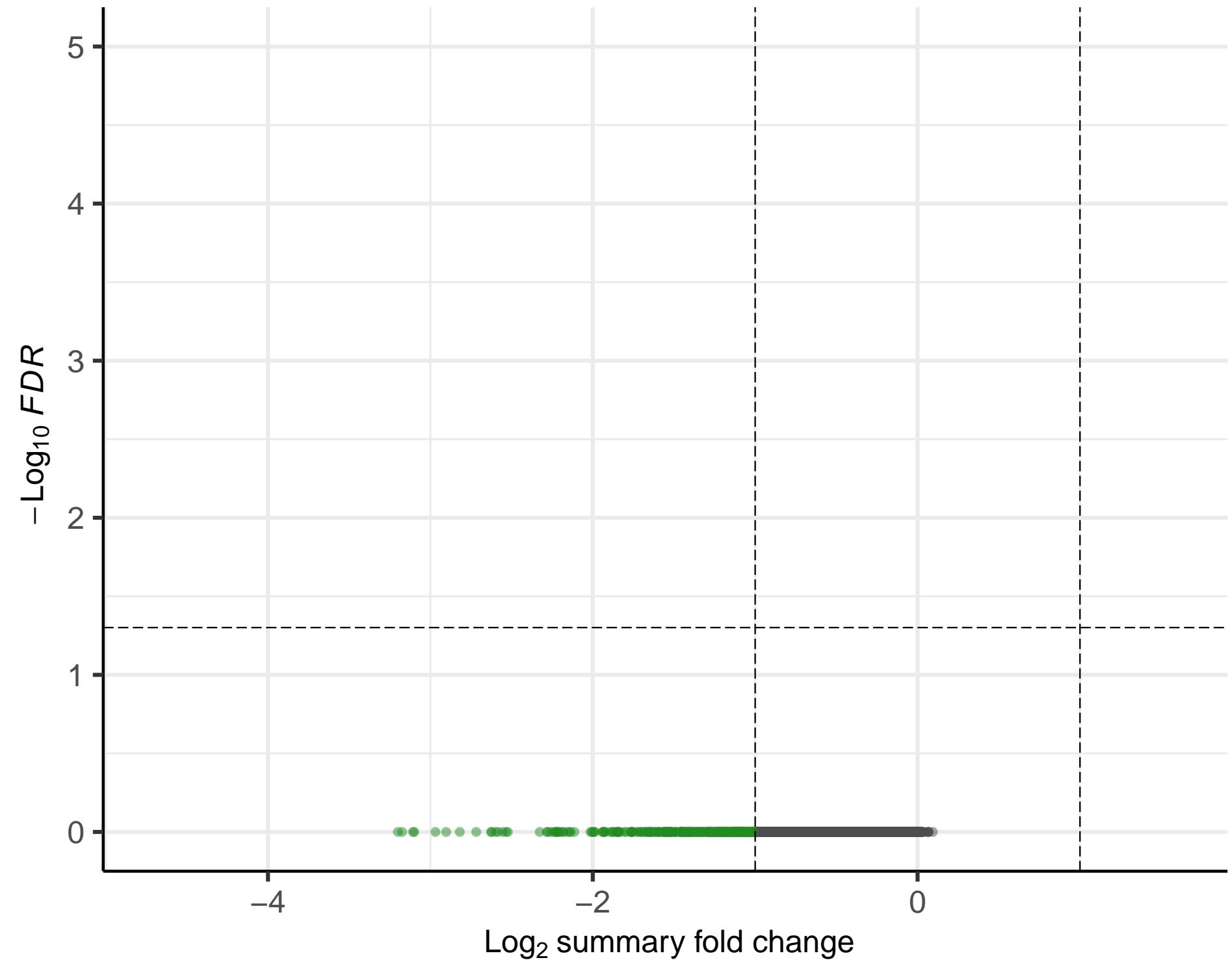
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Excit_09

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

● NS ● Log₂ FC

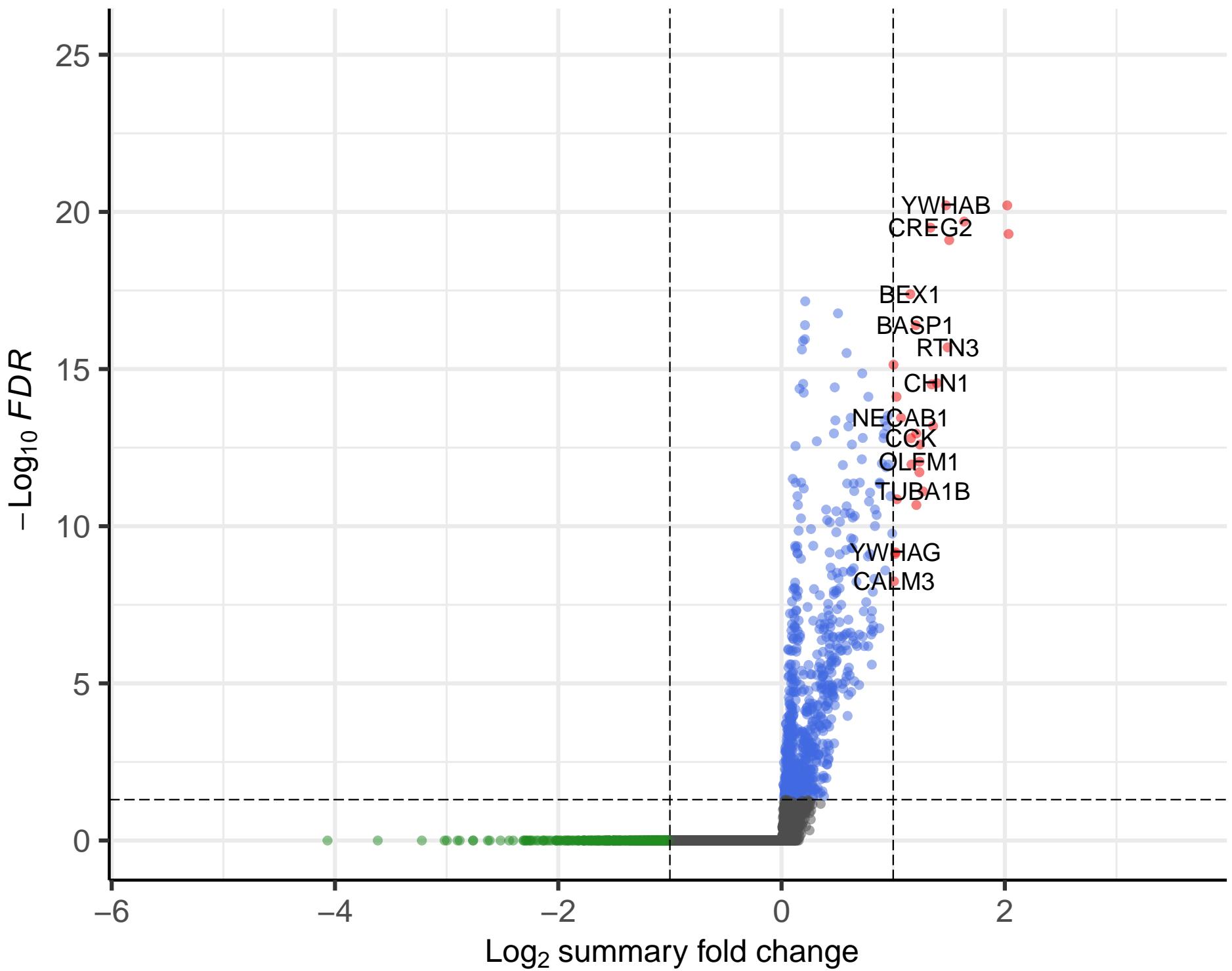


total = 36601 variables

Excit_10

n nuc=1079, n genes FDR<0.05=742

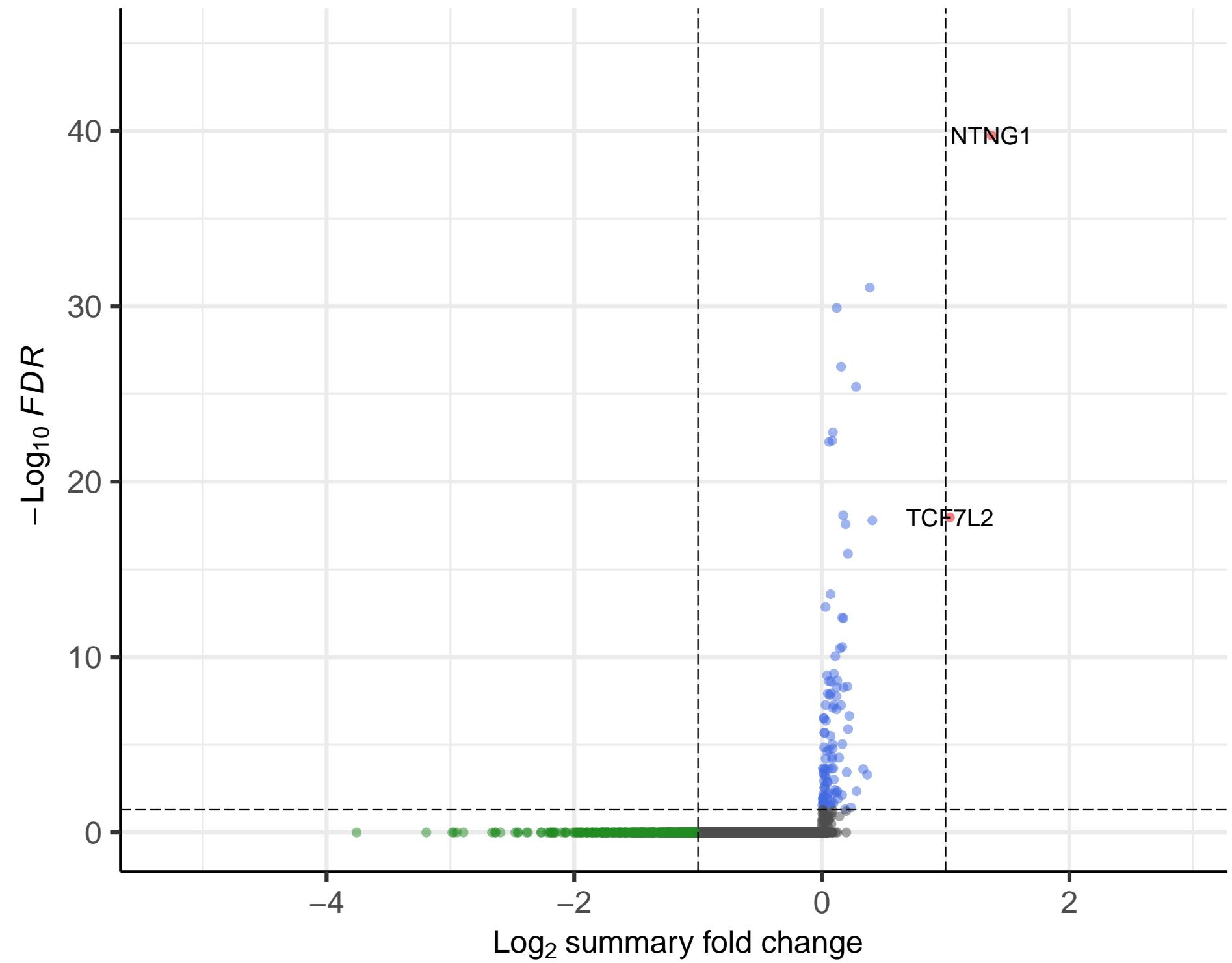
● NS ● Log₂ FC ● p-value ● p – value and log₂ FC



Excit_11

n nuc=482, n genes FDR<0.05=103

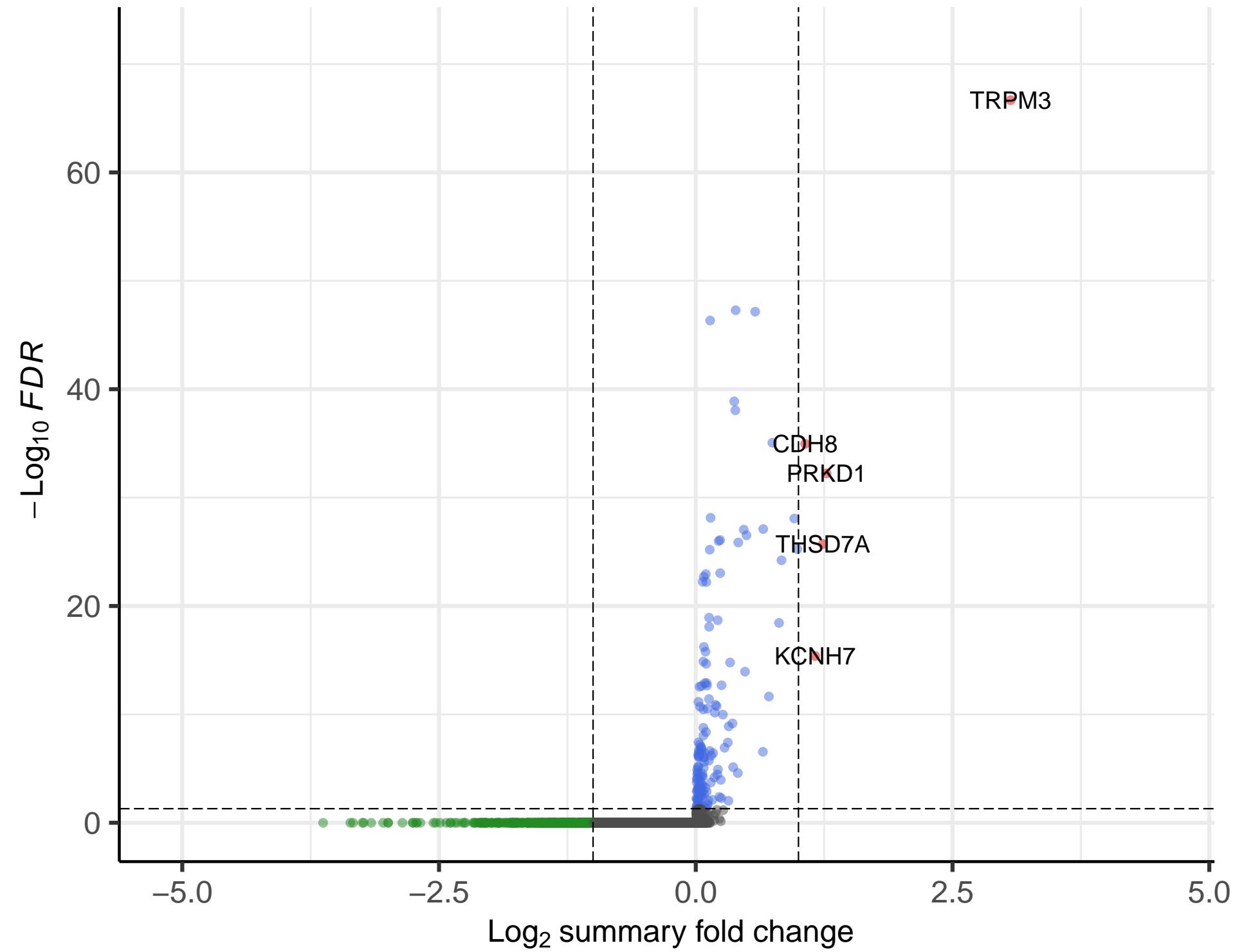
● NS ● Log₂ FC ● p-value ● p – value and log₂ FC



Excit_12

n nuc=420, n genes FDR<0.05=144

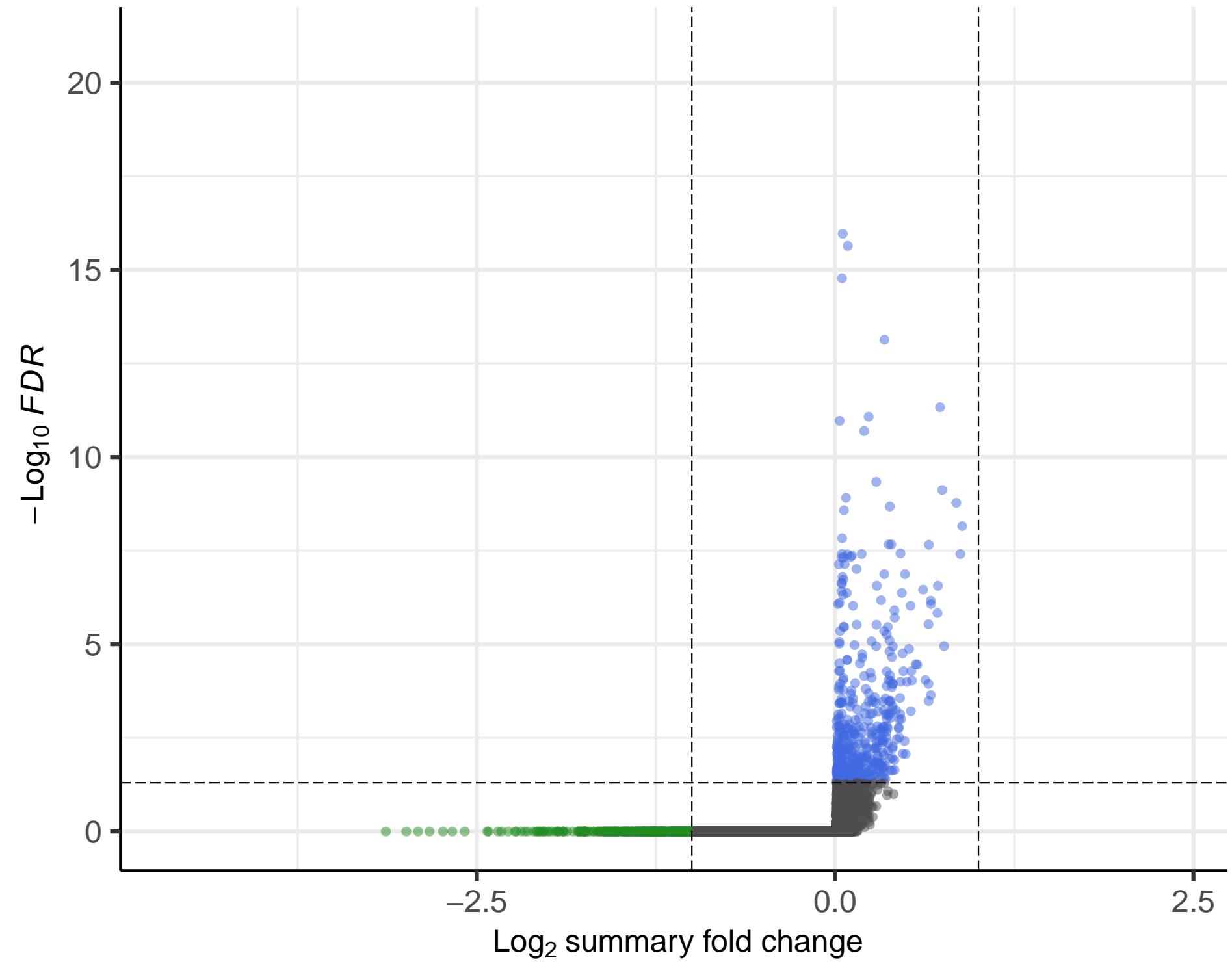
● NS ● Log₂ FC ● p-value ● p – value and log₂ FC



Excit_13

n nuc=1567, n genes FDR<0.05=442

● NS ● Log₂ FC ● p-value

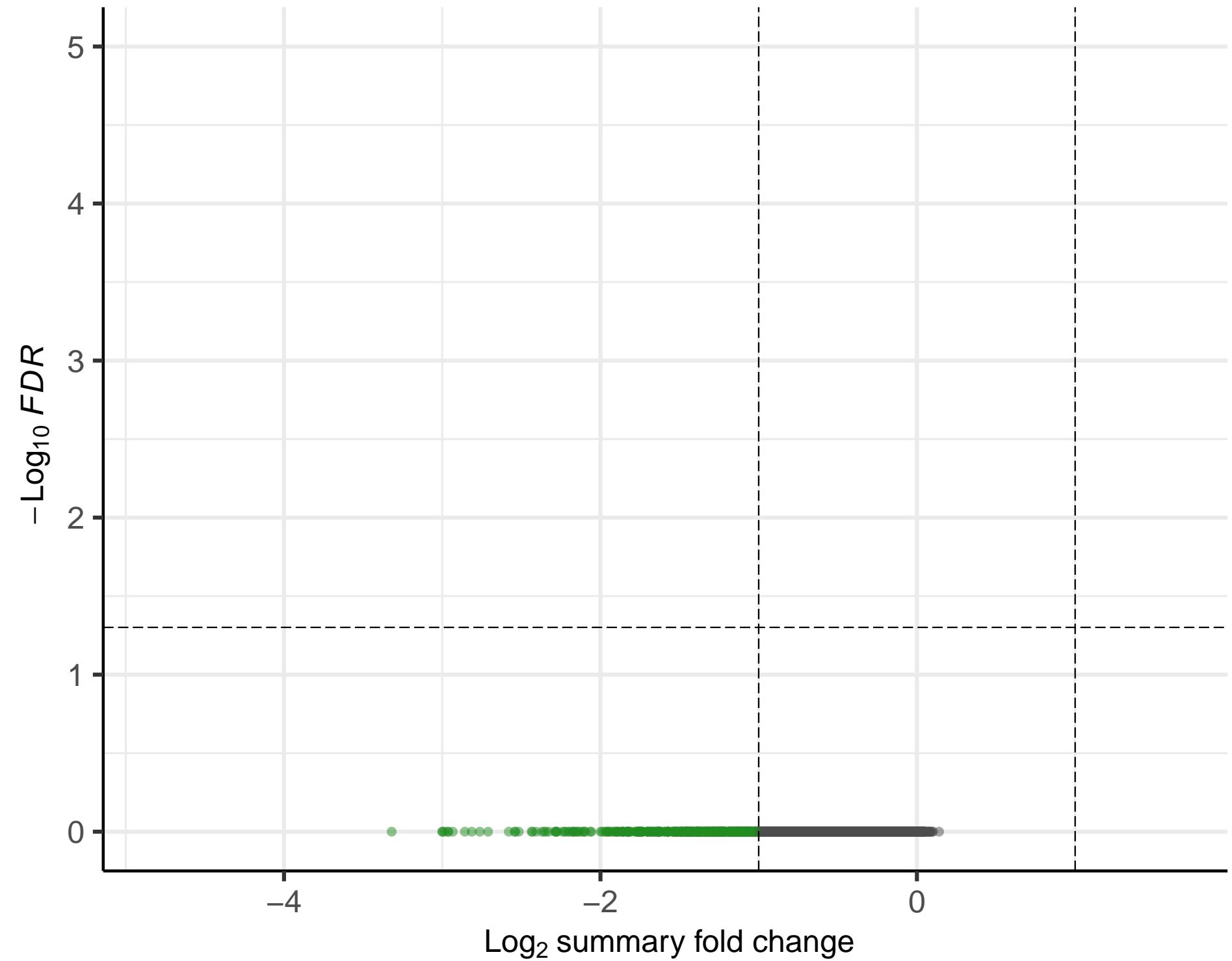


total = 36601 variables

Excit_14

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

● NS ● Log₂ FC

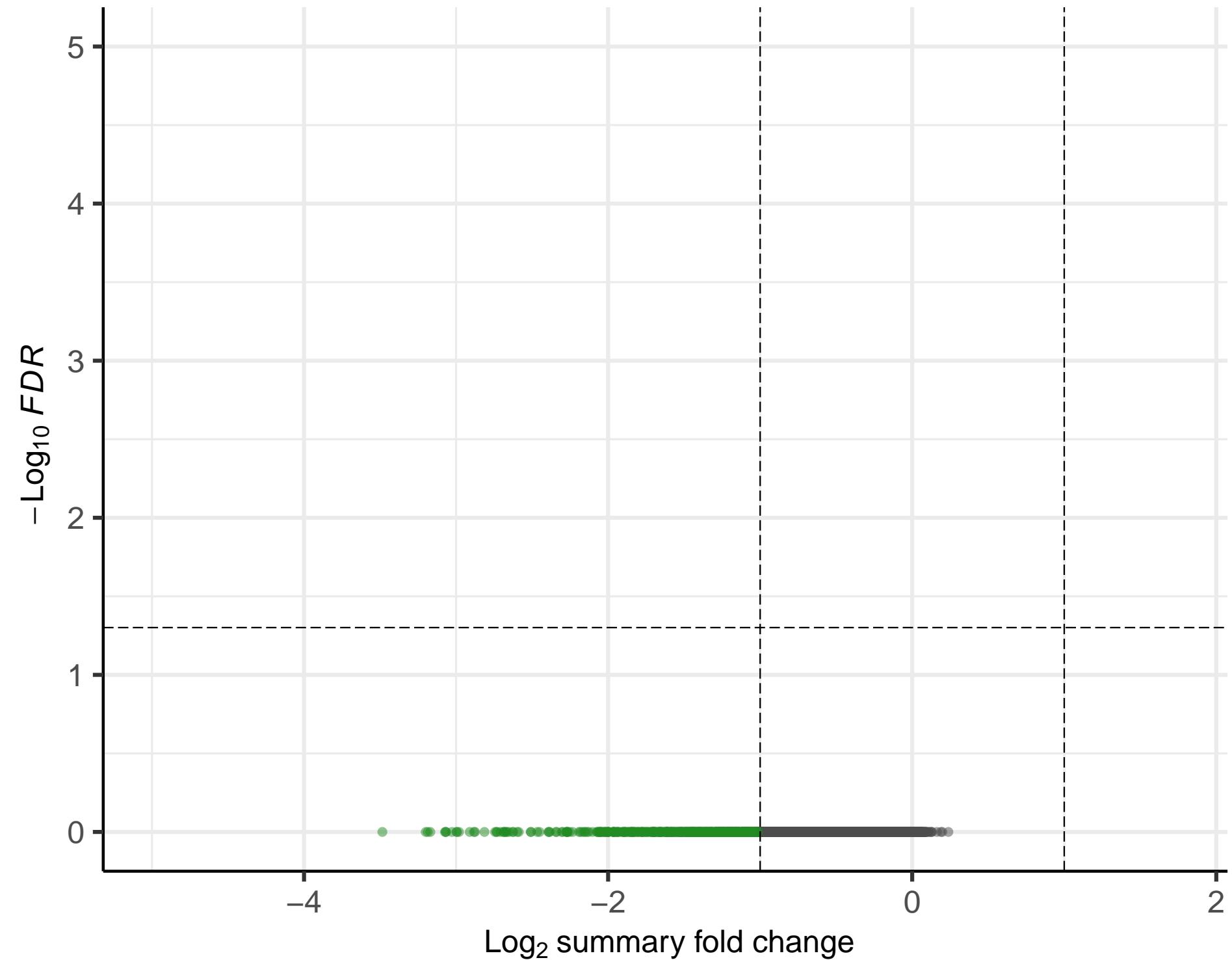


total = 36601 variables

Excit_15

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

● NS ● Log₂ FC

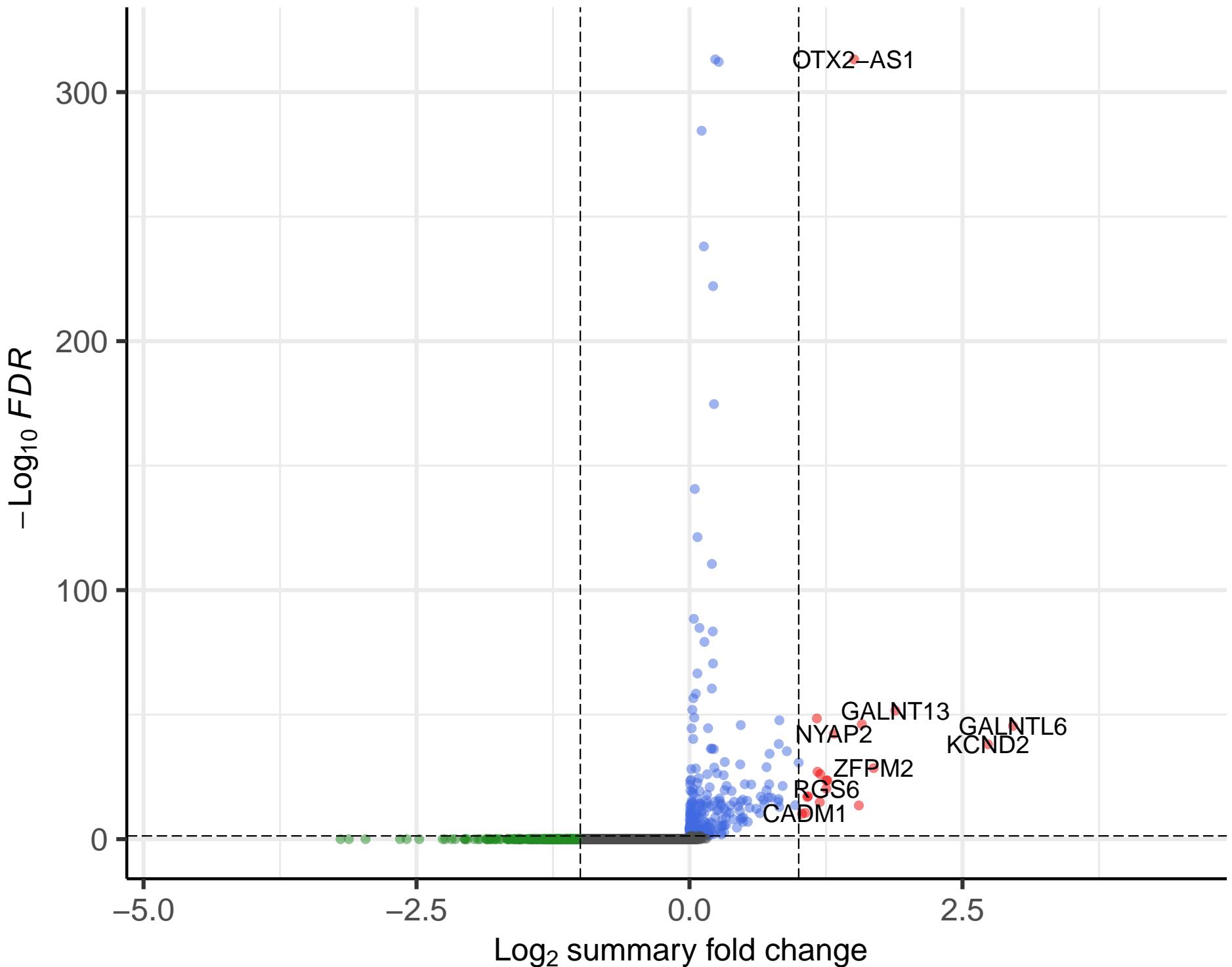


total = 36601 variables

Inhib_01

n nuc=5366, n genes FDR<0.05=240

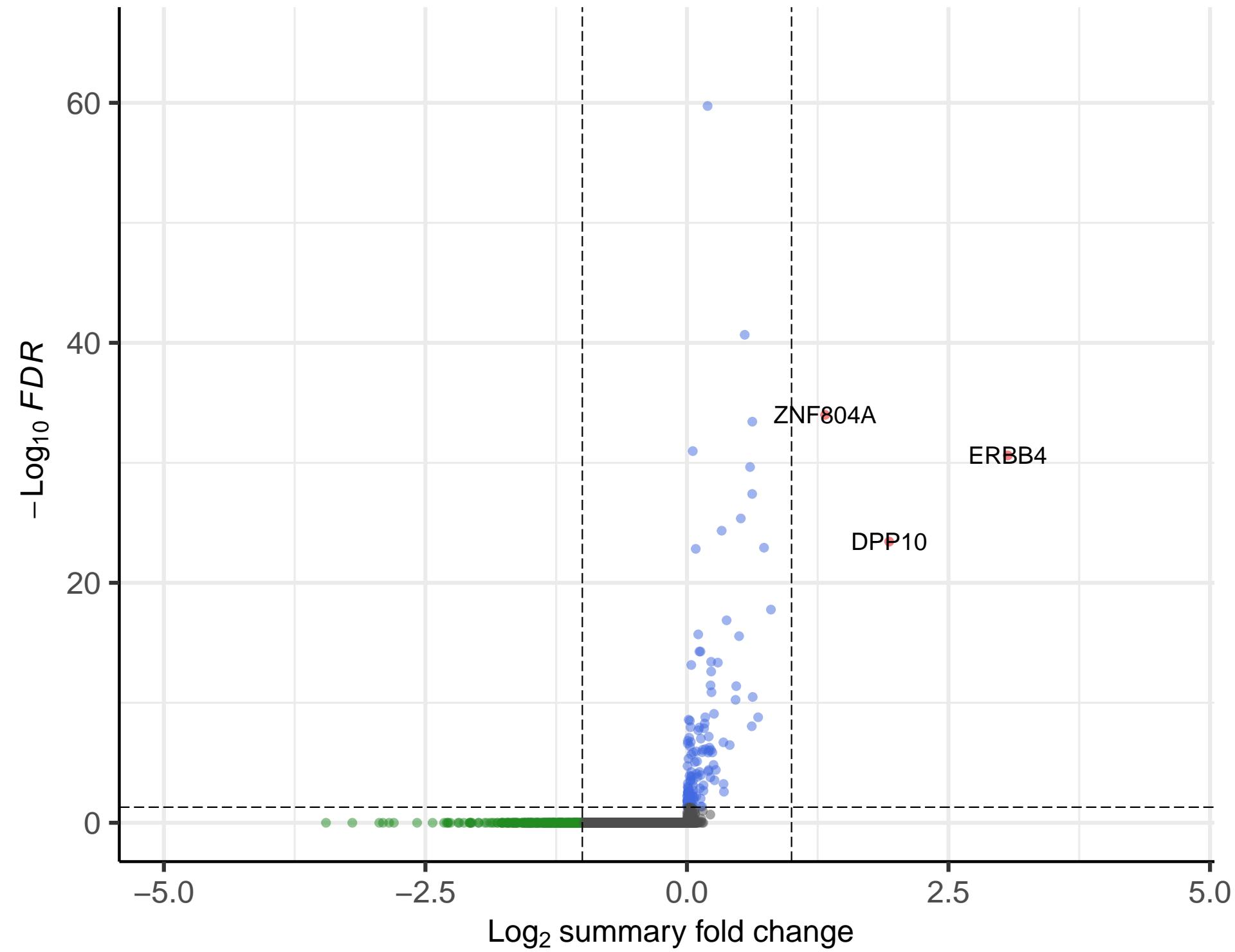
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Inhib_02

n nuc=1267, n genes FDR<0.05=131

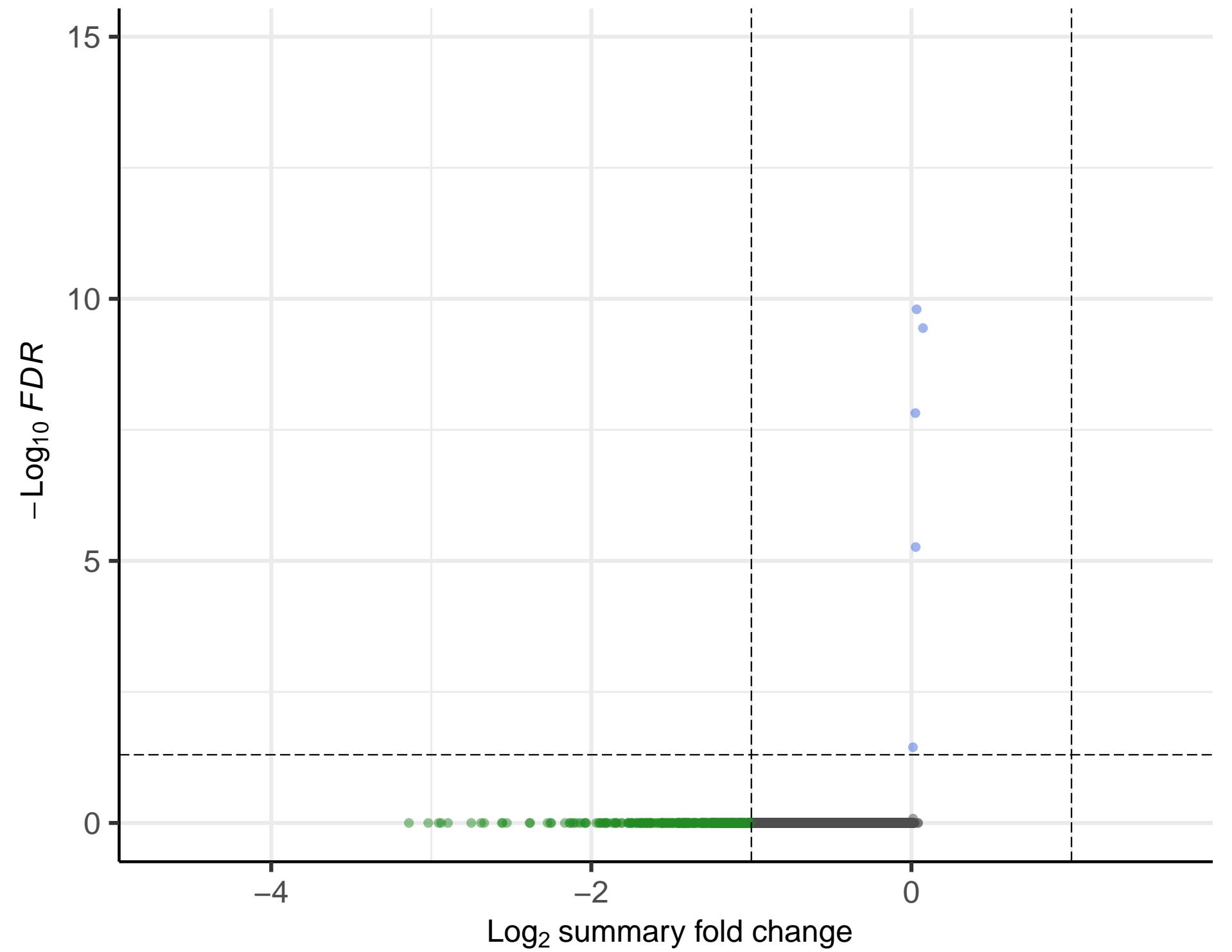
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Inhib_03

n nuc=1310, n genes FDR<0.05=5

● NS ● Log₂ FC ● p-value

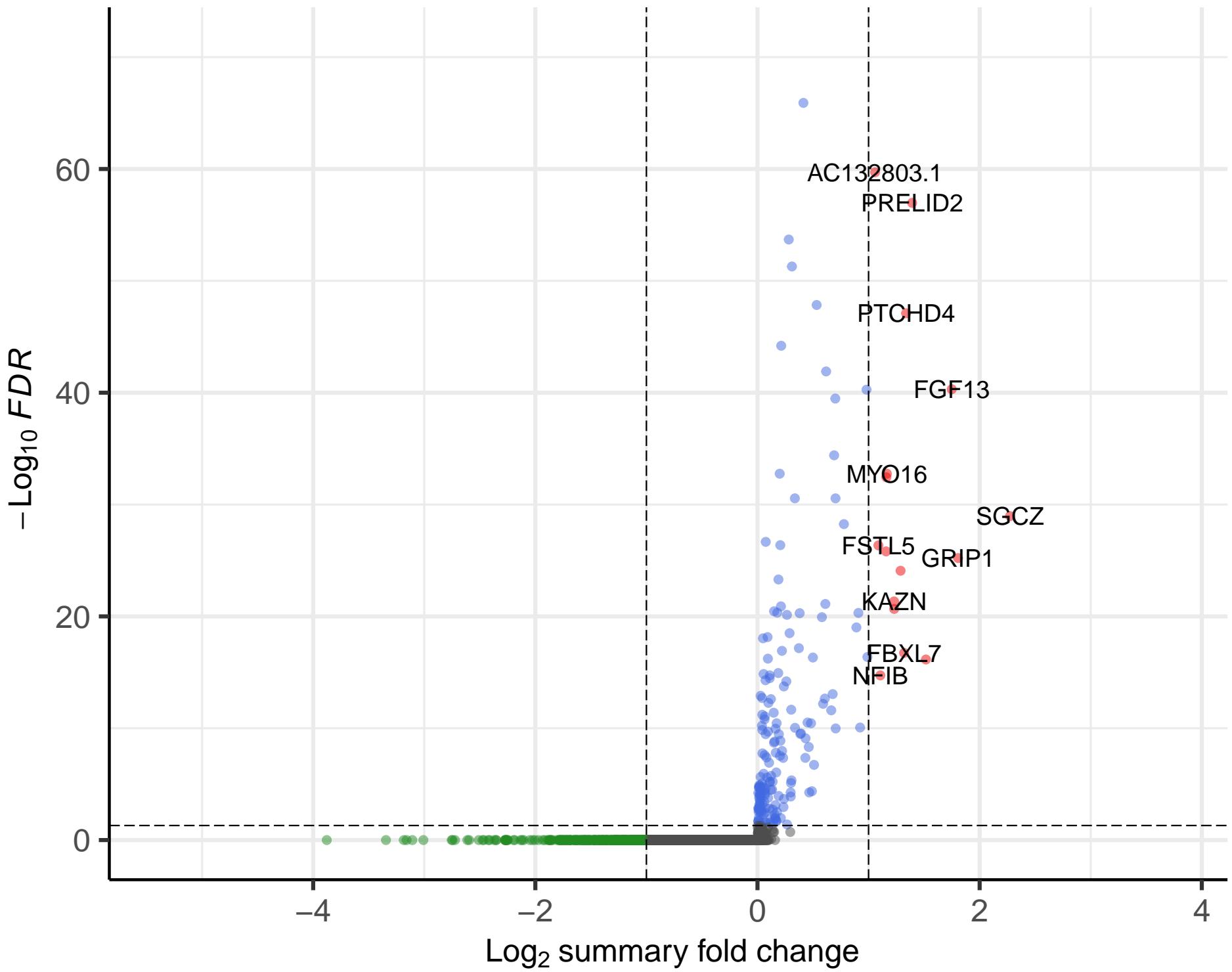


total = 36601 variables

Inhib_04

n nuc=565, n genes FDR<0.05=178

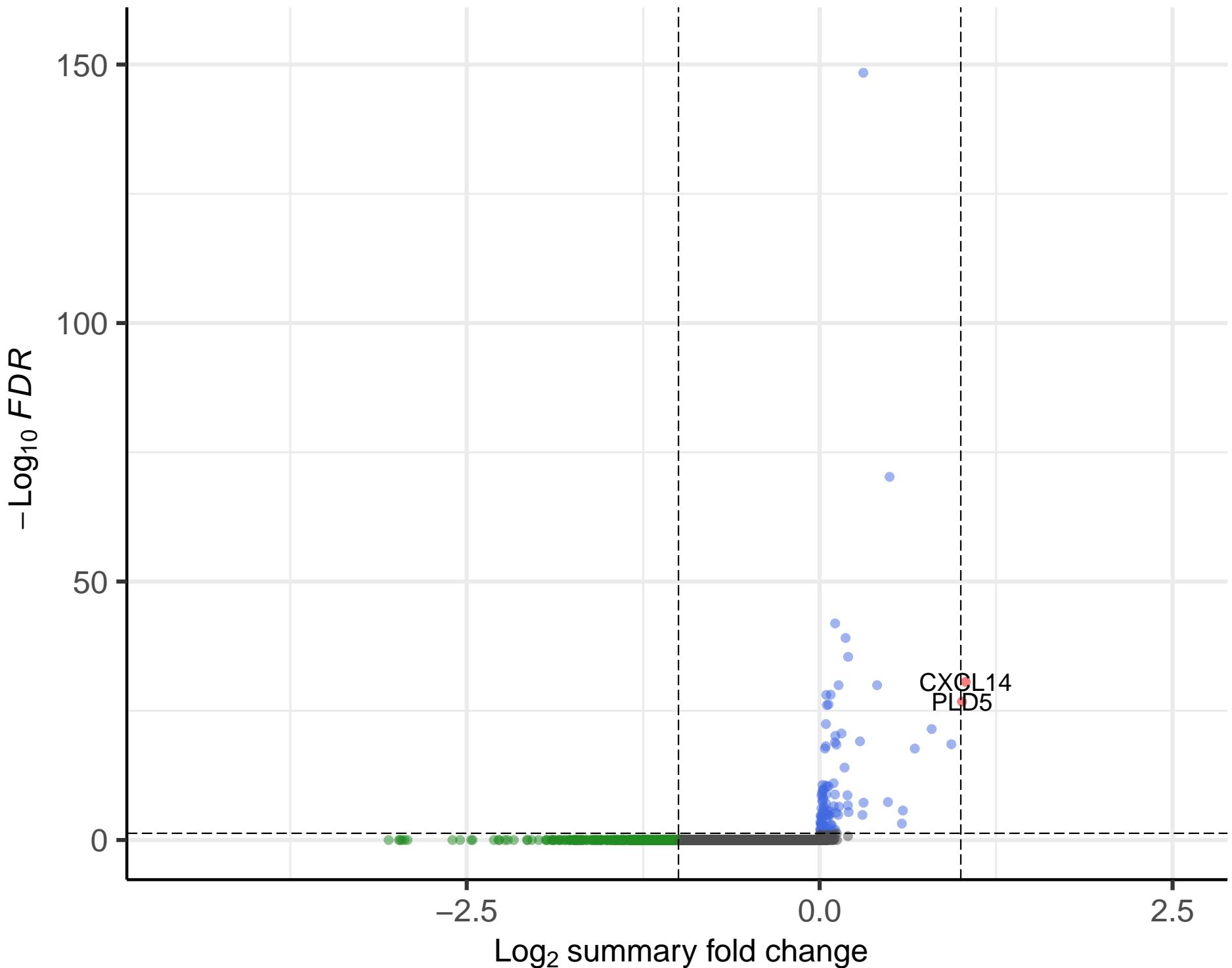
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Inhib_05

n nuc=1192, n genes FDR<0.05=97

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

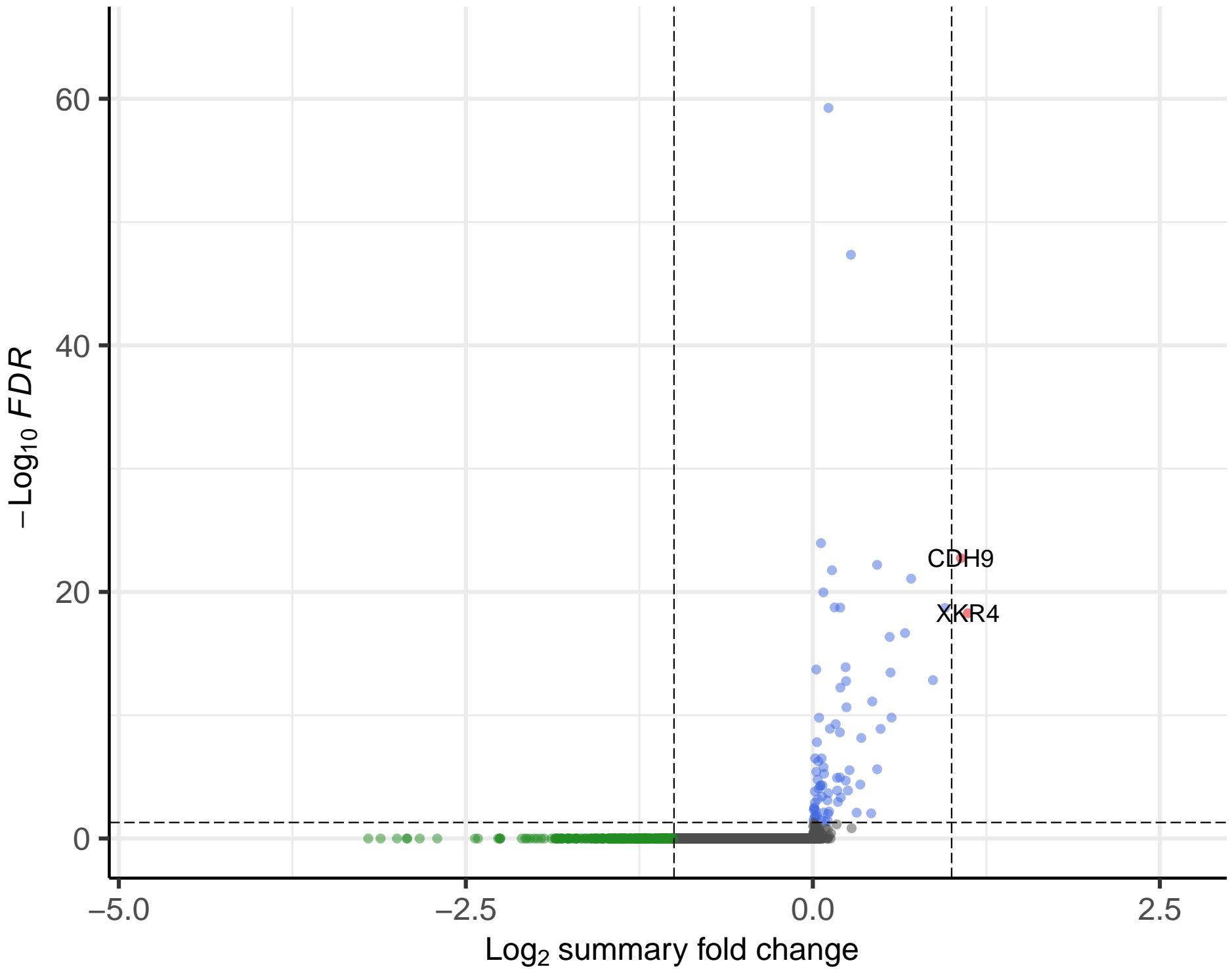


total = 36601 variables

Inhib_06

n nuc=1367, n genes FDR<0.05=74

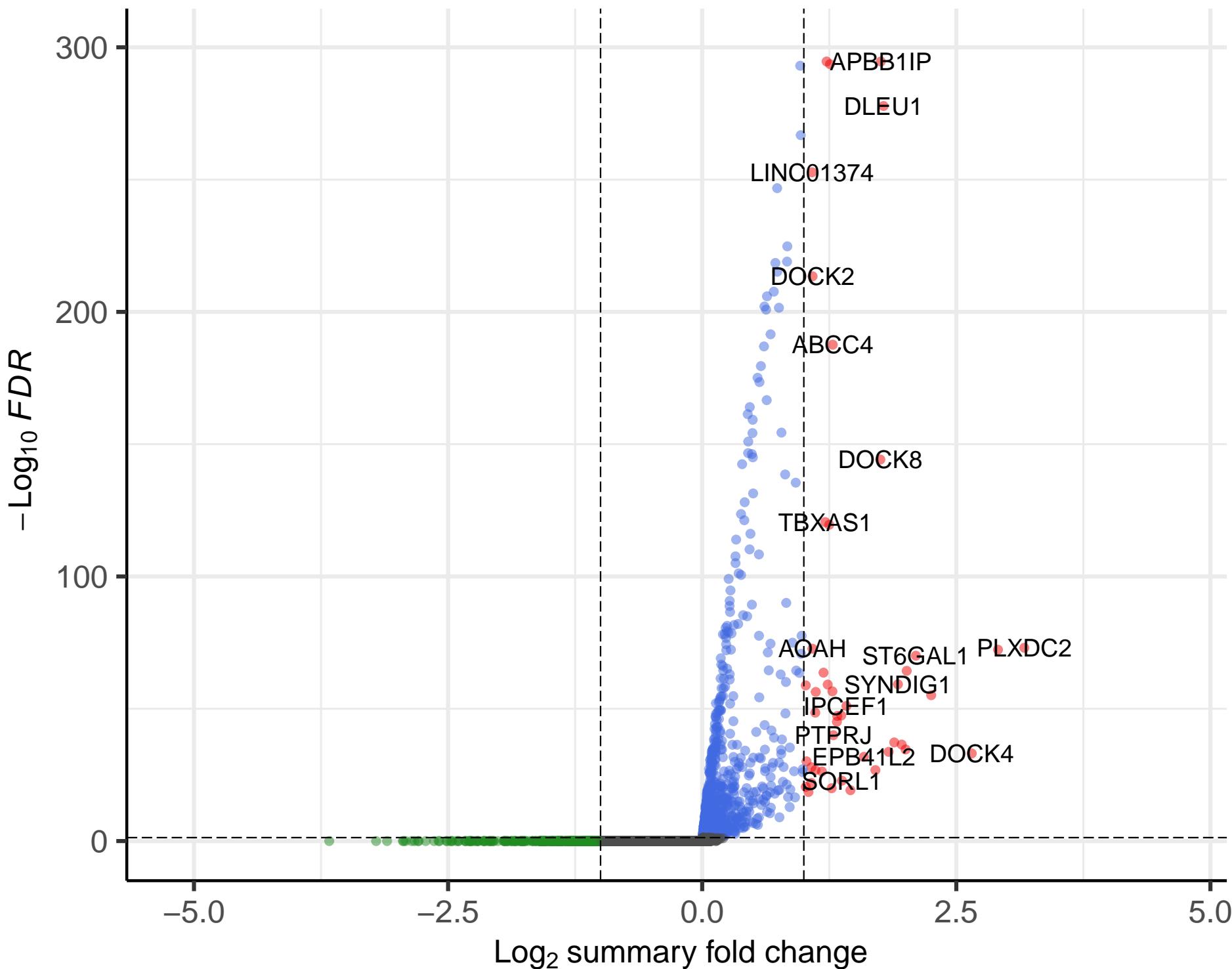
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Micro

n nuc=1601, n genes FDR<0.05=1016

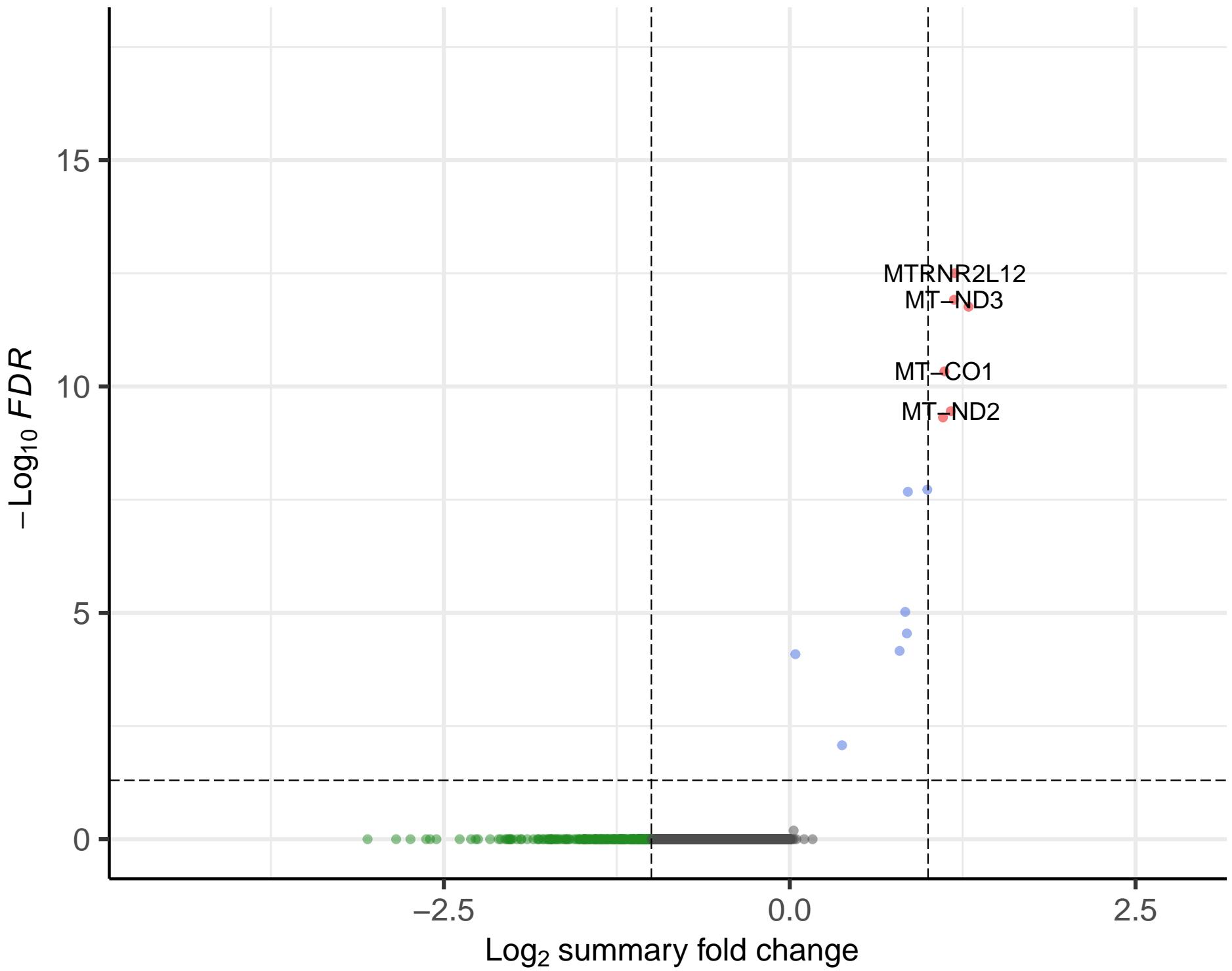
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Oligo_01

n nuc=1868, n genes FDR<0.05=13

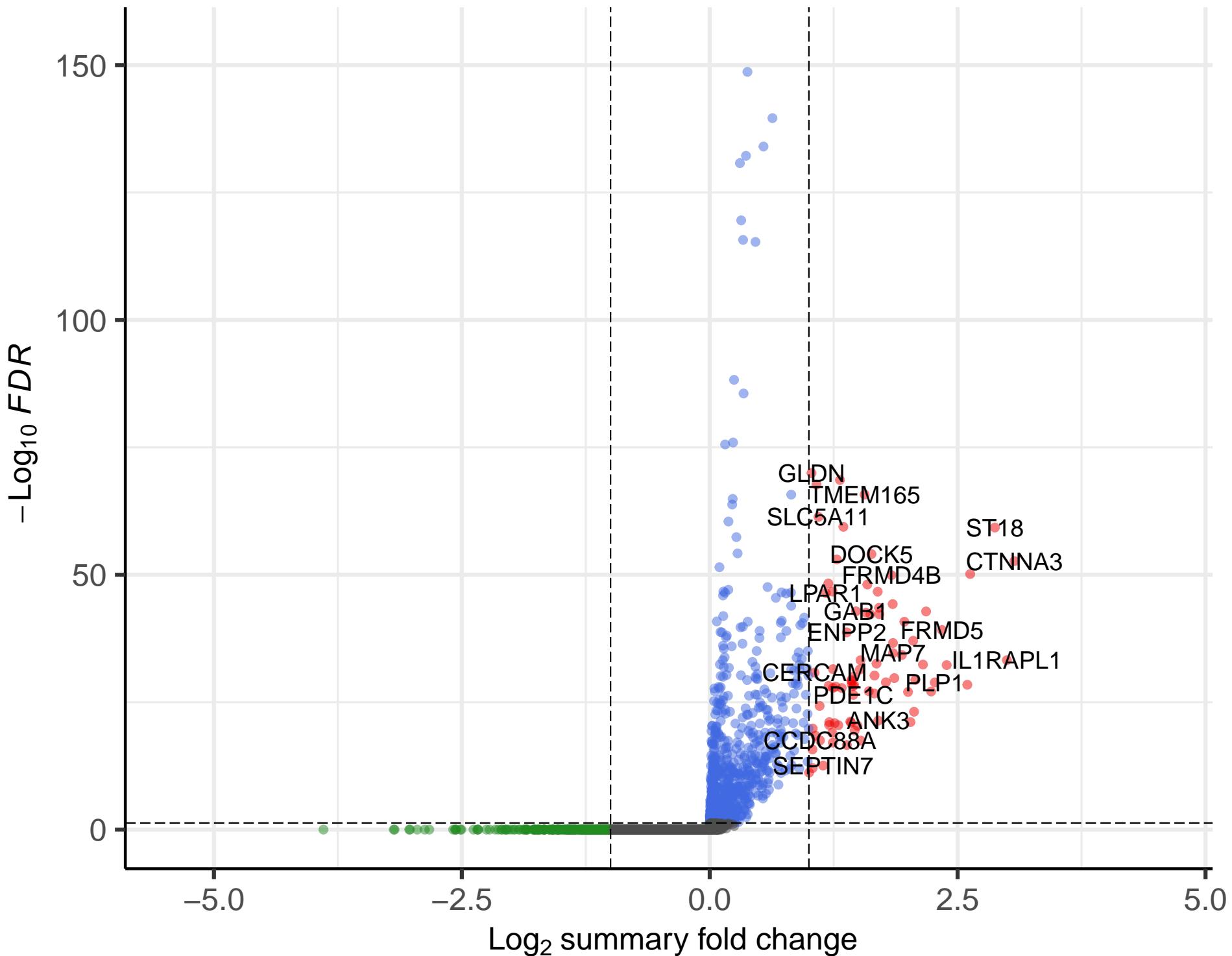
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Oligo_02

n nuc=4732, n genes FDR<0.05=889

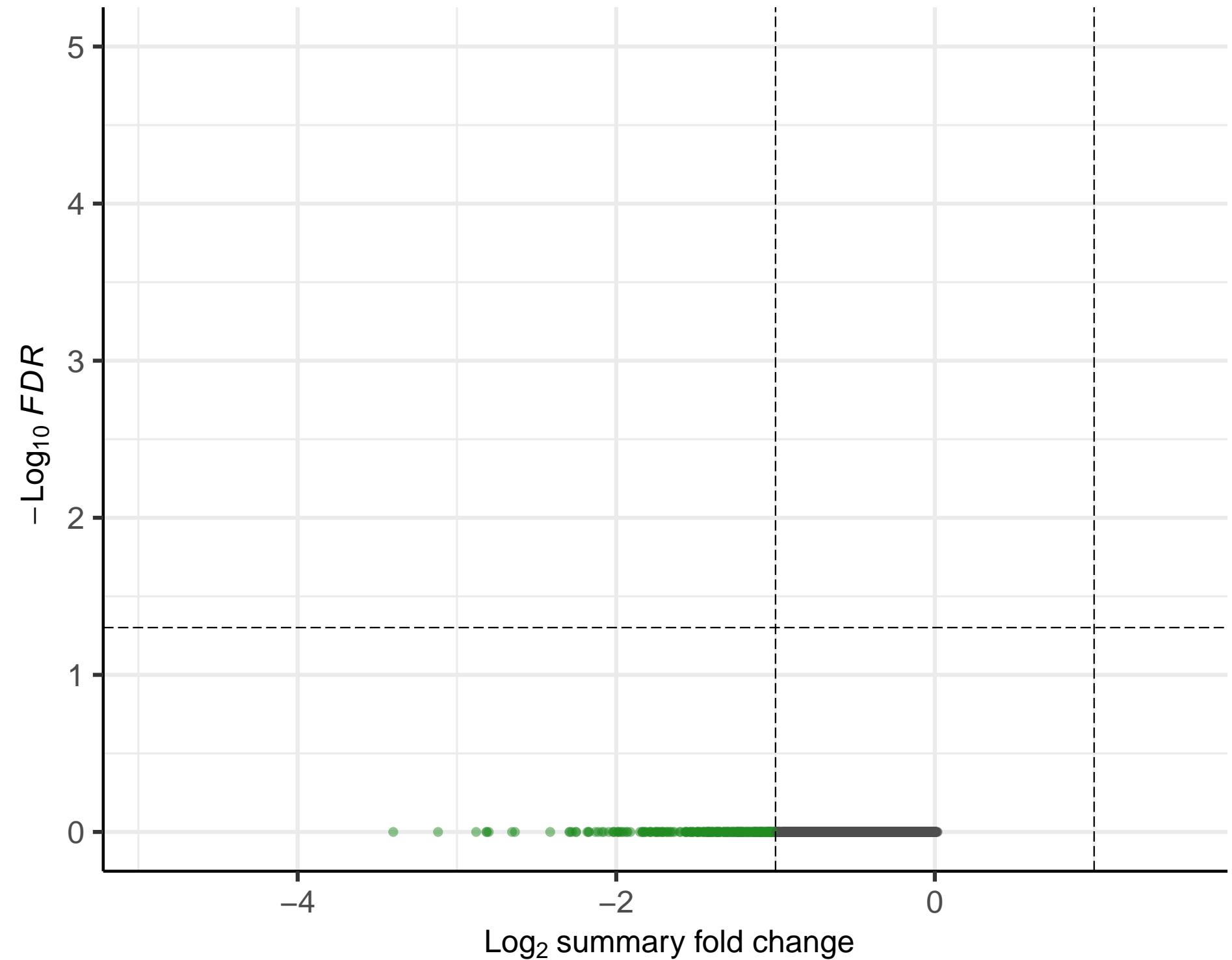
● NS ● Log₂ FC ● p-value ● p-value and log₂ FC



Oligo_03

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

● NS ● Log₂ FC



total = 36601 variables

OPC

n nuc=1940, n genes FDR<0.05=277

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

