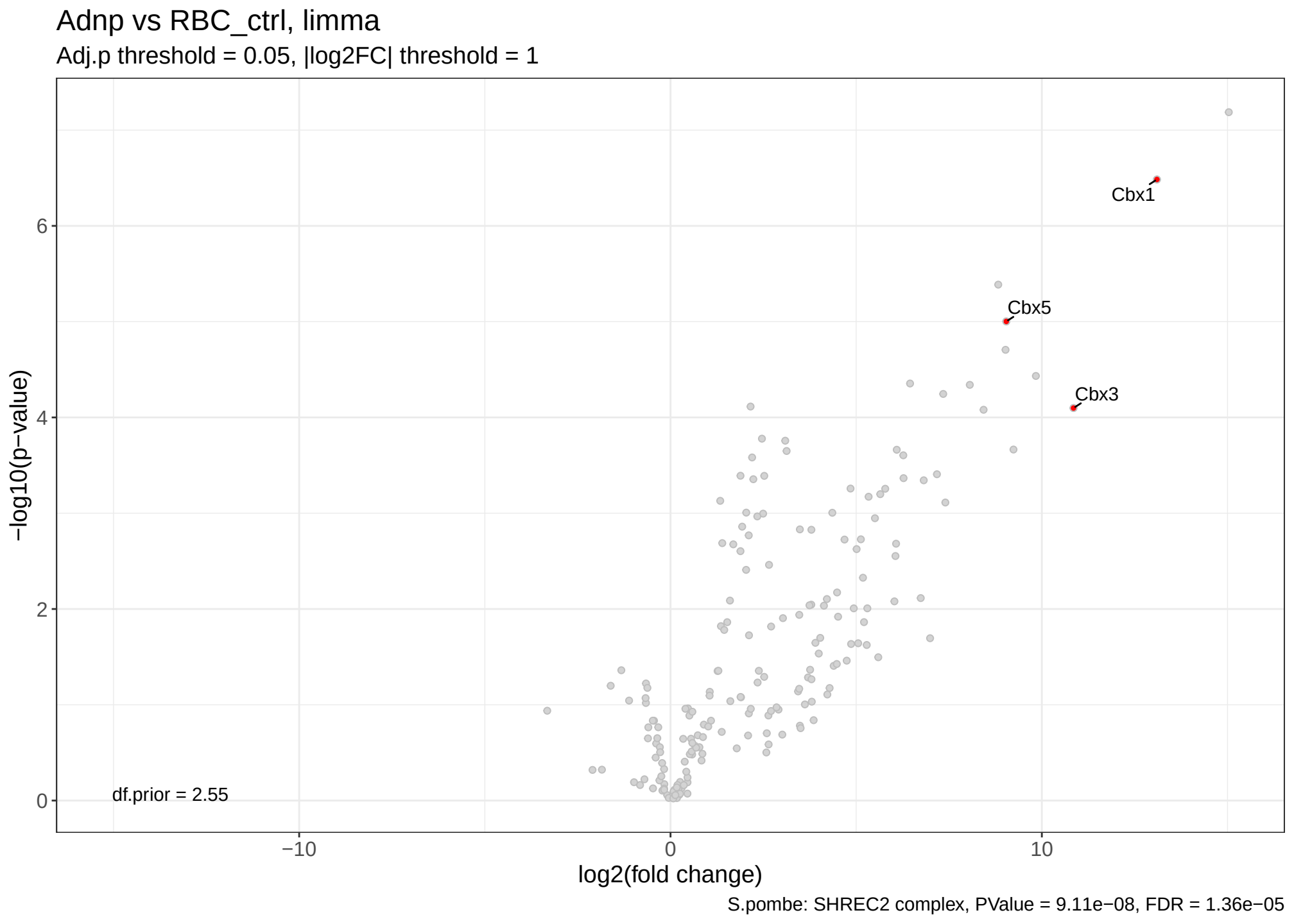
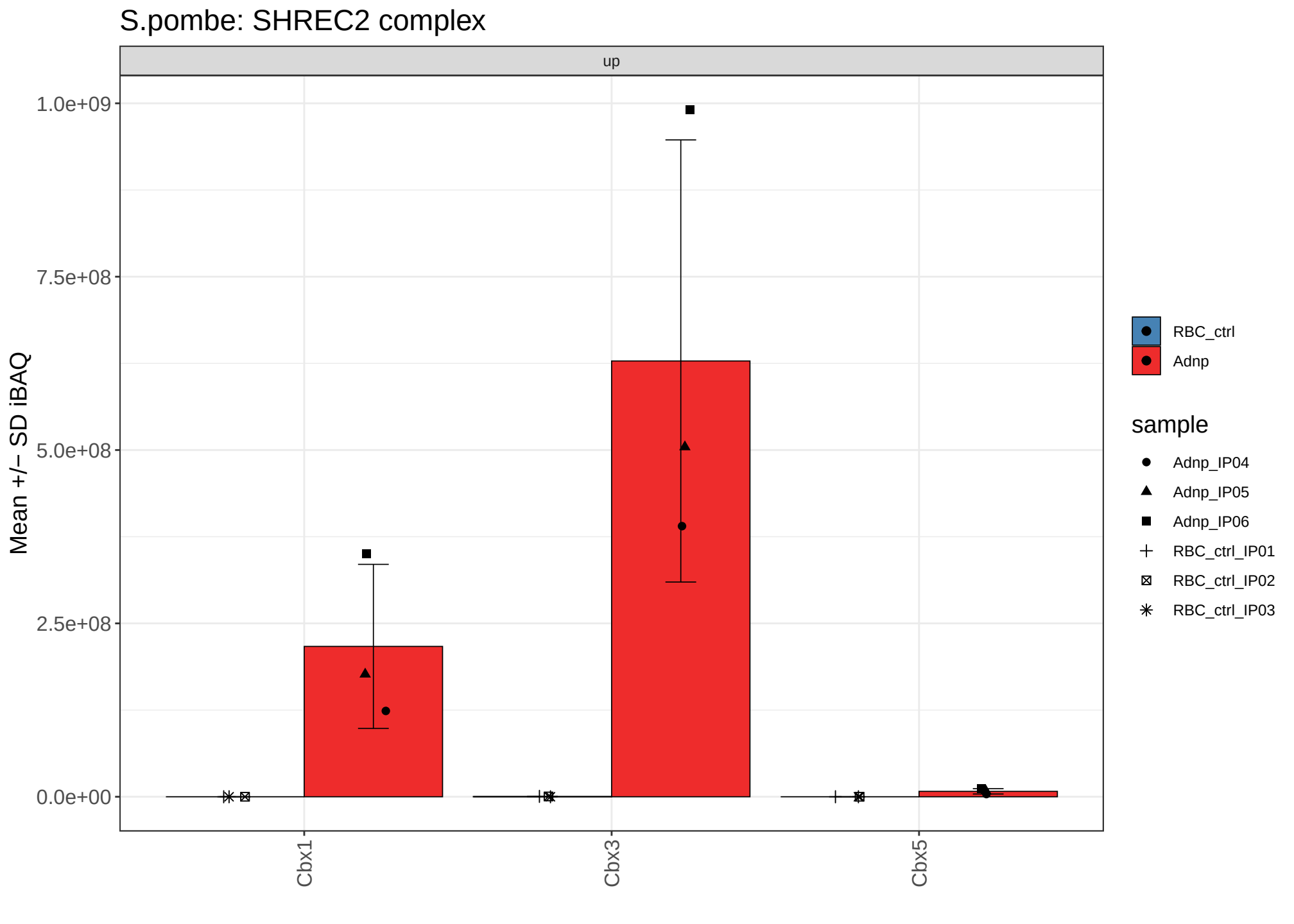
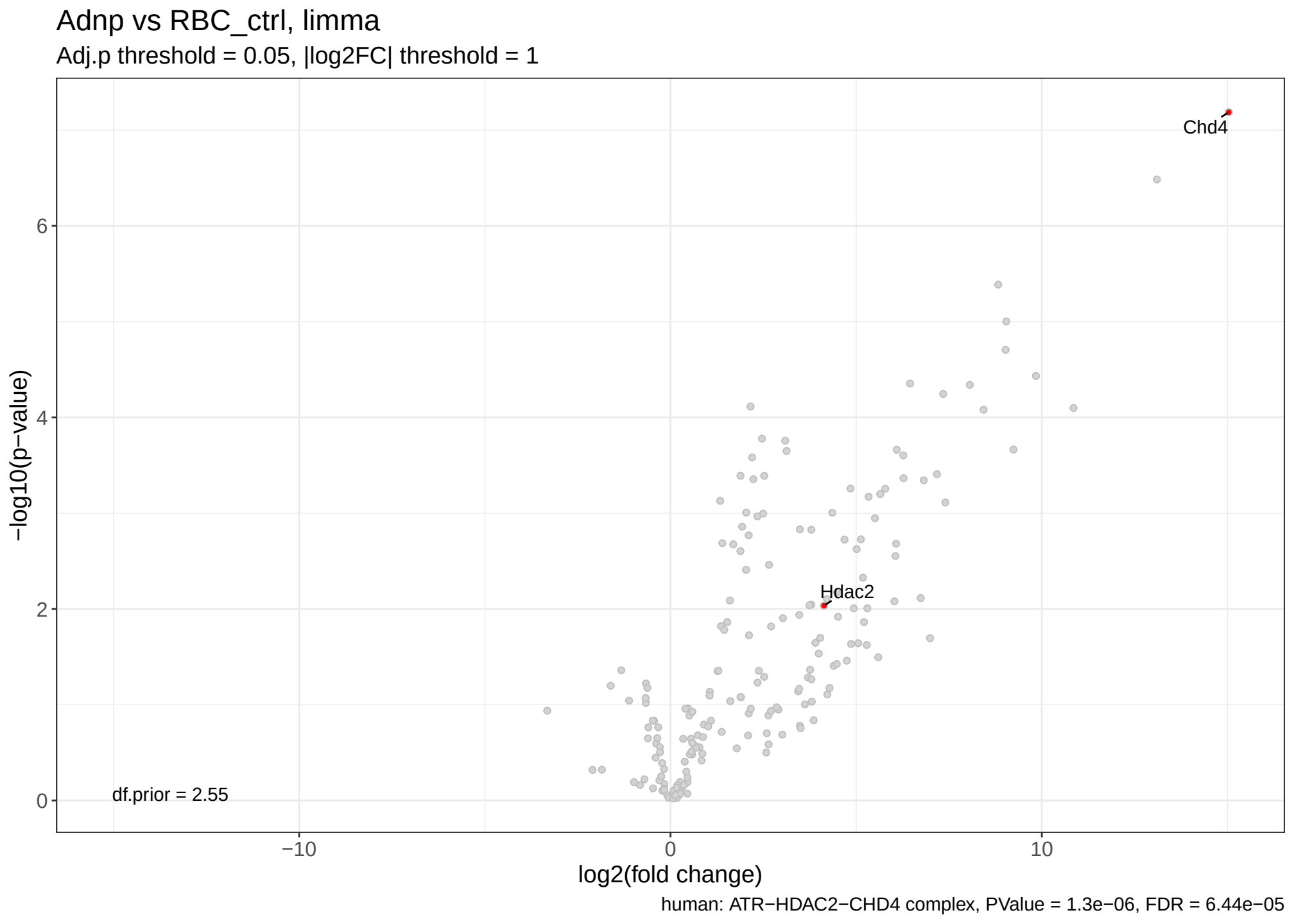
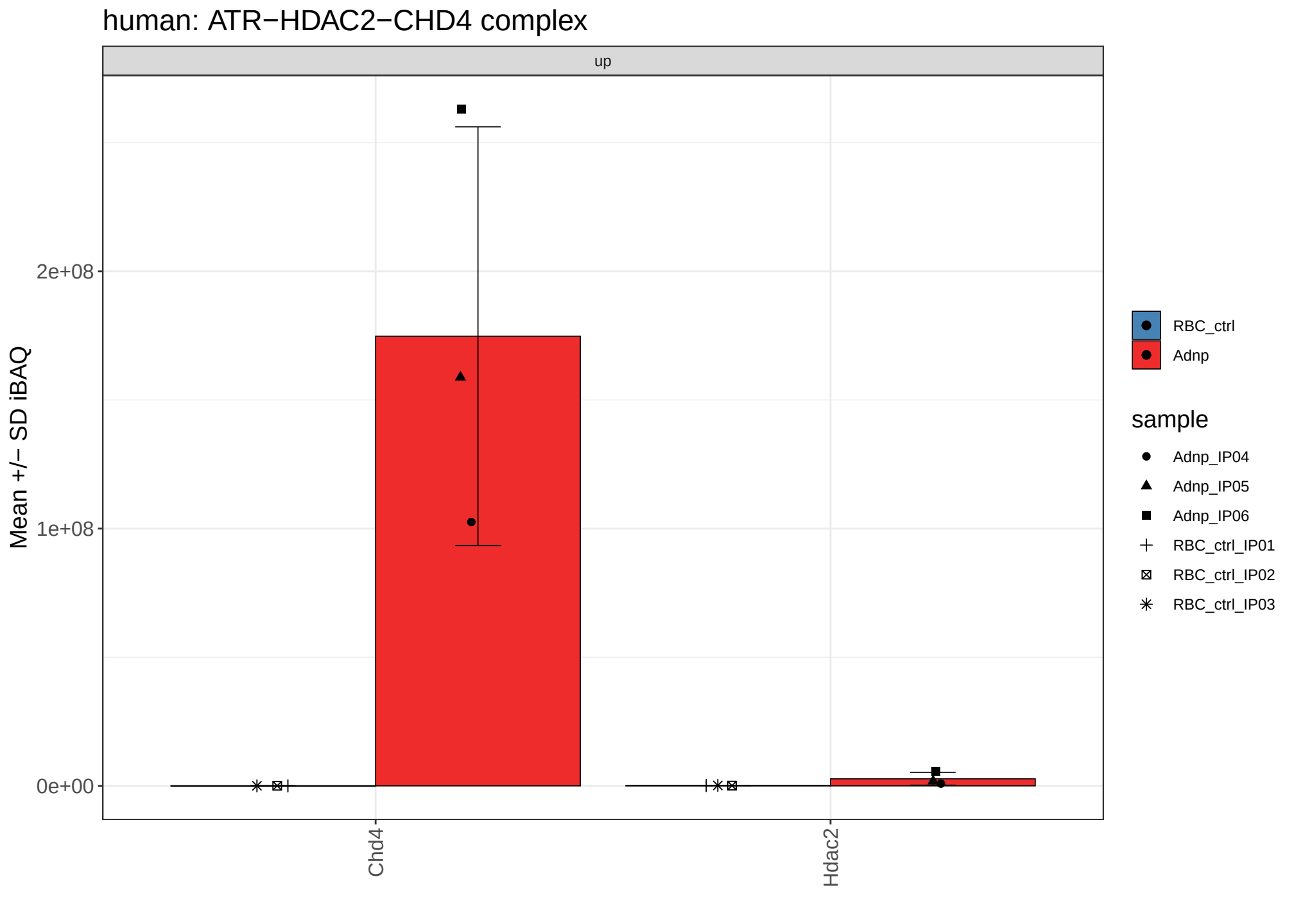
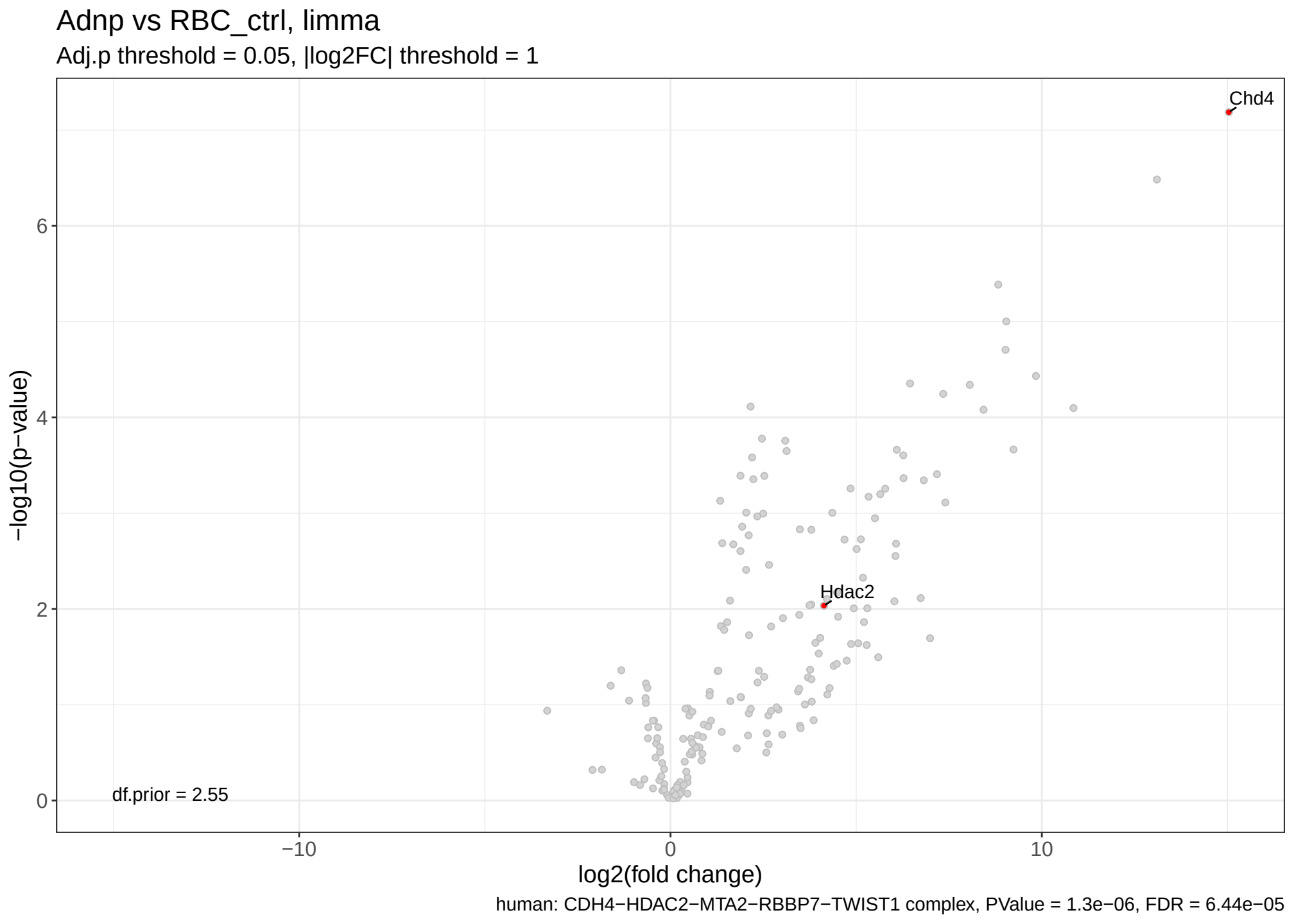
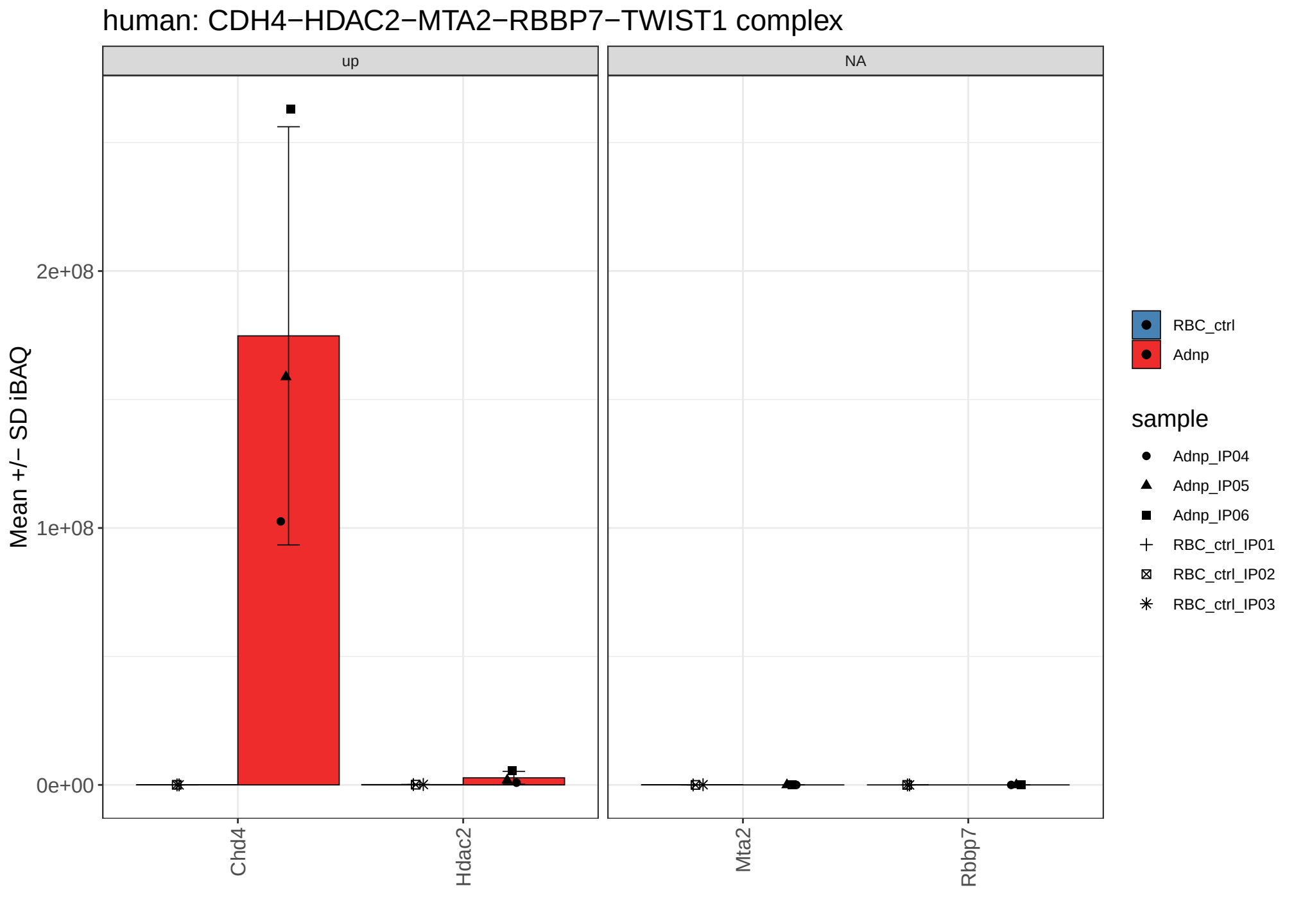

Adnp vs RBC_ctrl, limma
Adj.p threshold = 0.05, |log2FC| threshold = 1
Cbx1
Cbx5
Cbx3
df.prior = 2.55
−log10(p−value)
log2(fold change)
0
2
4
6
−10
0
10
S.pombe: SHREC2 complex, PValue = 9.11e−08, FDR = 1.36e−05

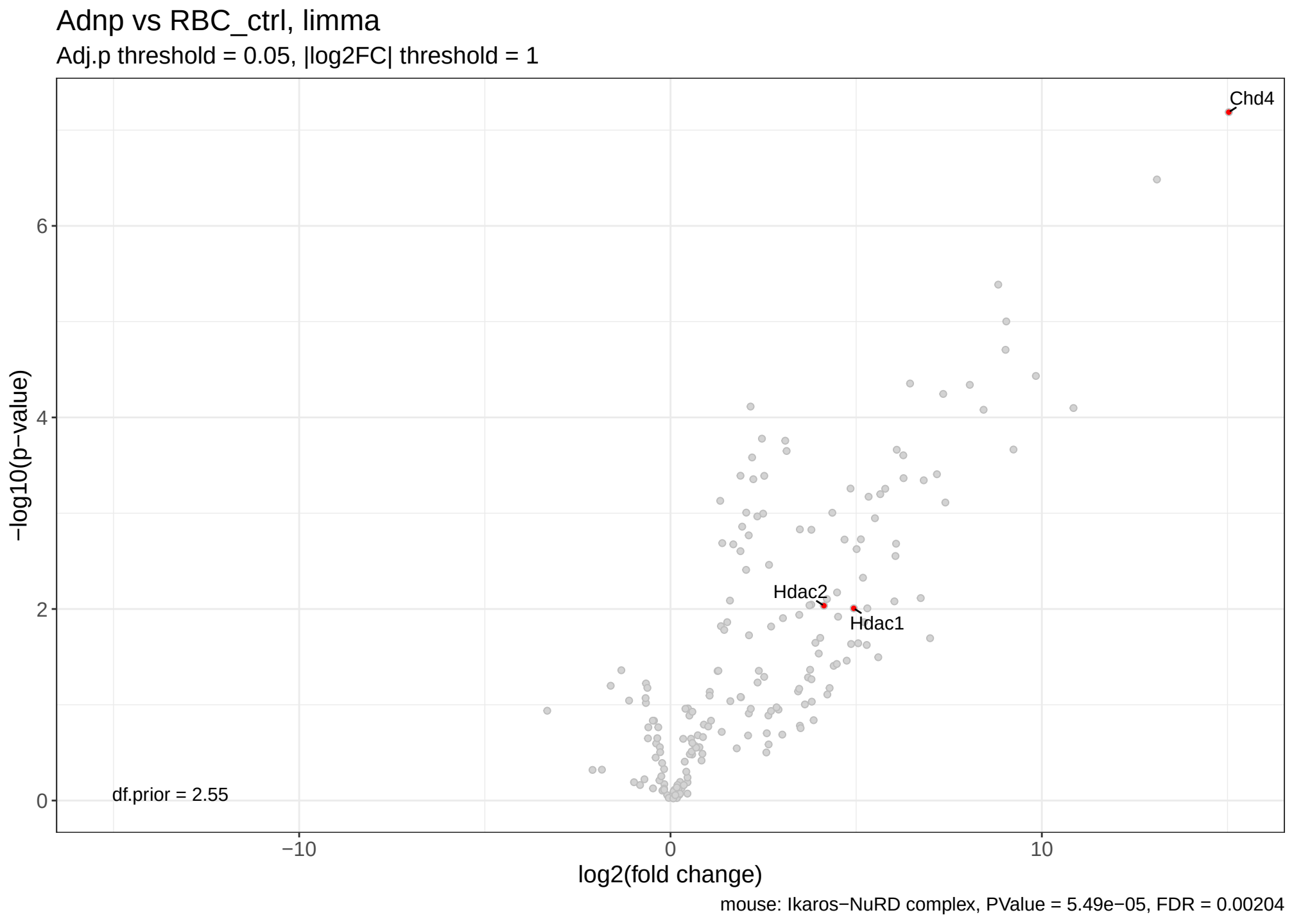

S.pombe: SHREC2 complex
up
Mean +/− SD iBAQ
1.0e+09
7.5e+08
5.0e+08
2.5e+08
0.0e+00
Cbx1
Cbx3
Cbx5
RBC_ctrl
Adnp
sample
Adnp_IP04
Adnp_IP05
Adnp_IP06
RBC_ctrl_IP01
RBC_ctrl_IP02
RBC_ctrl_IP03

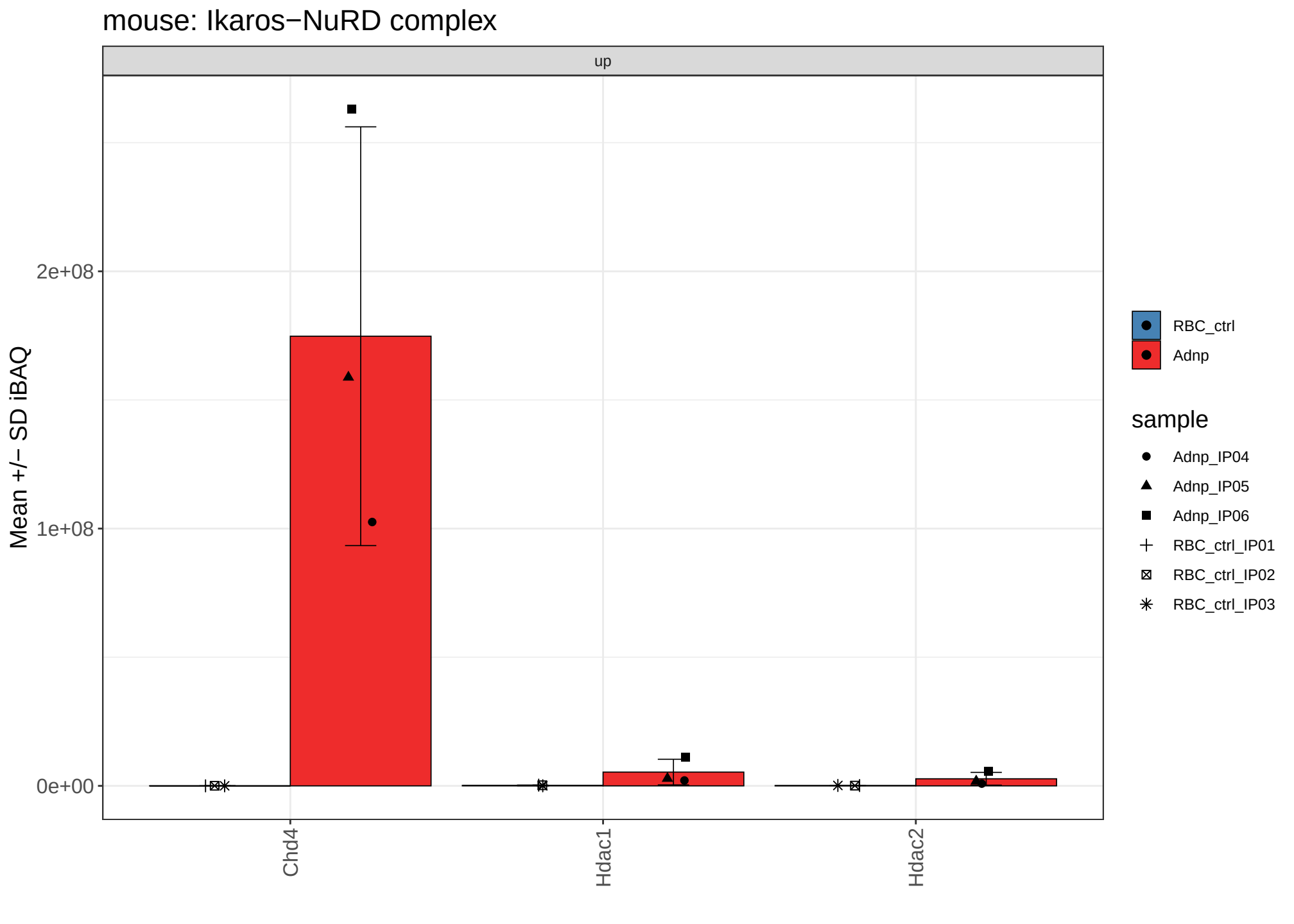

Adnp vs RBC_ctrl, limma
Adj.p threshold = 0.05, |log2FC| threshold = 1
Chd4
Hdac2
df.prior = 2.55
−log10(p−value)
log2(fold change)
0
2
4
6
−10
0
10
human: ATR−HDAC2−CHD4 complex, PValue = 1.3e−06, FDR = 6.44e−05

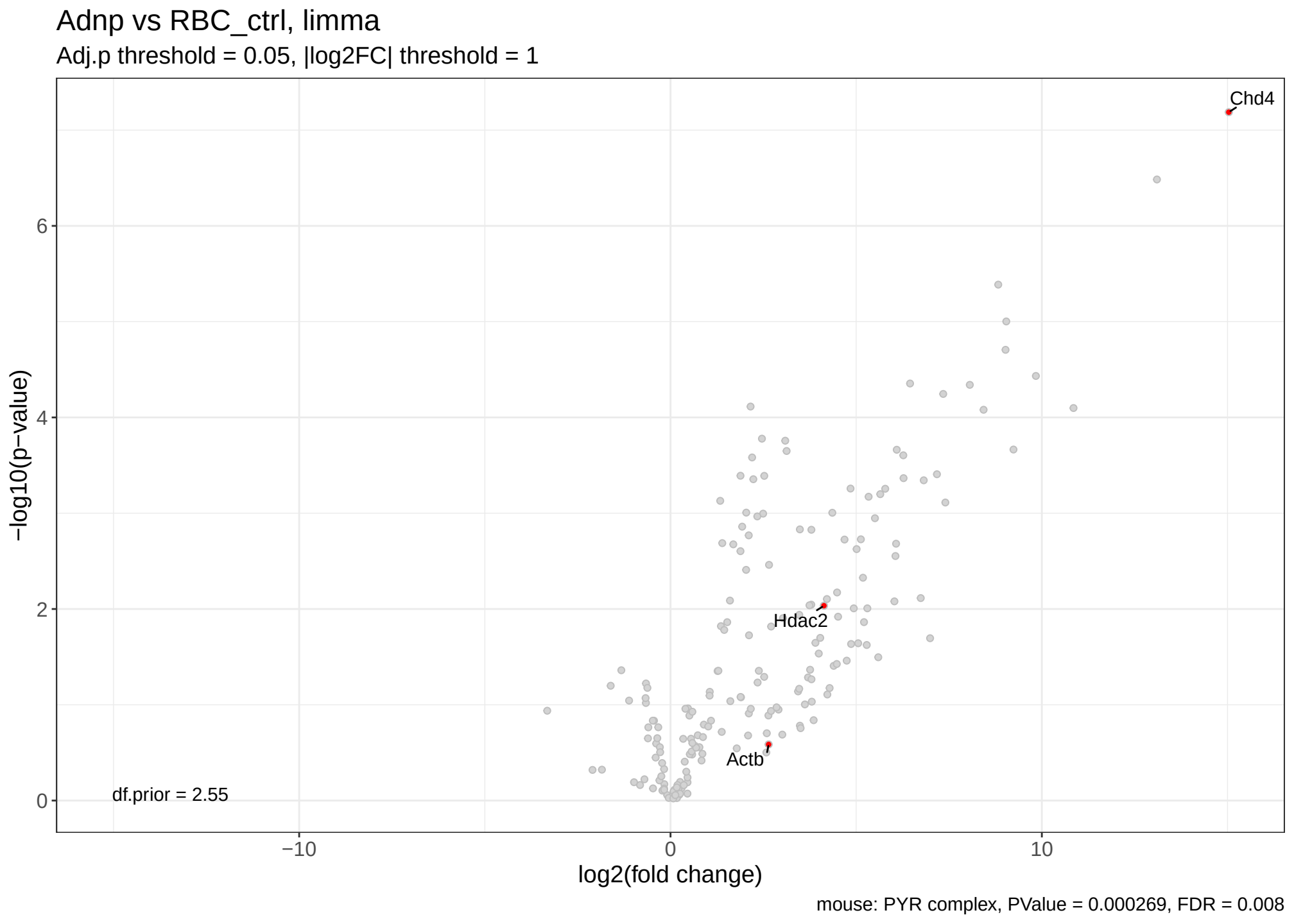

human: ATR−HDAC2−CHD4 complex
up
Mean +/− SD iBAQ
2e+08
1e+08
0e+00
Chd4
Hdac2
RBC_ctrl
Adnp
sample
Adnp_IP04
Adnp_IP05
Adnp_IP06
RBC_ctrl_IP01
RBC_ctrl_IP02
RBC_ctrl_IP03

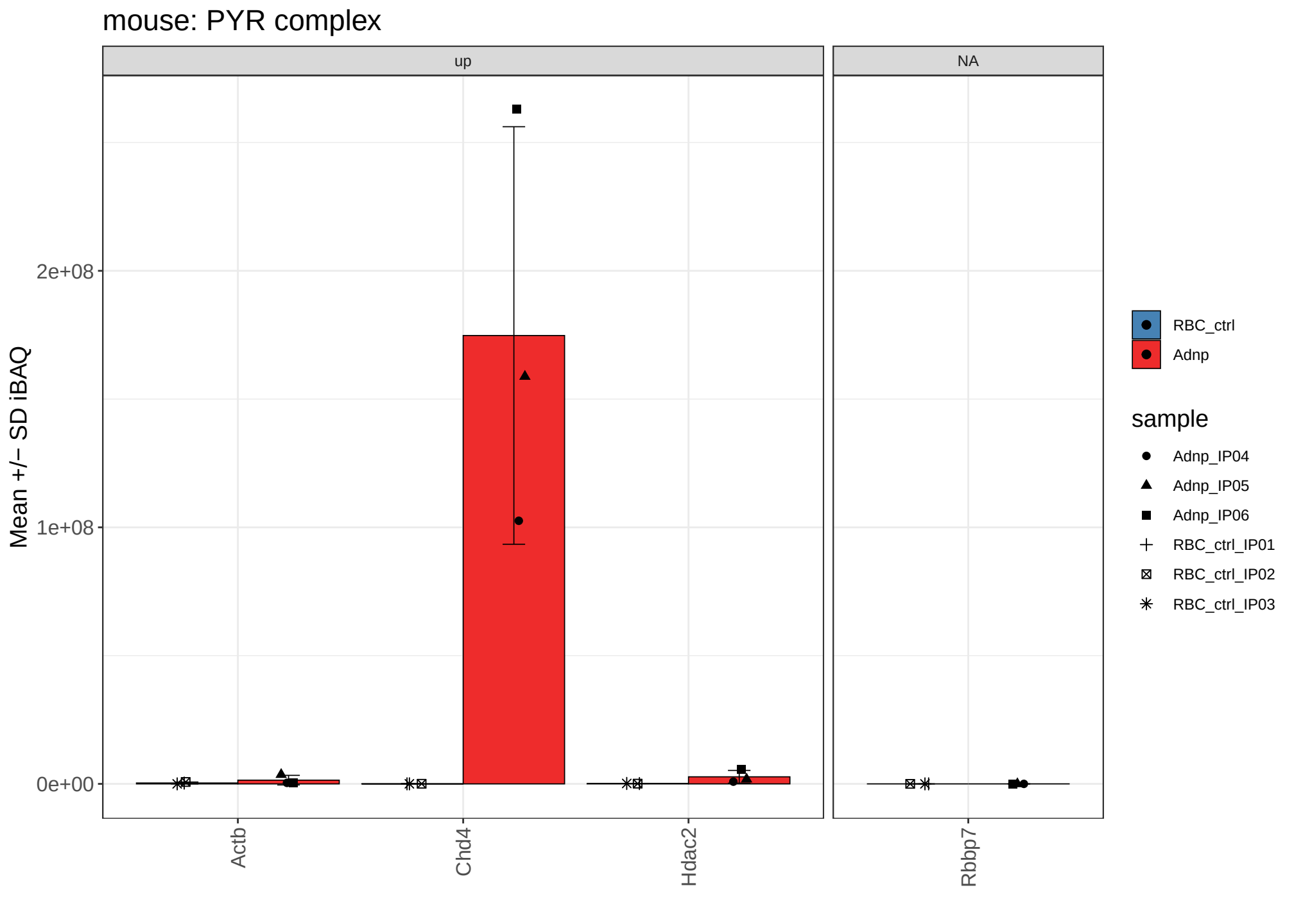

Adnp vs RBC_ctrl, limma
Adj.p threshold = 0.05, |log2FC| threshold = 1
Chd4
Hdac2
df.prior = 2.55
−log10(p−value)
log2(fold change)
human: CDH4−HDAC2−MTA2−RBBP7−TWIST1 complex, PValue = 1.3e−06, FDR = 6.44e−05

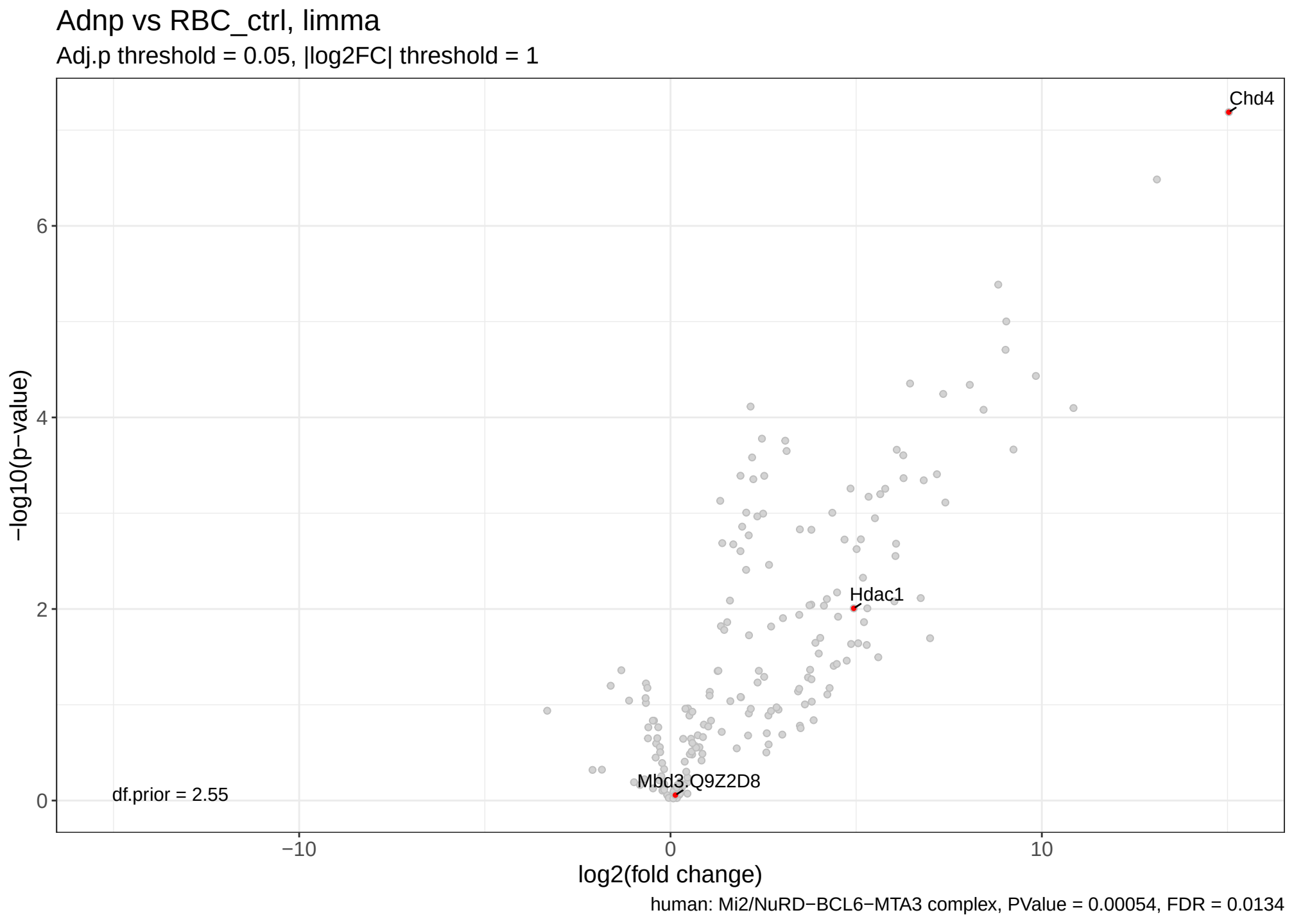

human: CDH4−HDAC2−MTA2−RBBP7−TWIST1 complex
up
NA
Mean +/− SD iBAQ
2e+08
1e+08
0e+00
Chd4
Hdac2
Mta2
Rbbp7
RBC_ctrl
Adnp
sample
Adnp_IP04
Adnp_IP05
Adnp_IP06
RBC_ctrl_IP01
RBC_ctrl_IP02
RBC_ctrl_IP03

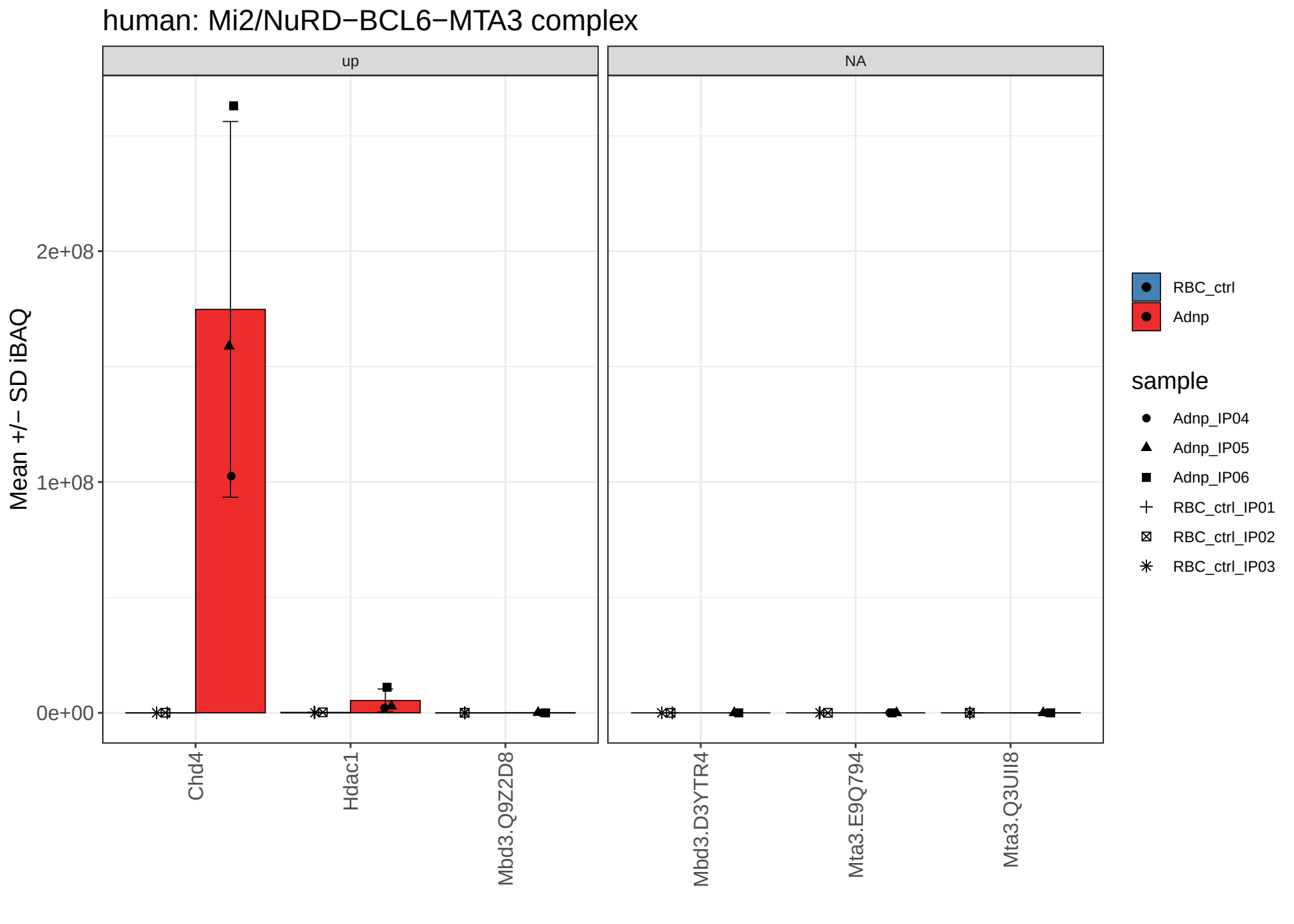

Adnp vs RBC_ctrl, limma
Adj.p threshold = 0.05, |log2FC| threshold = 1
Chd4
Hdac2
Hdac1
df.prior = 2.55
−log10(p−value)
log2(fold change)
0
2
4
6
−10
0
10
mouse: Ikaros−NuRD complex, PValue = 5.49e−05, FDR = 0.00204

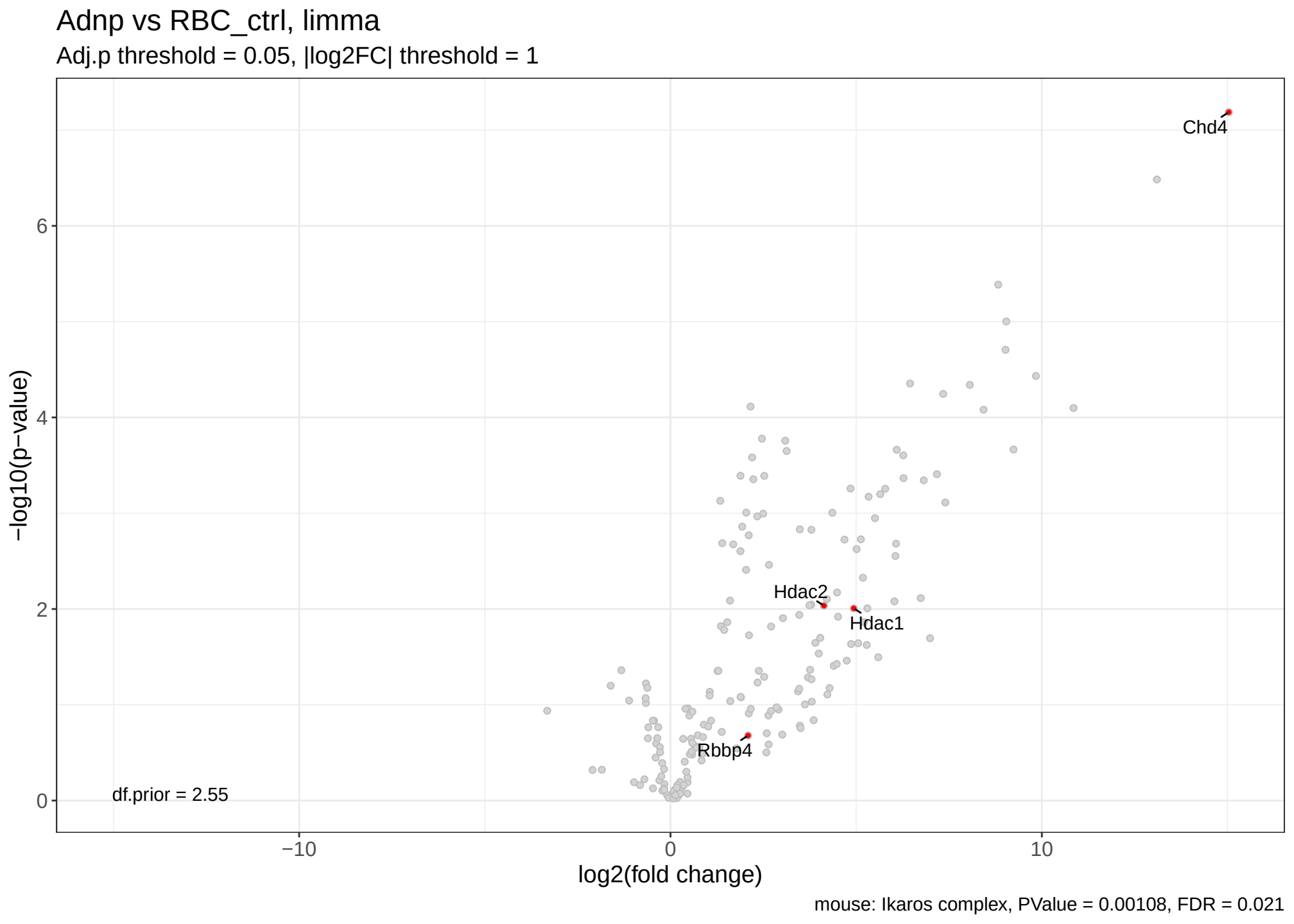

mouse: Ikaros−NuRD complex
up
Mean +/− SD iBAQ
2e+08
1e+08
0e+00
Chd4
Hdac1
Hdac2
RBC_ctrl
Adnp
sample
Adnp_IP04
Adnp_IP05
Adnp_IP06
RBC_ctrl_IP01
RBC_ctrl_IP02
RBC_ctrl_IP03

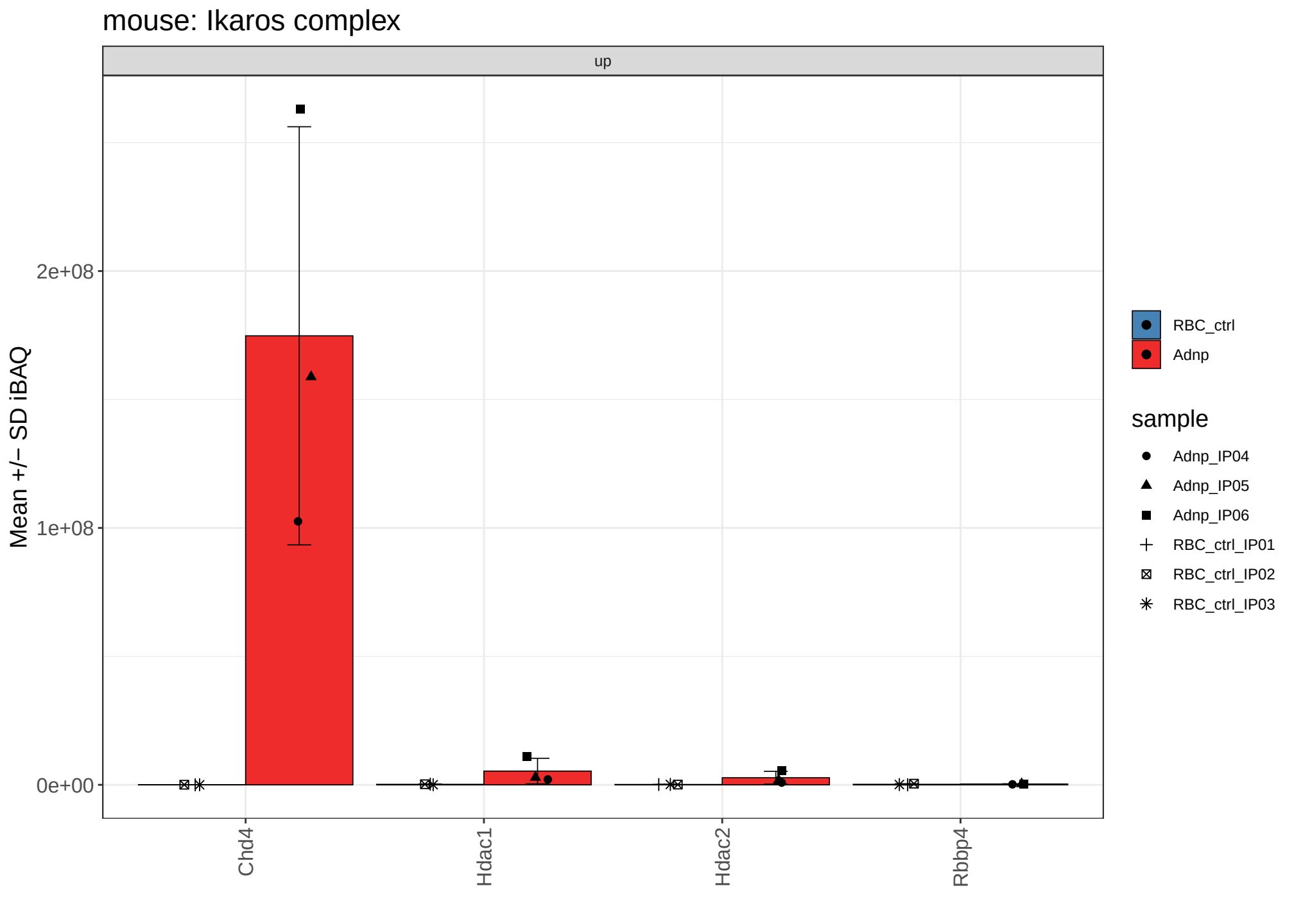

Adnp vs RBC_ctrl, limma
Adj.p threshold = 0.05, |log2FC| threshold = 1
Chd4
Hdac2
Actb
df.prior = 2.55
−log10(p−value)
log2(fold change)
0
2
4
6
−10
0
10
mouse: PYR complex, PValue = 0.000269, FDR = 0.008

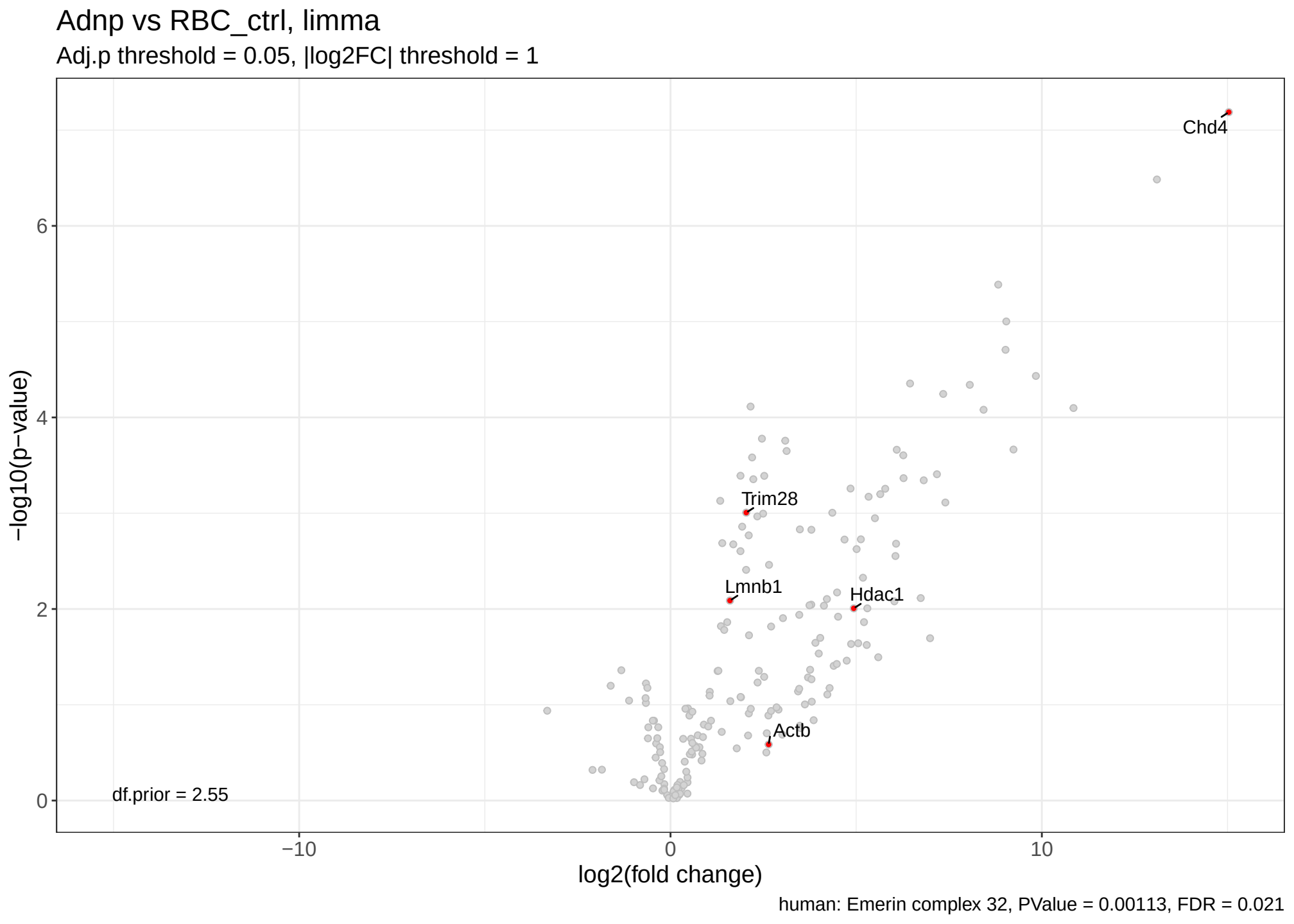

mouse: PYR complex
up
NA
Mean +/− SD iBAQ
2e+08
1e+08
0e+00
Actb
Chd4
Hdac2
Rbbp7
RBC_ctrl
Adnp
sample
Adnp_IP04
Adnp_IP05
Adnp_IP06
RBC_ctrl_IP01
RBC_ctrl_IP02
RBC_ctrl_IP03

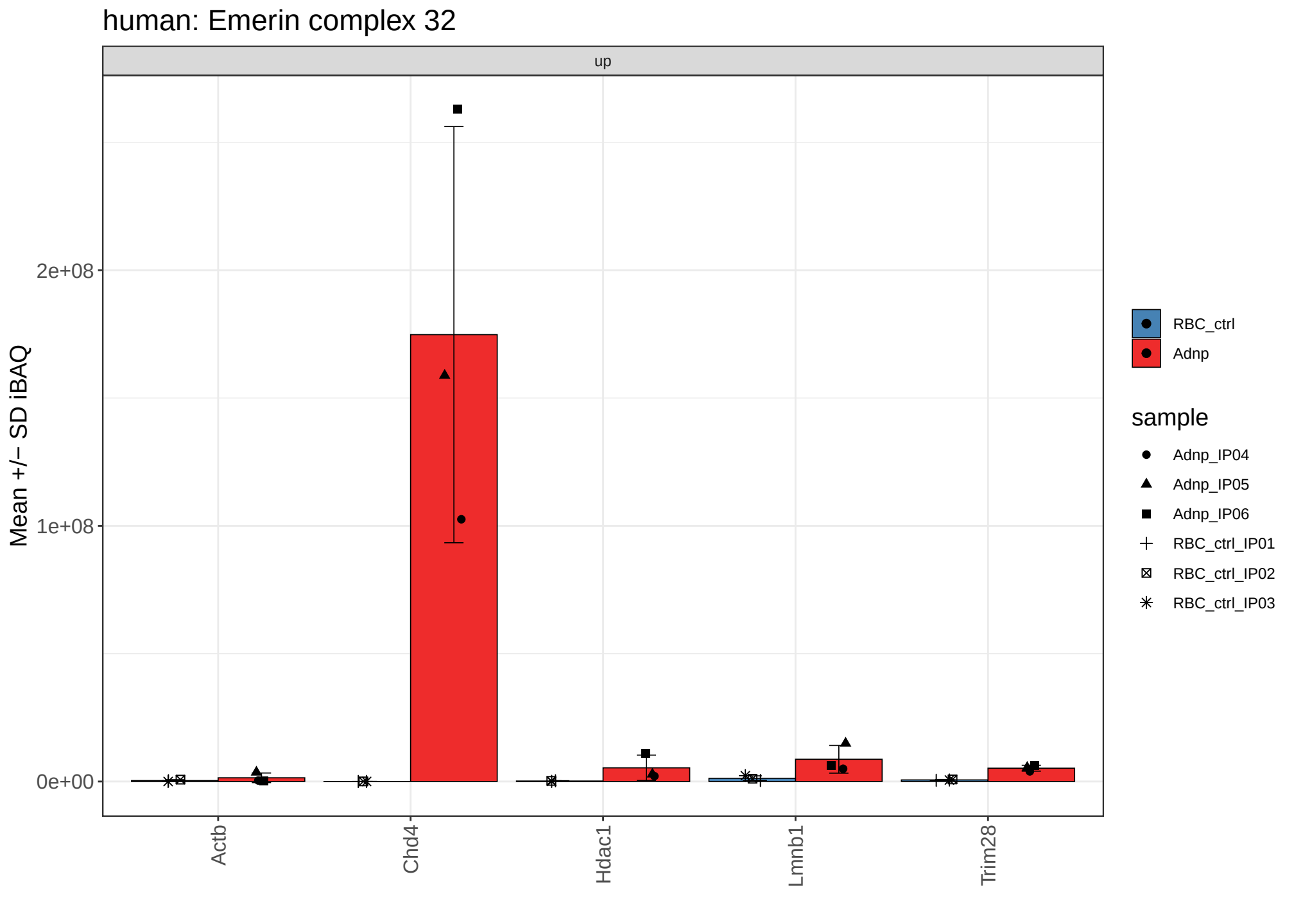

Adnp vs RBC_ctrl, limma
Adj.p threshold = 0.05, |log2FC| threshold = 1
Chd4
Hdac1
Mbd3.Q9Z2D8
df.prior = 2.55
−log10(p−value)
log2(fold change)
human: Mi2/NuRD−BCL6−MTA3 complex, PValue = 0.00054, FDR = 0.0134

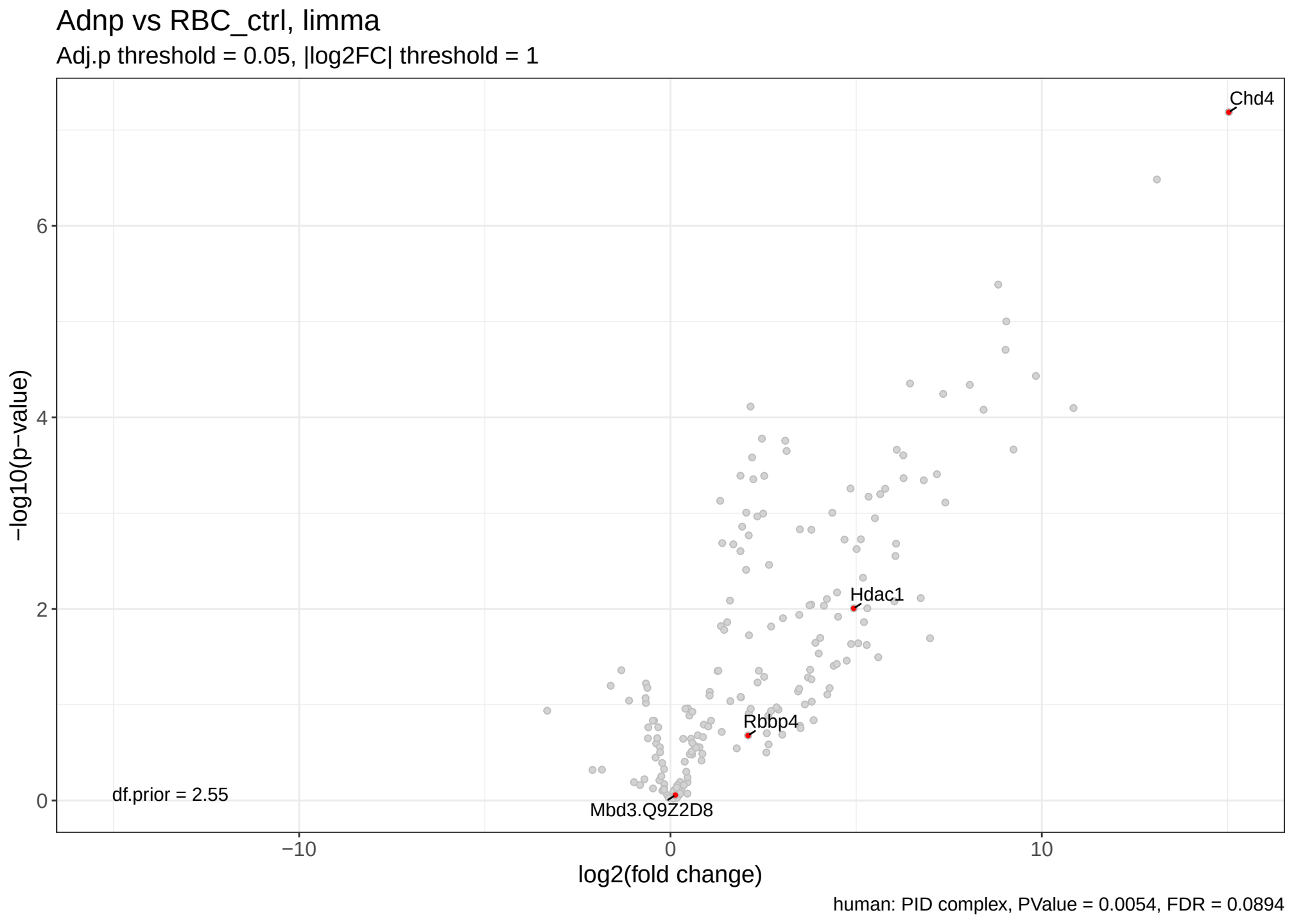

human: Mi2/NuRD−BCL6−MTA3 complex
up
NA
Mean +/− SD iBAQ
2e+08
1e+08
0e+00
Chd4
Hdac1
Mbd3.Q9Z2D8
Mbd3.D3YTR4
Mta3.E9Q794
Mta3.Q3UII8
RBC_ctrl
Adnp
sample
Adnp_IP04
Adnp_IP05
Adnp_IP06
RBC_ctrl_IP01
RBC_ctrl_IP02
RBC_ctrl_IP03

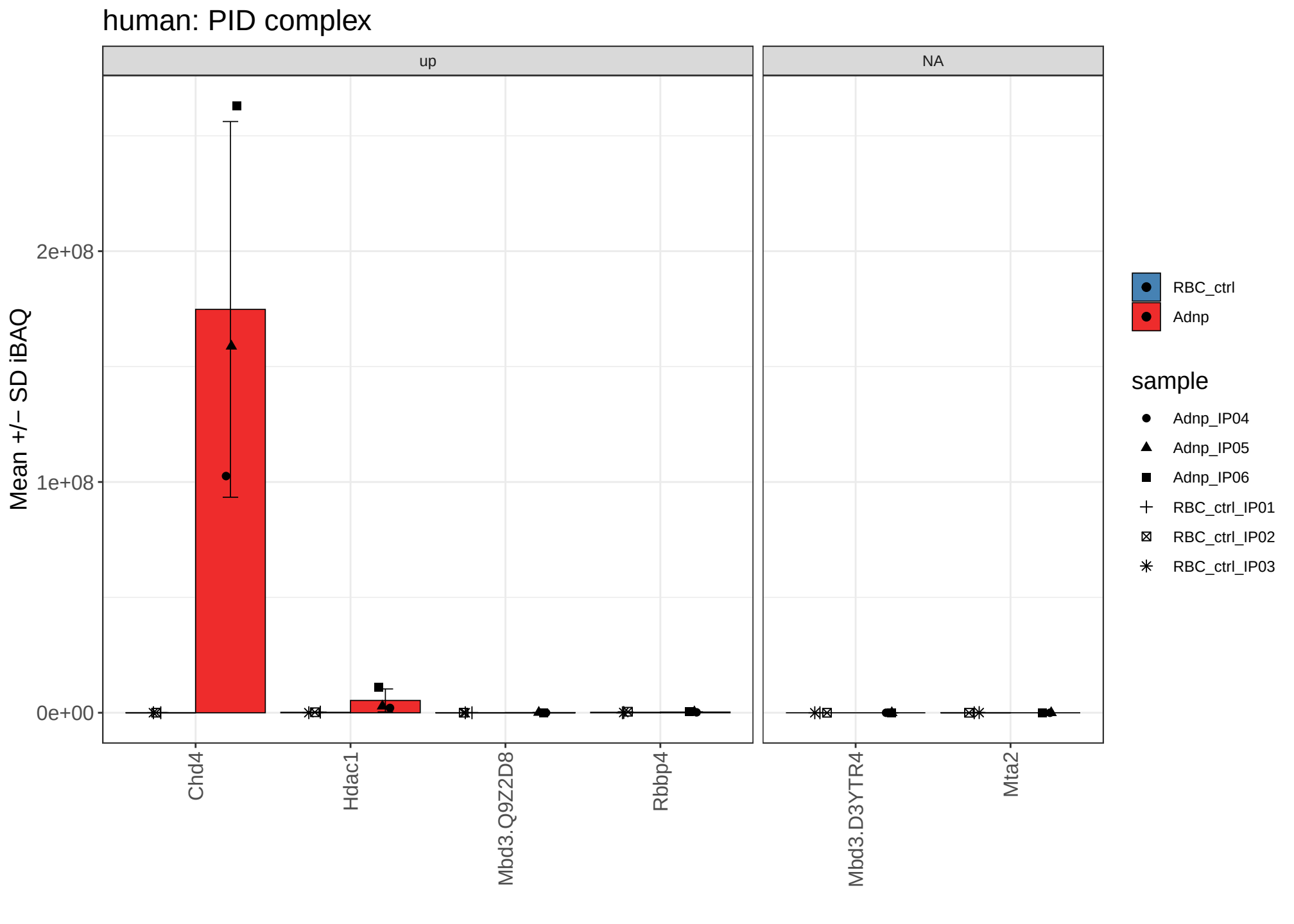

Adnp vs RBC_ctrl, limma
Adj.p threshold = 0.05, |log2FC| threshold = 1
Chd4
Hdac2
Hdac1
Rbbp4
df.prior = 2.55
−log10(p−value)
log2(fold change)
0
2
4
6
−10
0
10
mouse: Ikaros complex, PValue = 0.00108, FDR = 0.021


mouse: Ikaros complex
up
Mean +/− SD iBAQ
2e+08
1e+08
0e+00
Chd4
Hdac1
Hdac2
Rbbp4
RBC_ctrl
Adnp
sample
Adnp_IP04
Adnp_IP05
Adnp_IP06
RBC_ctrl_IP01
RBC_ctrl_IP02
RBC_ctrl_IP03


Adnp vs RBC_ctrl, limma
Adj.p threshold = 0.05, |log2FC| threshold = 1
Chd4
Trim28
Lmnb1
Hdac1
Actb
df.prior = 2.55
−log10(p−value)
log2(fold change)
6
4
2
0
−10
0
10
human: Emerin complex 32, PValue = 0.00113, FDR = 0.021


human: Emerin complex 32
up
Mean +/− SD iBAQ
2e+08
1e+08
0e+00
Actb
Chd4
Hdac1
Lmnb1
Trim28
RBC_ctrl
Adnp
sample
Adnp_IP04
Adnp_IP05
Adnp_IP06
RBC_ctrl_IP01
RBC_ctrl_IP02
RBC_ctrl_IP03


Adnp vs RBC_ctrl, limma
Adj.p threshold = 0.05, |log2FC| threshold = 1
Chd4
Hdac1
Rbbp4
Mbd3.Q9Z2D8
df.prior = 2.55
−log10(p−value)
log2(fold change)
6
4
2
0
−10
0
10
human: PID complex, PValue = 0.0054, FDR = 0.0894


human: PID complex
up
NA
Mean +/− SD iBAQ
2e+08
1e+08
0e+00
Chd4
Hdac1
Mbd3.Q9Z2D8
Rbbp4
Mbd3.D3YTR4
Mta2
RBC_ctrl
Adnp
sample
Adnp_IP04
Adnp_IP05
Adnp_IP06
RBC_ctrl_IP01
RBC_ctrl_IP02
RBC_ctrl_IP03