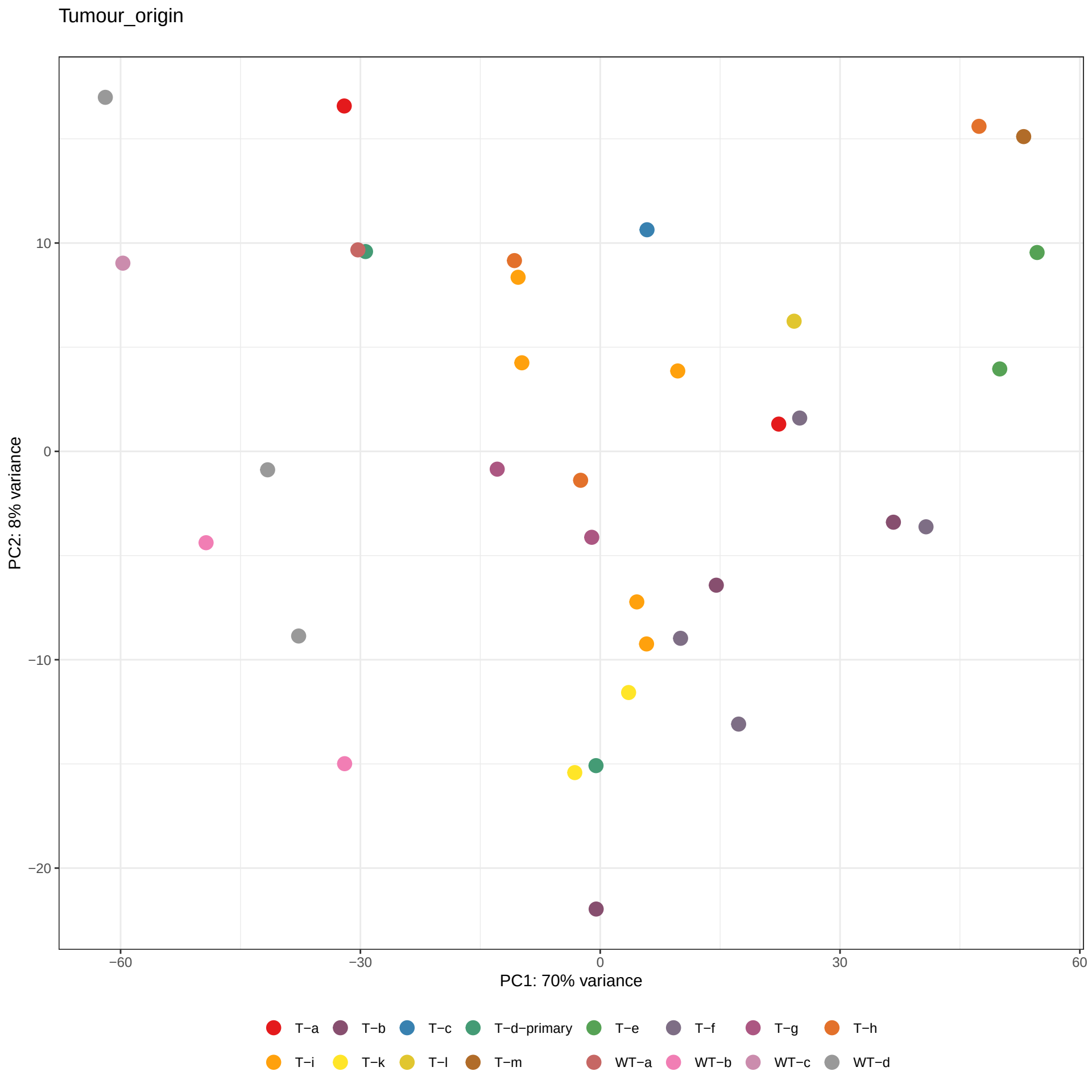
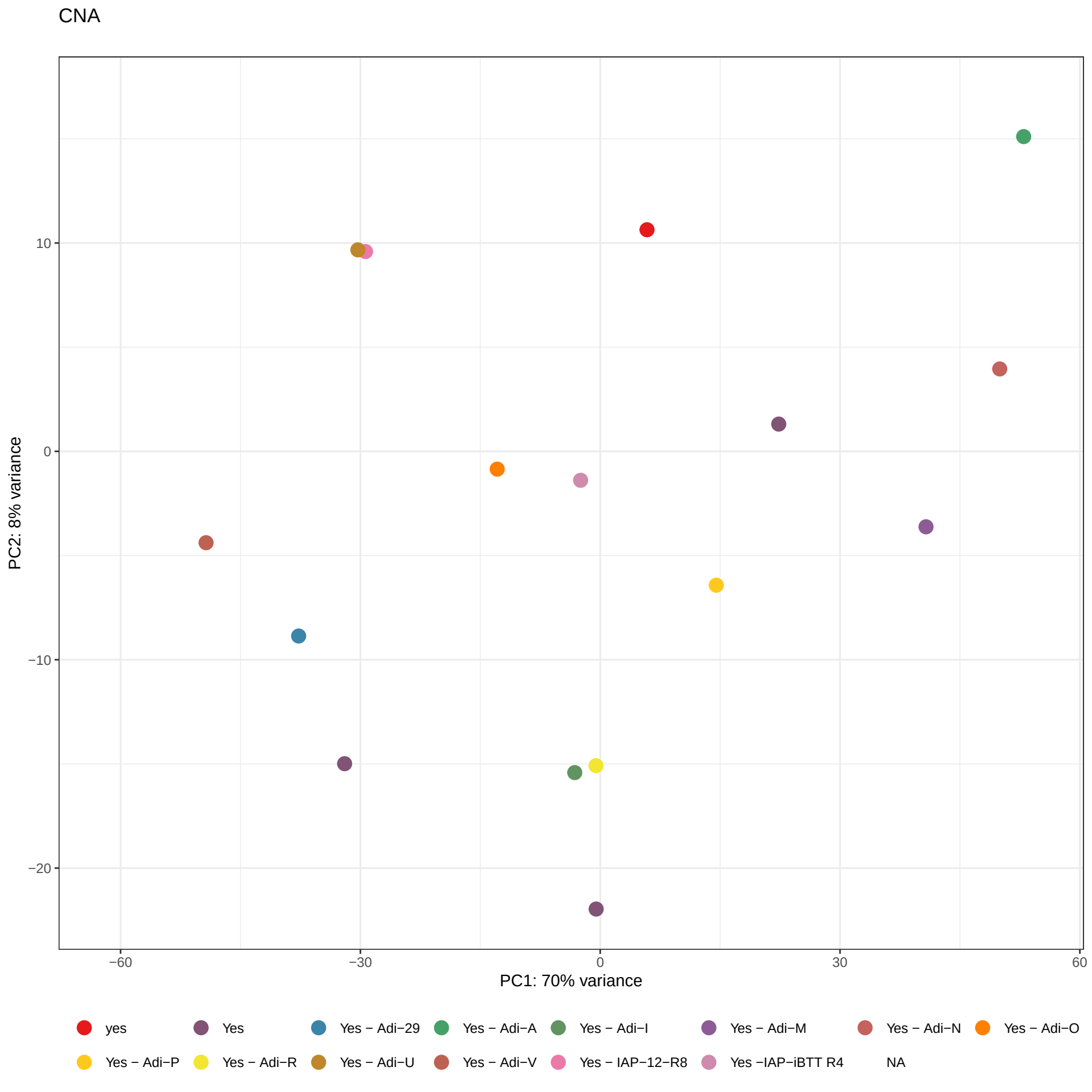
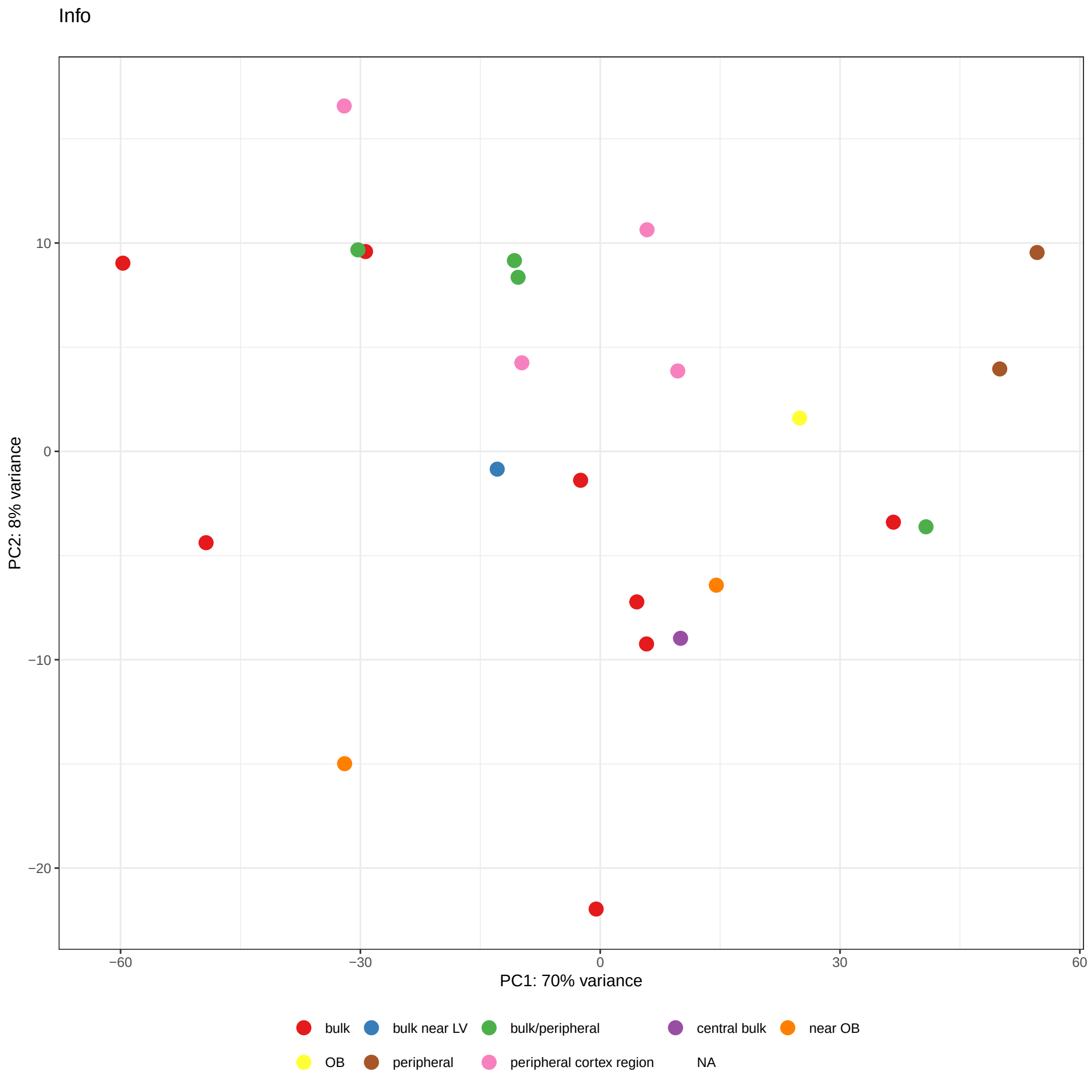
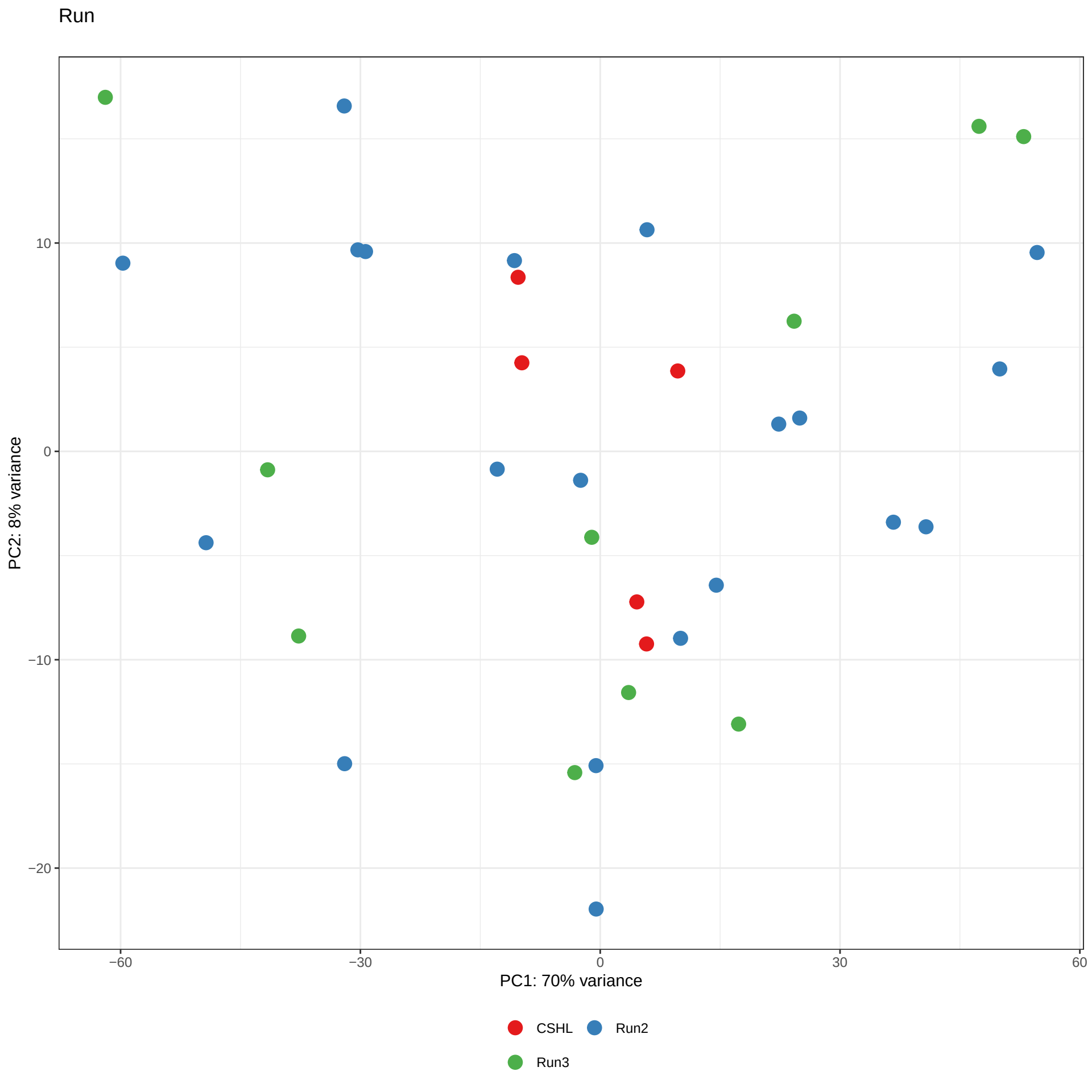
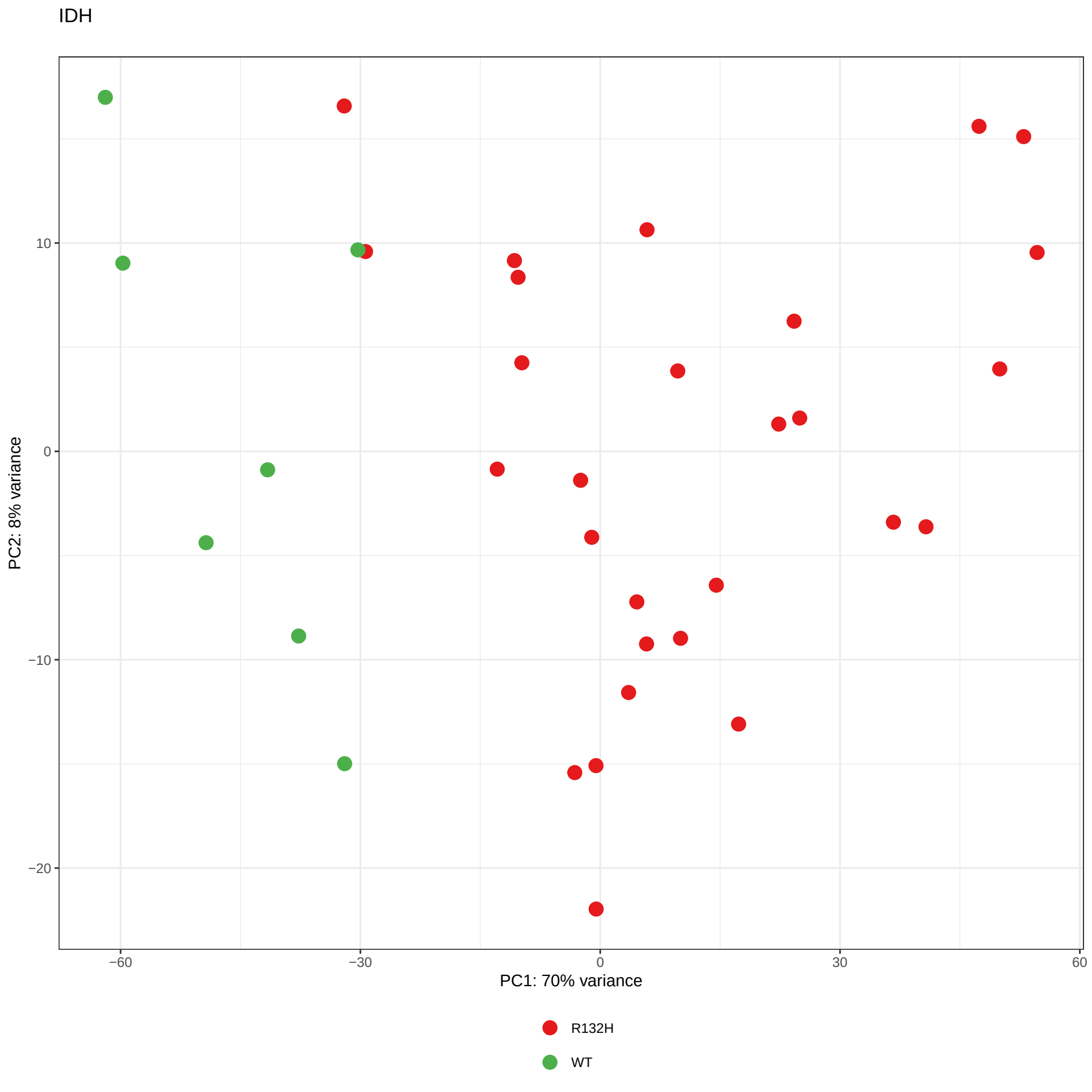
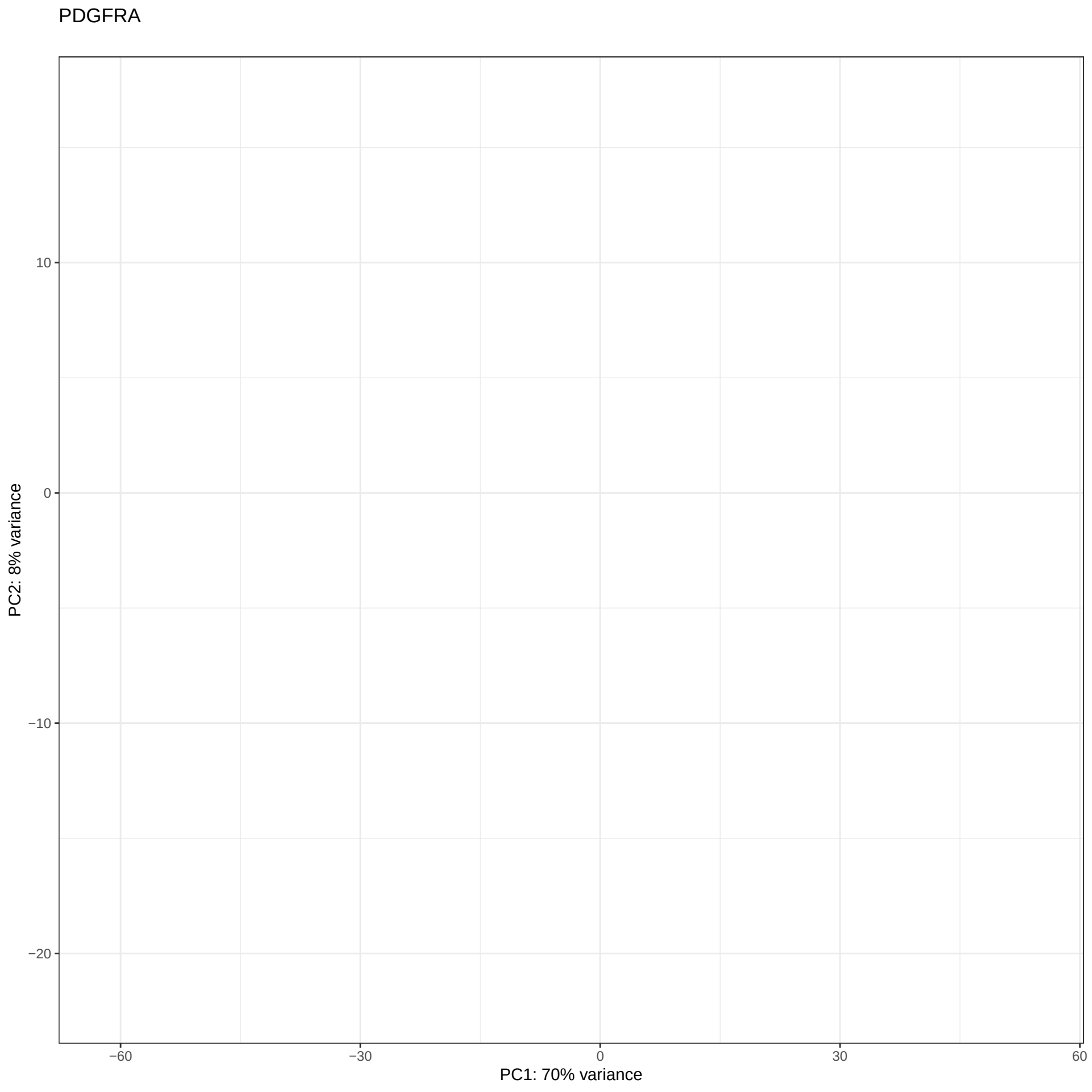

Tumour_origin
PC2: 8% variance
10
0
−10
−20
−60
−30
0
30
60
PC1: 70% variance
T−a
T−b
T−c
T−d−primary
T−e
T−f
T−g
T−h
T−i
T−k
T−l
T−m
WT−a
WT−b
WT−c
WT−d

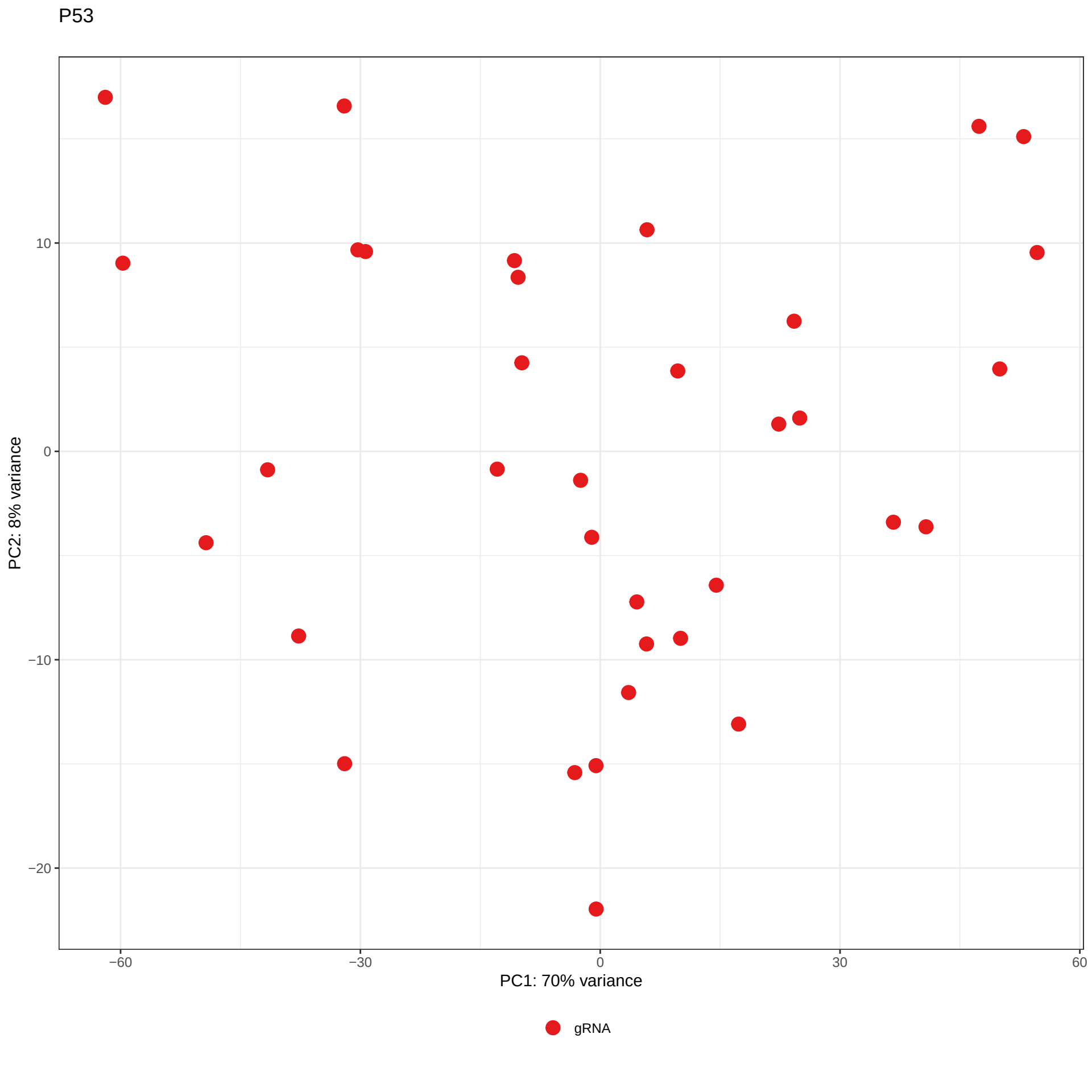

CNA
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance
yes
Yes
Yes − Adi−29
Yes − Adi−A
Yes − Adi−I
Yes − Adi−M
Yes − Adi−N
Yes − Adi−O
Yes − Adi−P
Yes − Adi−R
Yes − Adi−U
Yes − Adi−V
Yes − IAP−12−R8
Yes −IAP−iBTT R4
NA

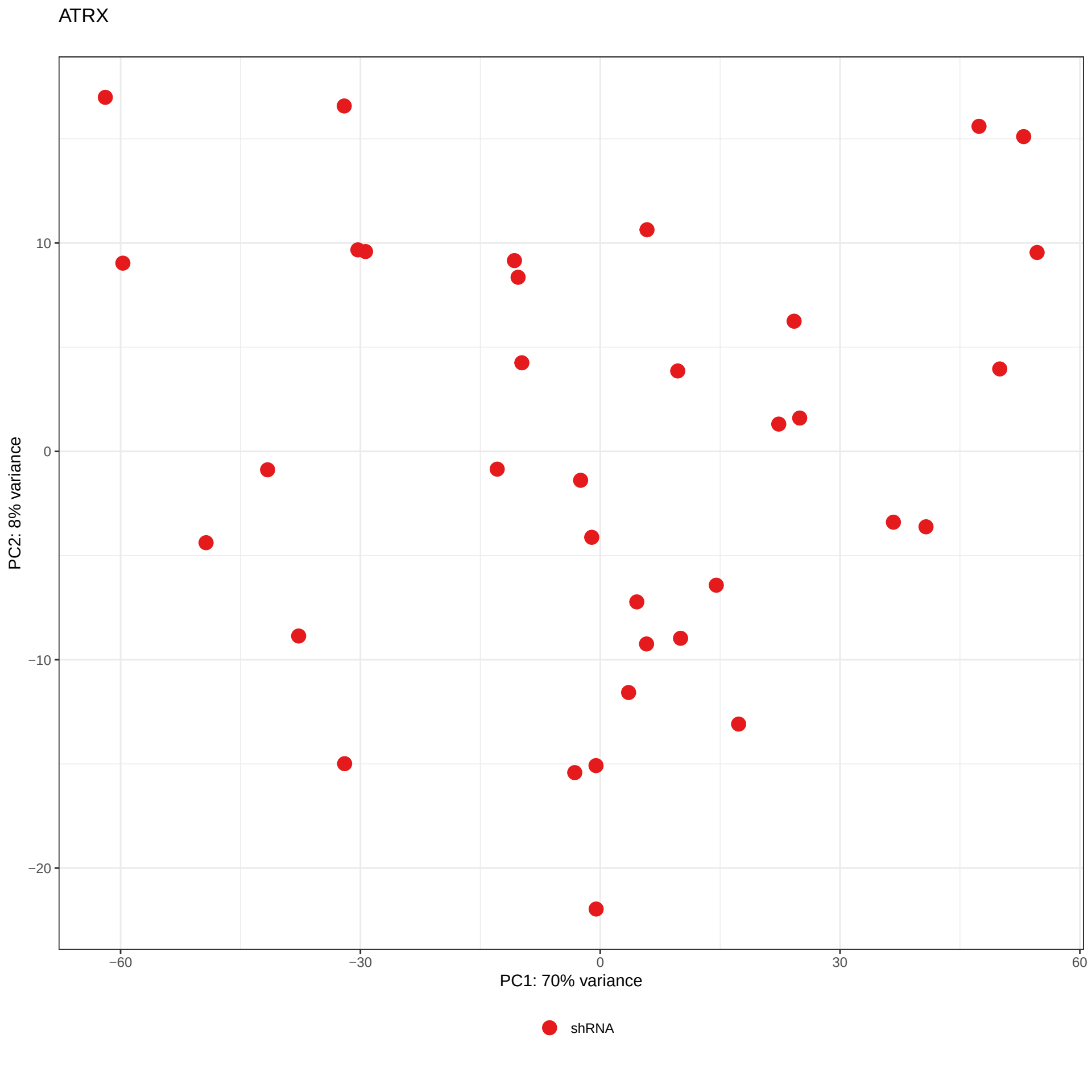

Info
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance
bulk
bulk near LV
bulk/peripheral
central bulk
near OB
OB
peripheral
peripheral cortex region
NA

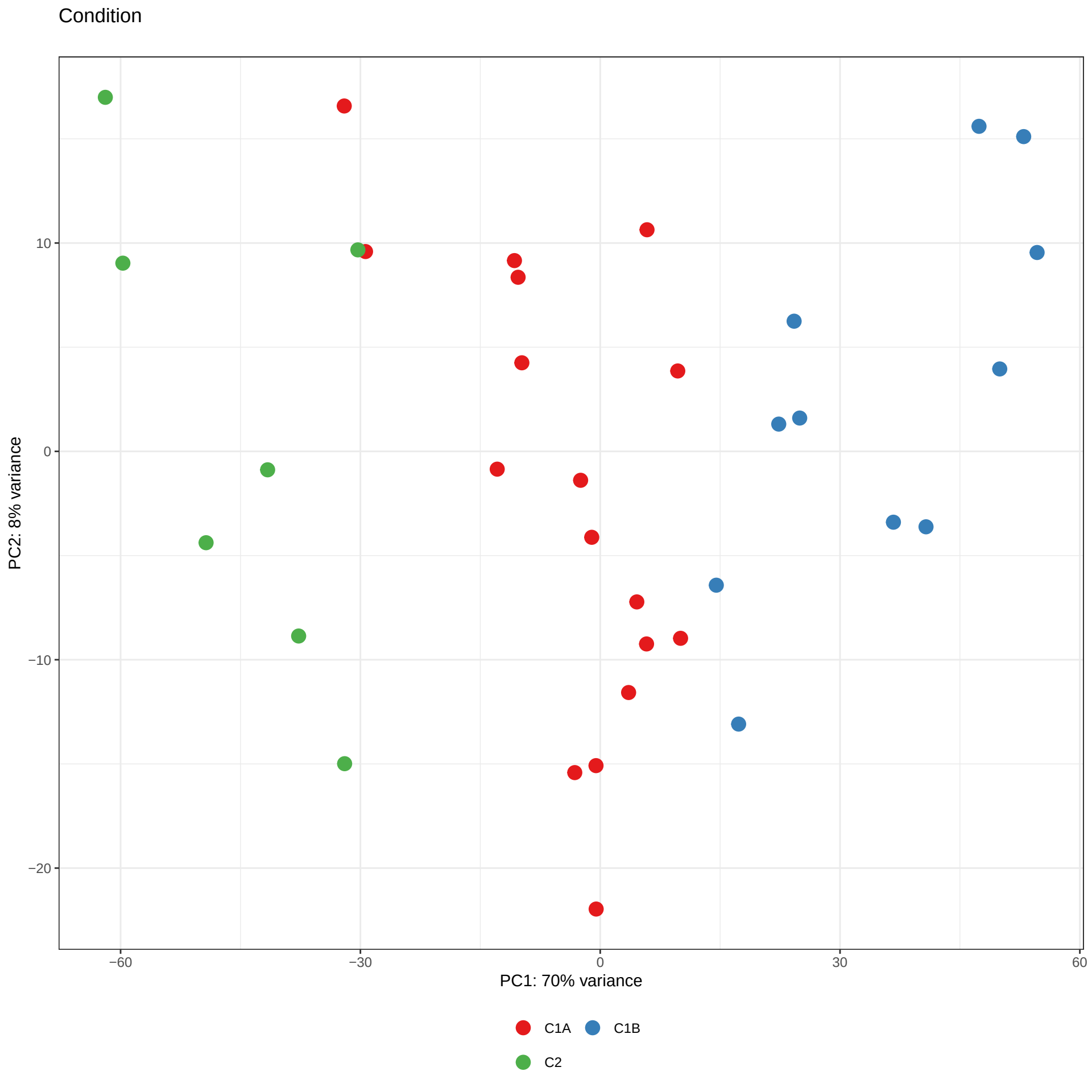

Run
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance
CSHL
Run2
Run3

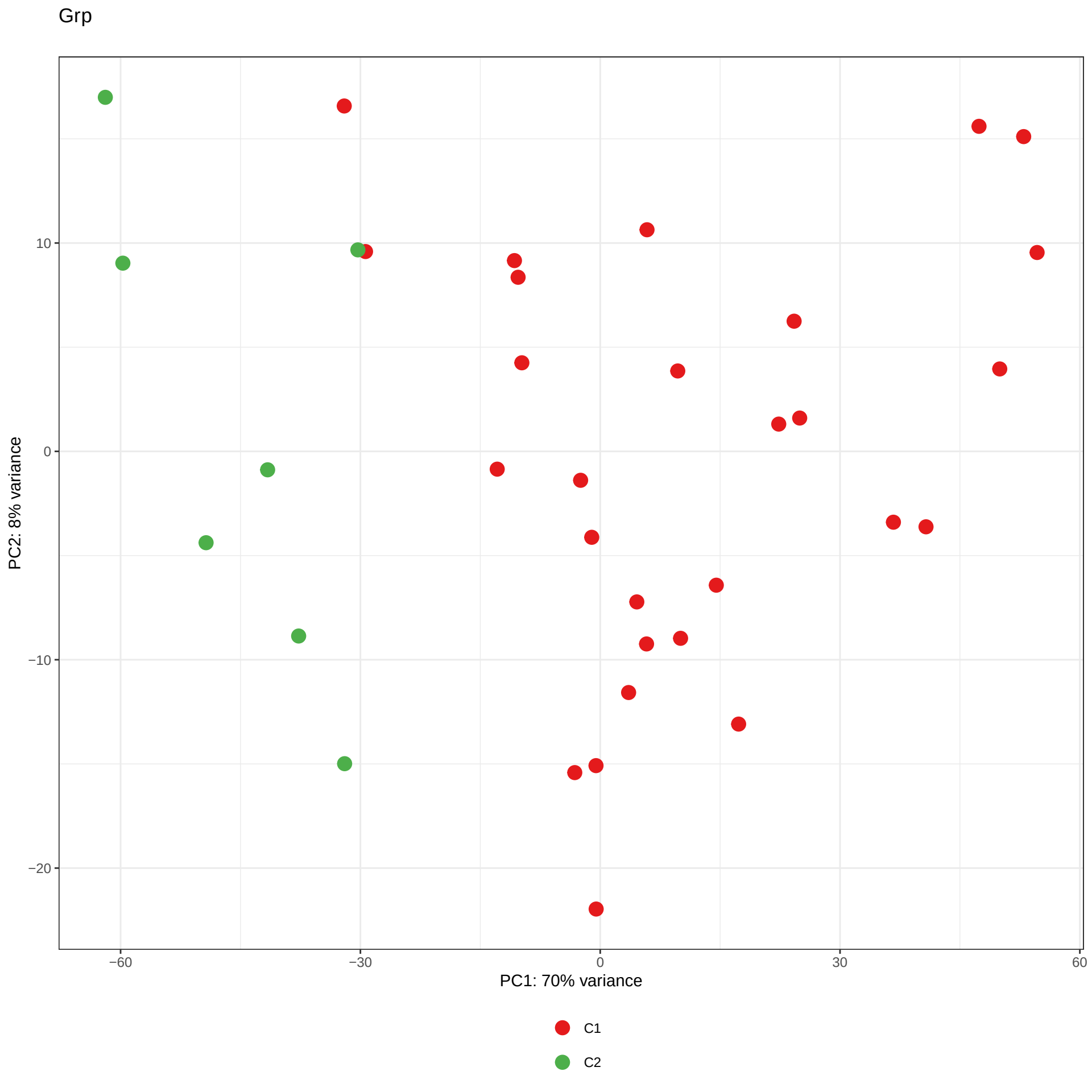

IDH
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance
R132H
WT

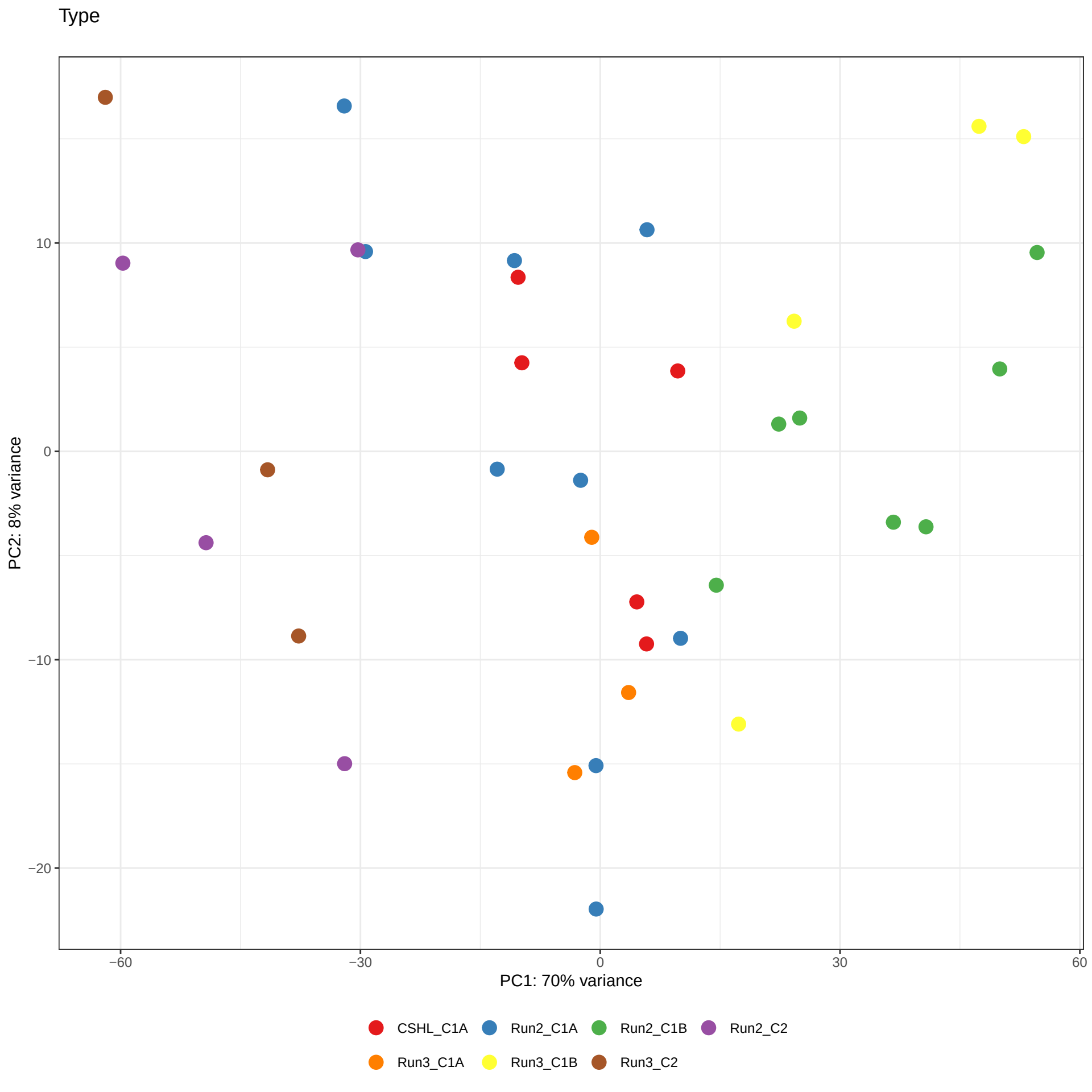

PDGFRA
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance

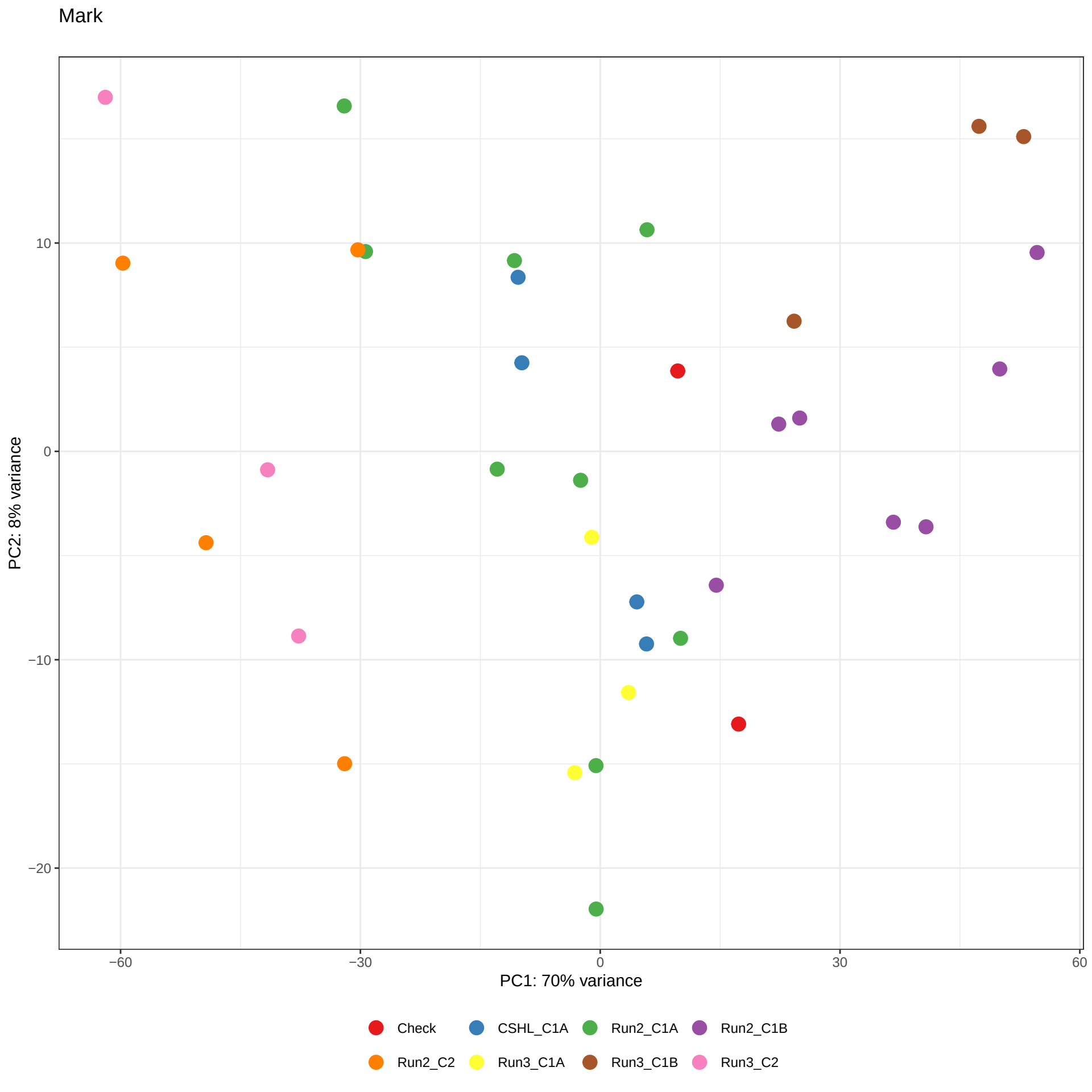

P53
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance
gRNA

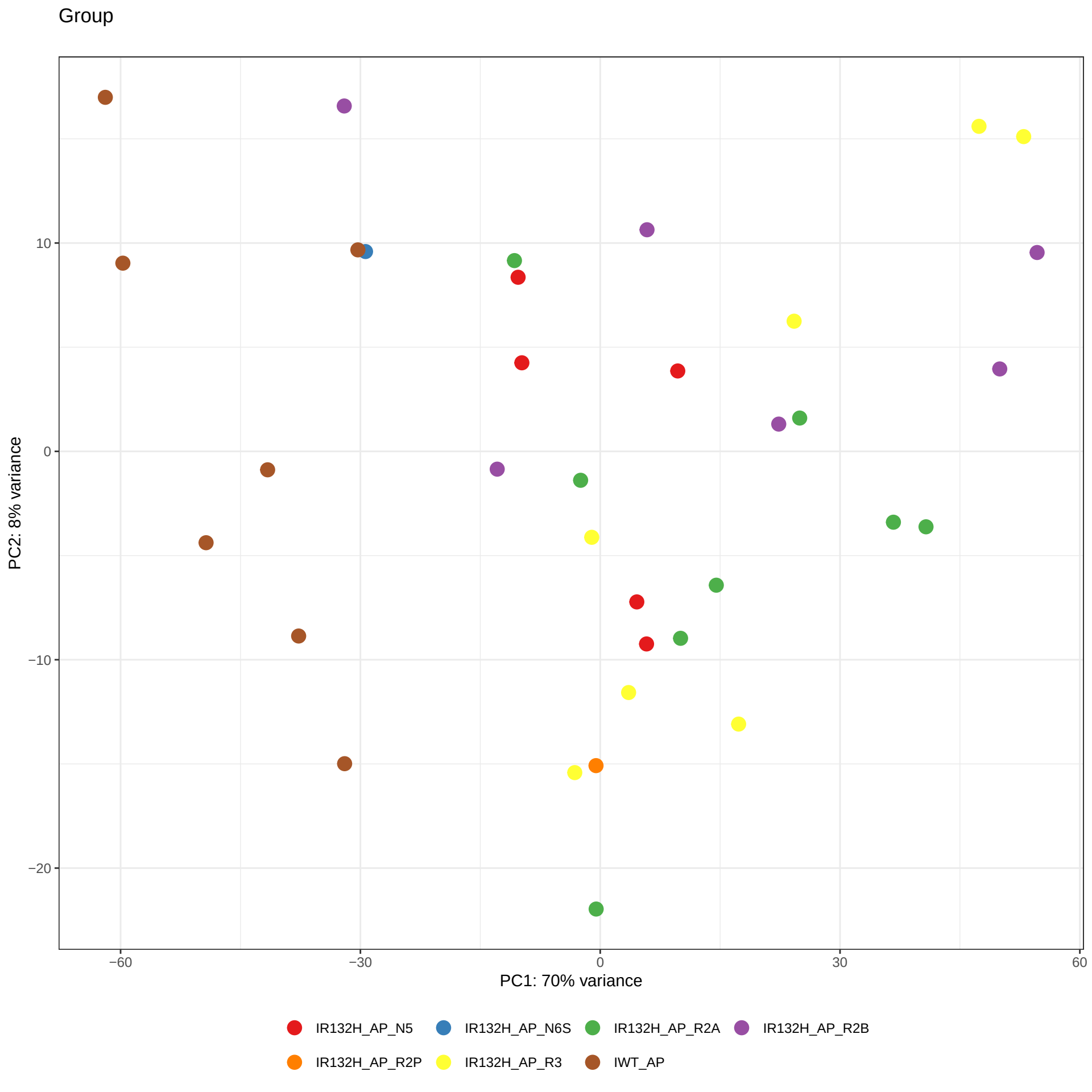

ATRX
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance
shRNA


Condition
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance
C1A
C1B
C2


Grp
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance
C1
C2


Type
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance
CSHL_C1A
Run2_C1A
Run2_C1B
Run2_C2
Run3_C1A
Run3_C1B
Run3_C2


Mark
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance
Check
CSHL_C1A
Run2_C1A
Run2_C1B
Run2_C2
Run3_C1A
Run3_C1B
Run3_C2


Group
10
0
−10
−20
PC2: 8% variance
−60
−30
0
30
60
PC1: 70% variance
IR132H_AP_N5
IR132H_AP_N6S
IR132H_AP_R2A
IR132H_AP_R2B
IR132H_AP_R2P
IR132H_AP_R3
IWT_AP