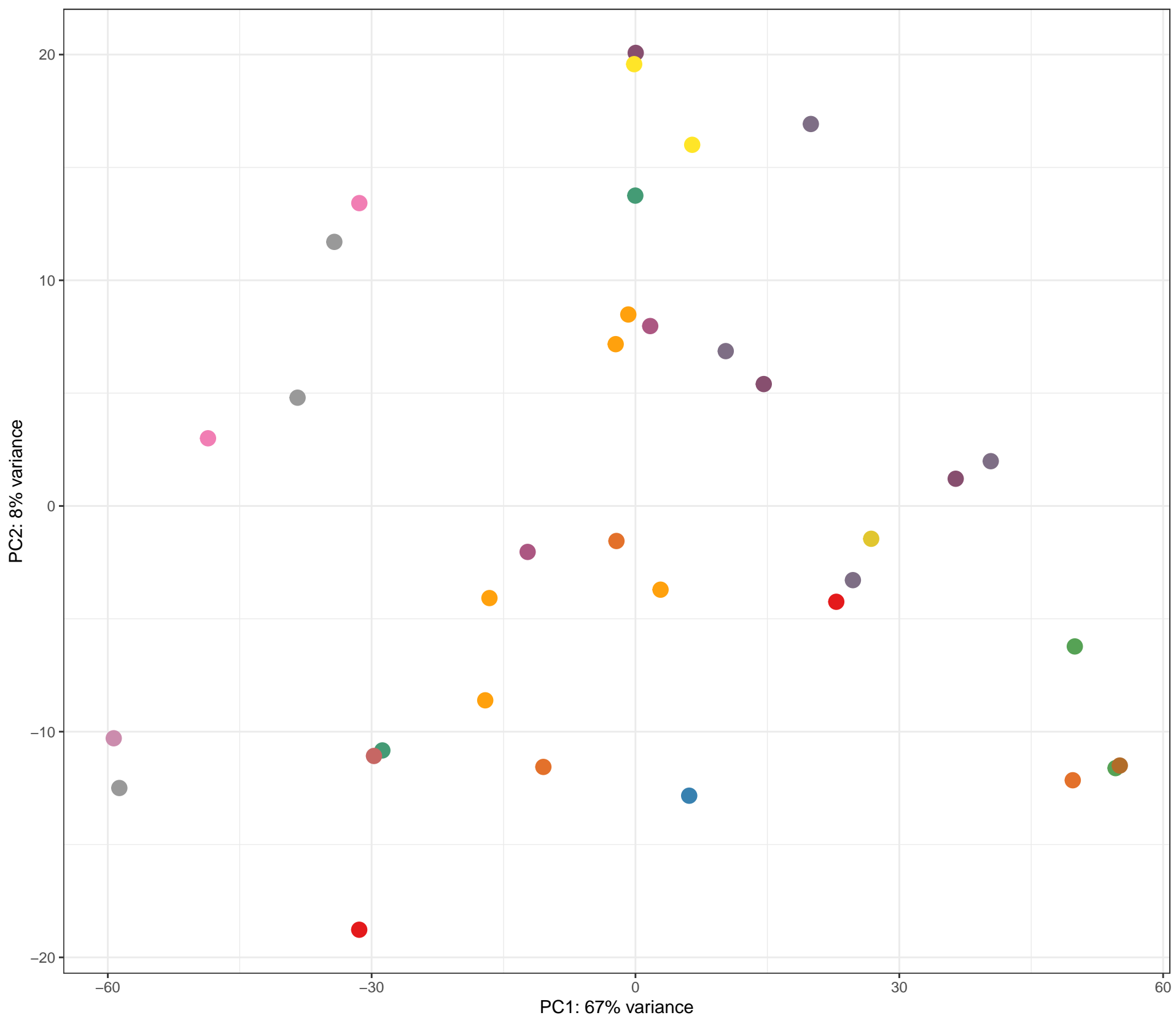
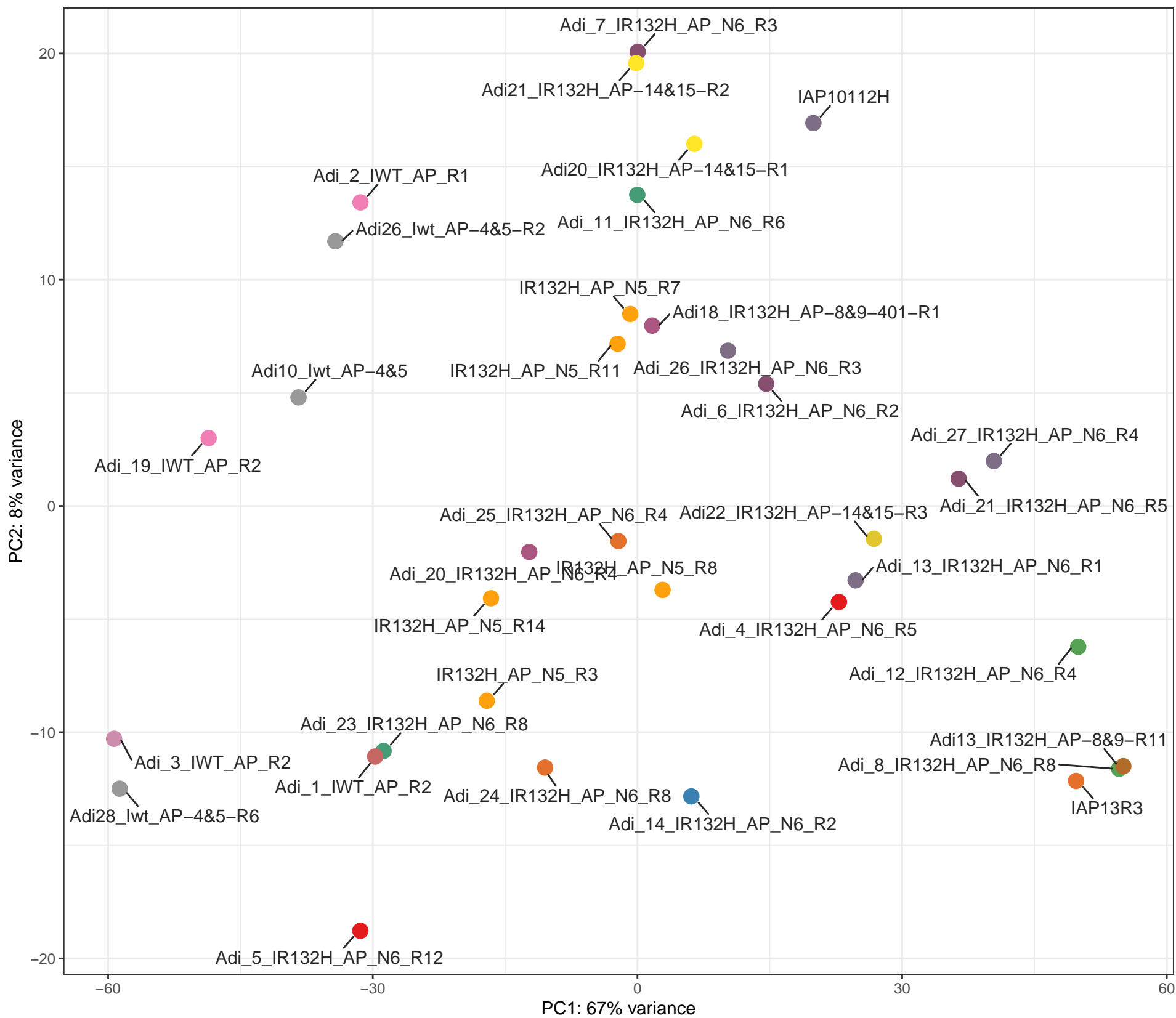


Tumour\_origin

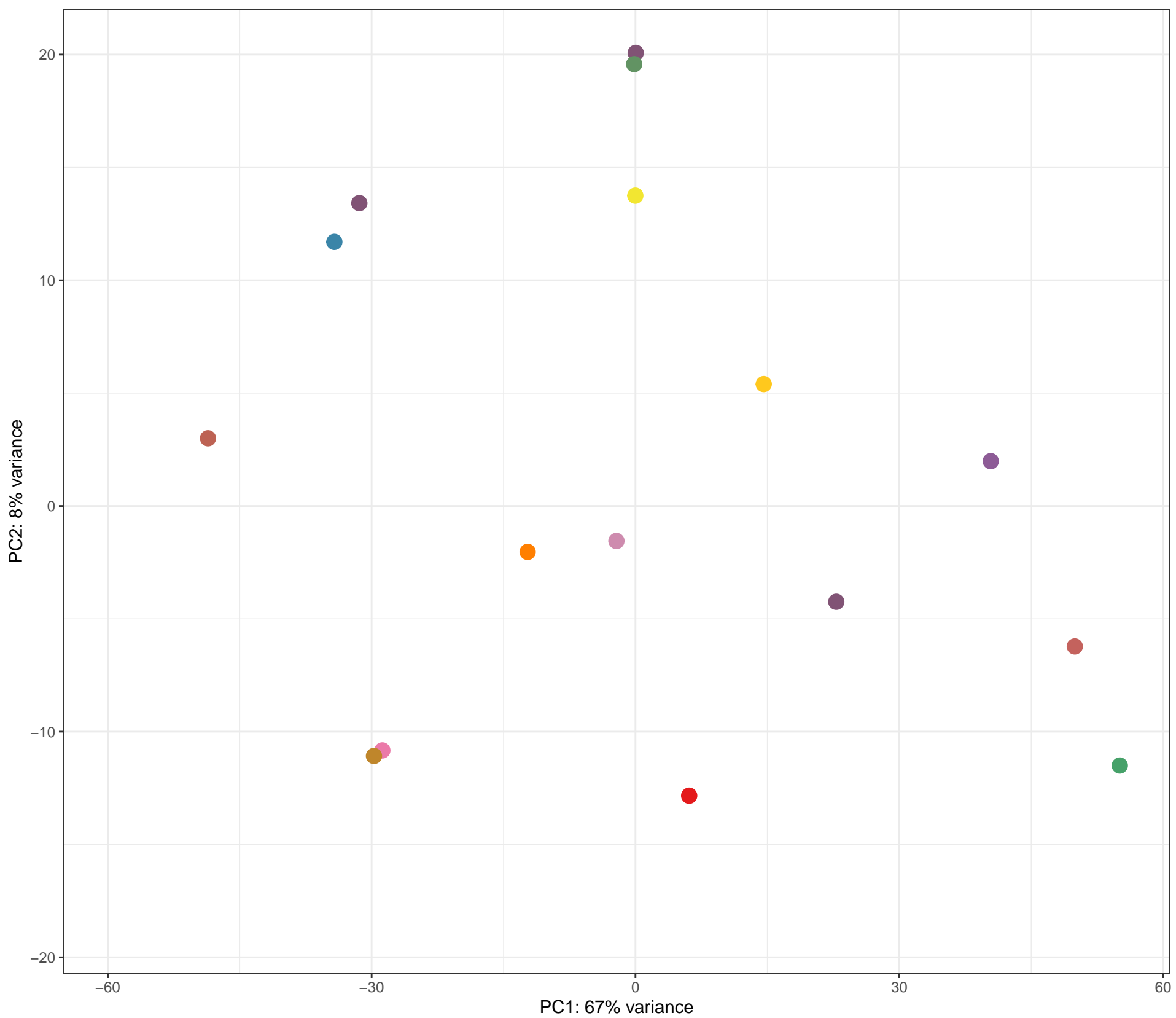


● T-a    ● T-b    ● T-c    ● T-d-primary    ● T-e    ● T-f    ● T-g    ● T-h  
● T-i    ● T-k    ● T-l    ● T-m    ● WT-a    ● WT-b    ● WT-c    ● WT-d

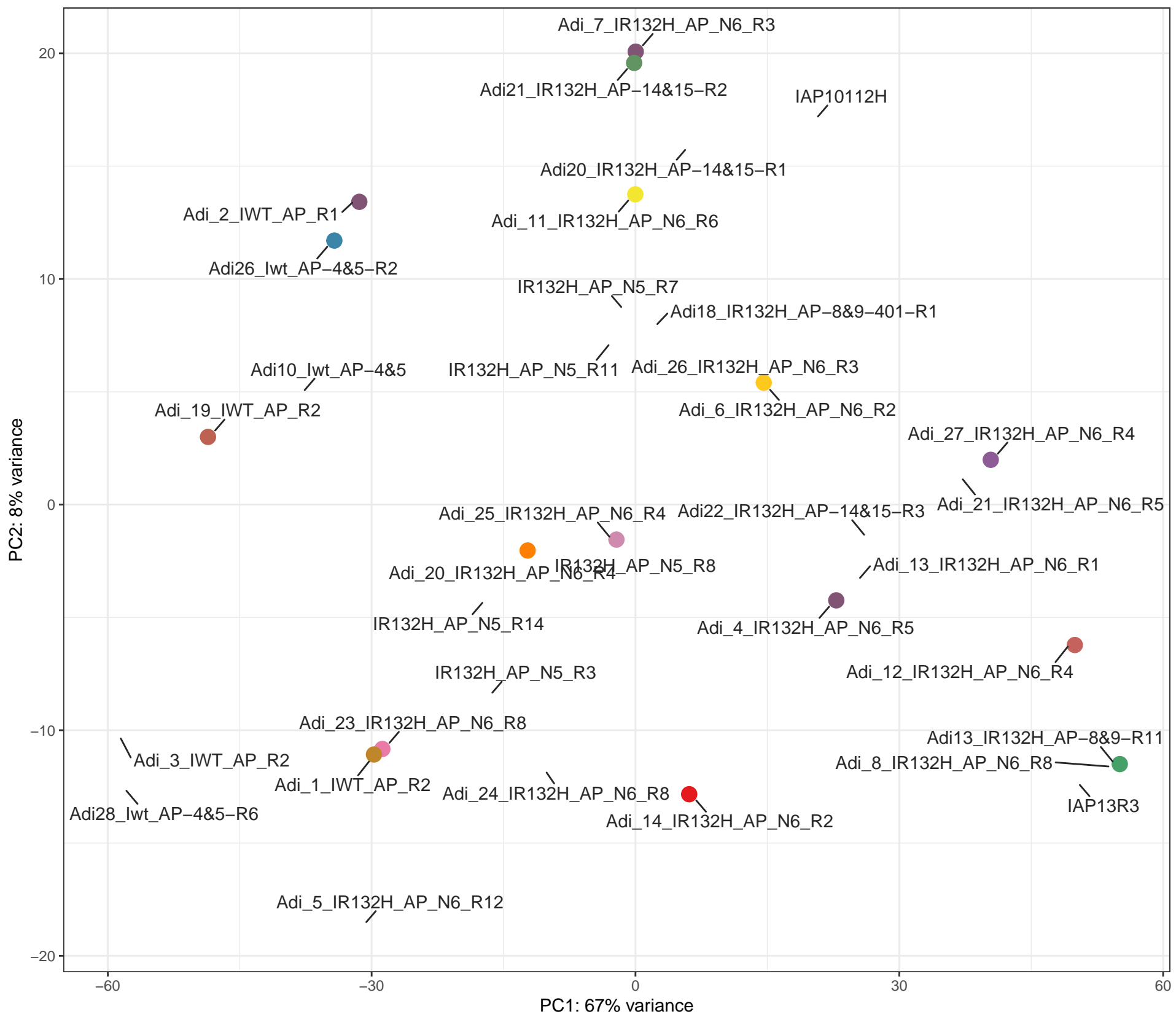
# Tumour\_origin



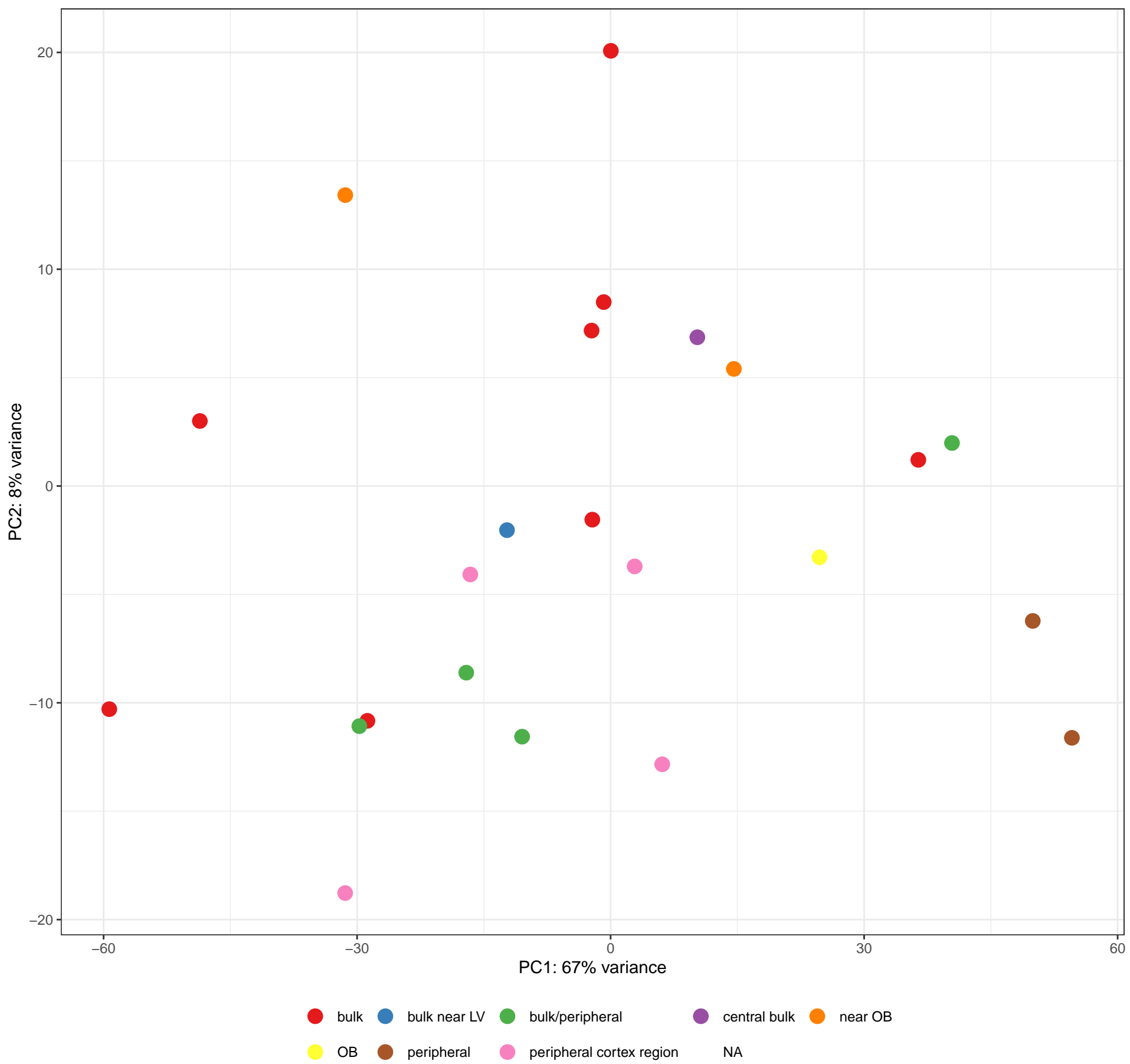
CNA

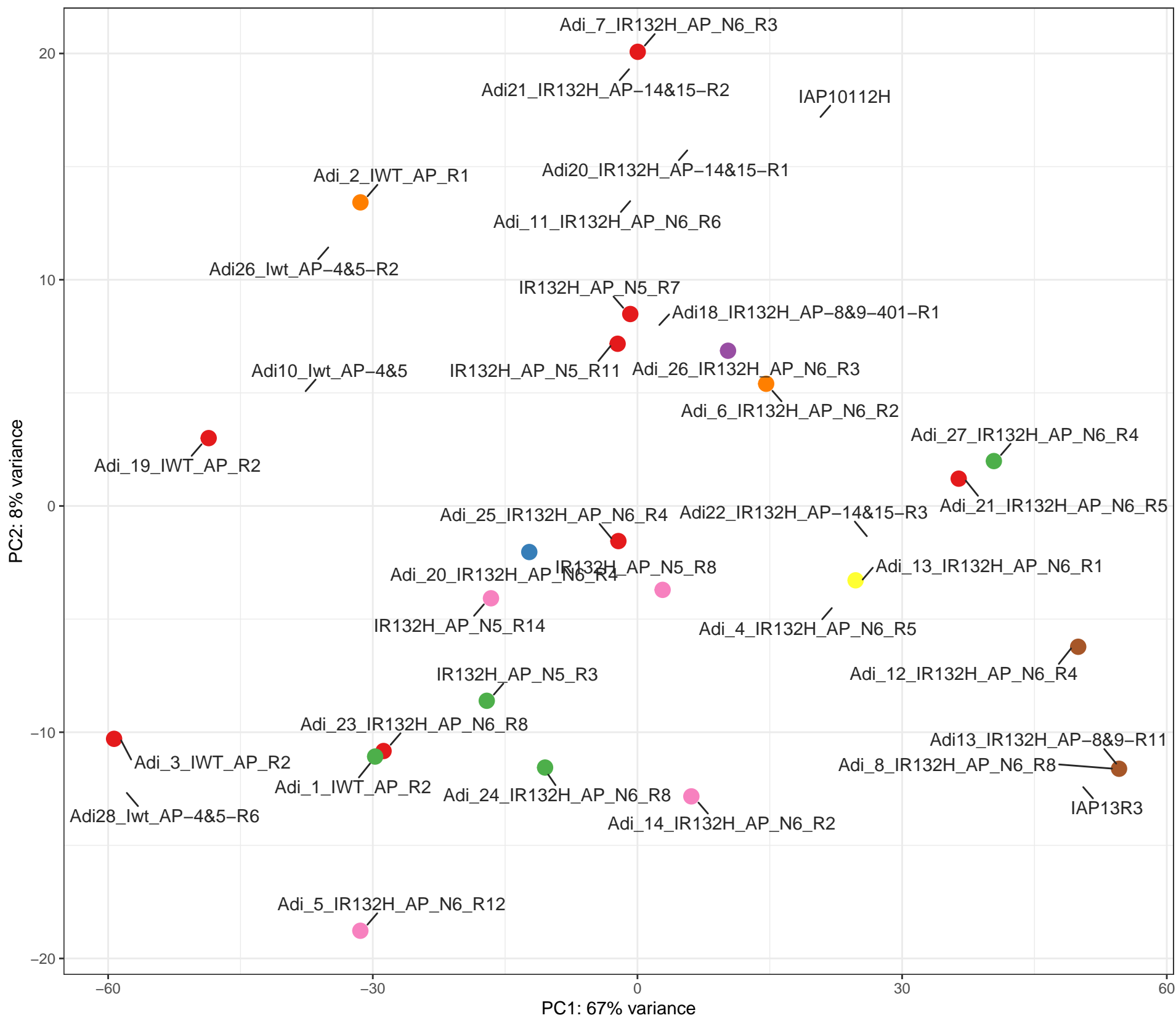


● yes
 ● Yes
 ● Yes - Adi-29
 ● Yes - Adi-A
 ● Yes - Adi-I
 ● Yes - Adi-M
 ● Yes - Adi-N
 ● Yes - Adi-O
 ● Yes - Adi-P
 ● Yes - Adi-R
 ● Yes - Adi-U
 ● Yes - Adi-V
 ● Yes - IAP-12-R8
 ● Yes - IAP-iBTT R4
 ● NA

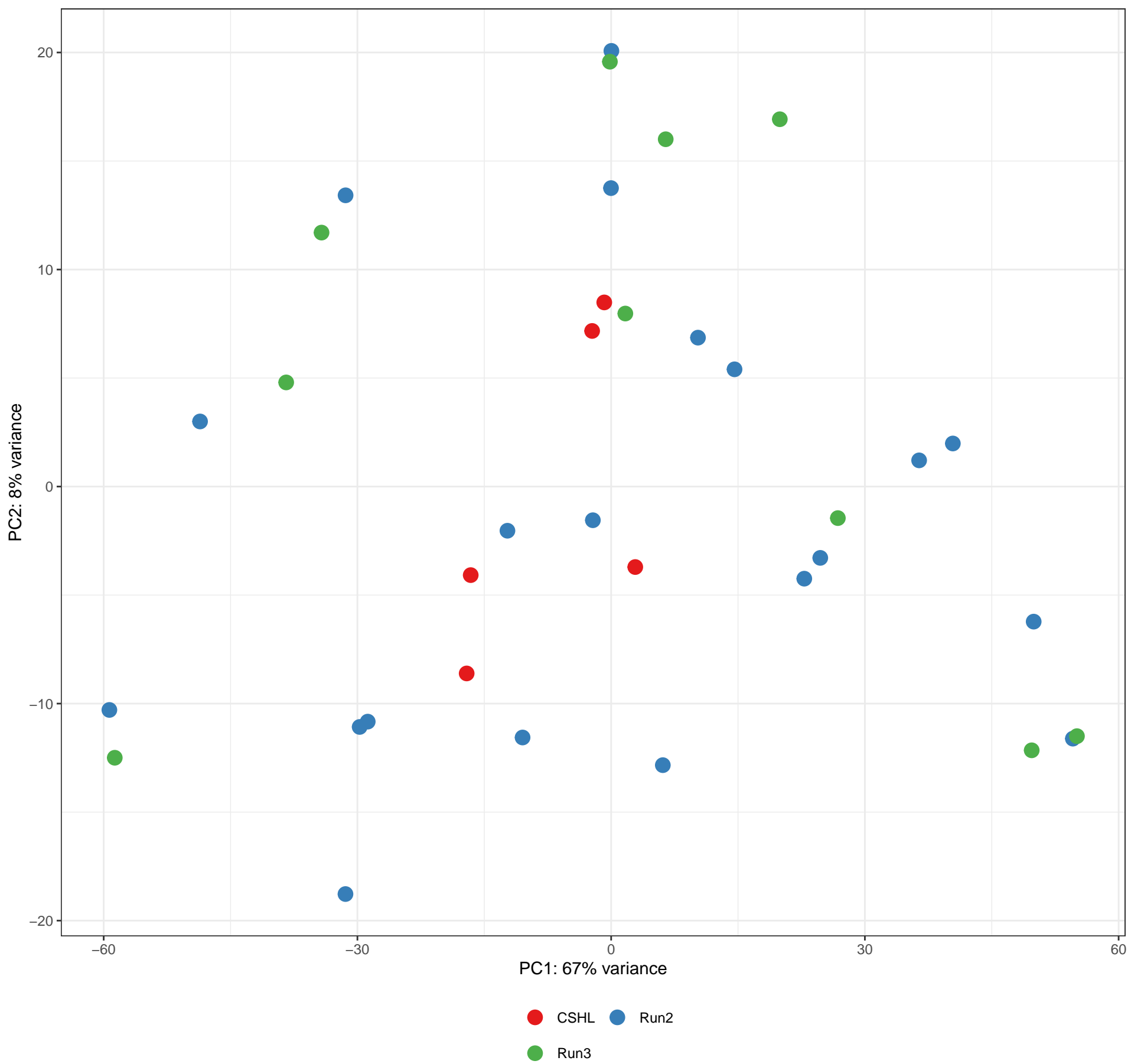


Info

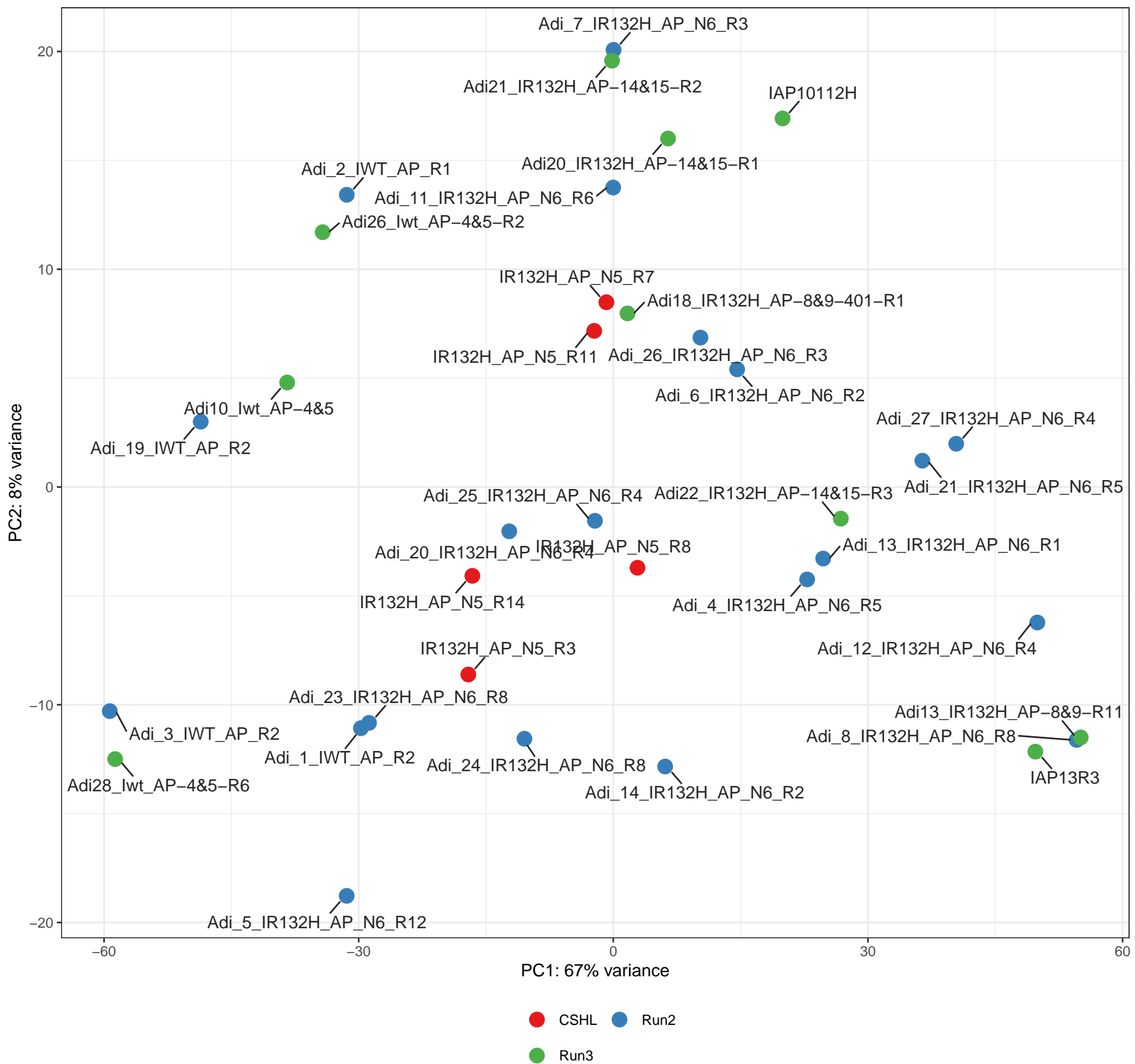




Run

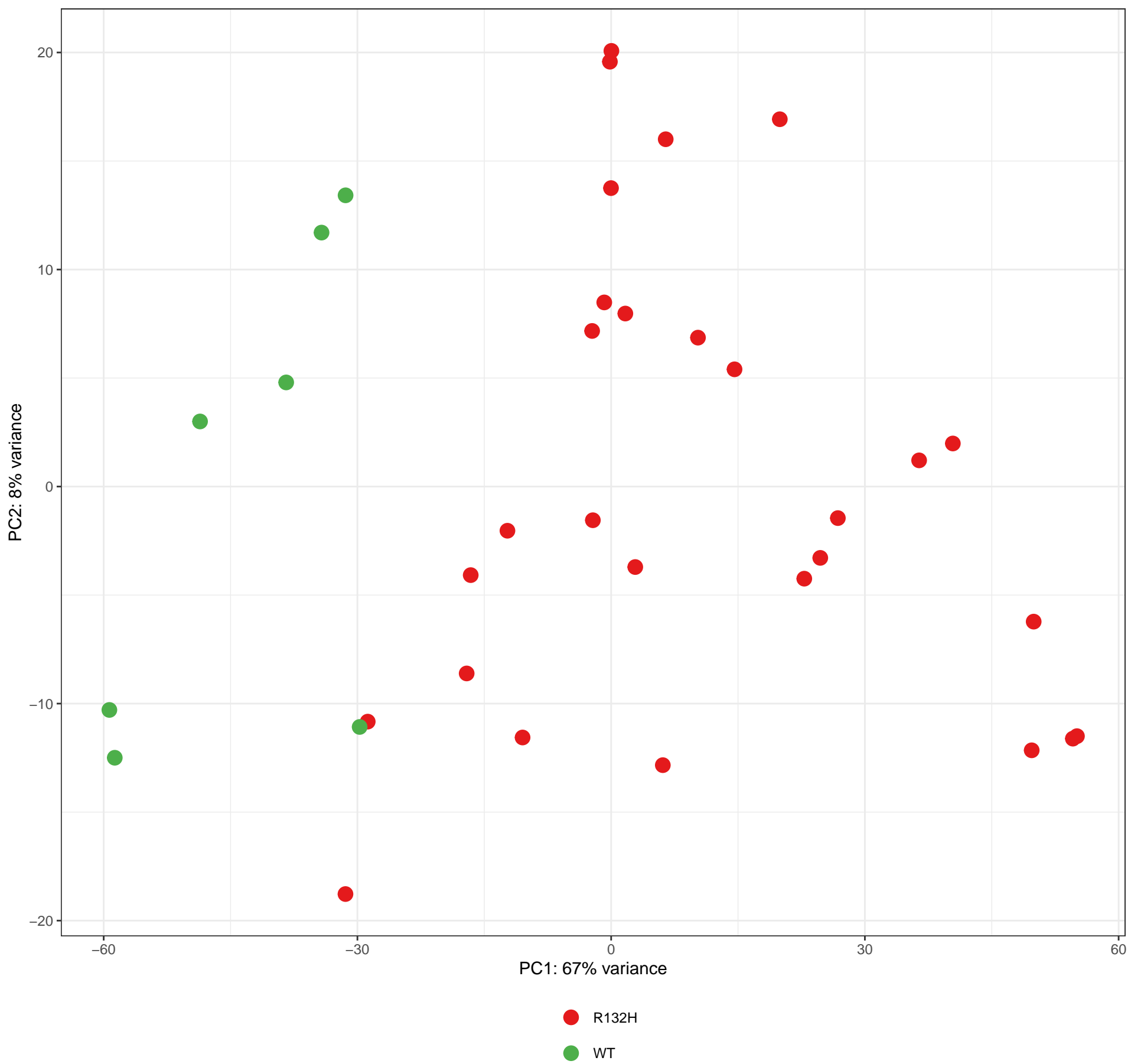


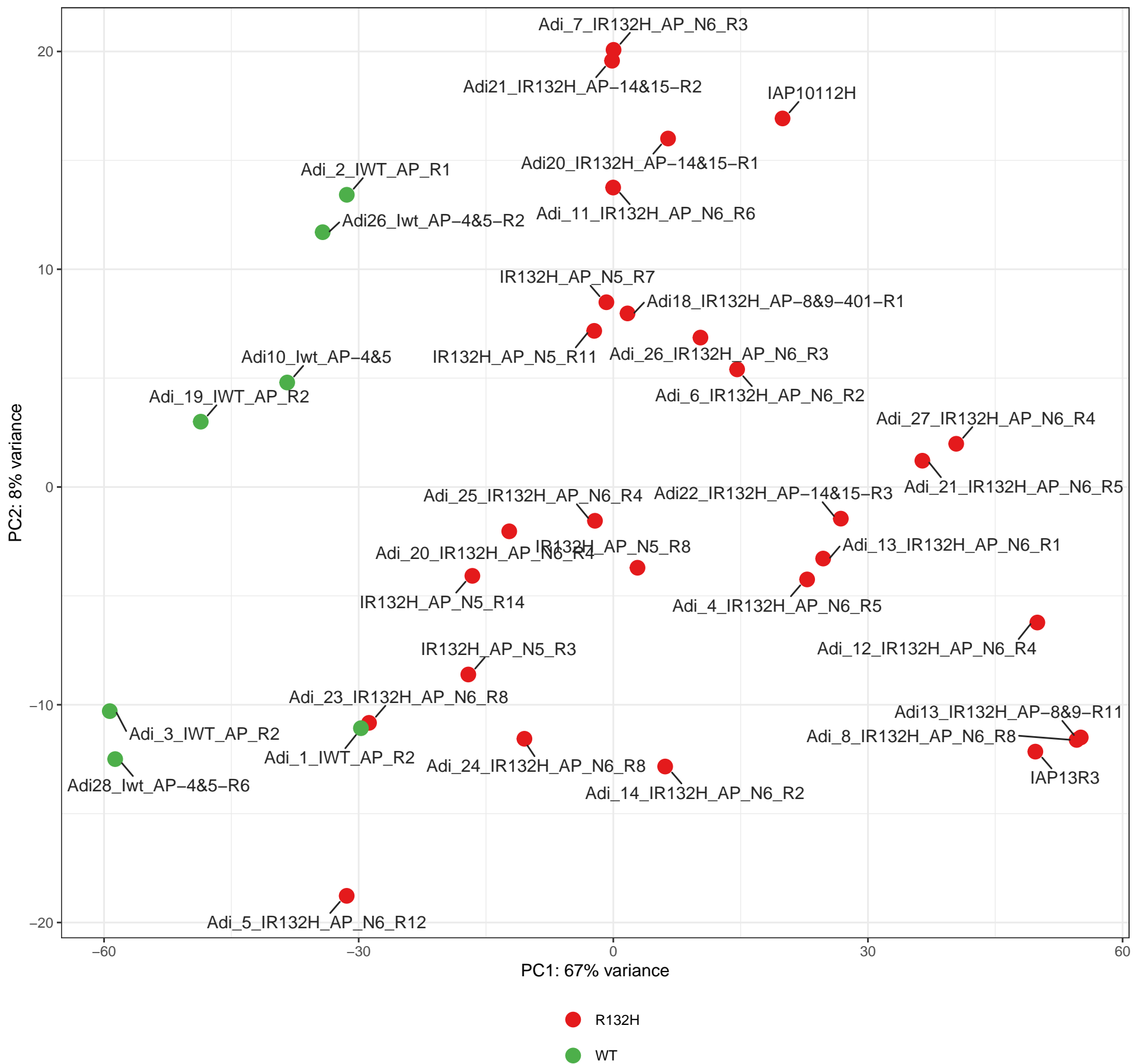
Run



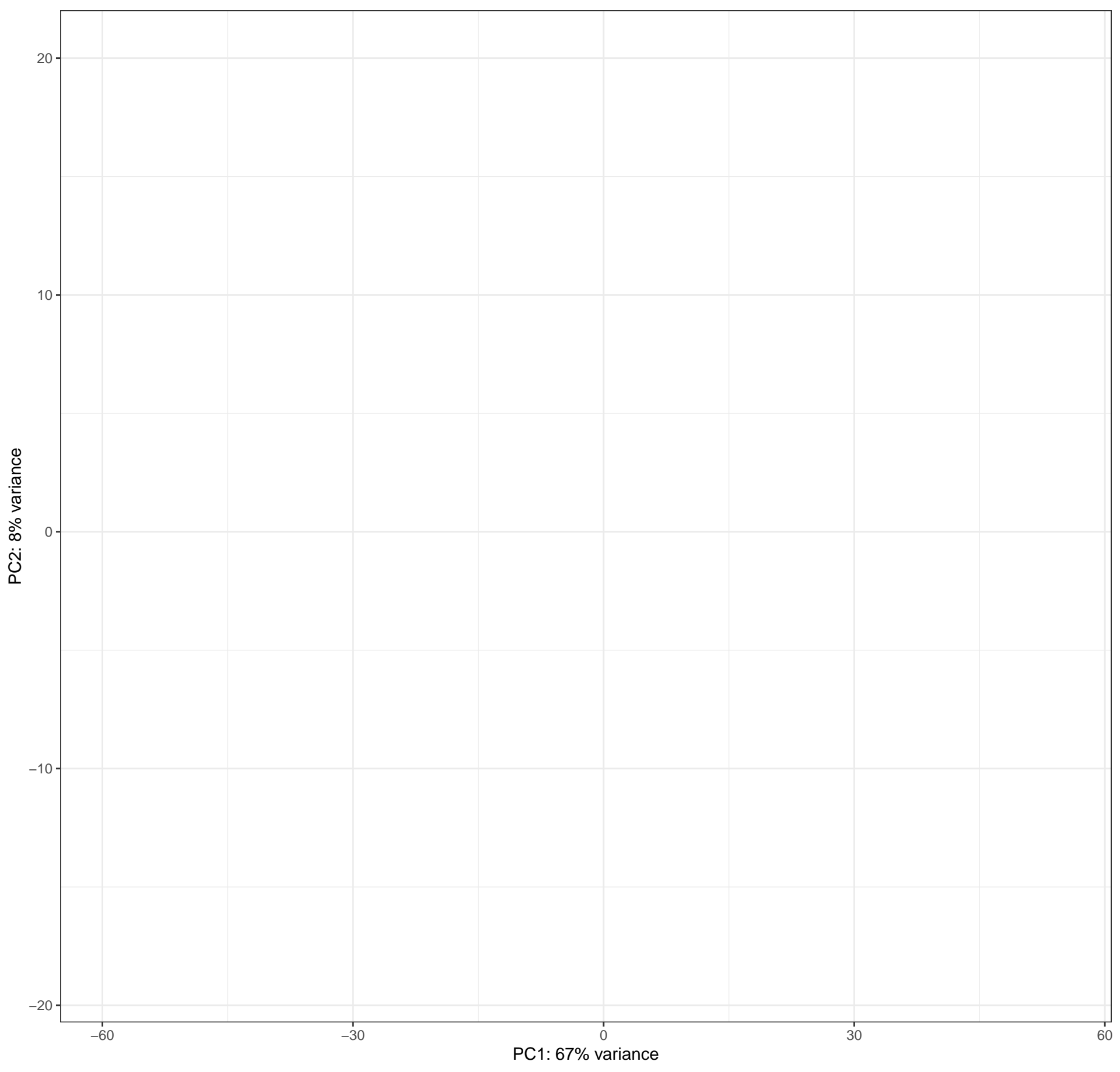


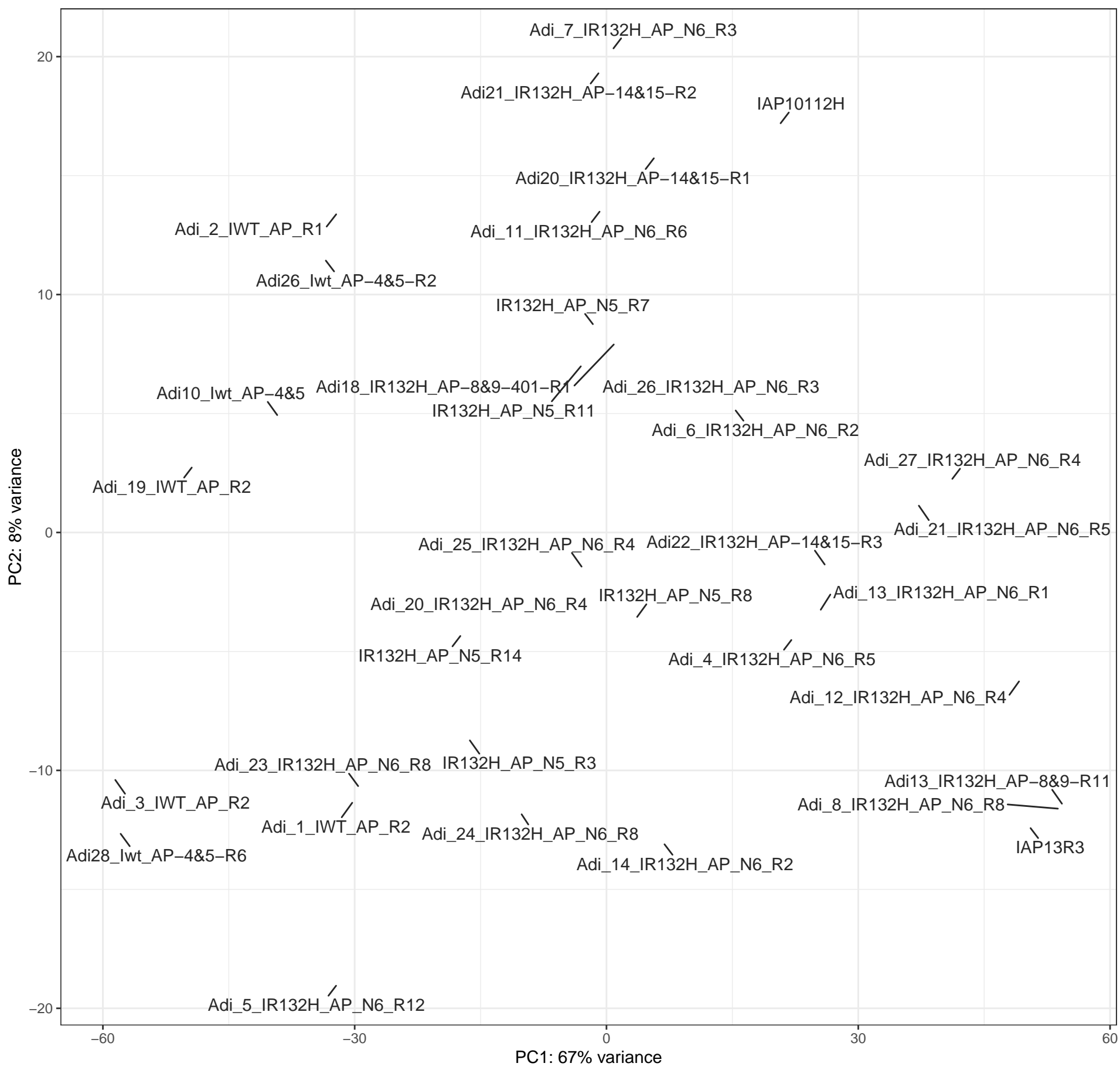
IDH



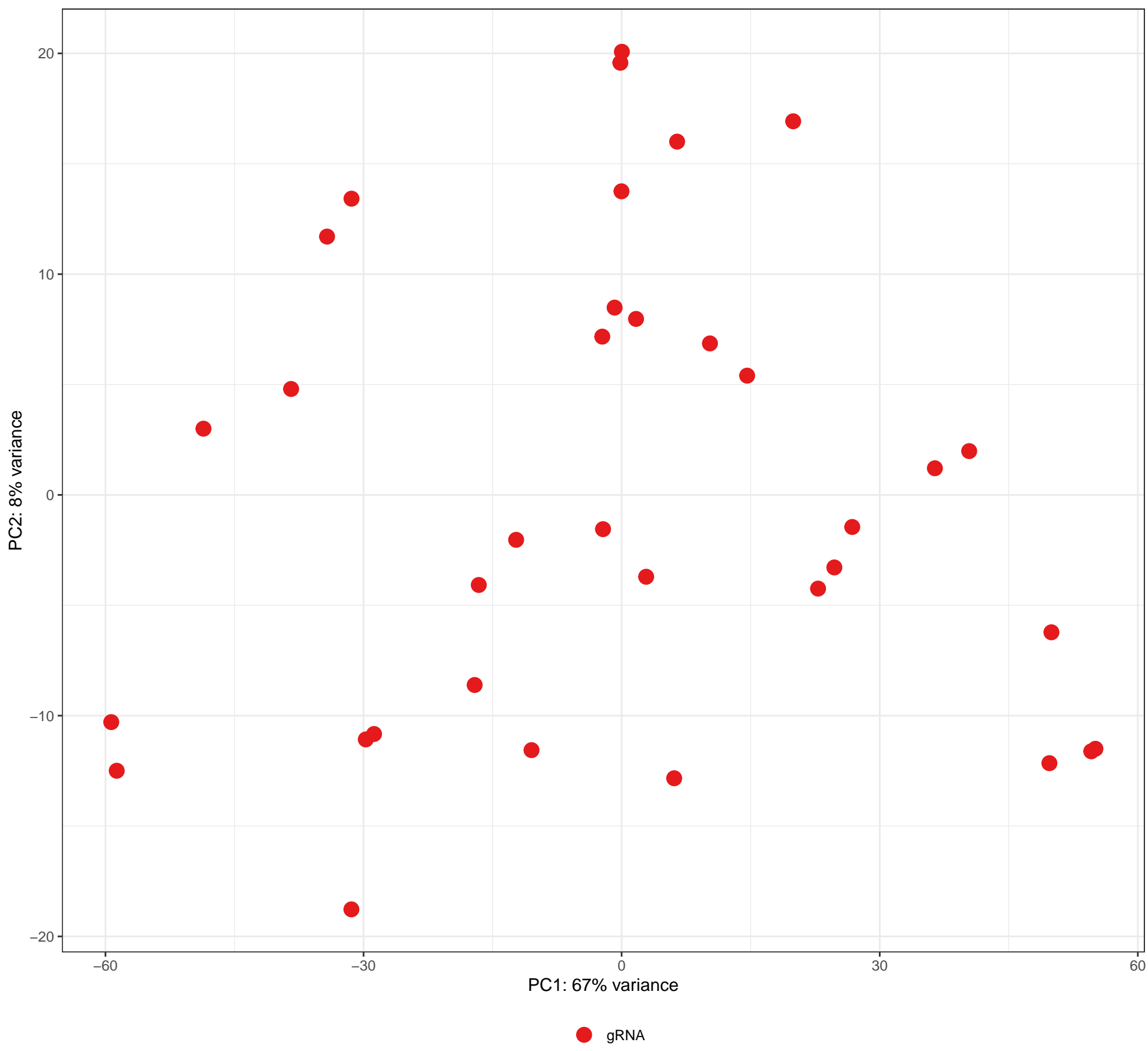


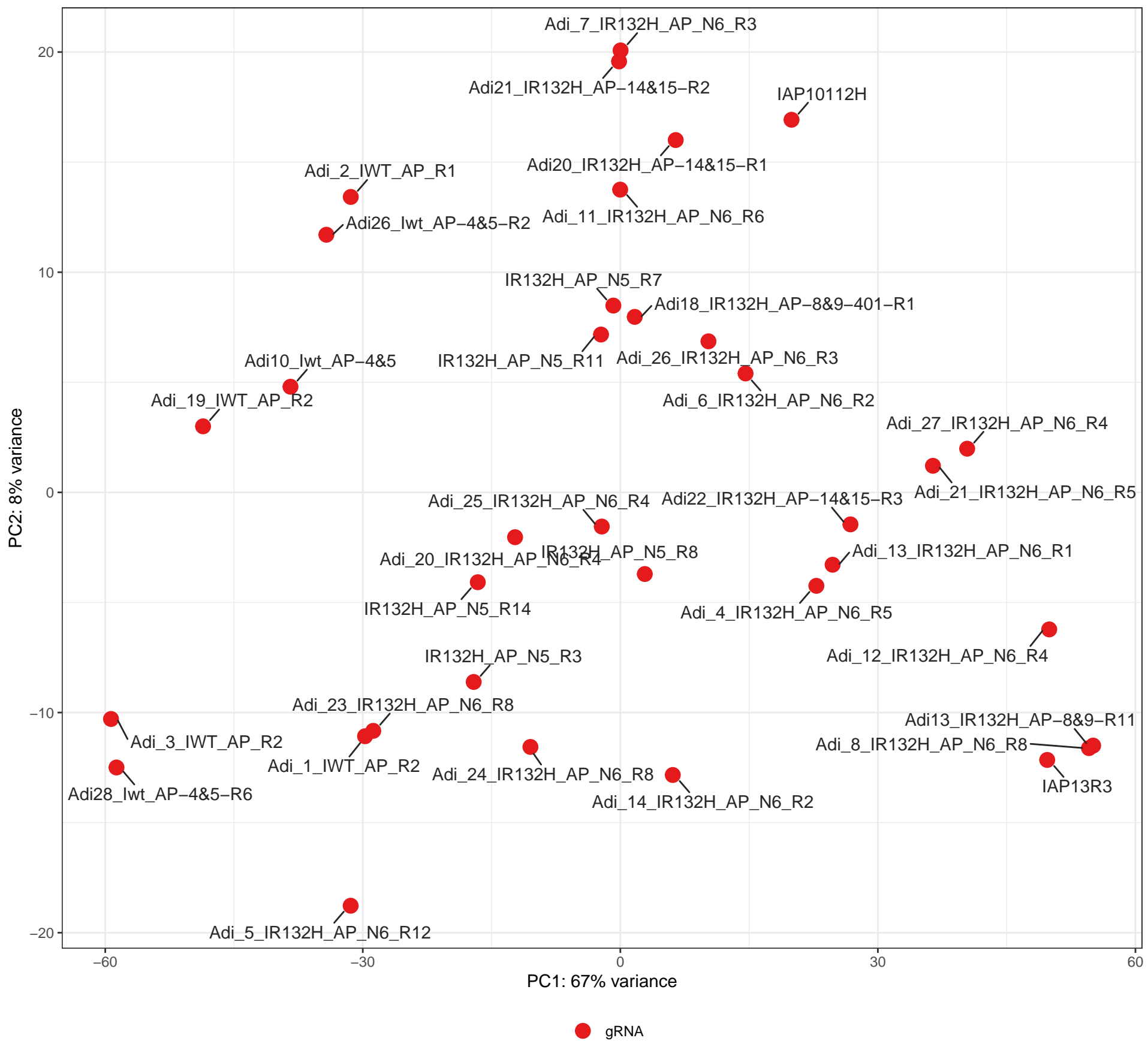
PDGFRA



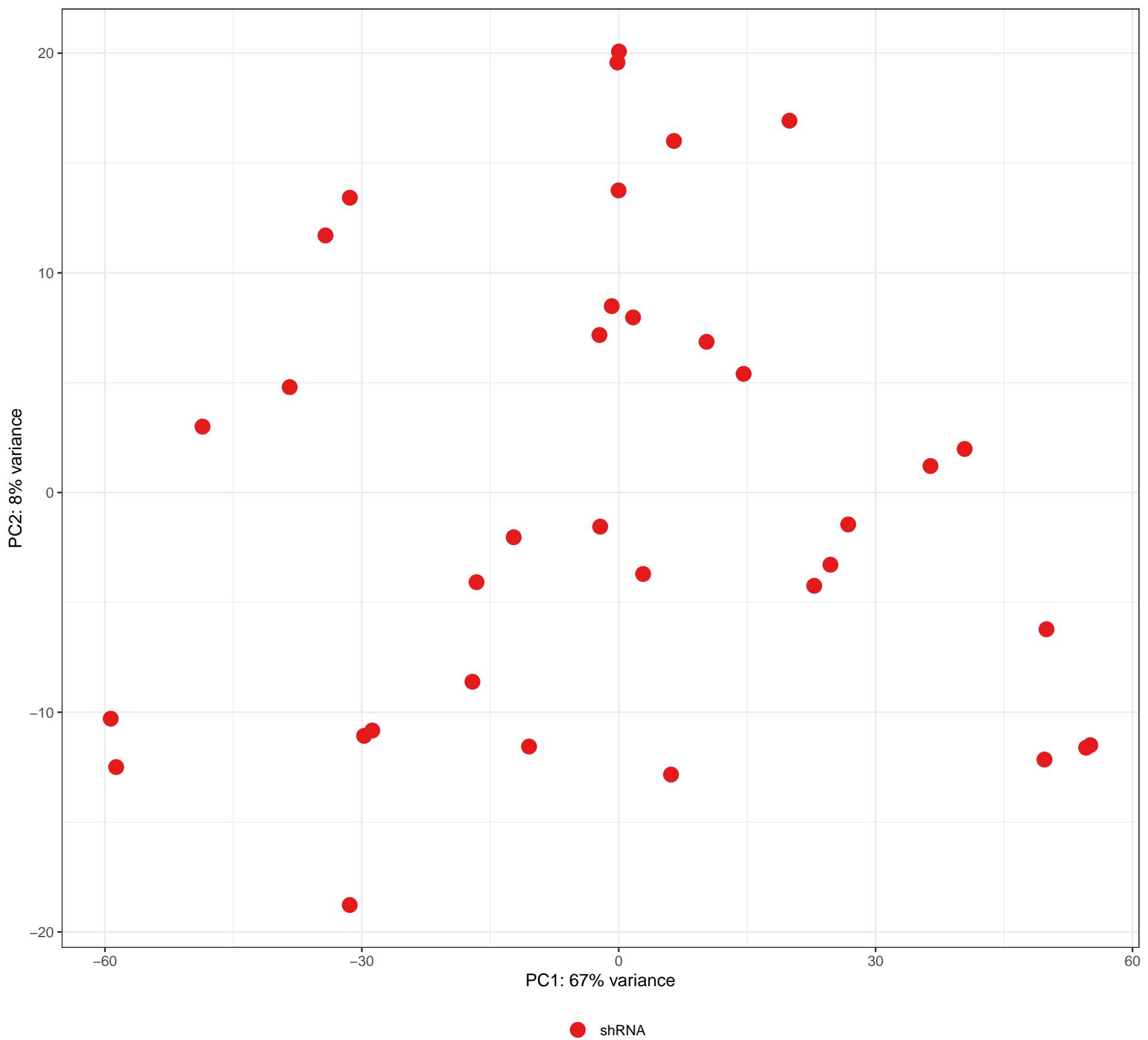


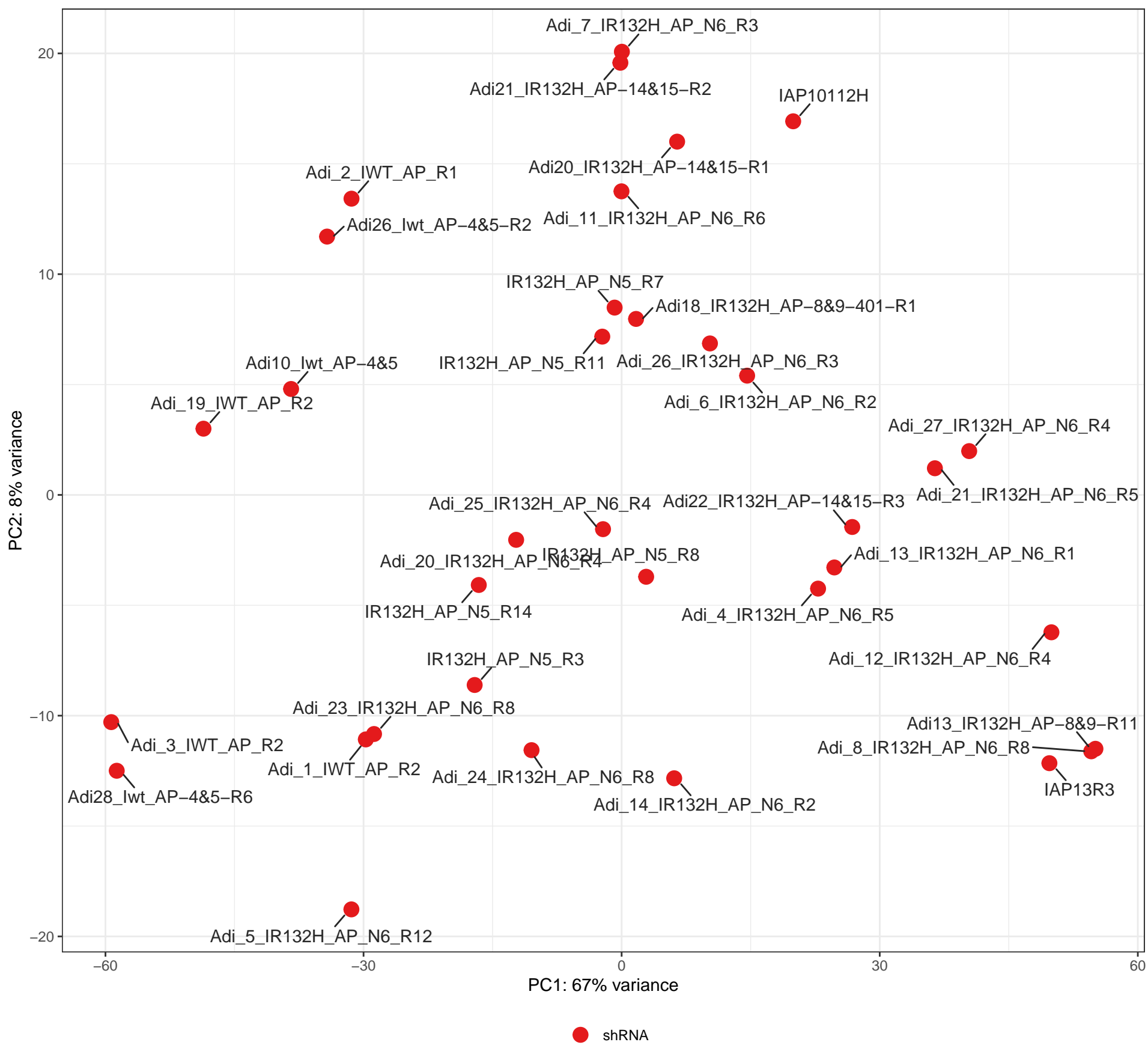
P53





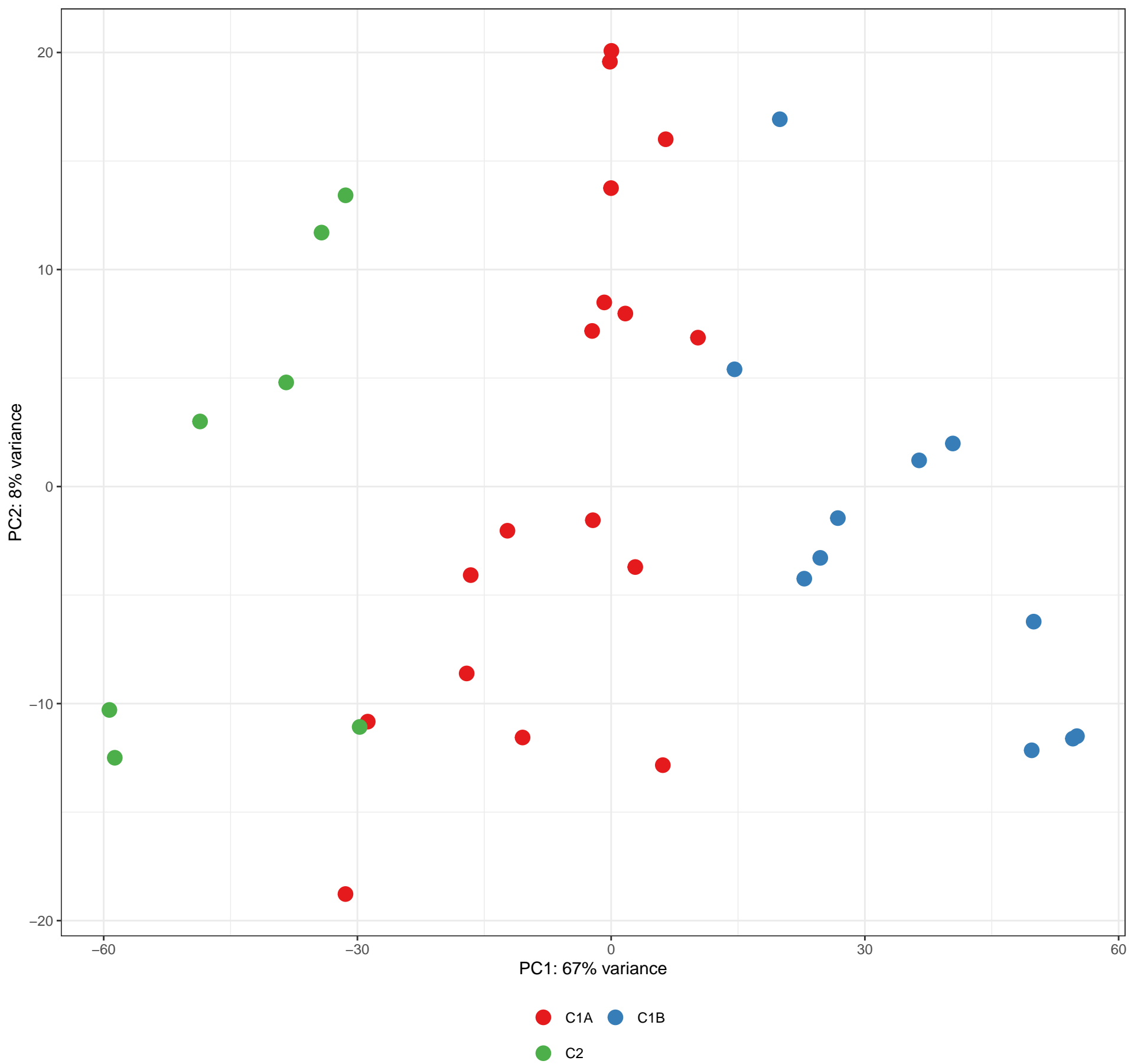
ATRX



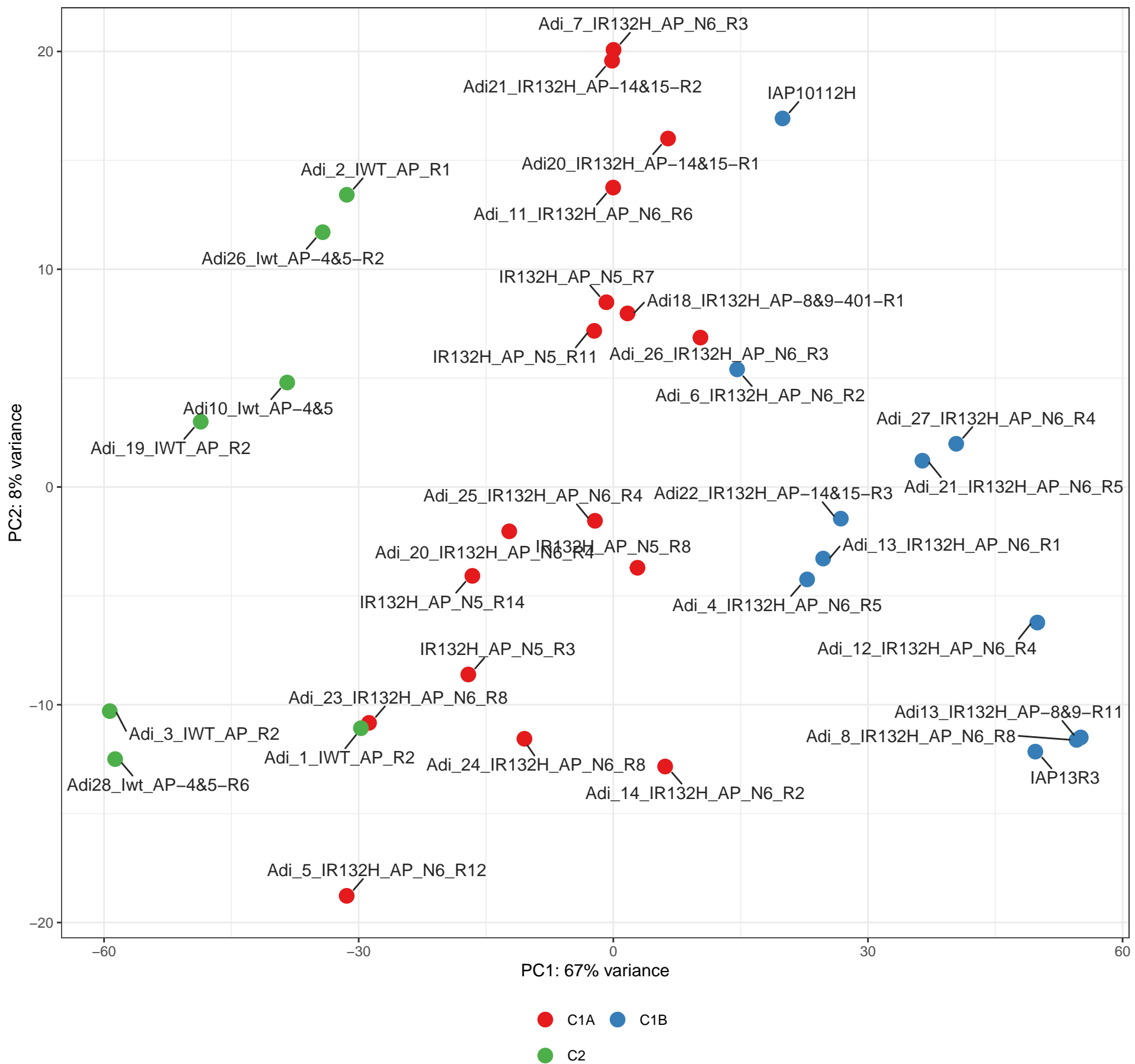




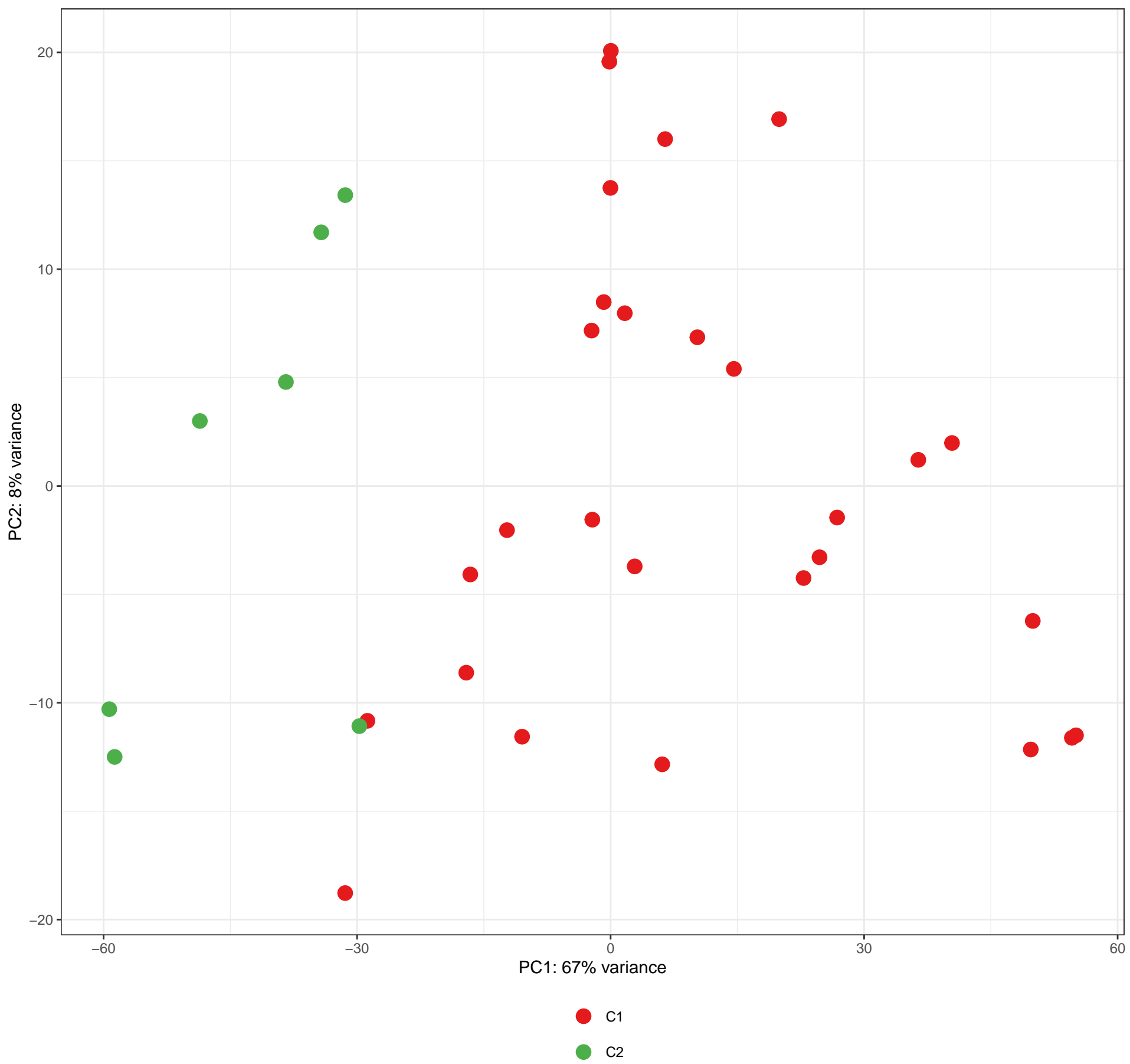
Condition



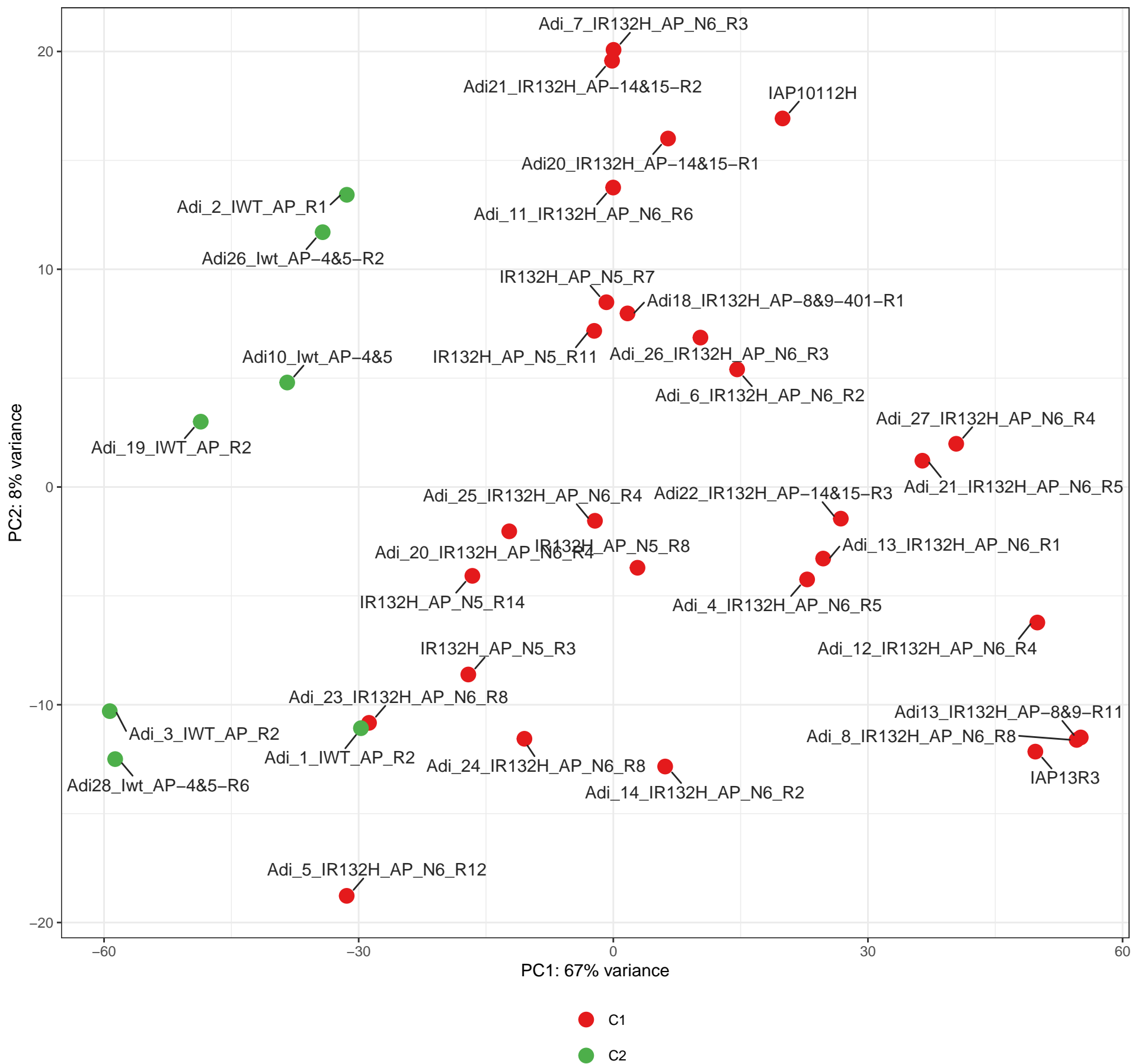
# Condition

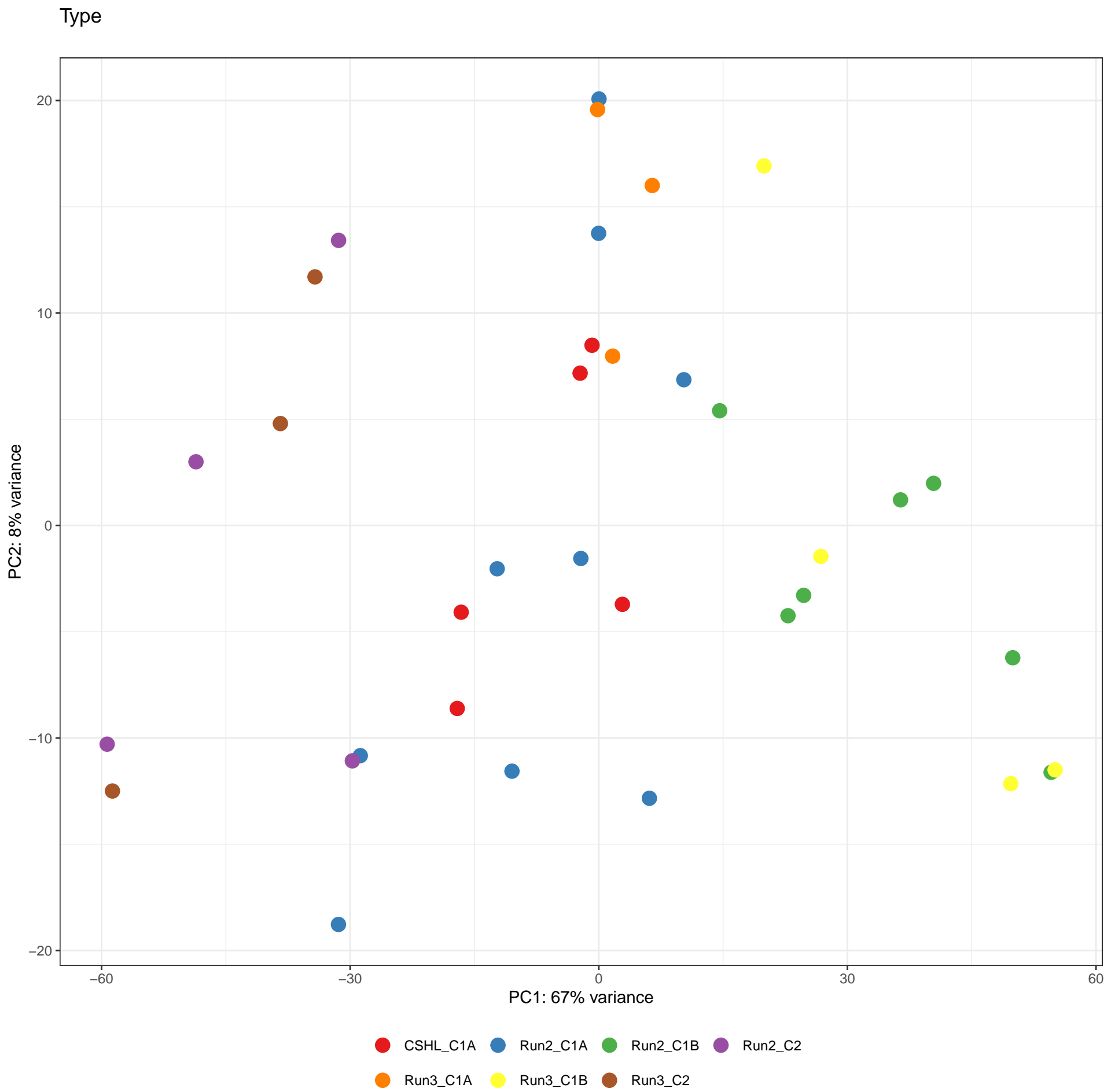


Grp

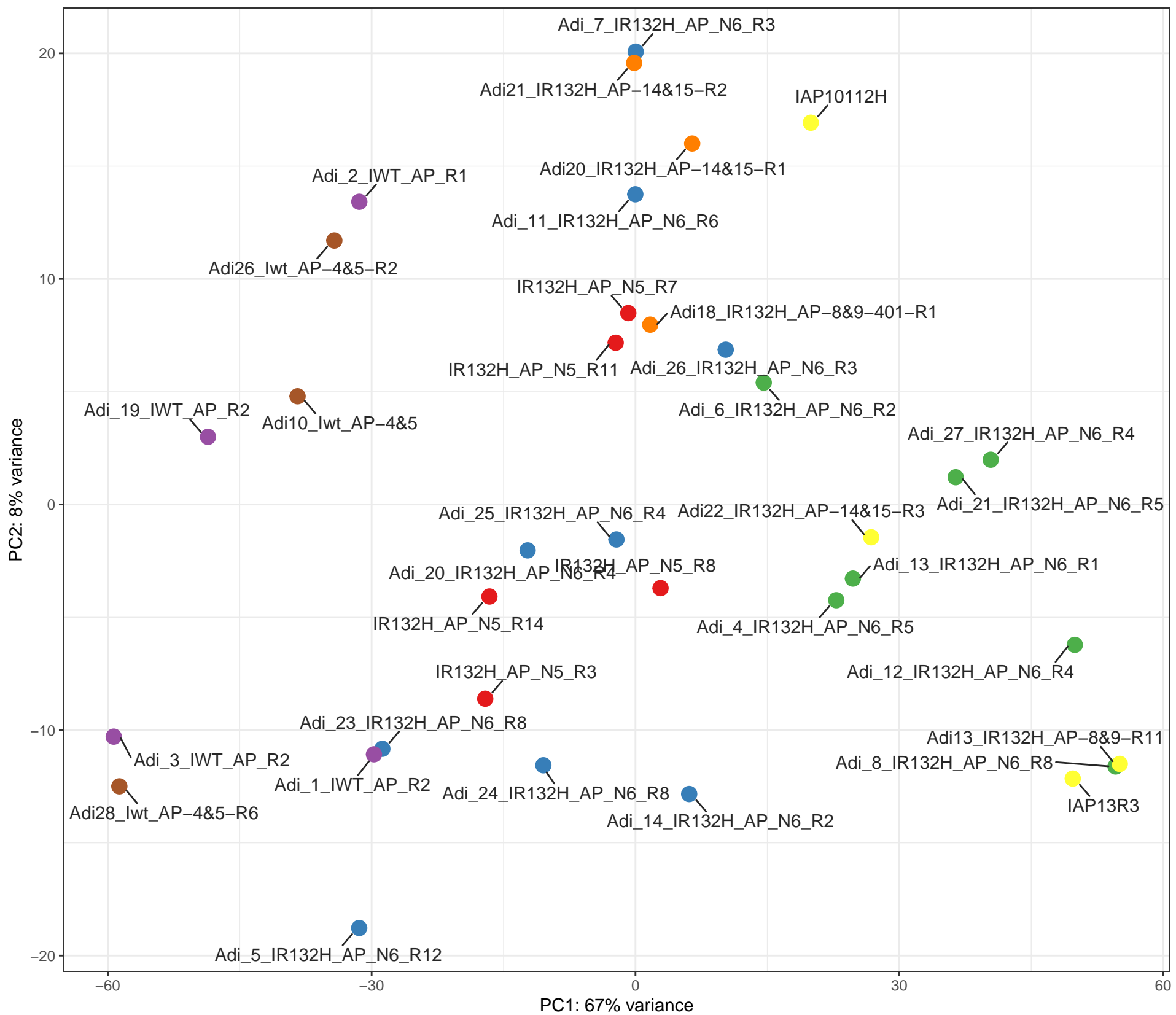


Grp



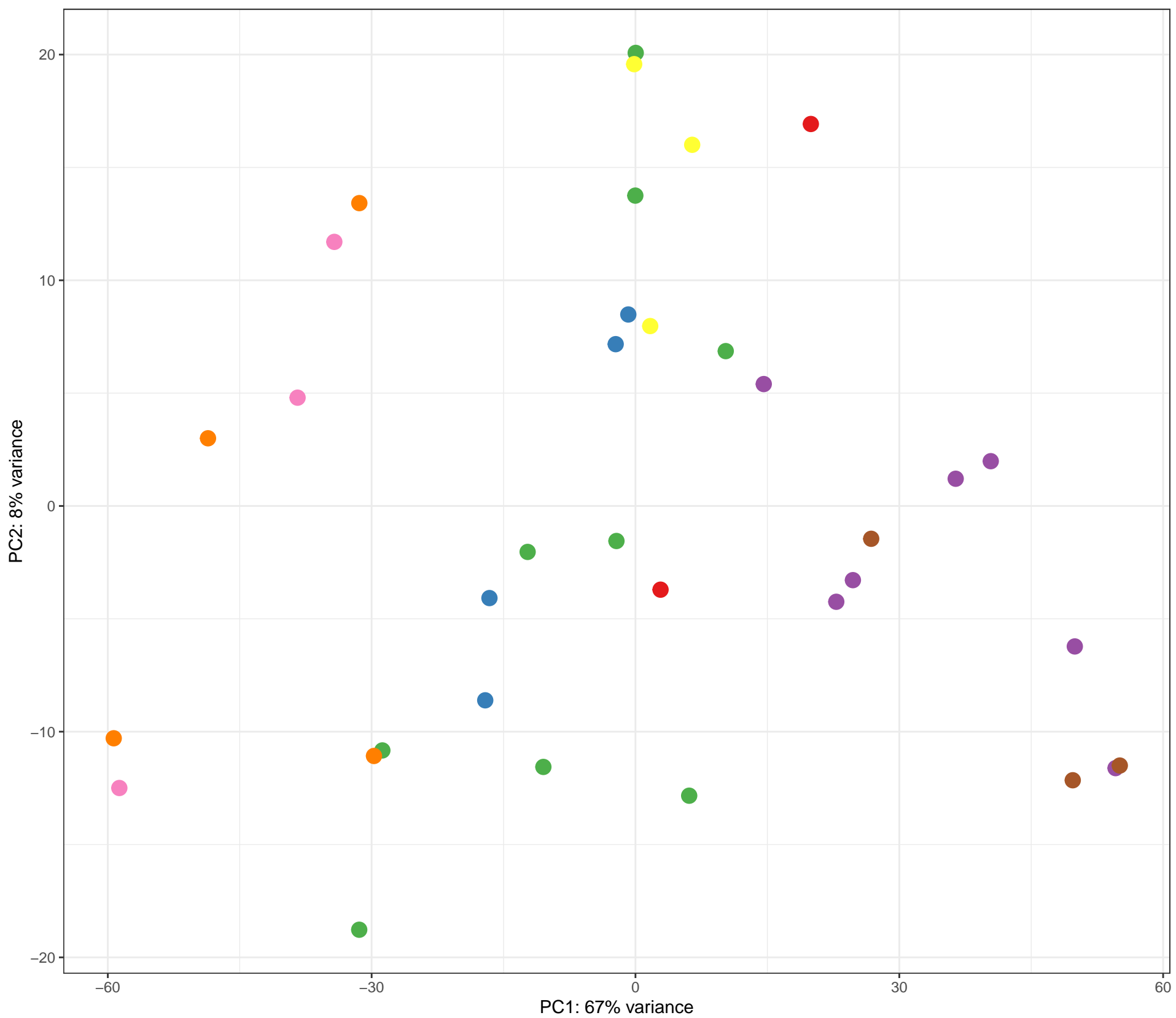


Type

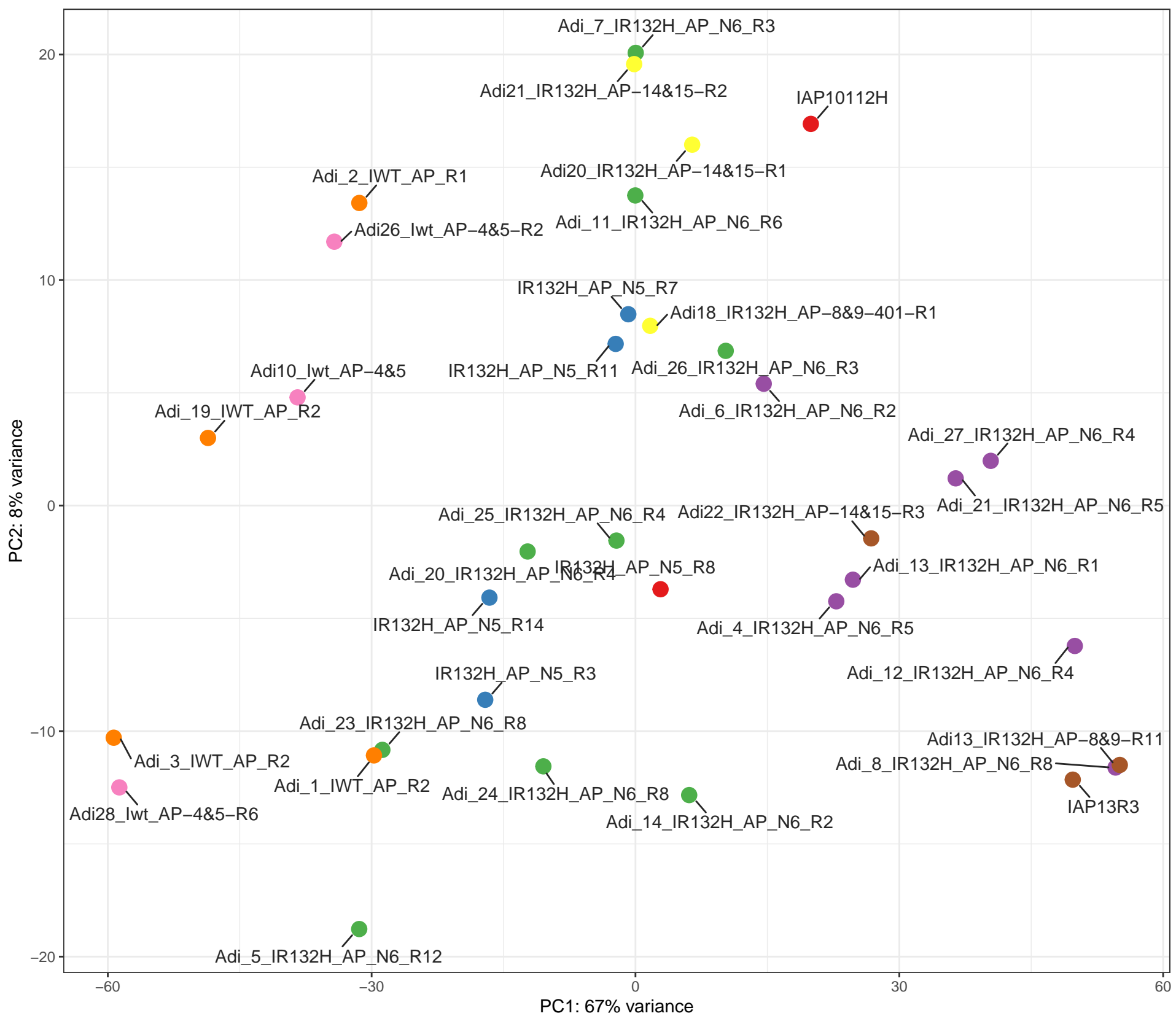


● CSHL\_C1A ● Run2\_C1A ● Run2\_C1B ● Run2\_C2  
● Run3\_C1A ● Run3\_C1B ● Run3\_C2

Mark

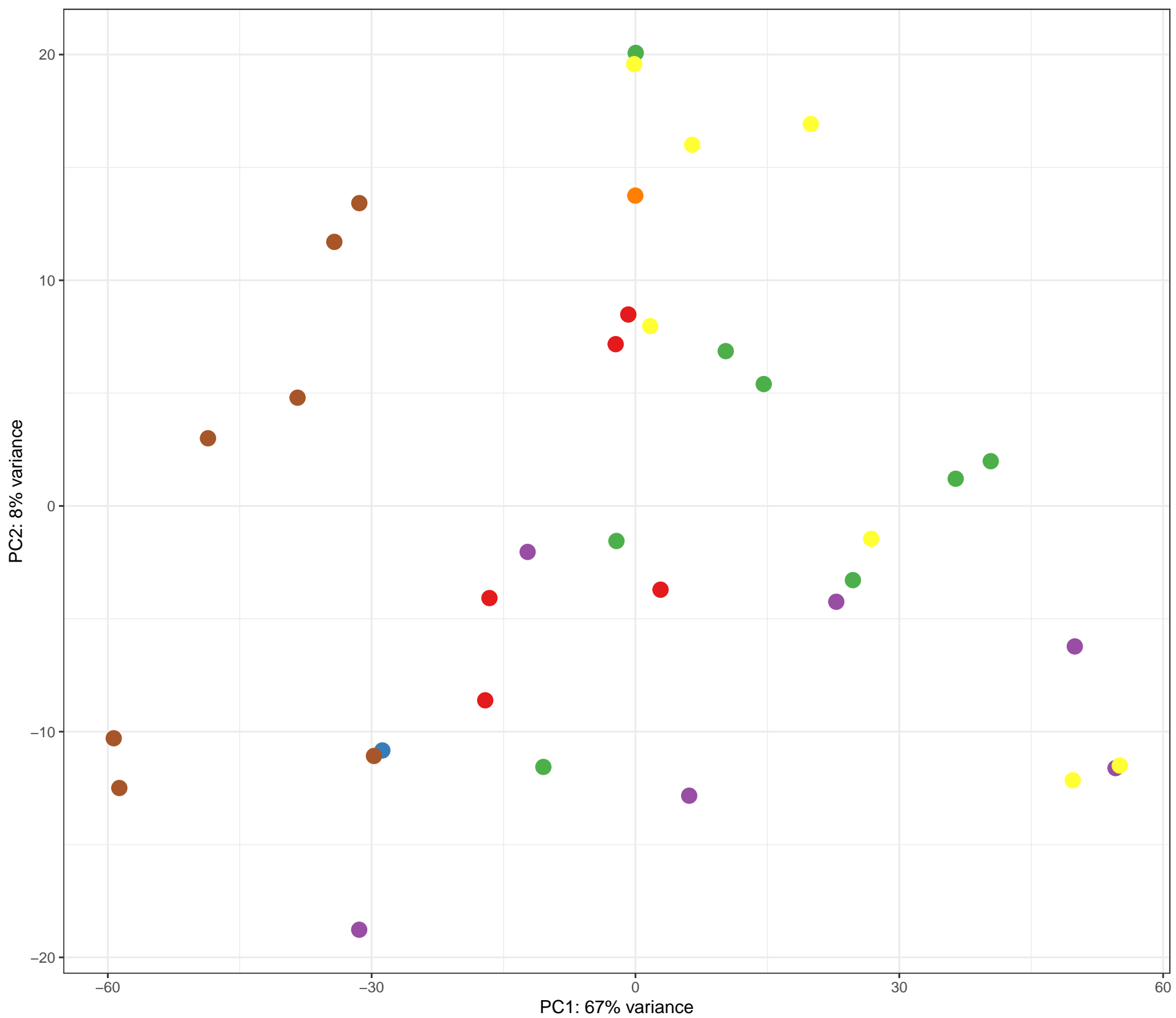


Mark





Group



IR132H\_AP\_N5 IR132H\_AP\_N6S IR132H\_AP\_R2A IR132H\_AP\_R2B  
IR132H\_AP\_R2P IR132H\_AP\_R3 IWT\_AP

Group

