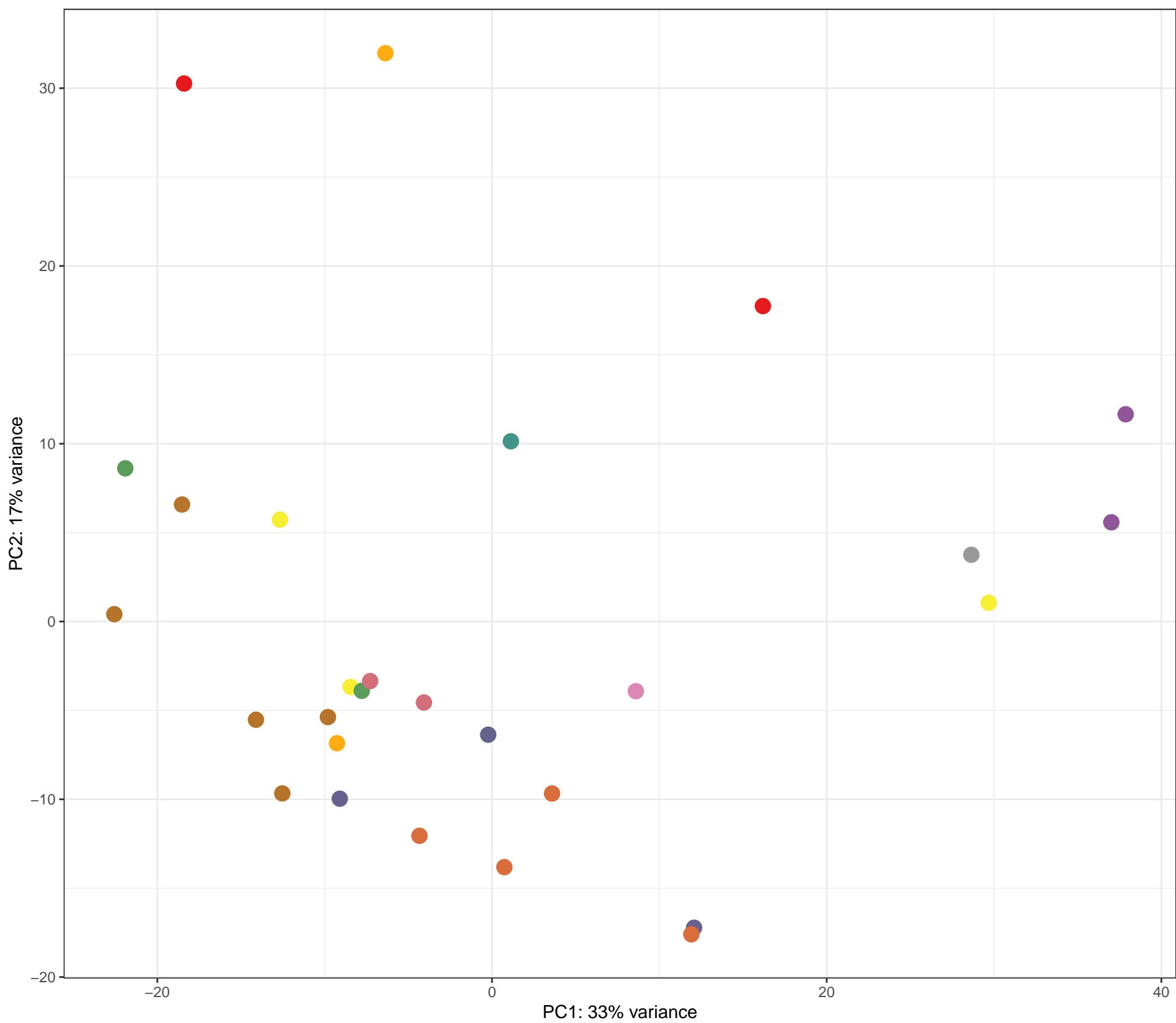
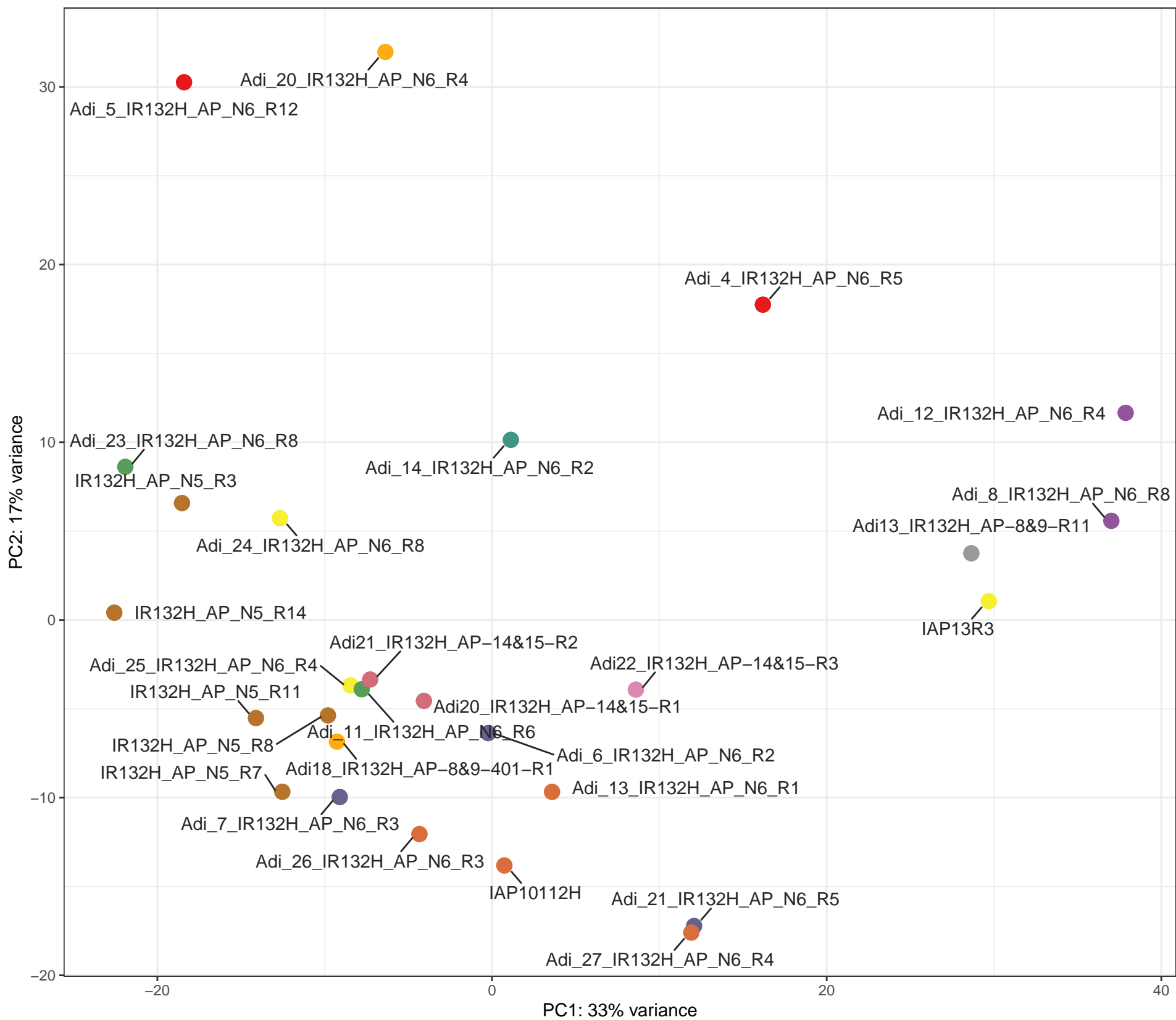


Tumour\_origin

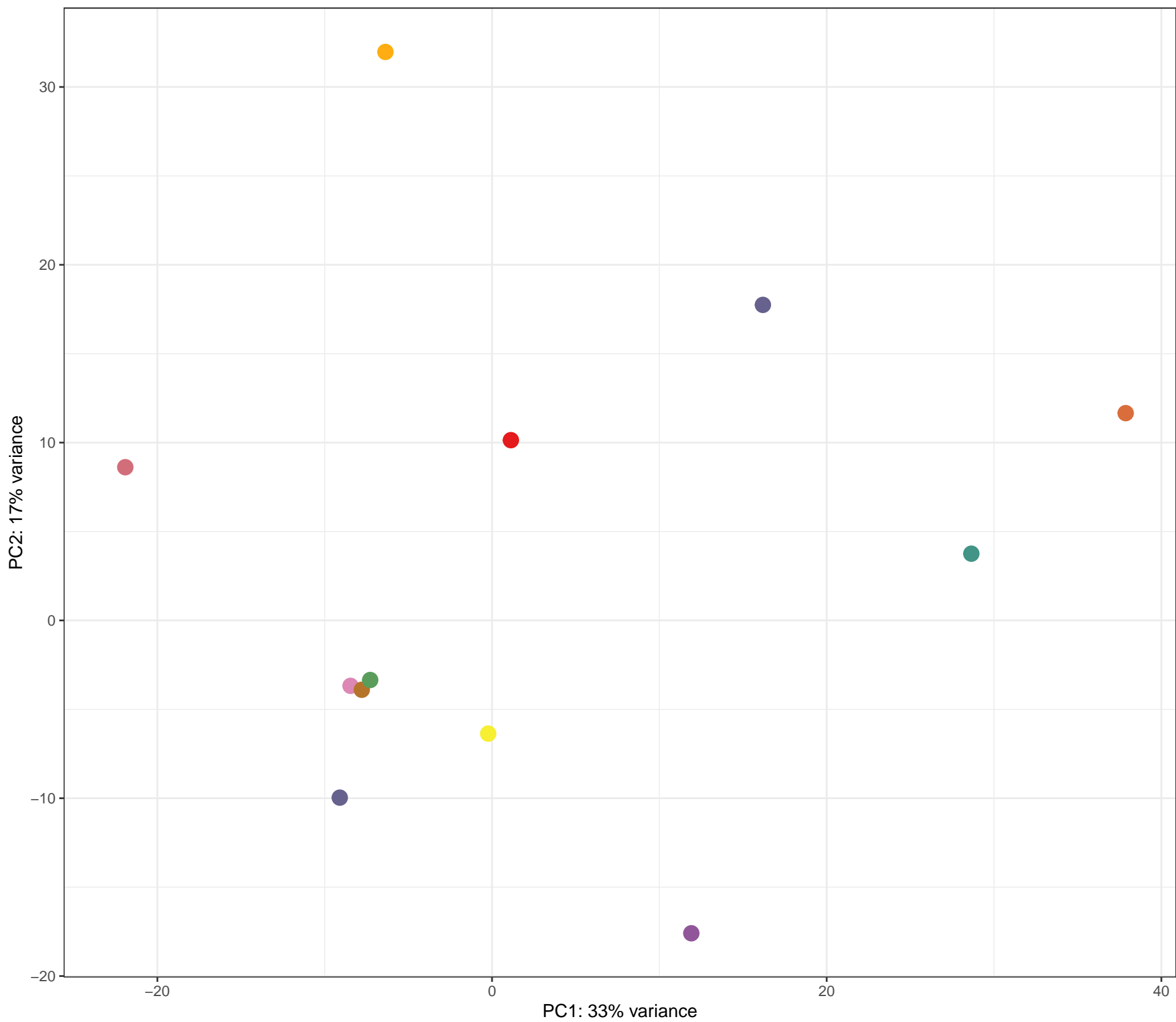


● T-a    ● T-b    ● T-c    ● T-d-primary    ● T-e    ● T-f  
● T-g    ● T-h    ● T-i    ● T-k    ● T-l    ● T-m

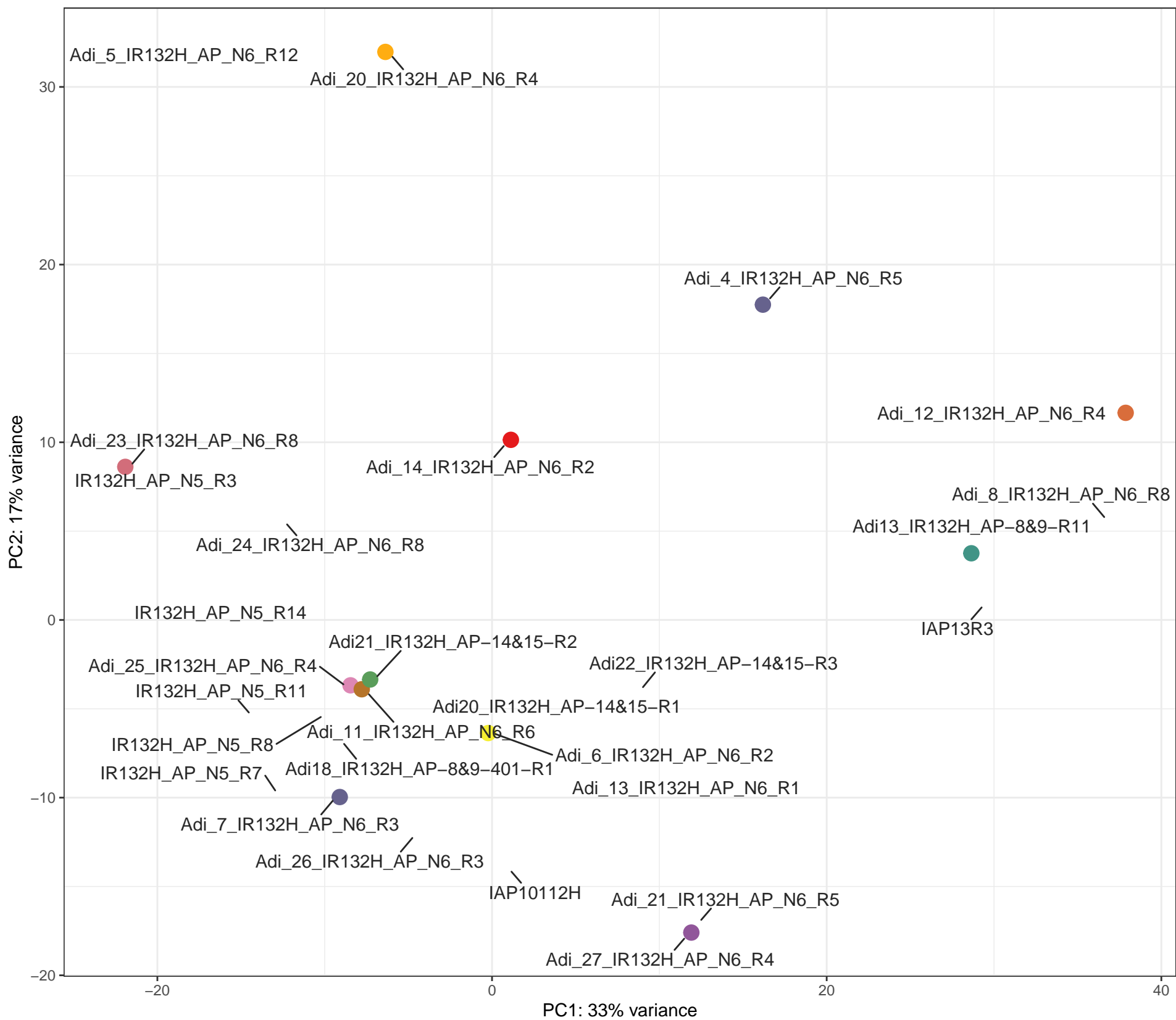
# Tumour\_origin



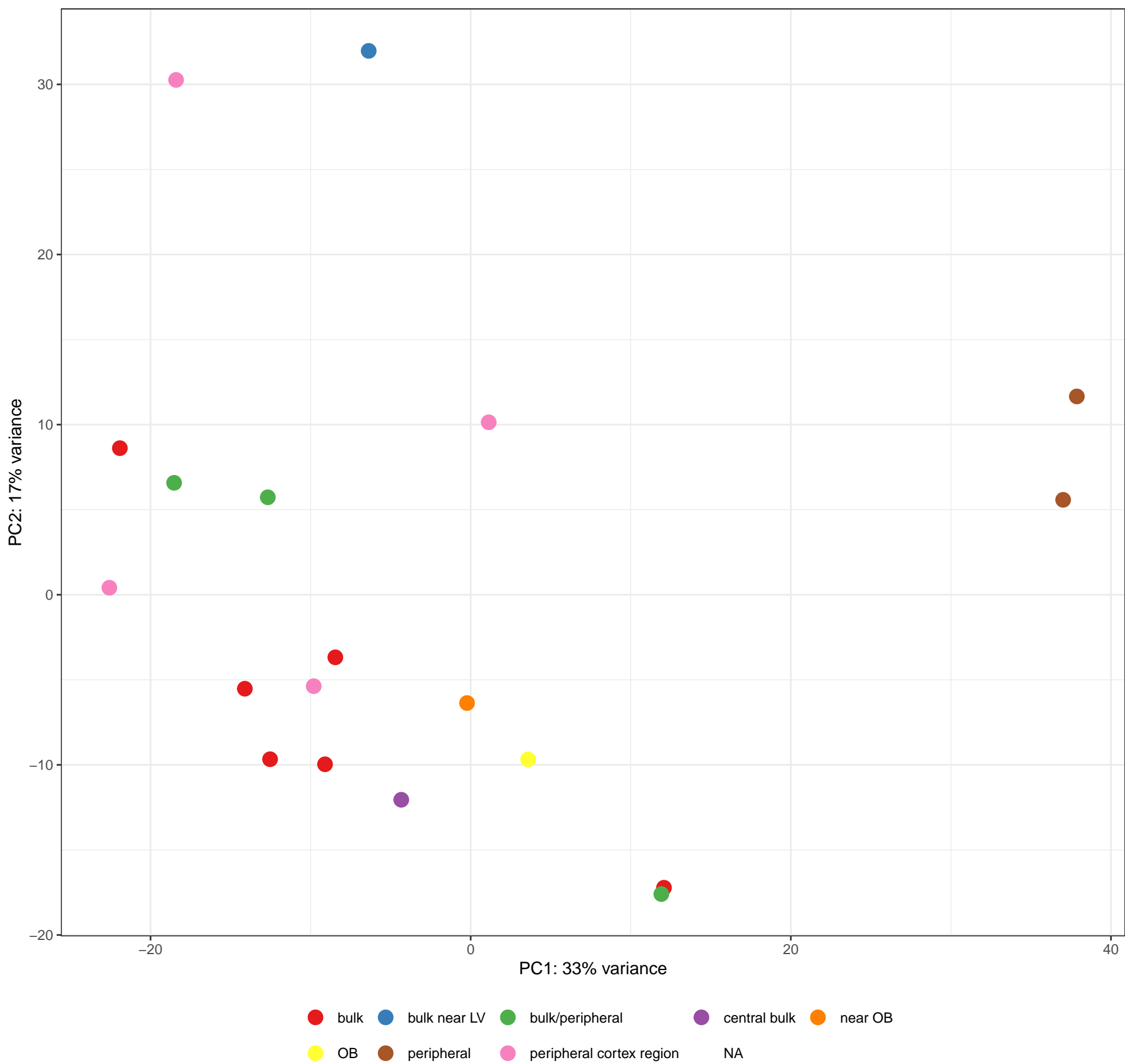
CNA

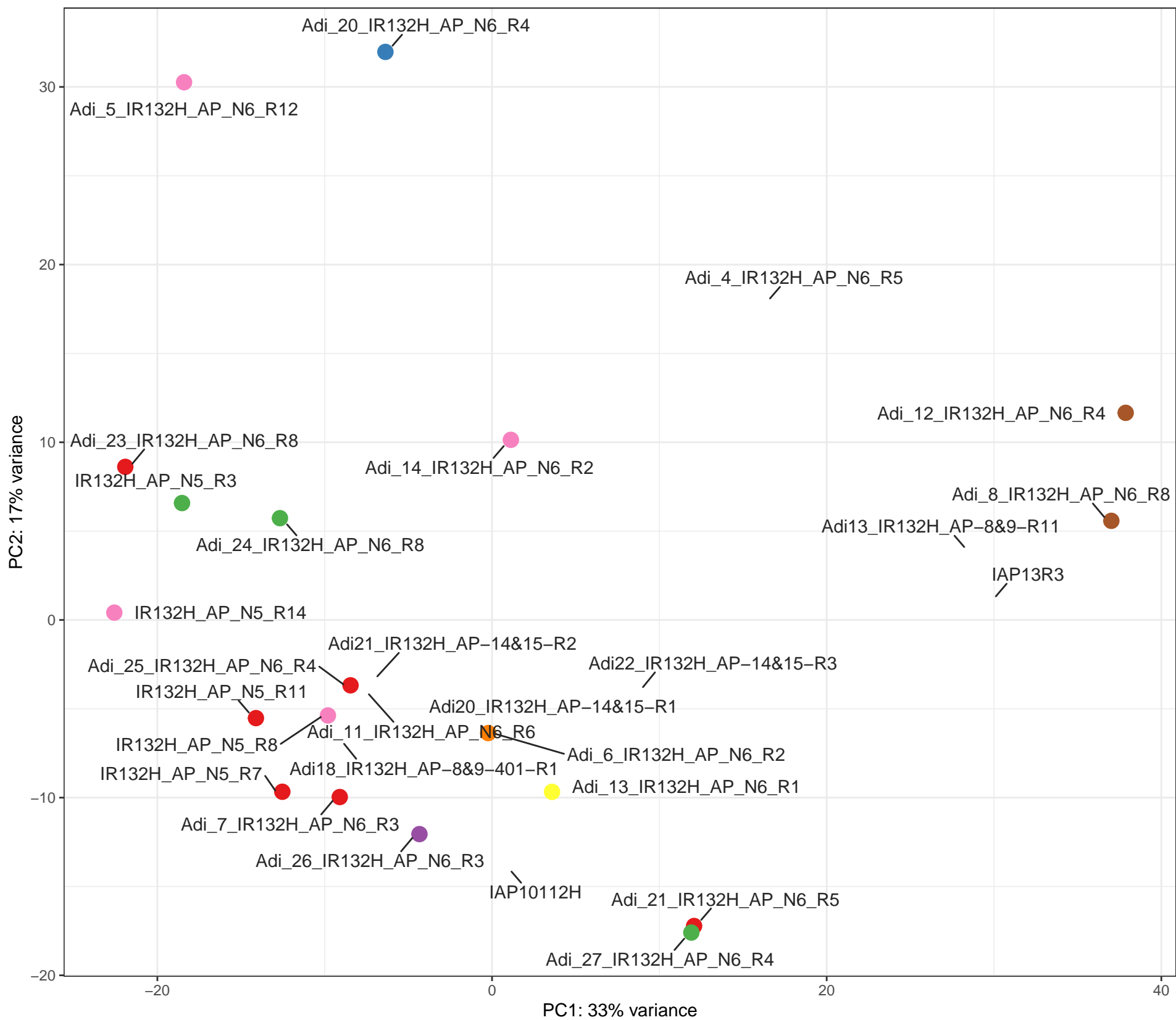


# CNA

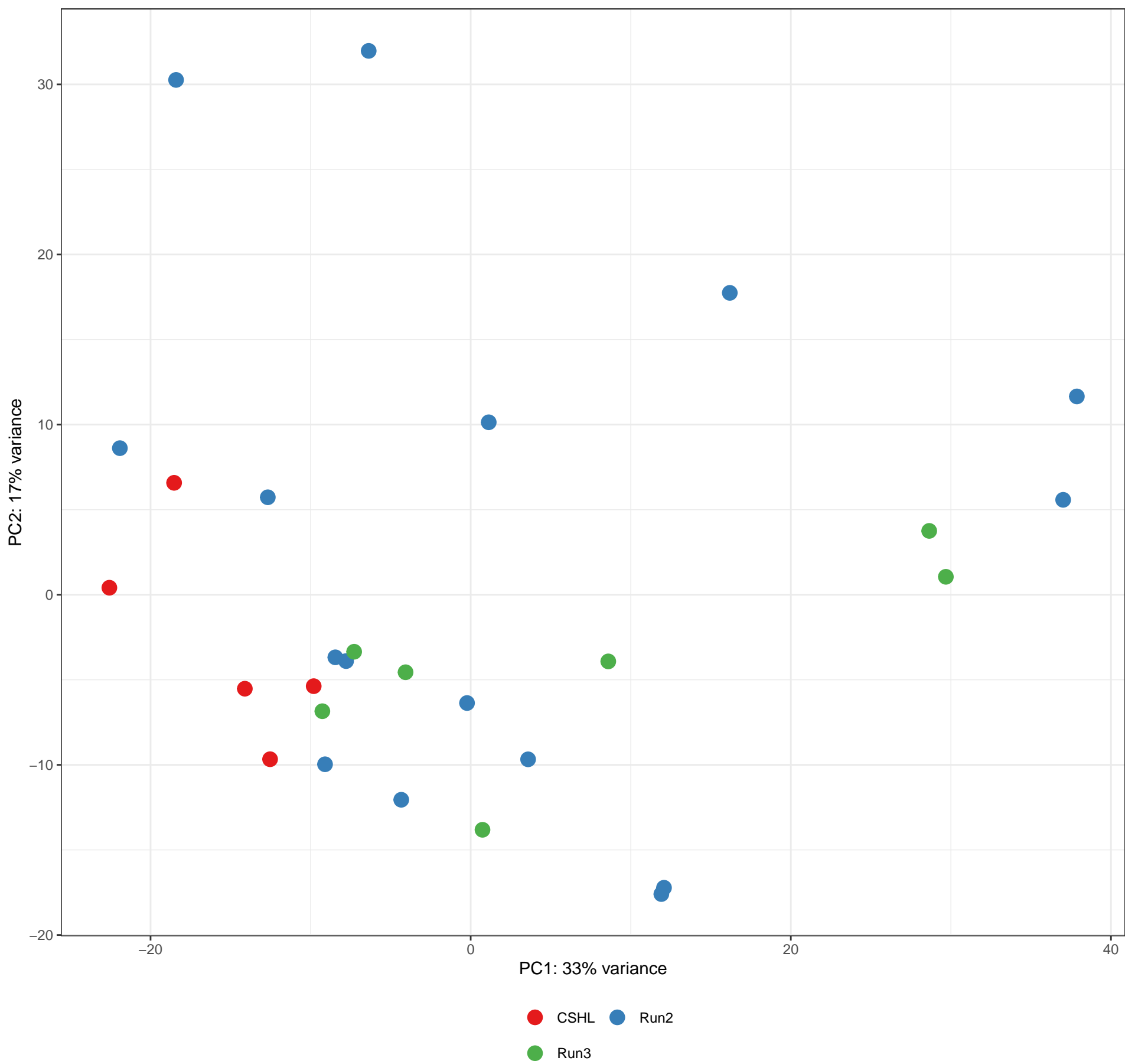


Info

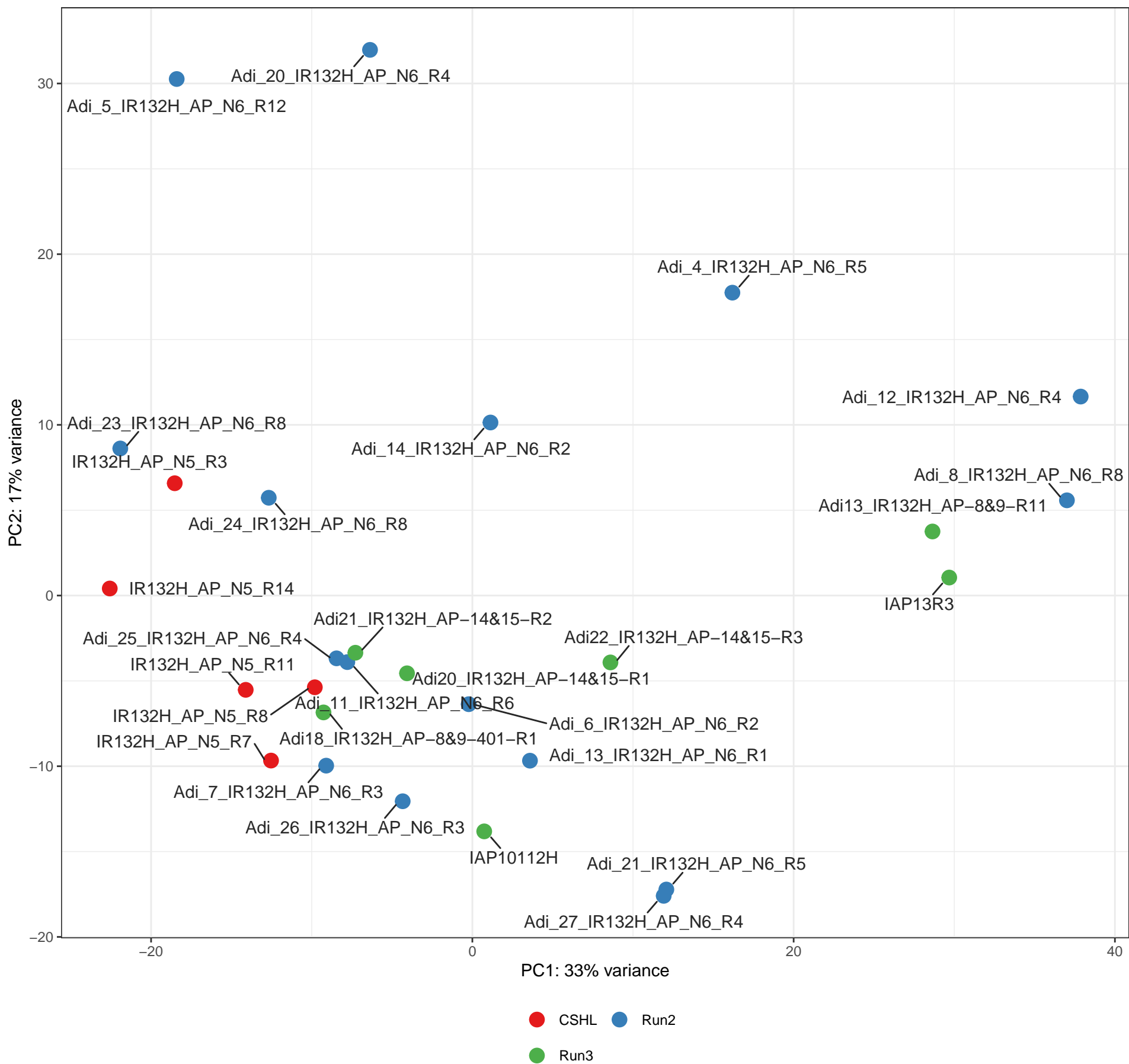




Run

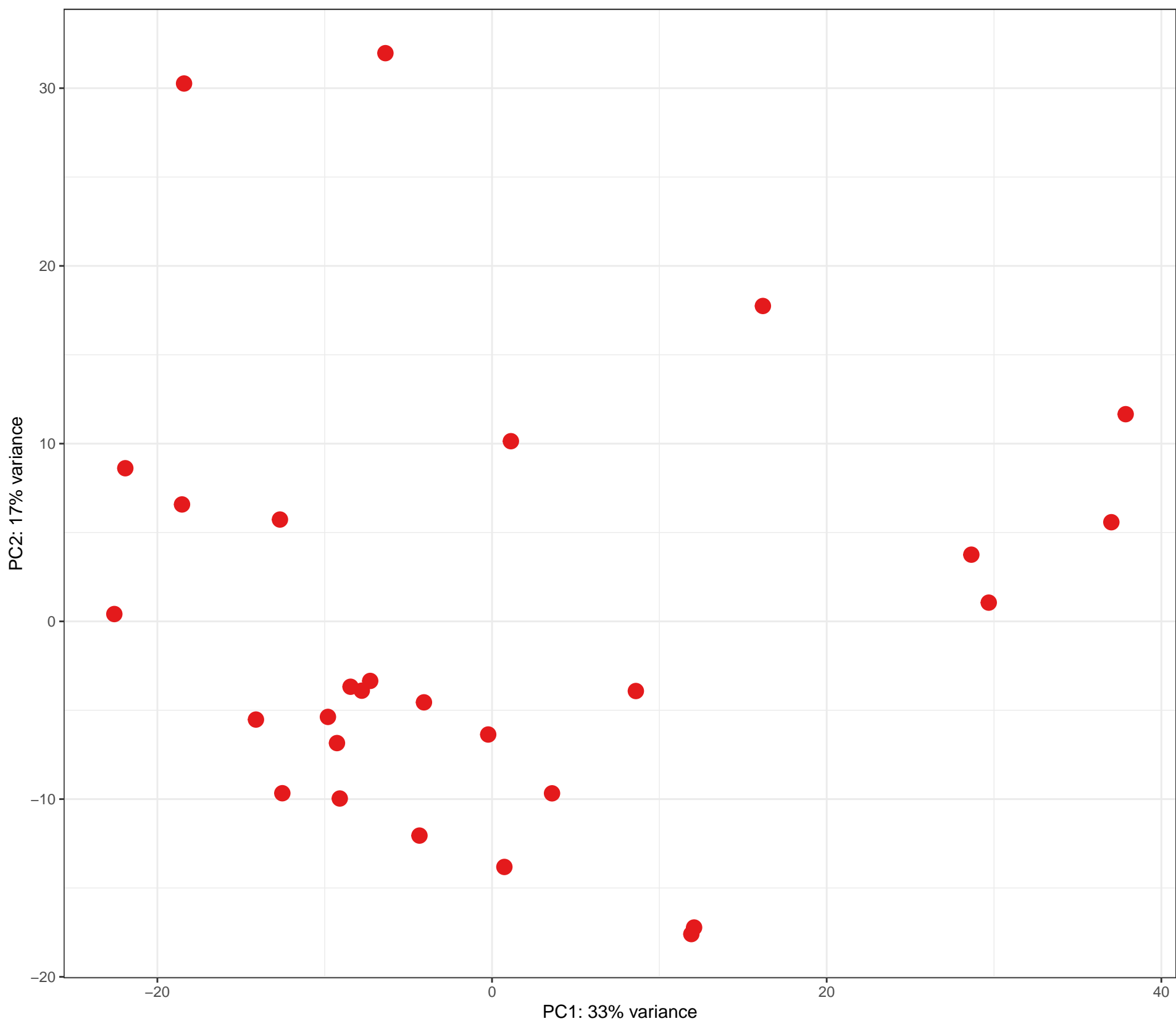


Run



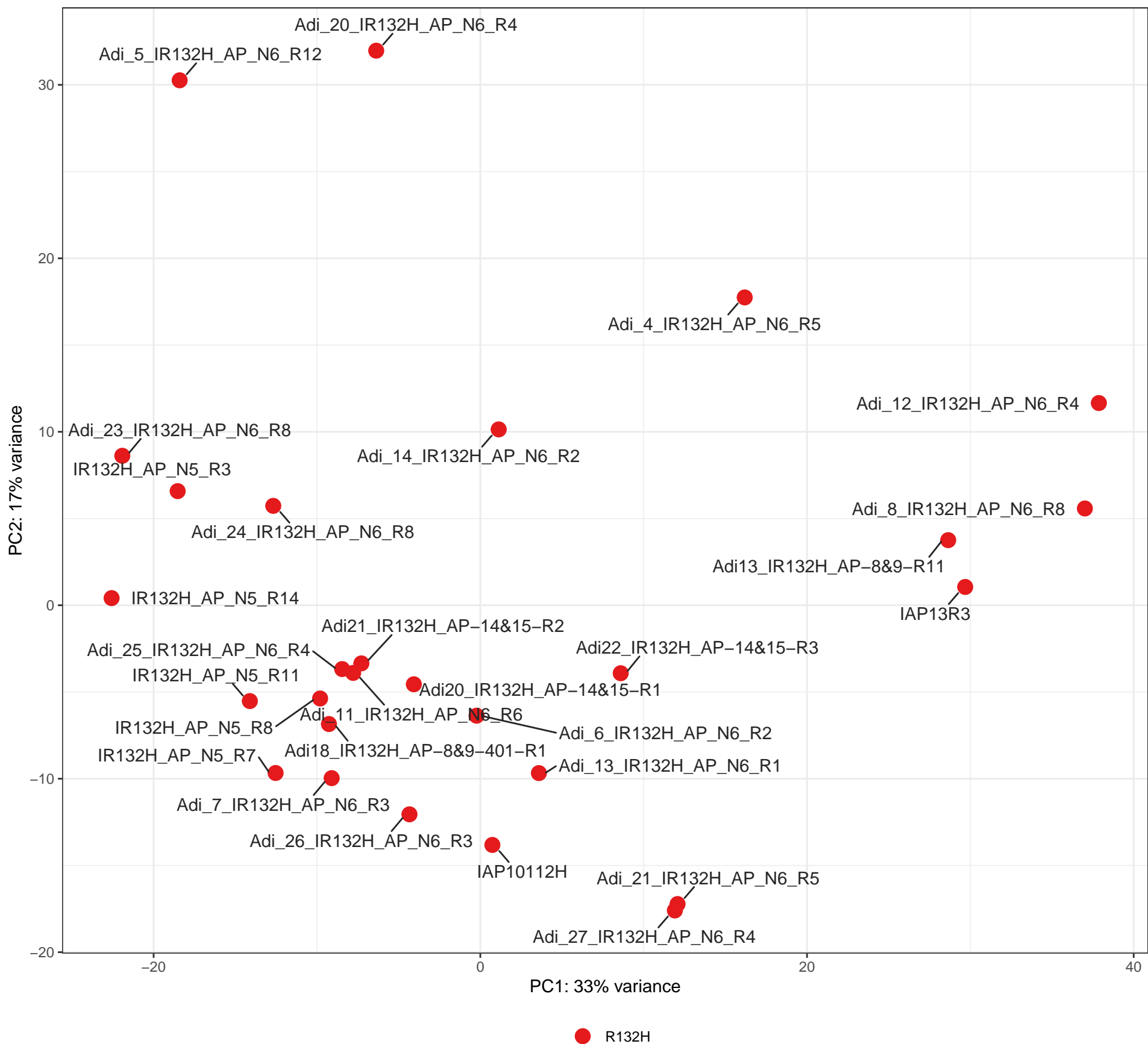


IDH

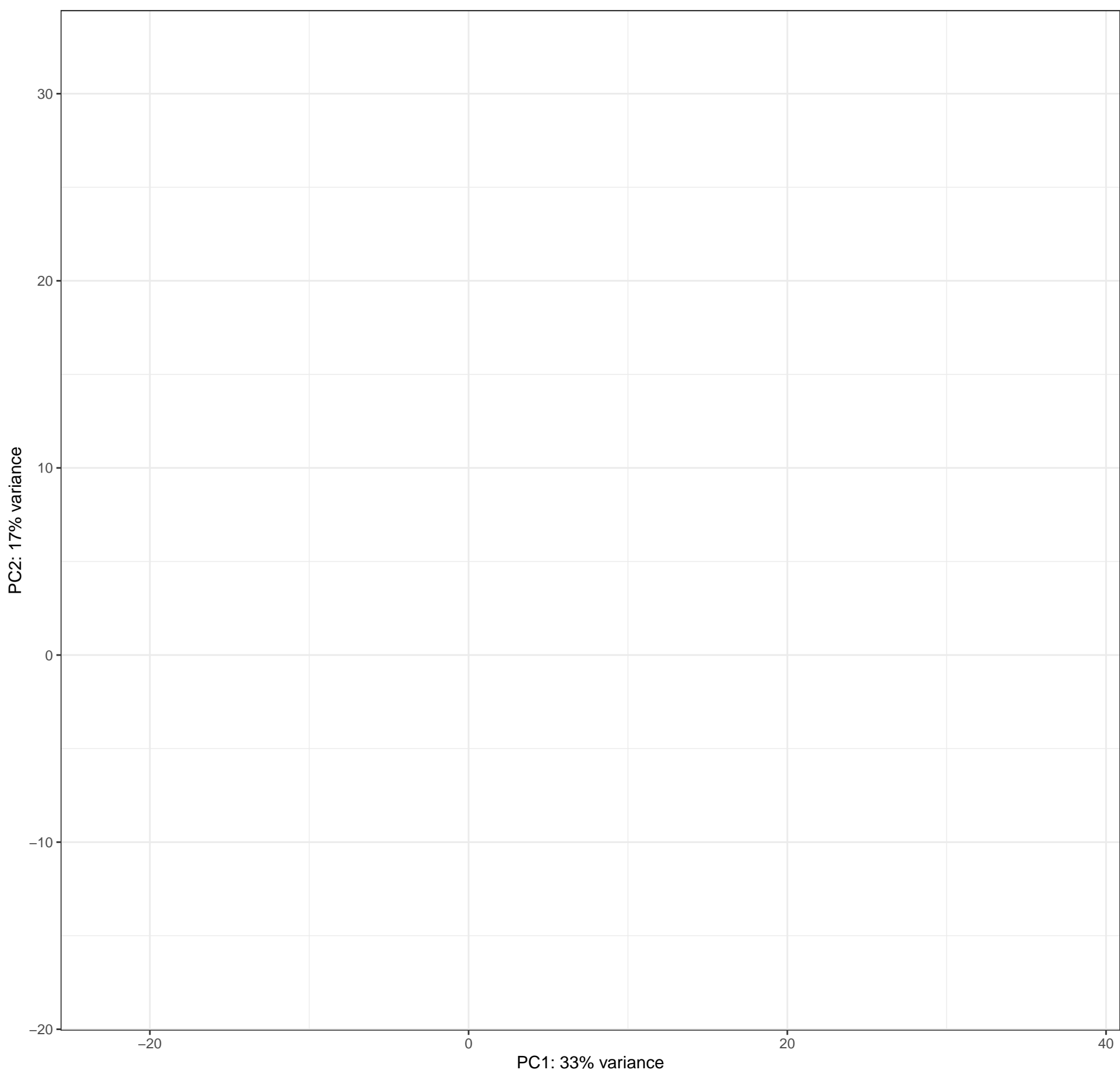


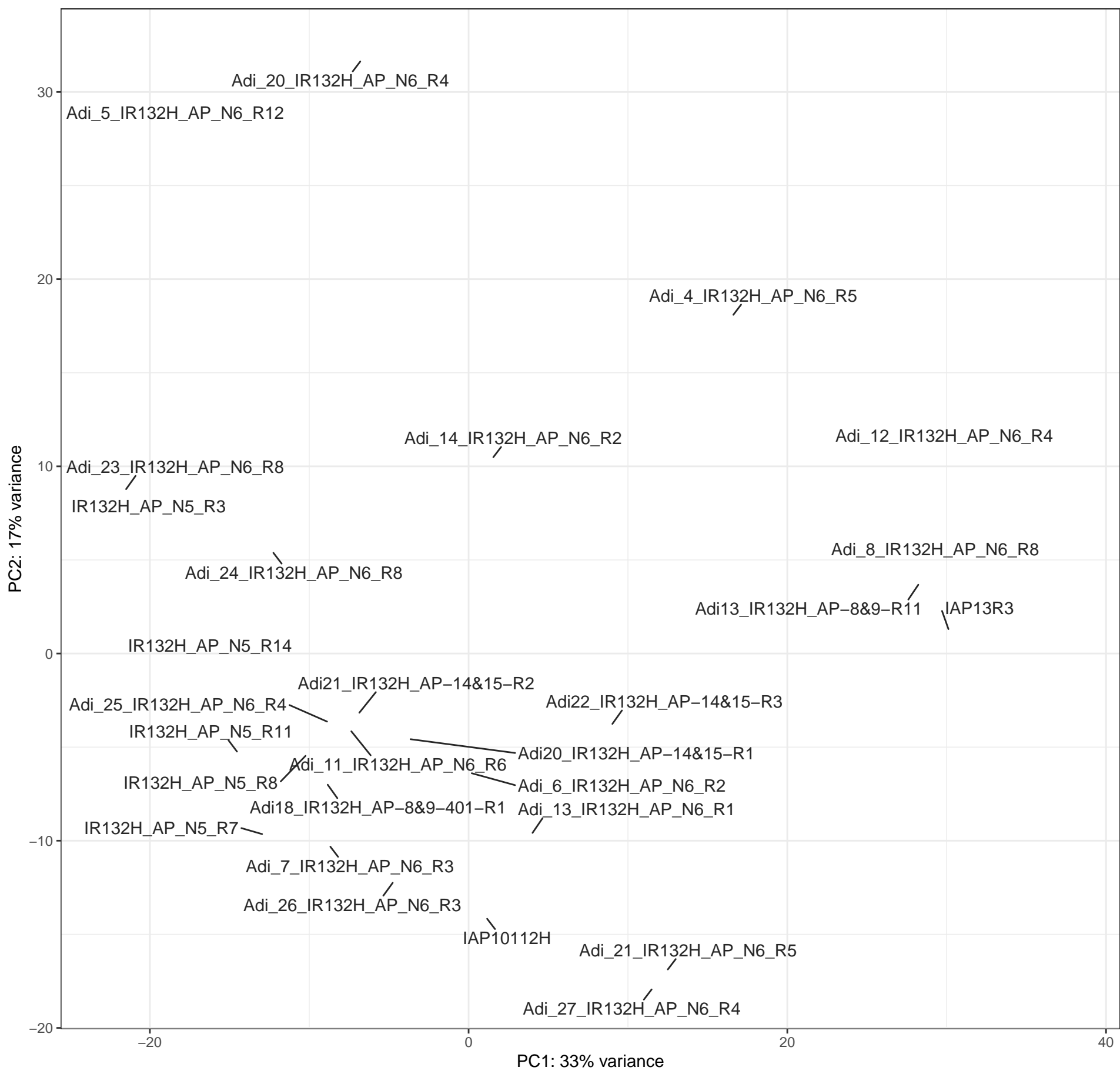
● R132H

IDH

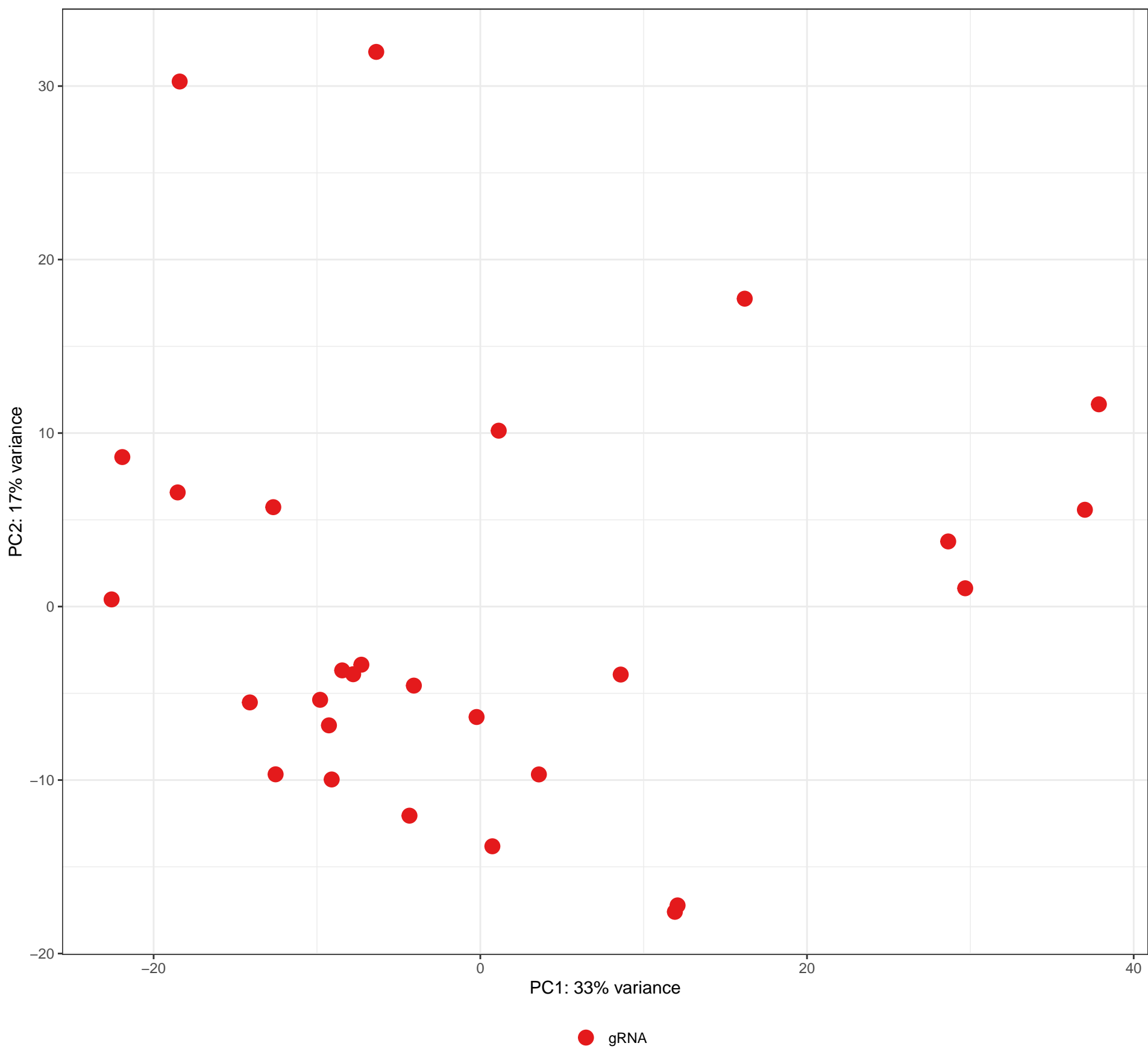


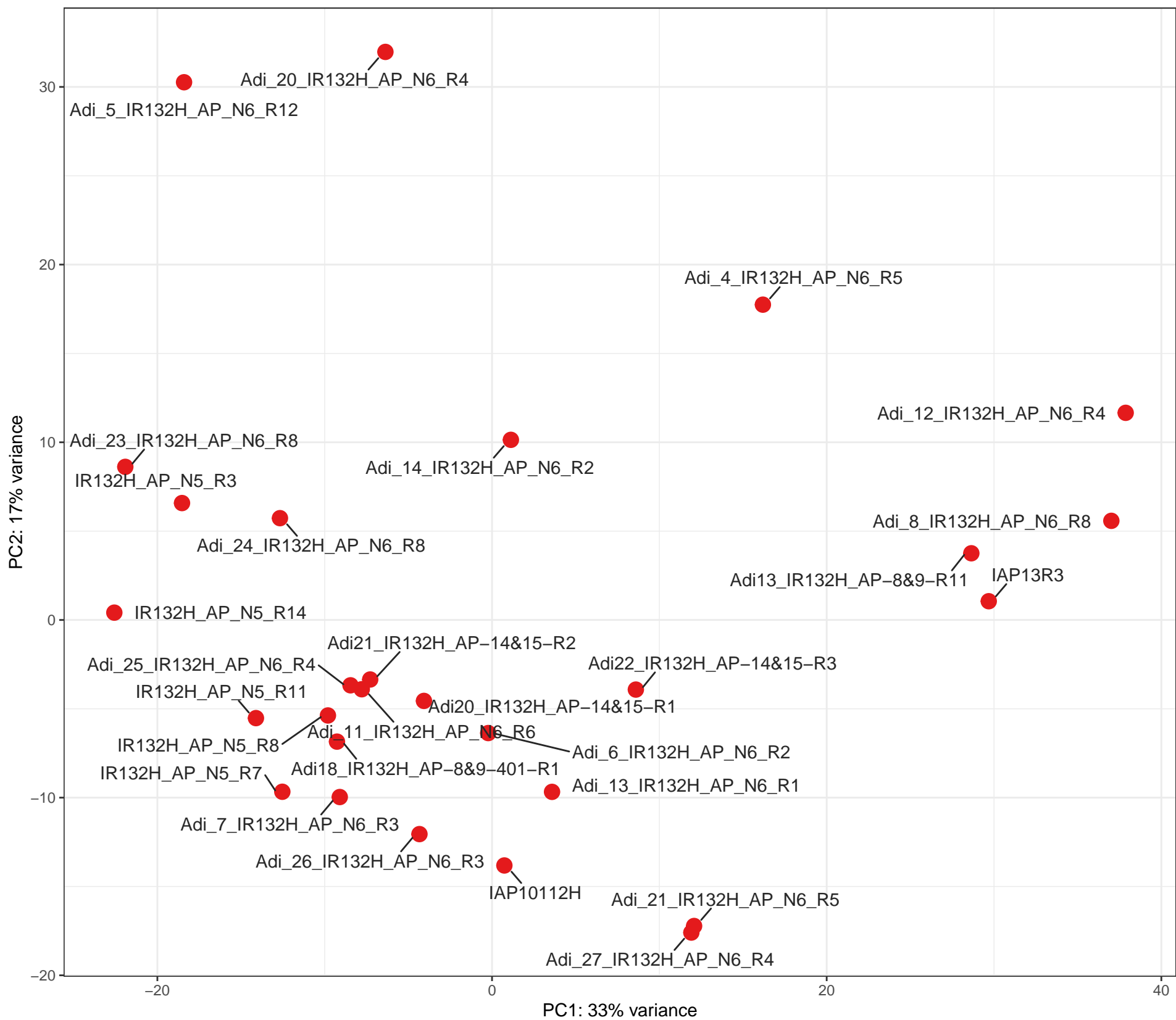
PDGFRA





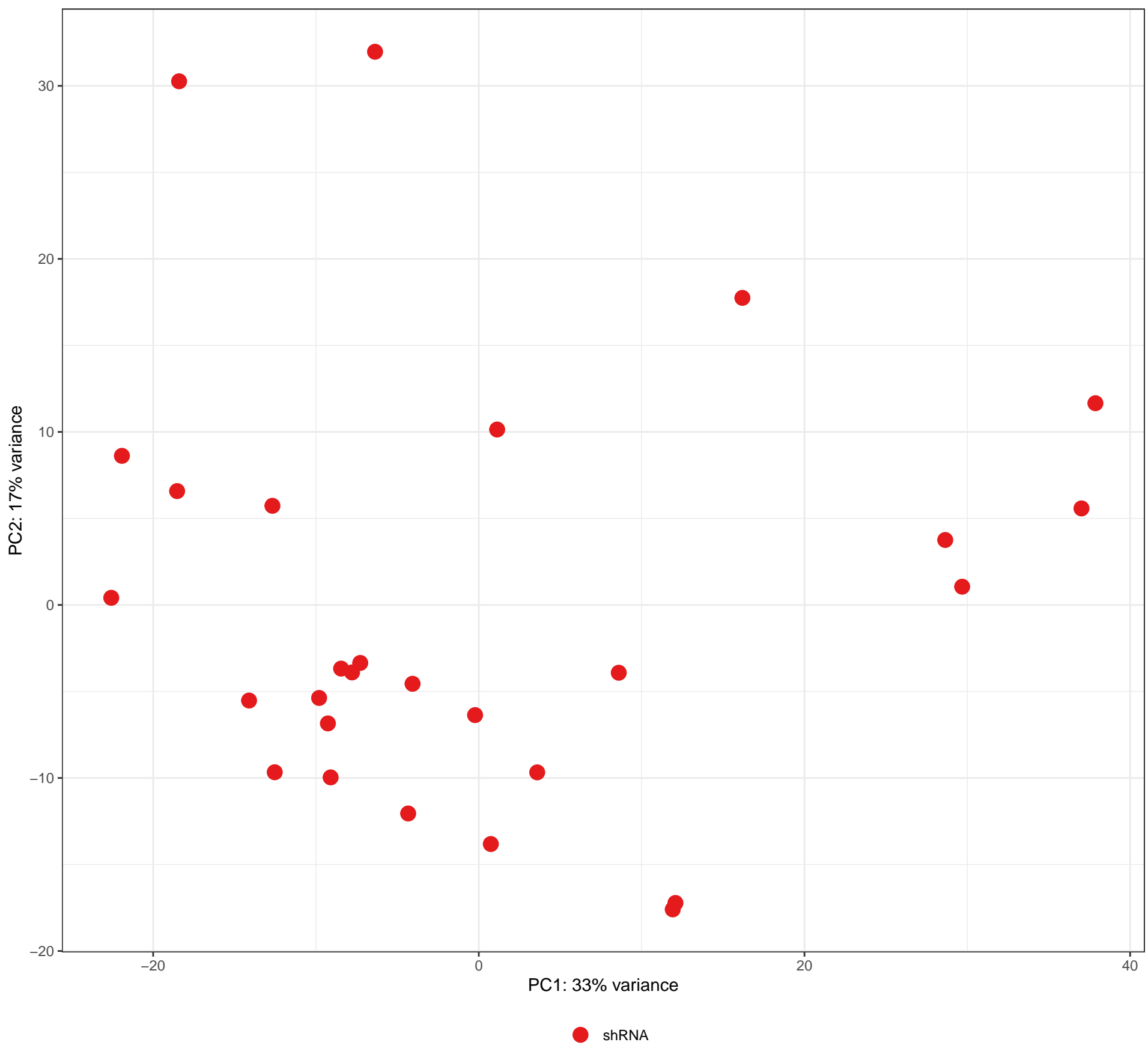
P53

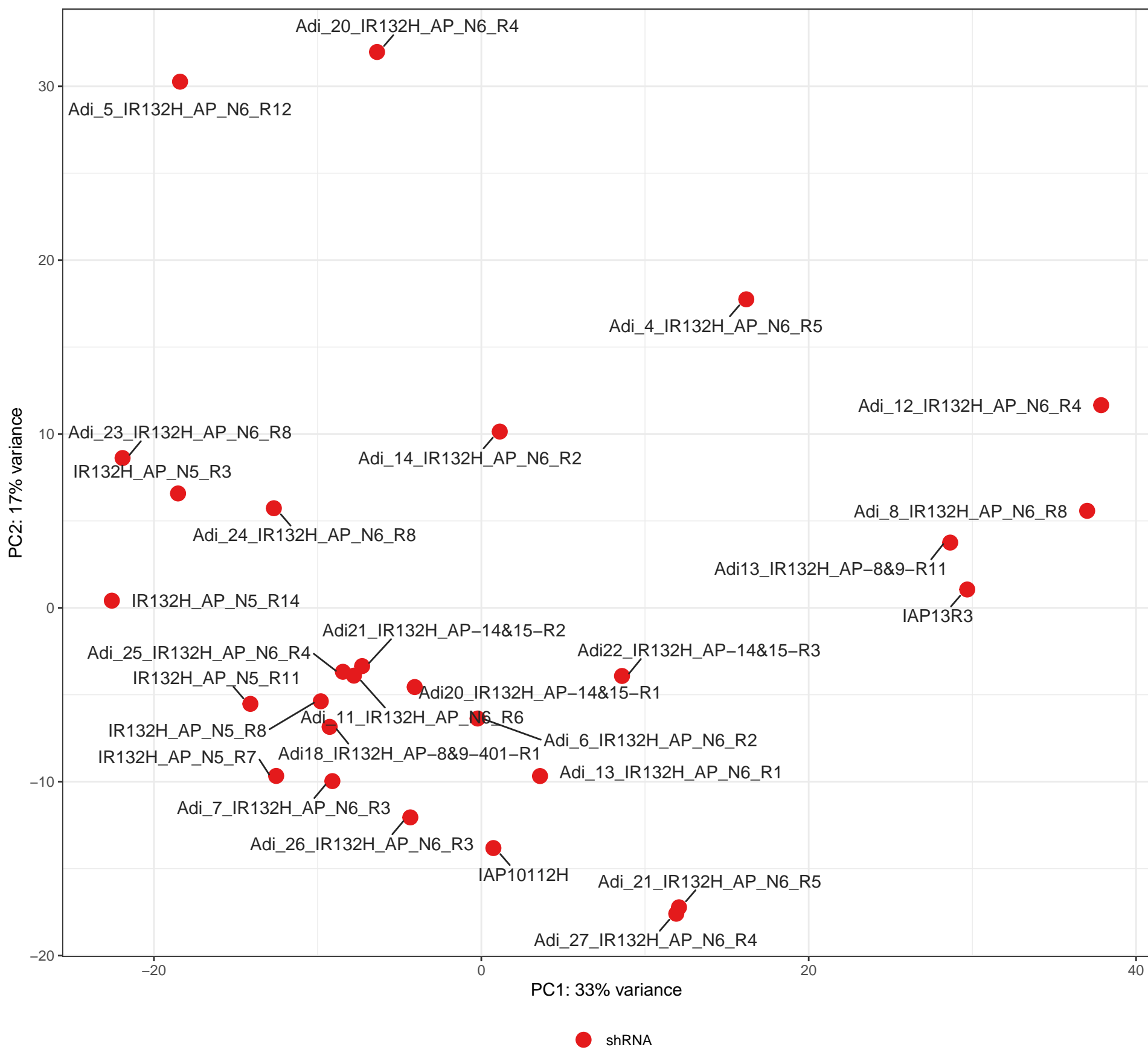




● gRNA

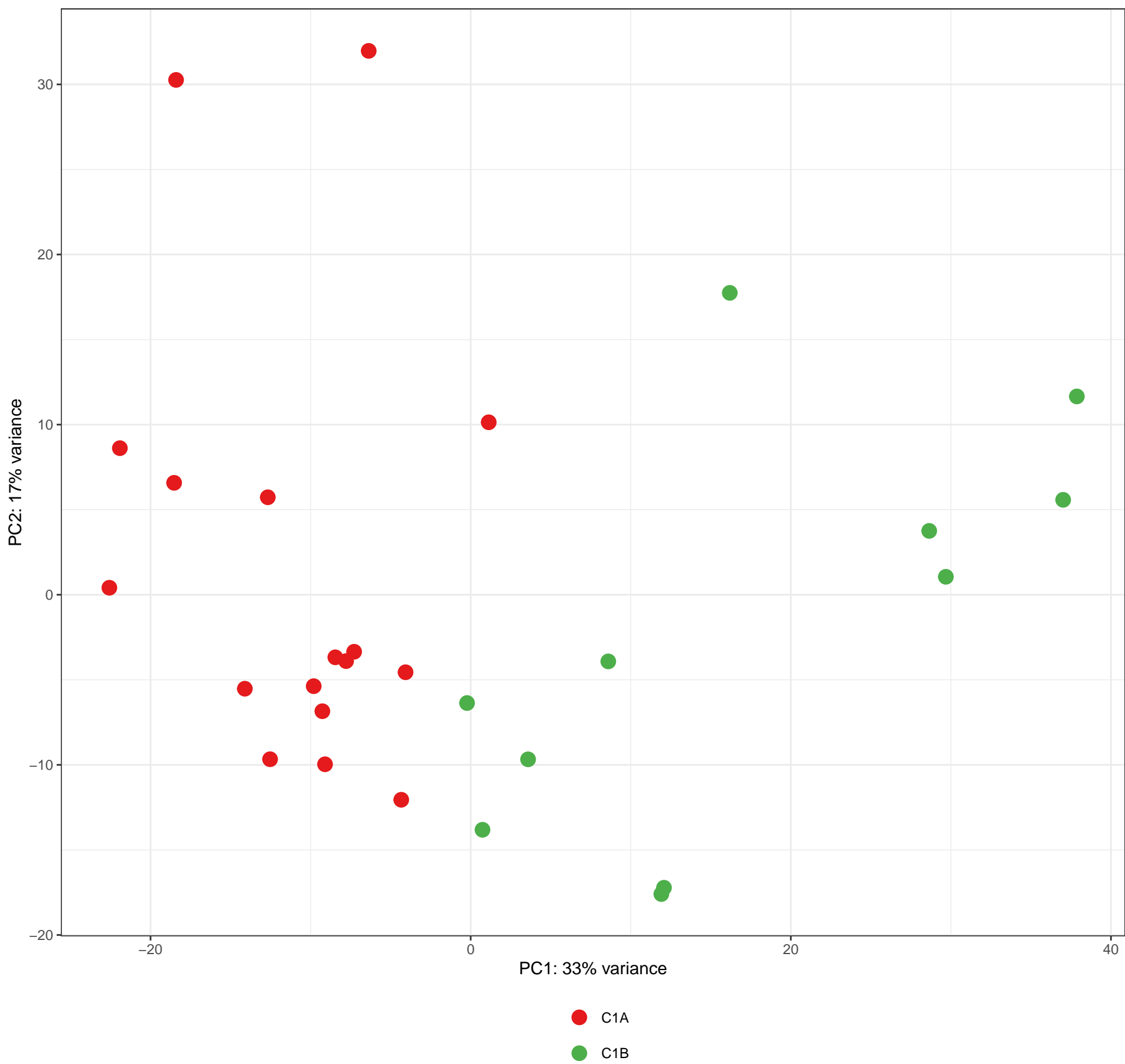
ATRX



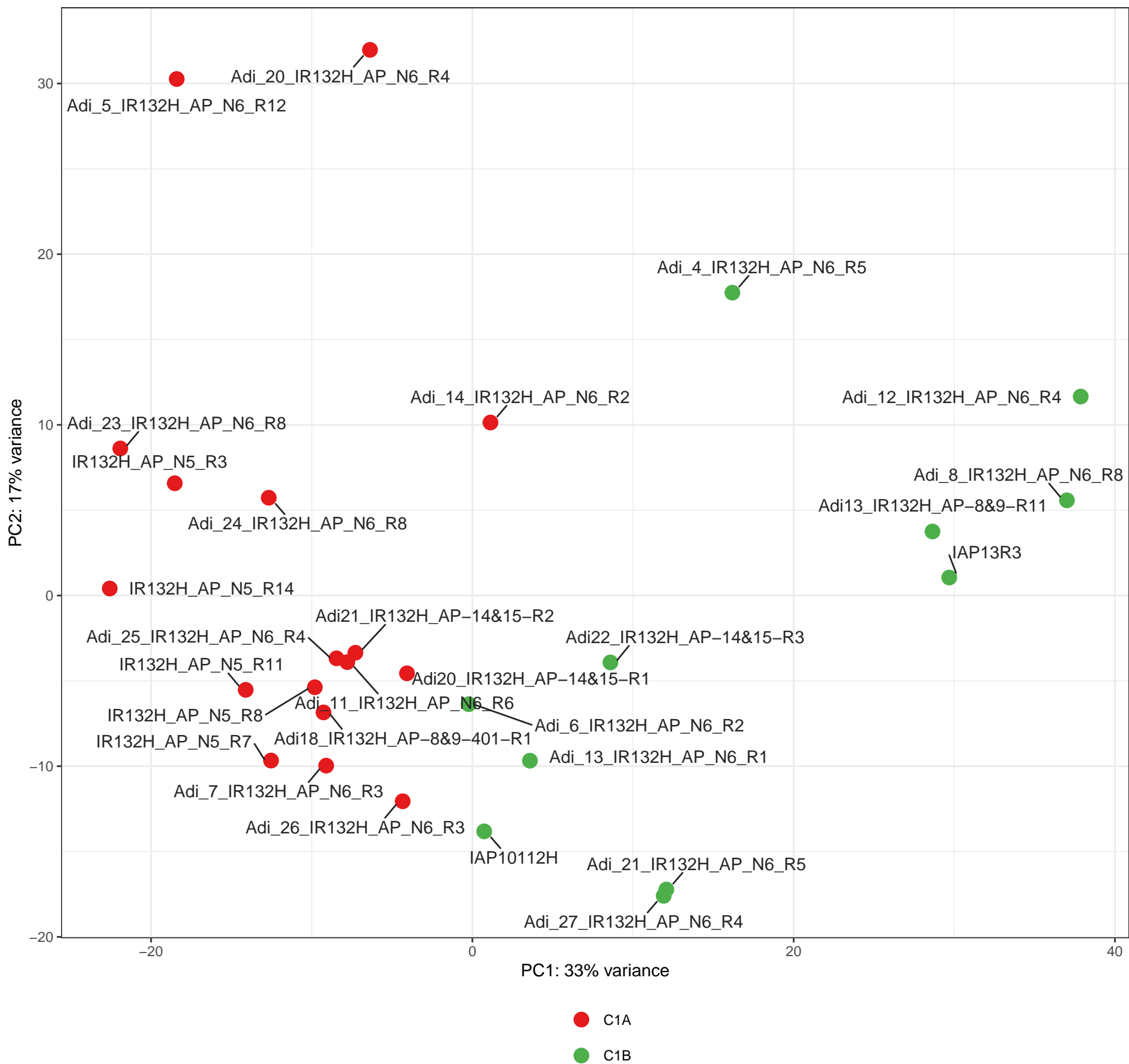




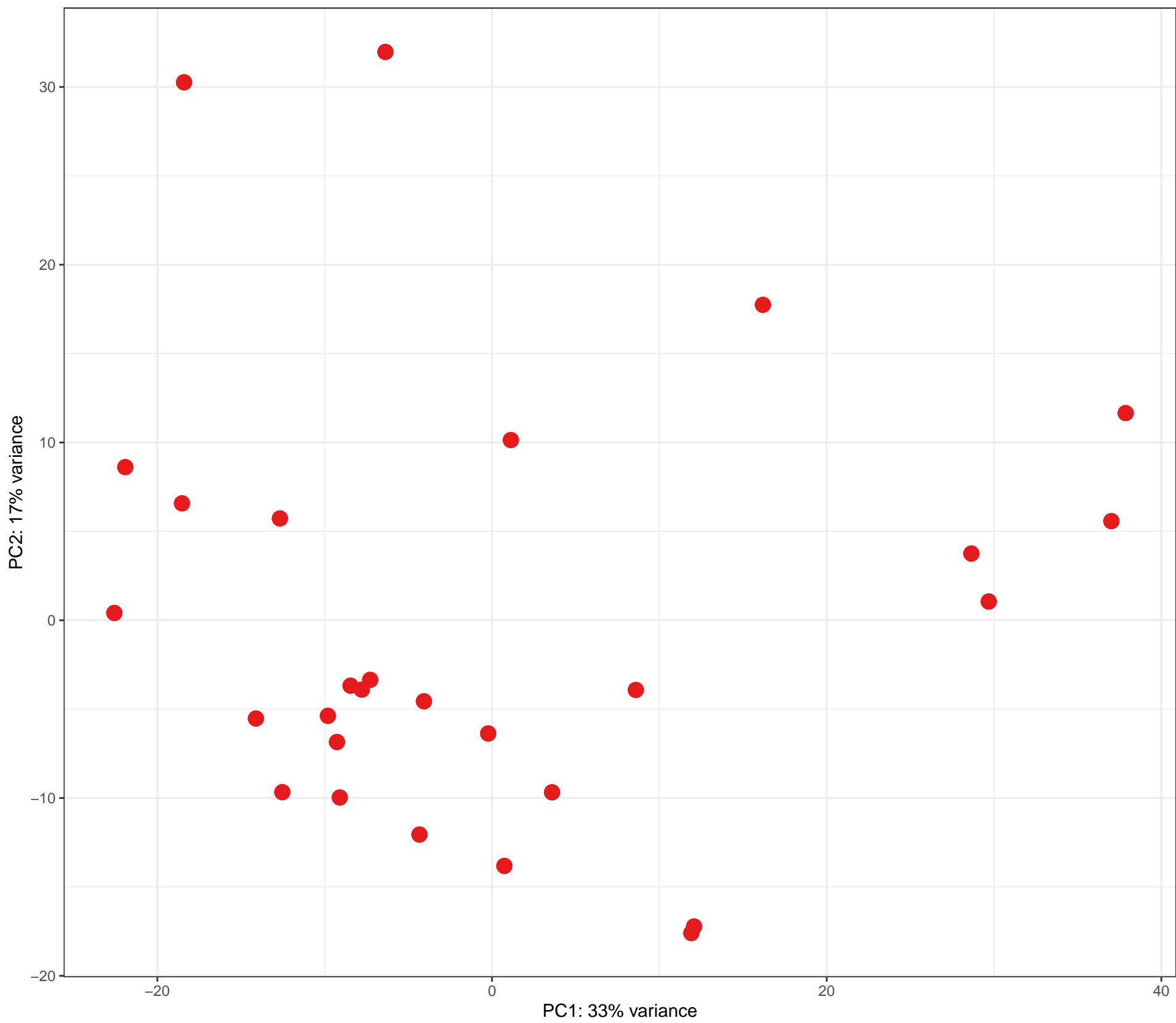
Condition



Condition

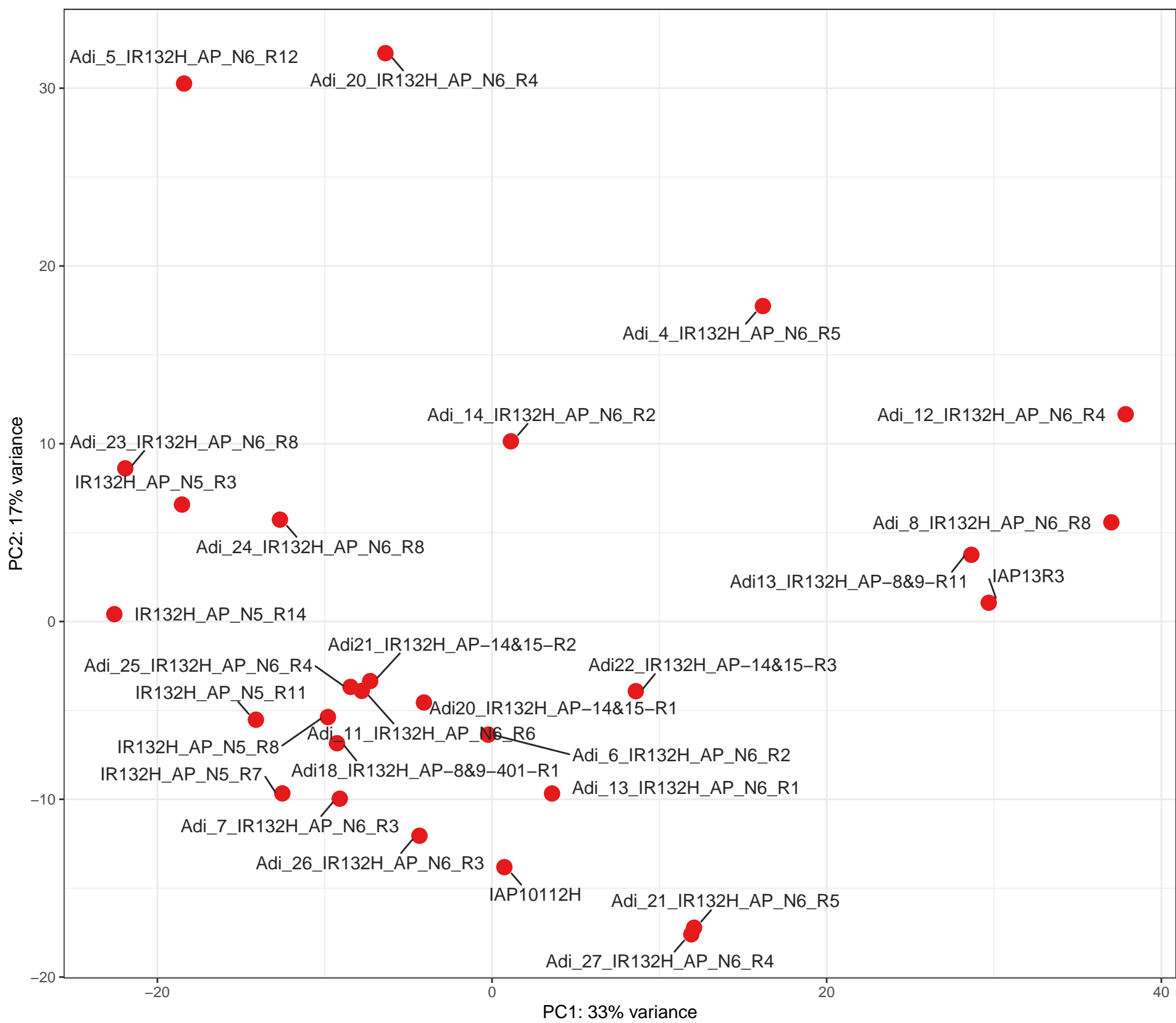


Grp



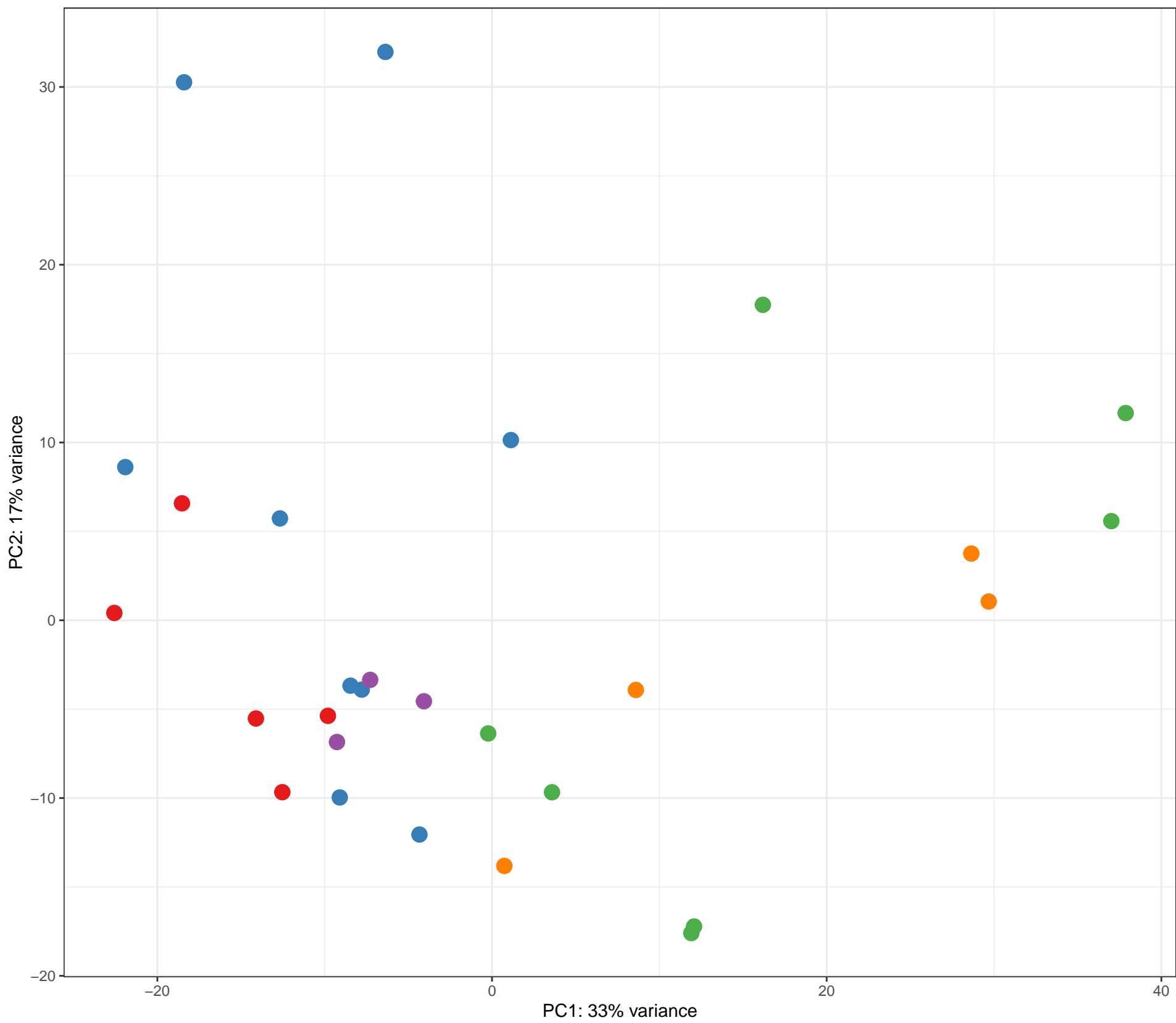
● C1

Grp



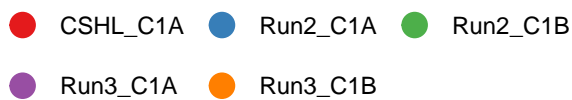
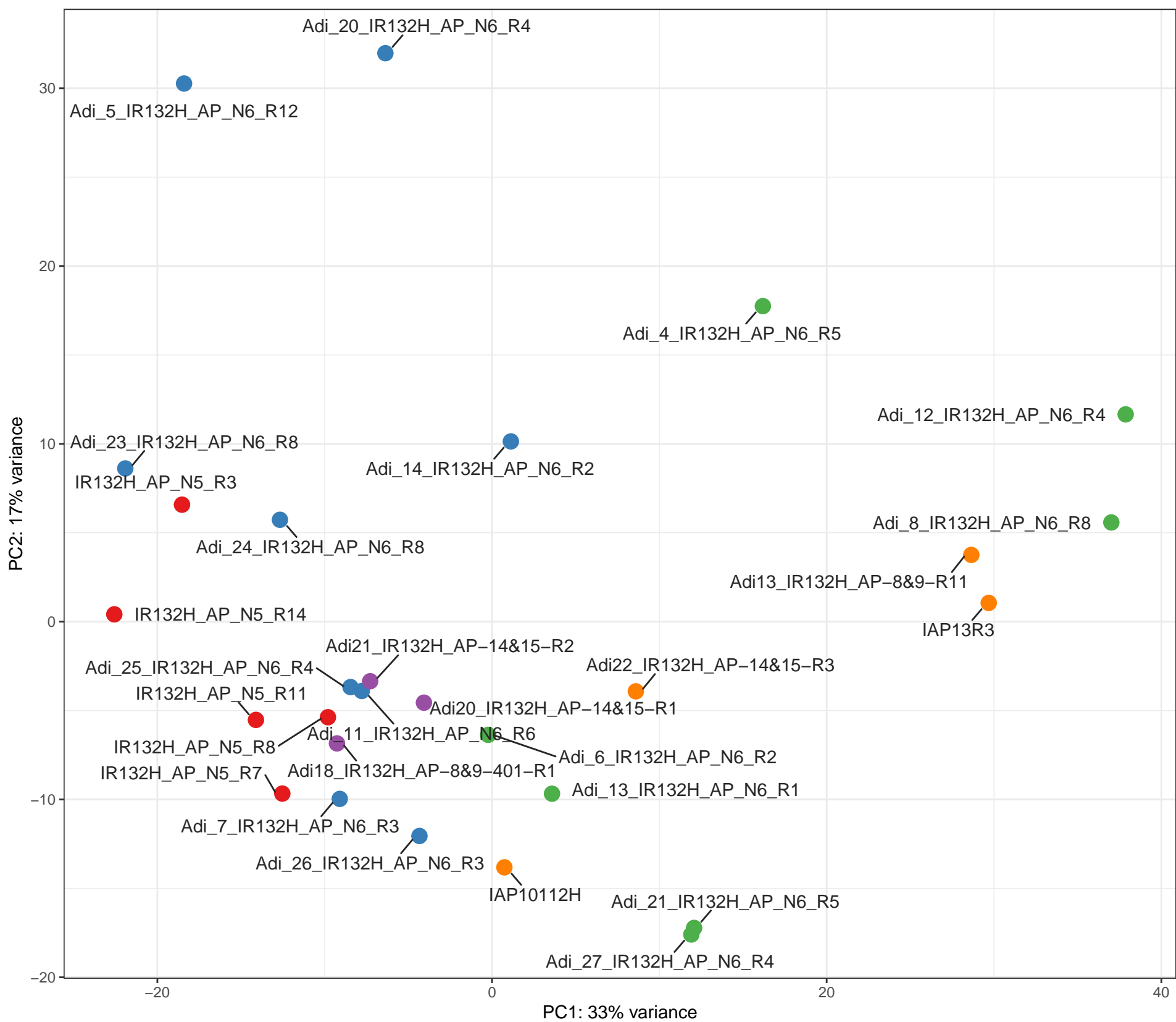
● C1

Type

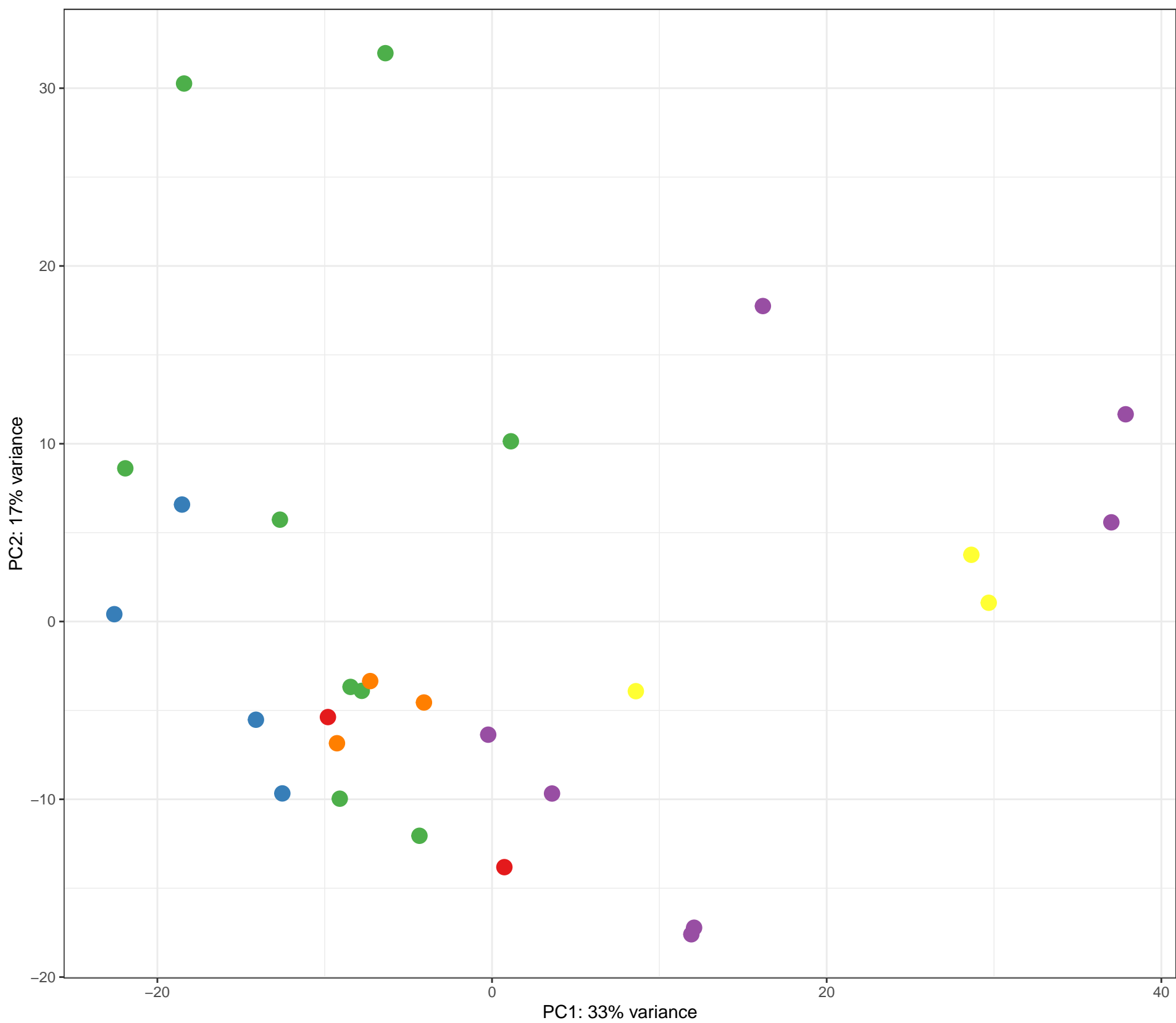


CSHL\_C1A Run2\_C1A Run2\_C1B  
Run3\_C1A Run3\_C1B

Type

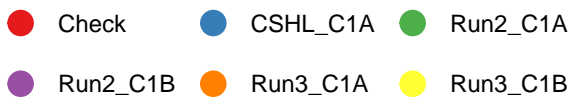
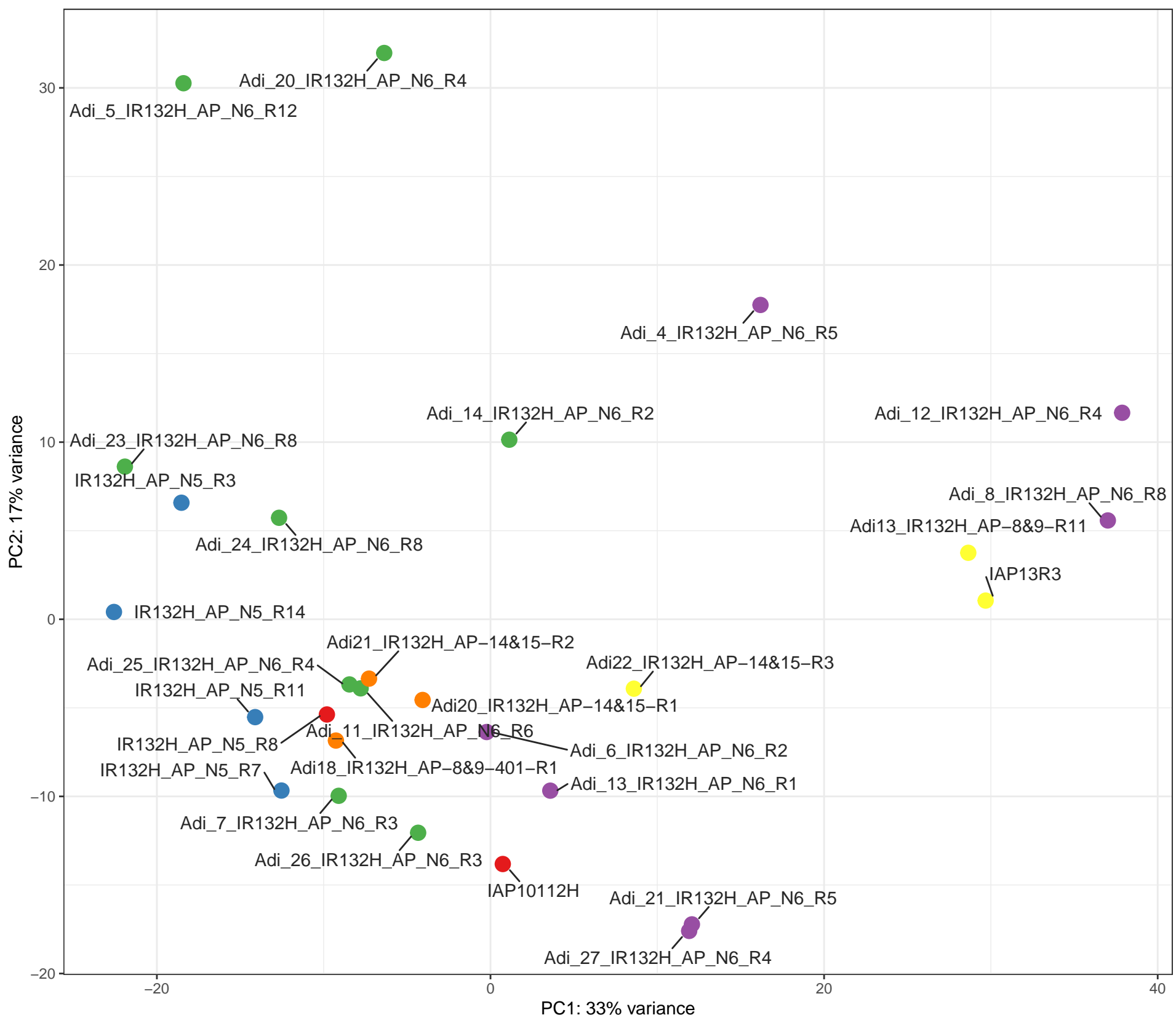


Mark



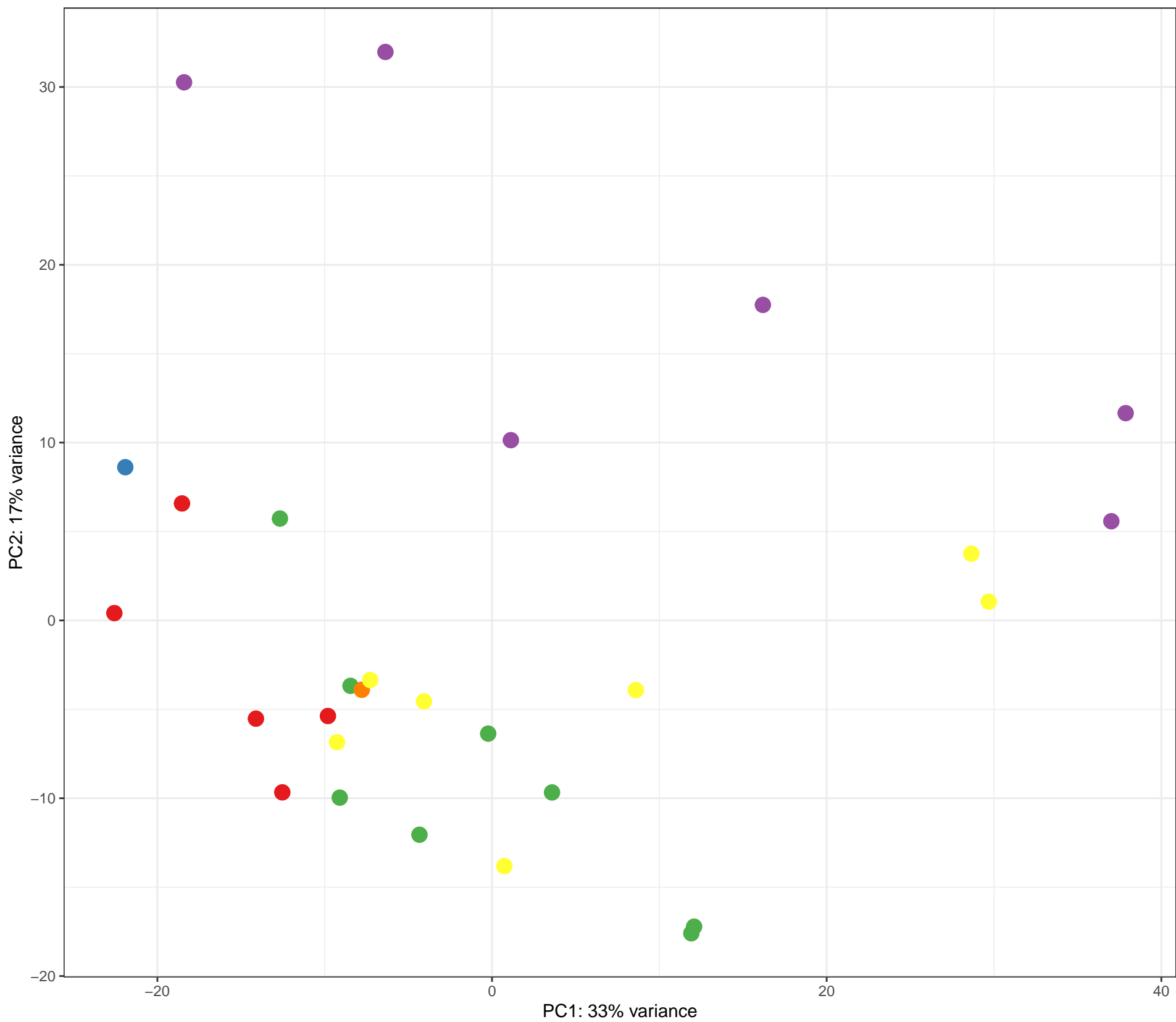
Check CSHL\_C1A Run2\_C1A  
Run2\_C1B Run3\_C1A Run3\_C1B

Mark





Group



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

Group

