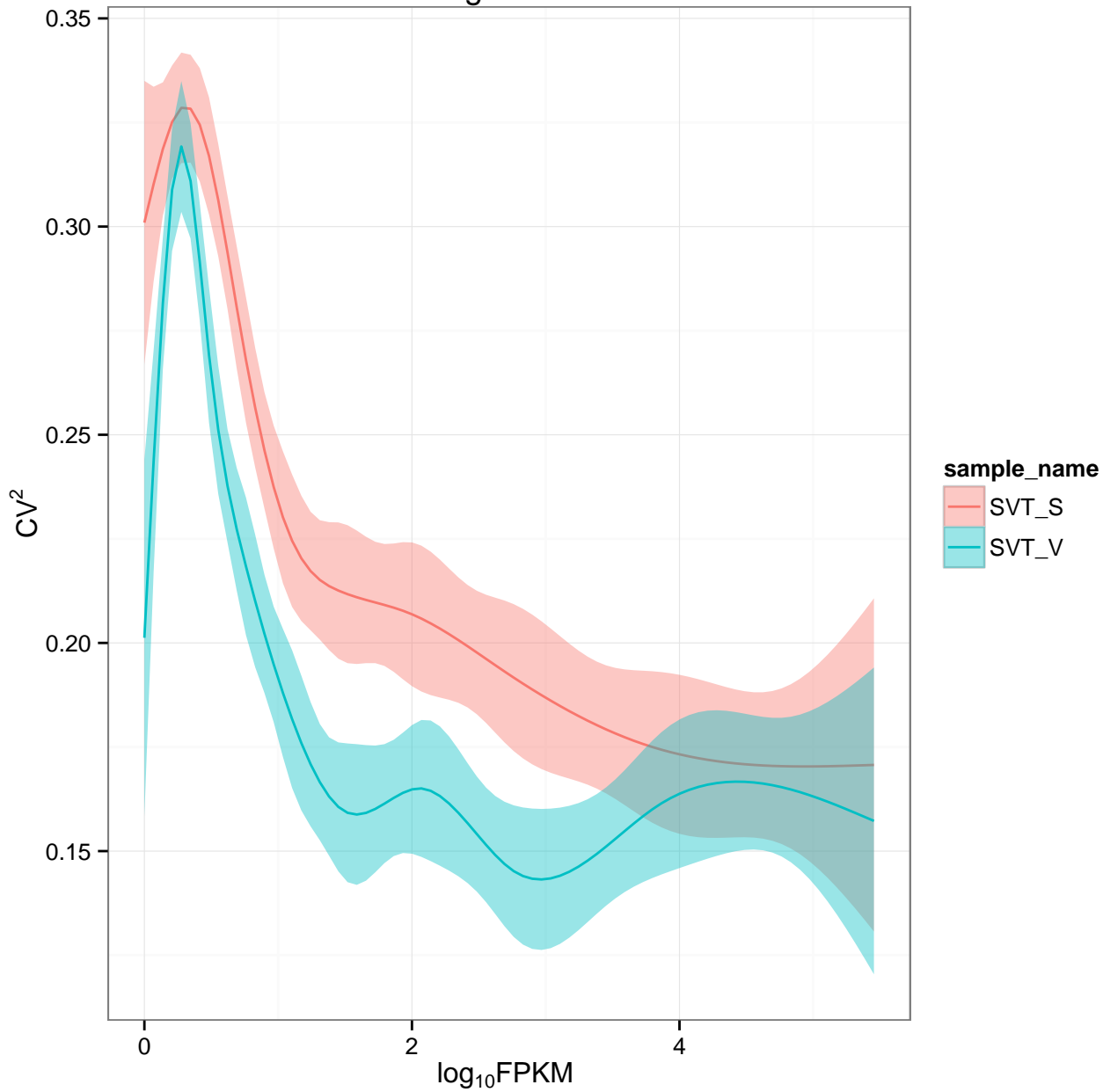
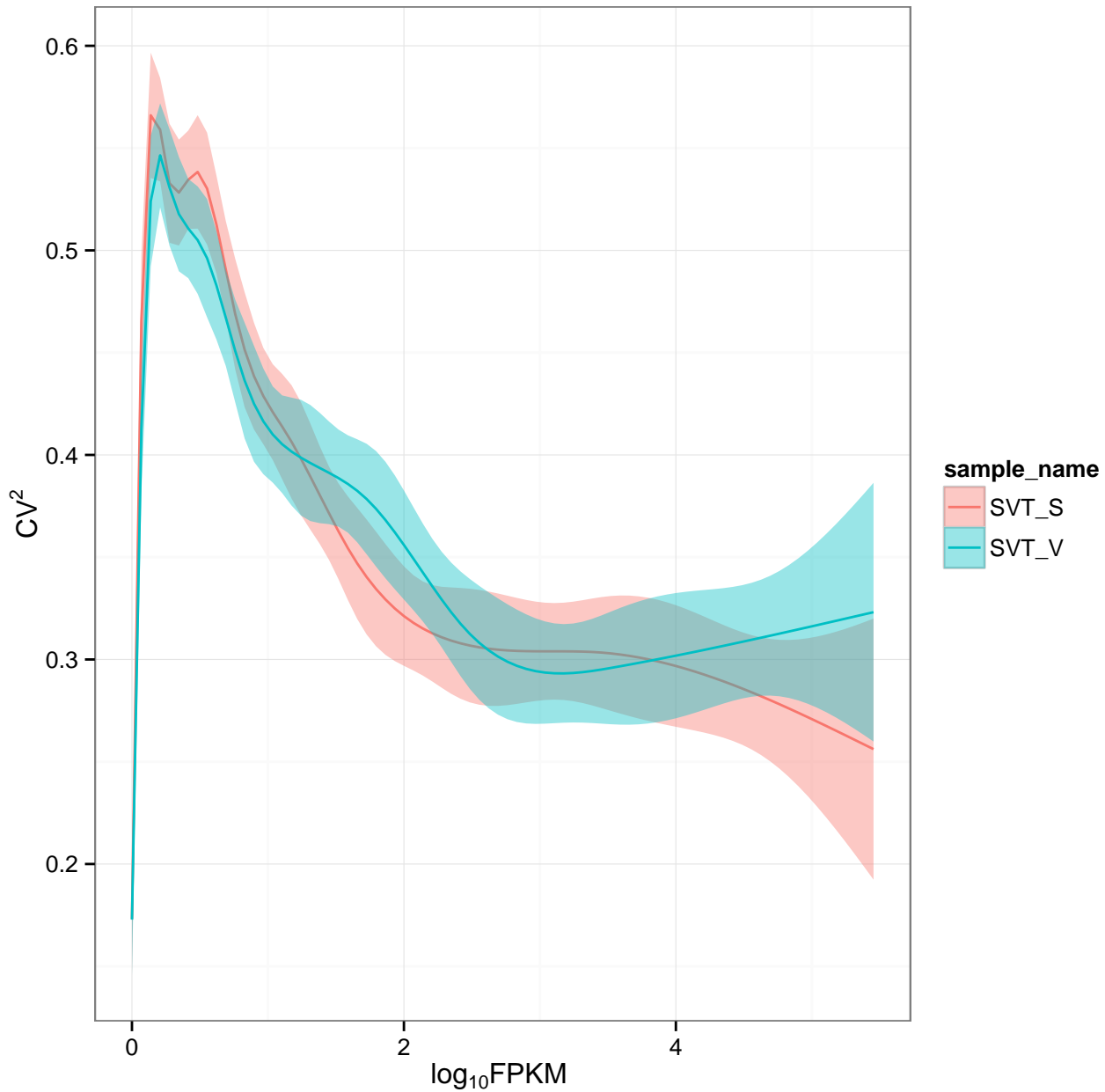


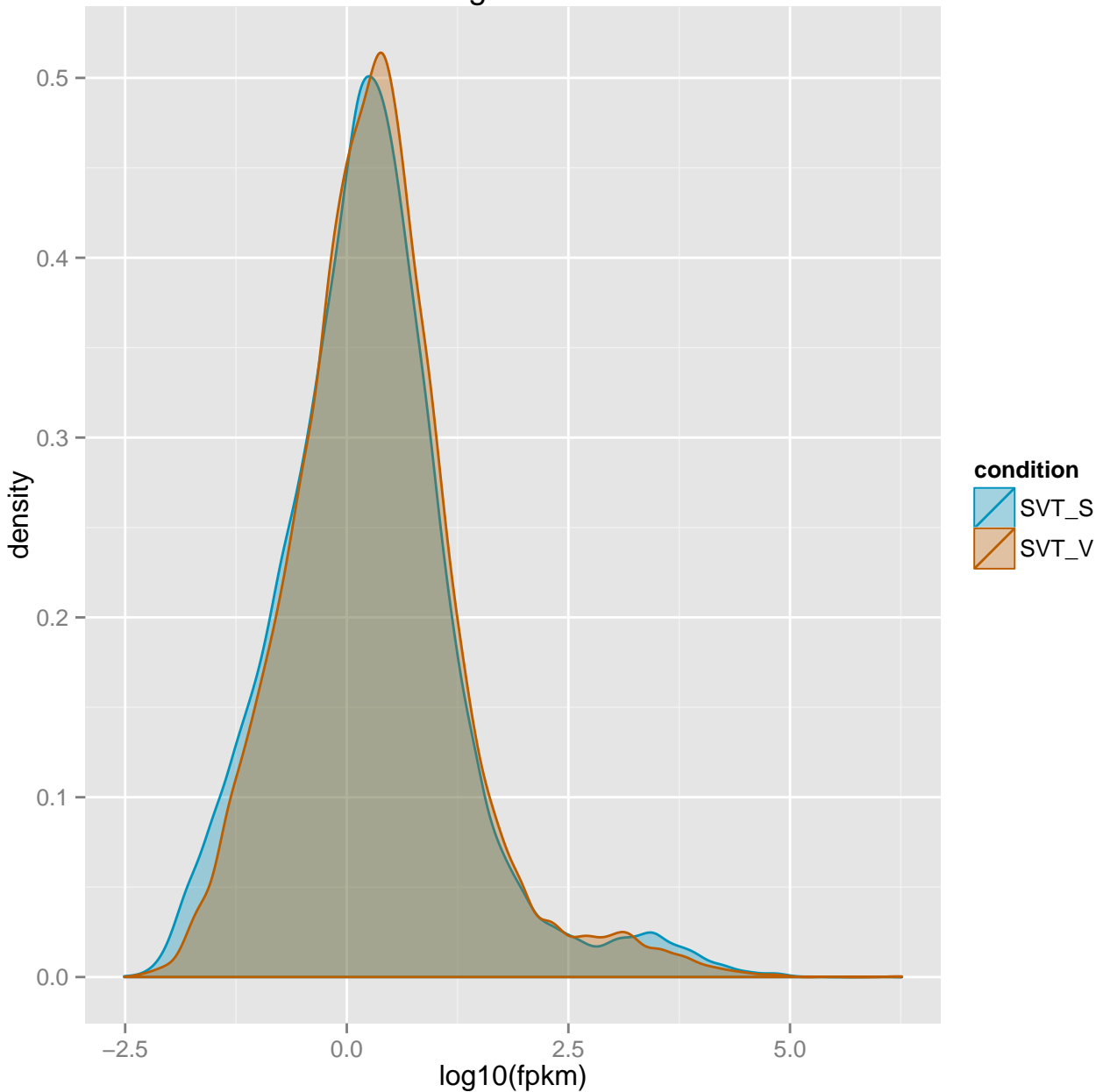
# genes



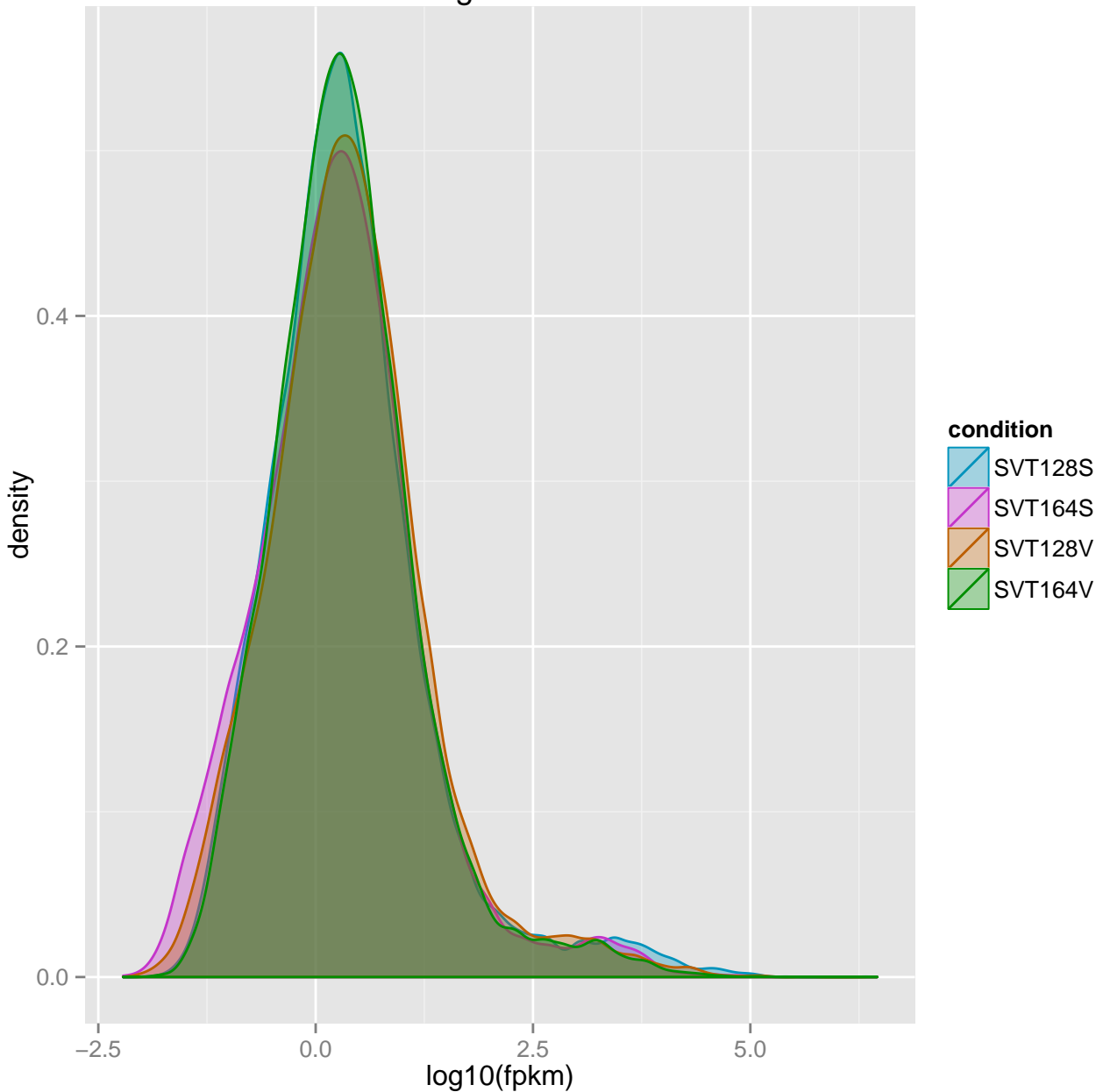
# isoforms

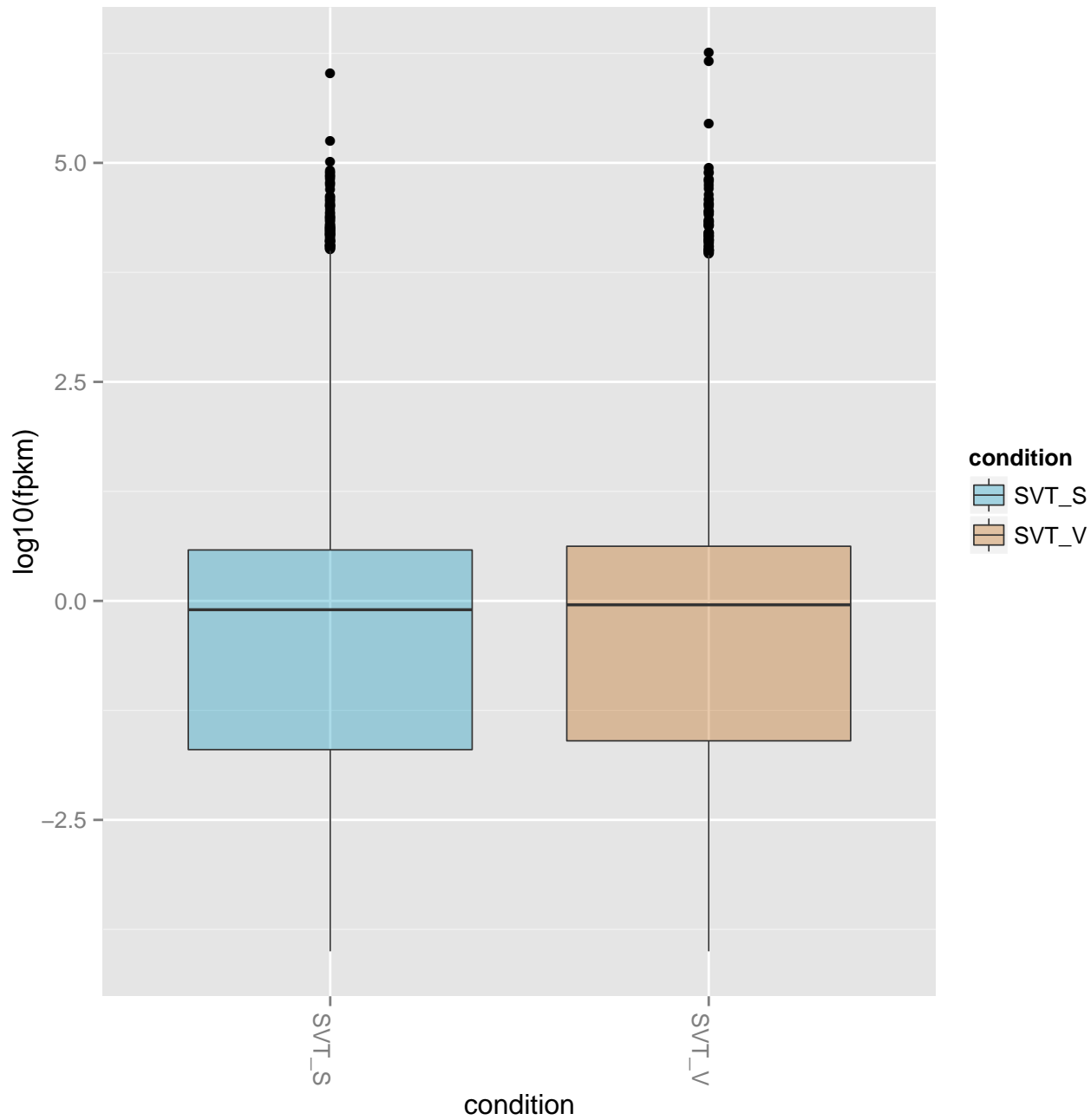


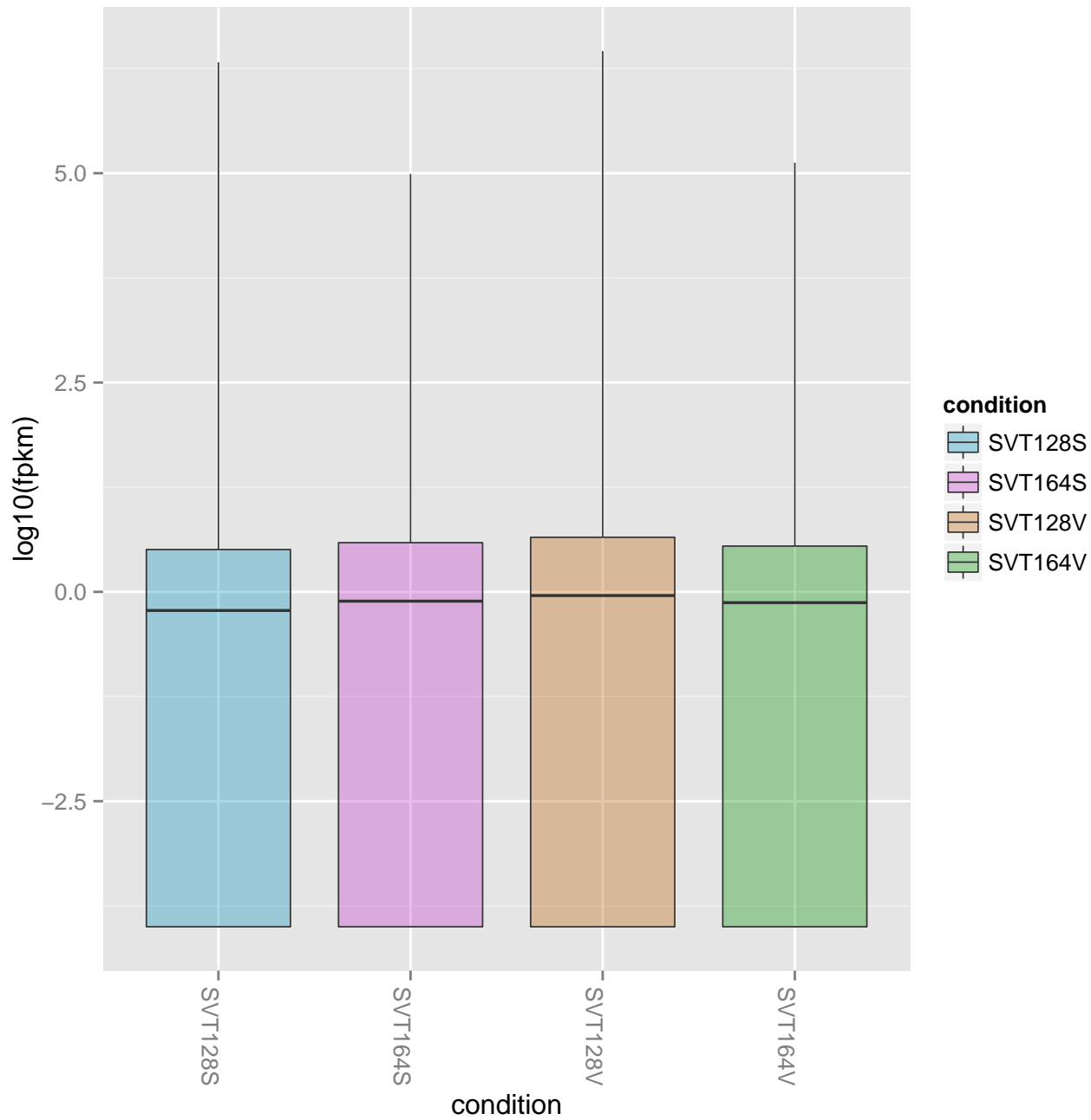
# genes

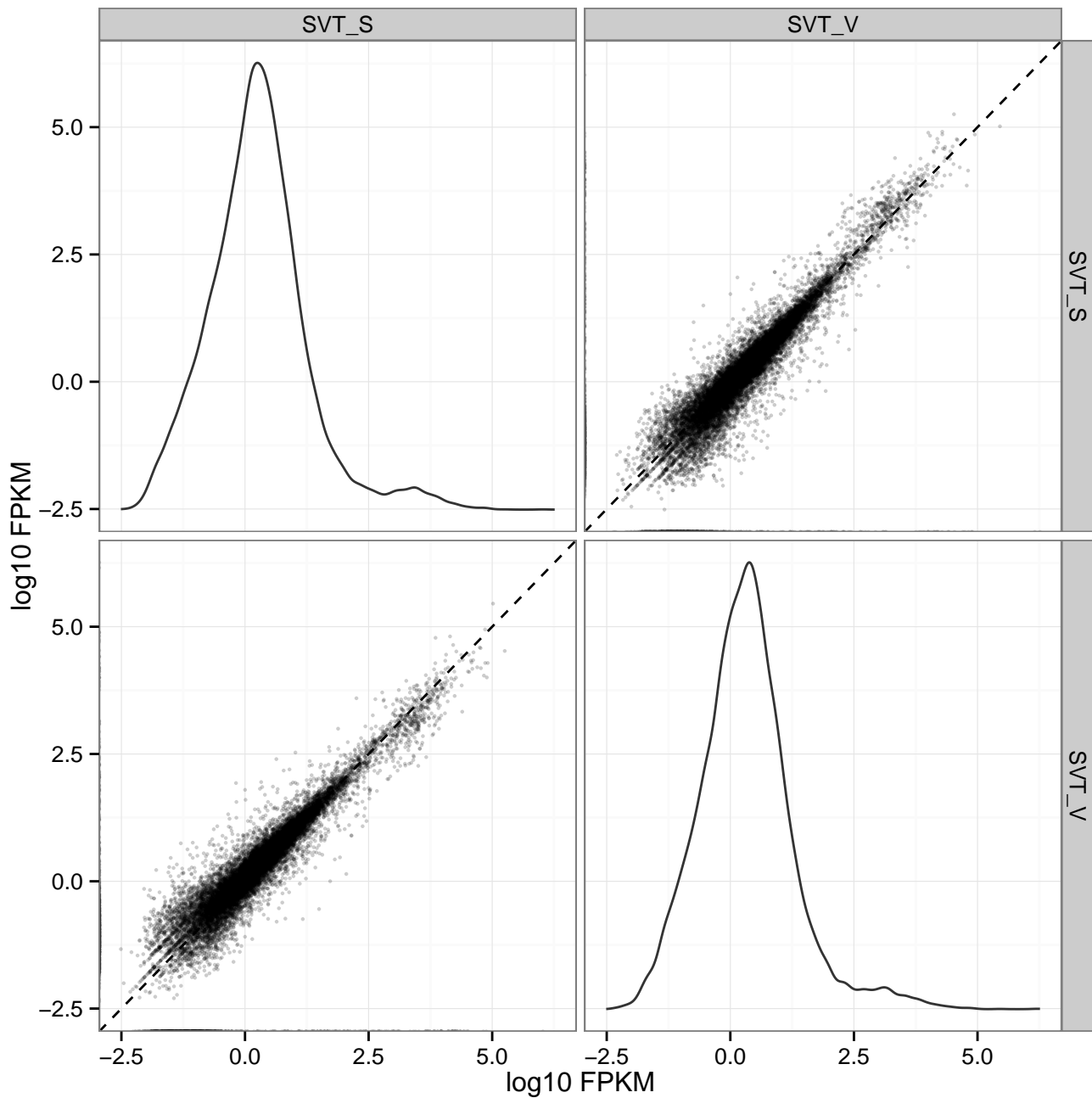


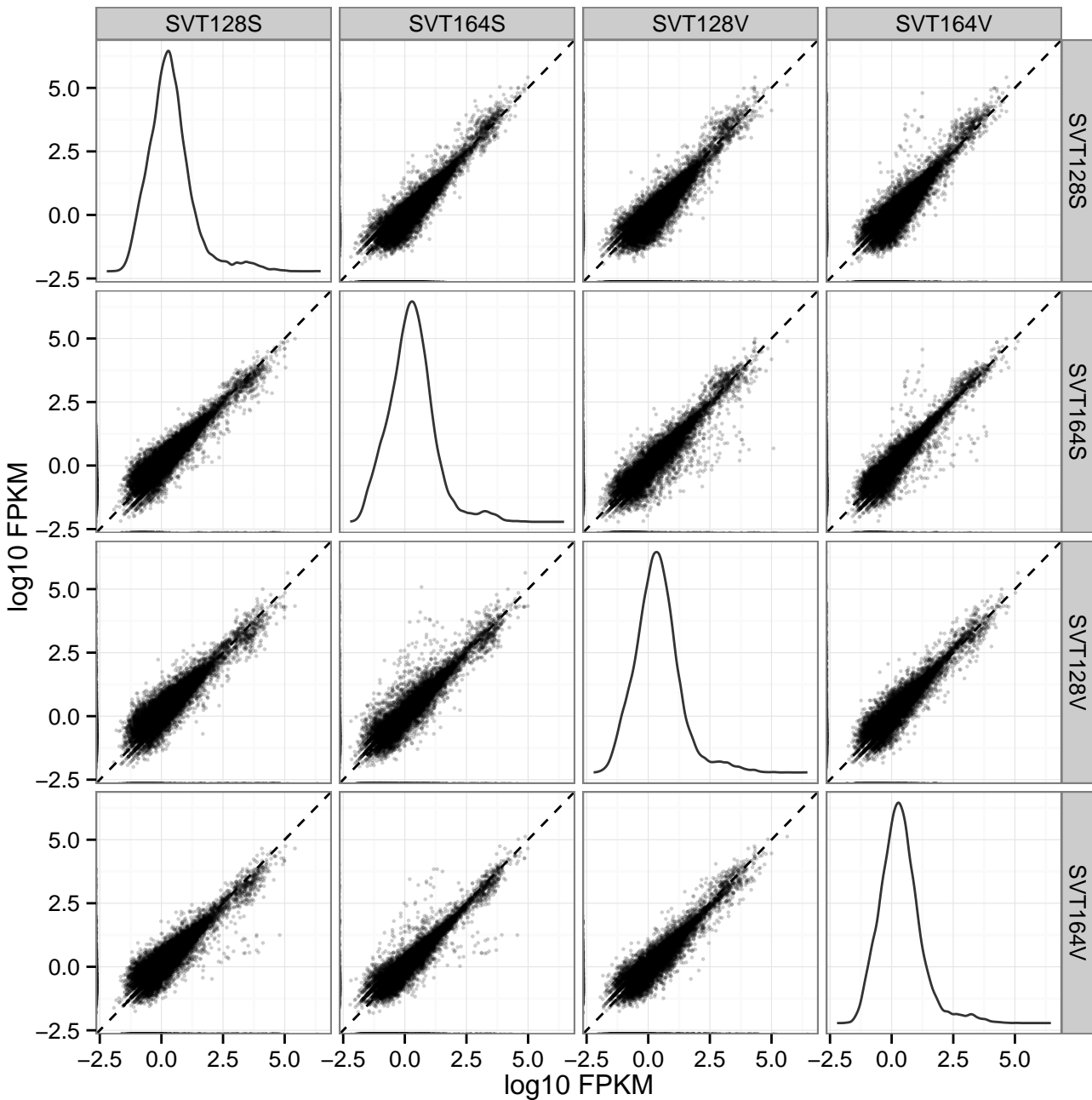
# genes



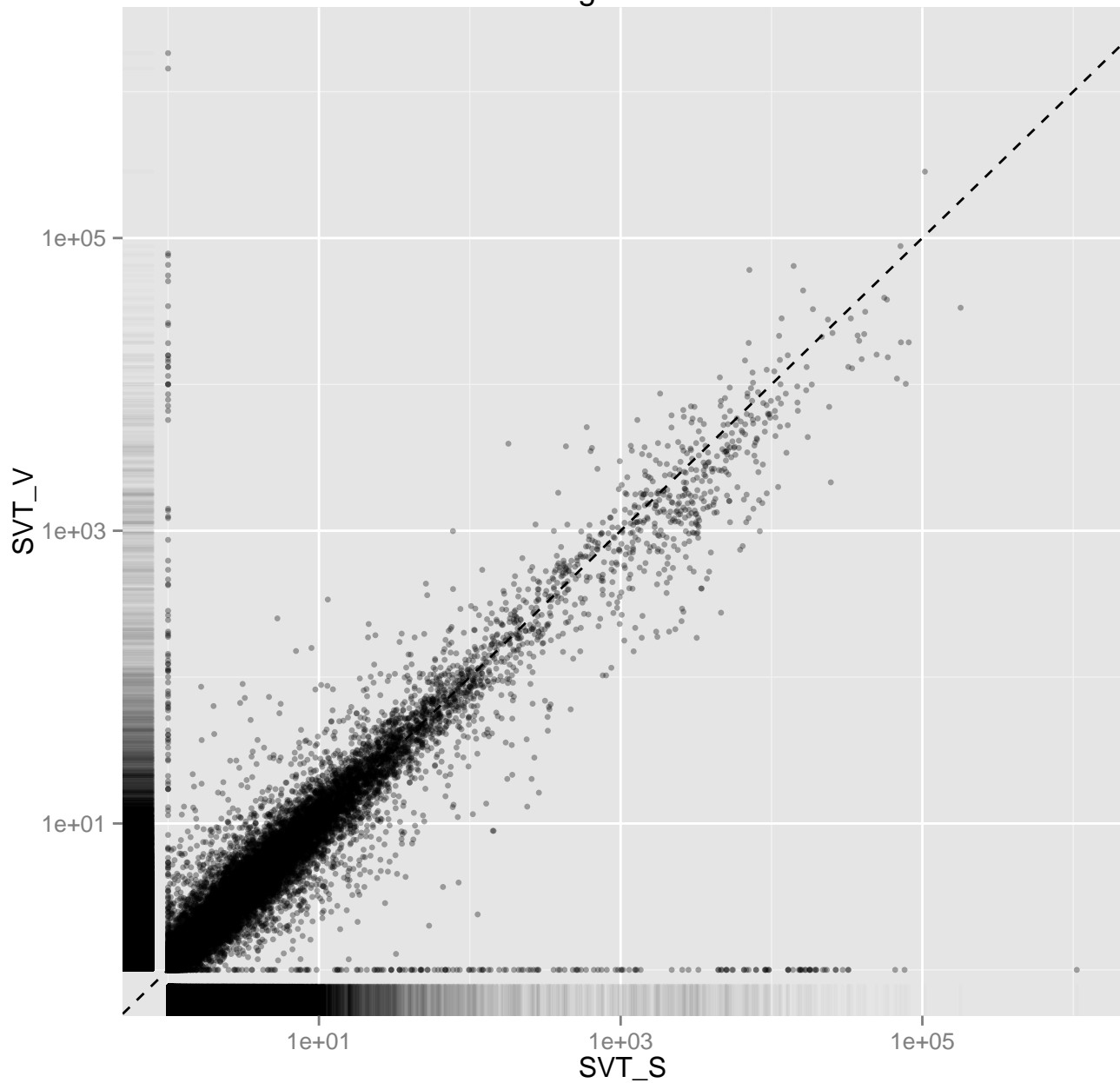


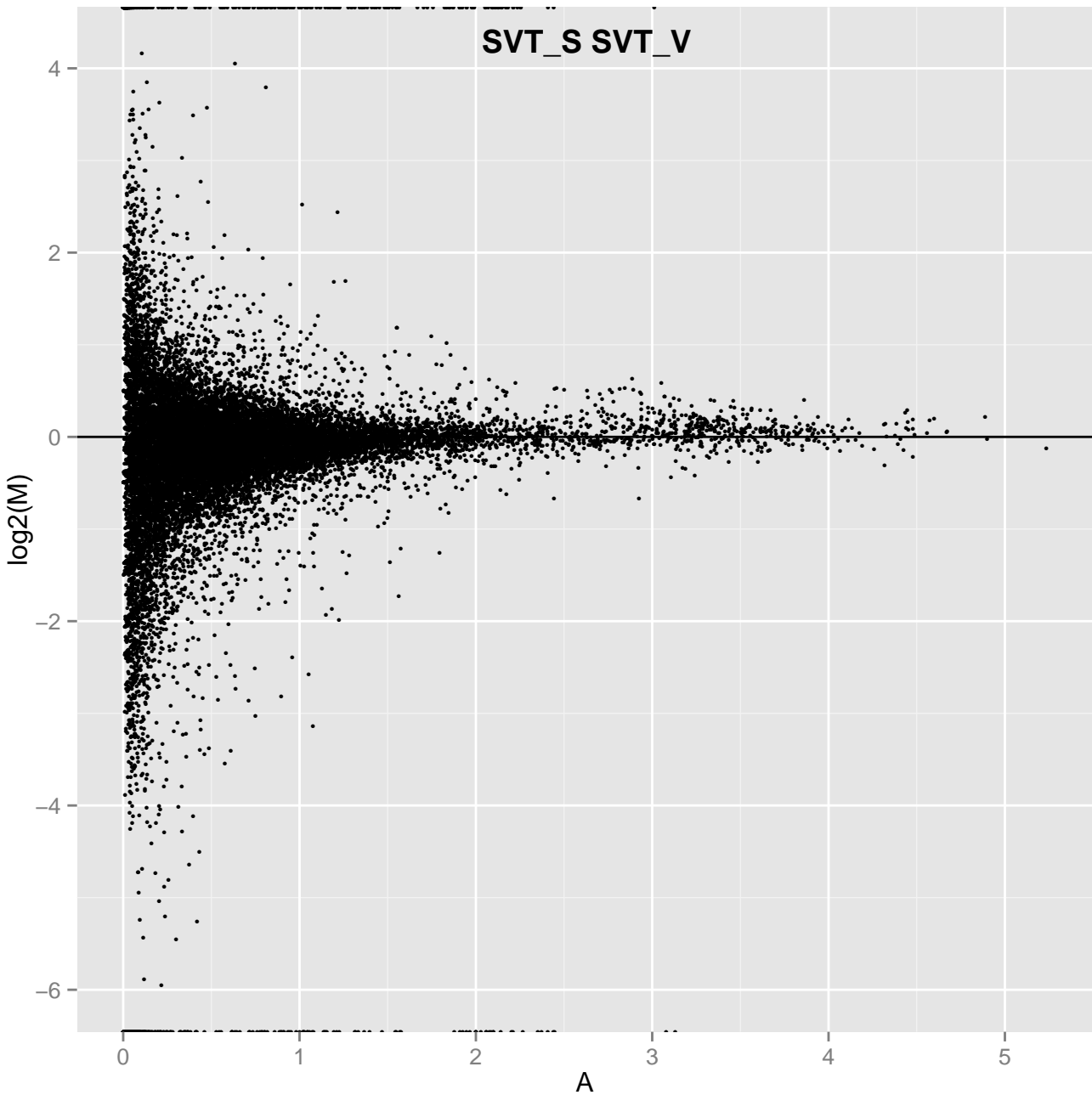






genes





normalized count values SVT\_S SVT\_V

log2(M)

0

2

4

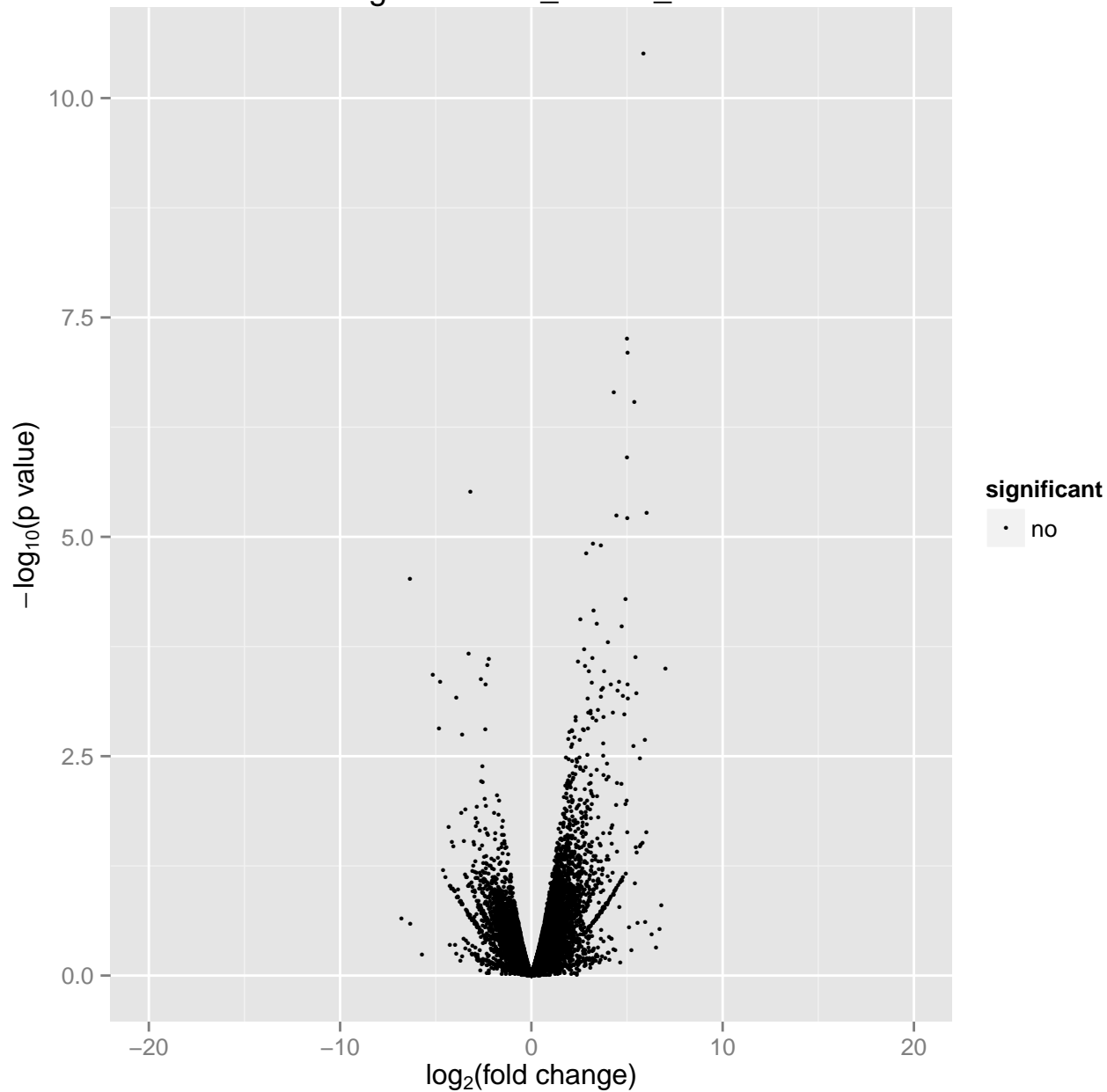
A

2

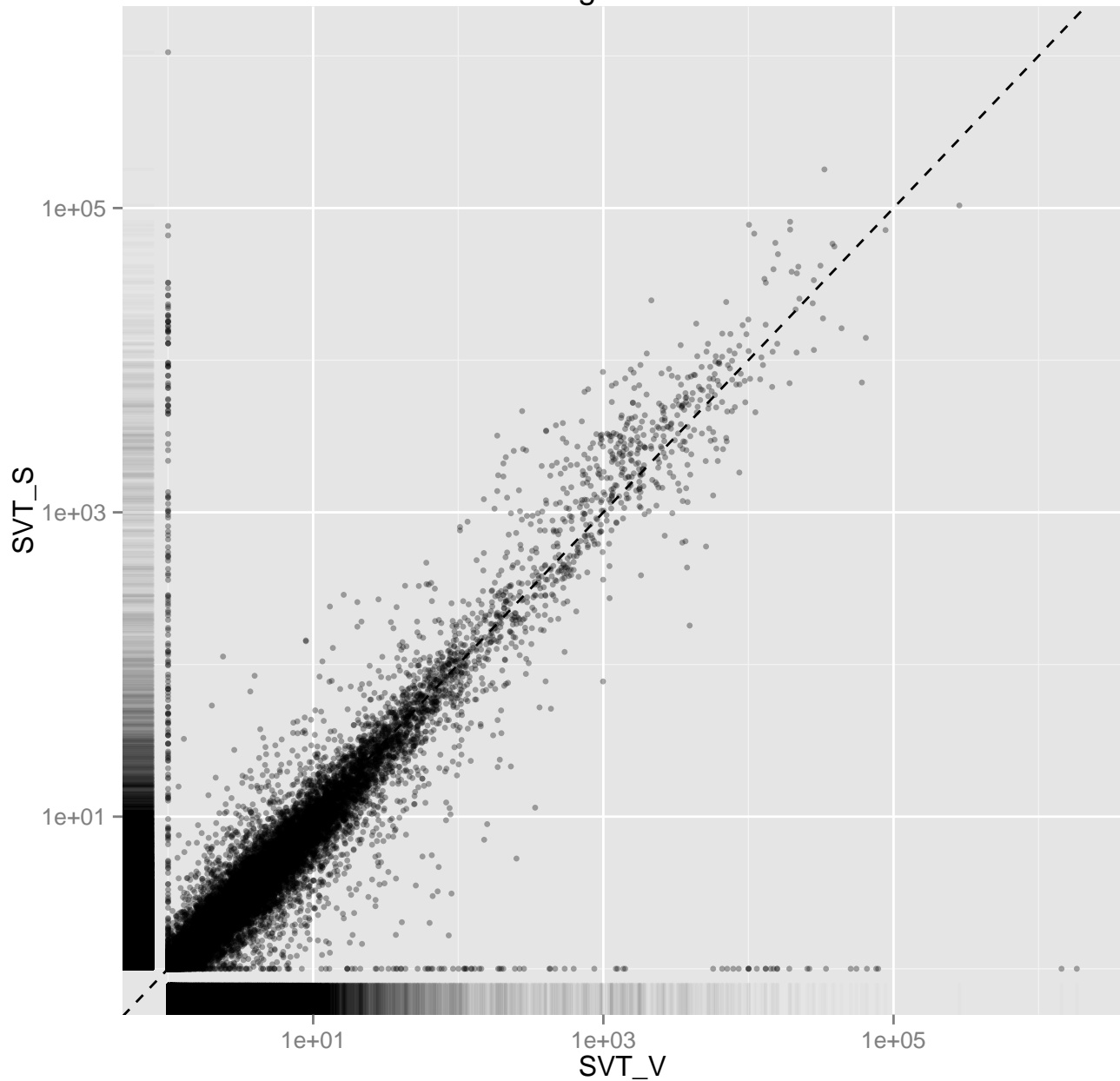
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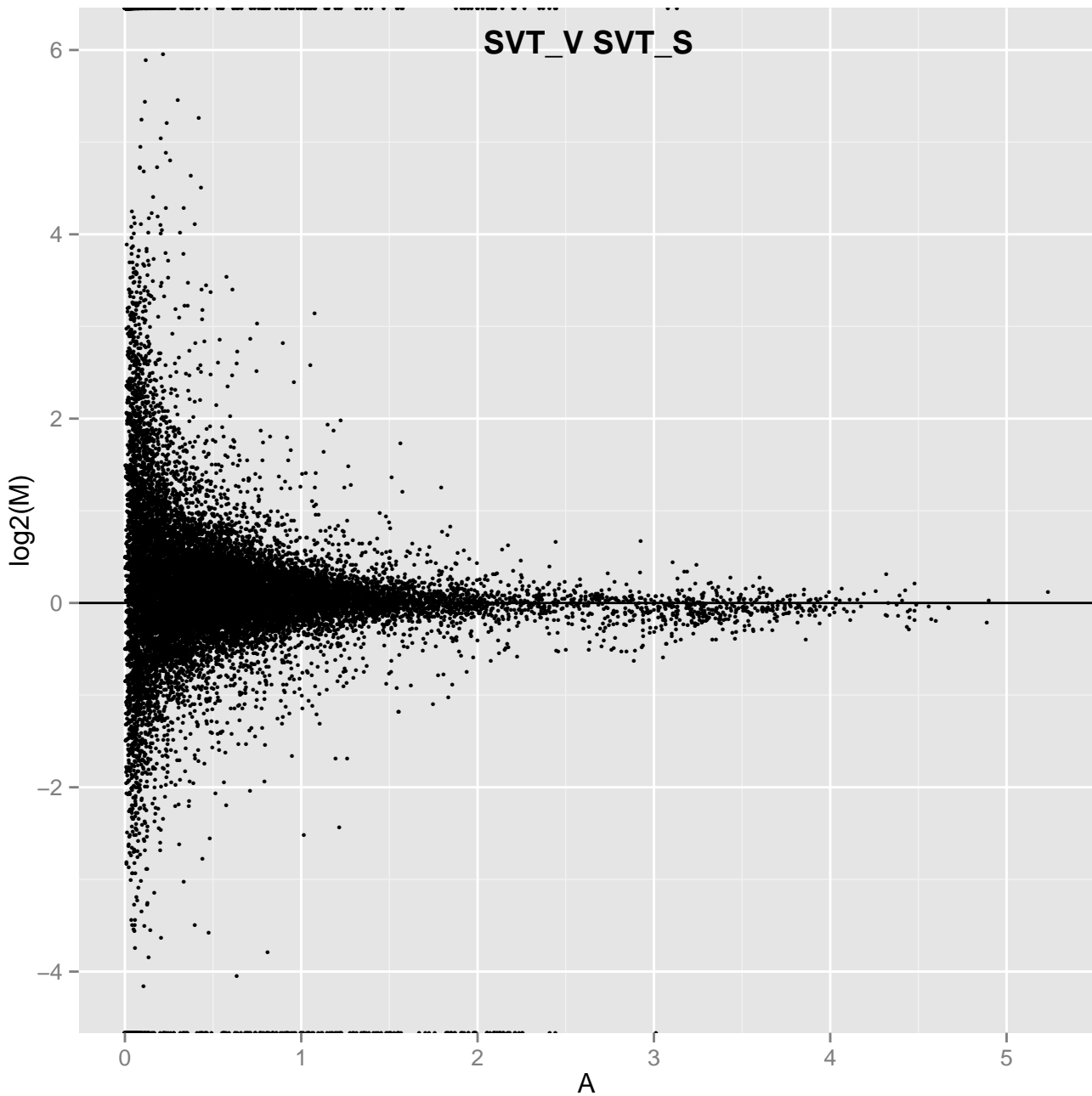
-2

genes: SVT\_V/SVT\_S



genes





normalized count values SVT\_V SVT\_S

$\log_2(M)$

2

0

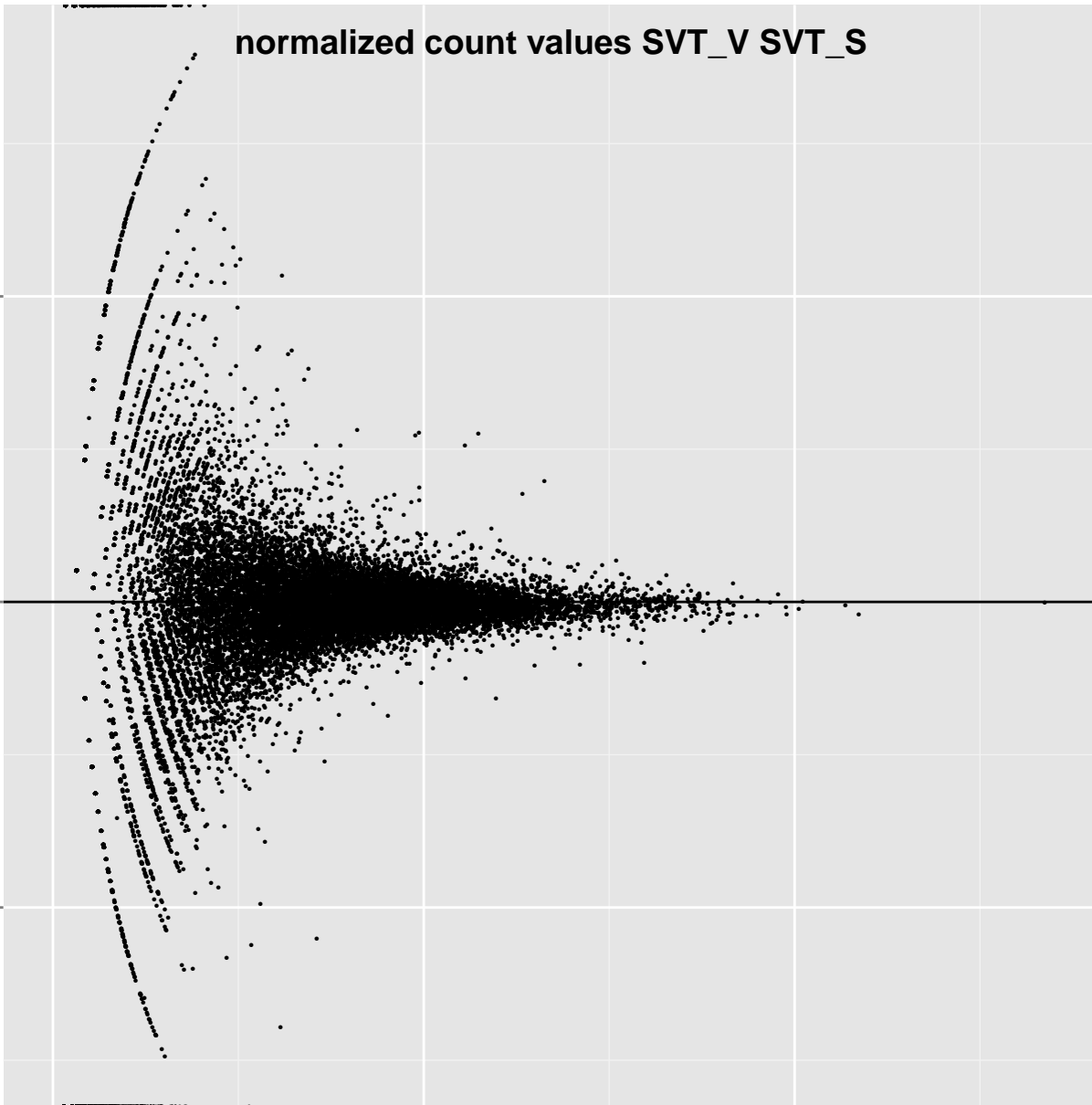
-2

0

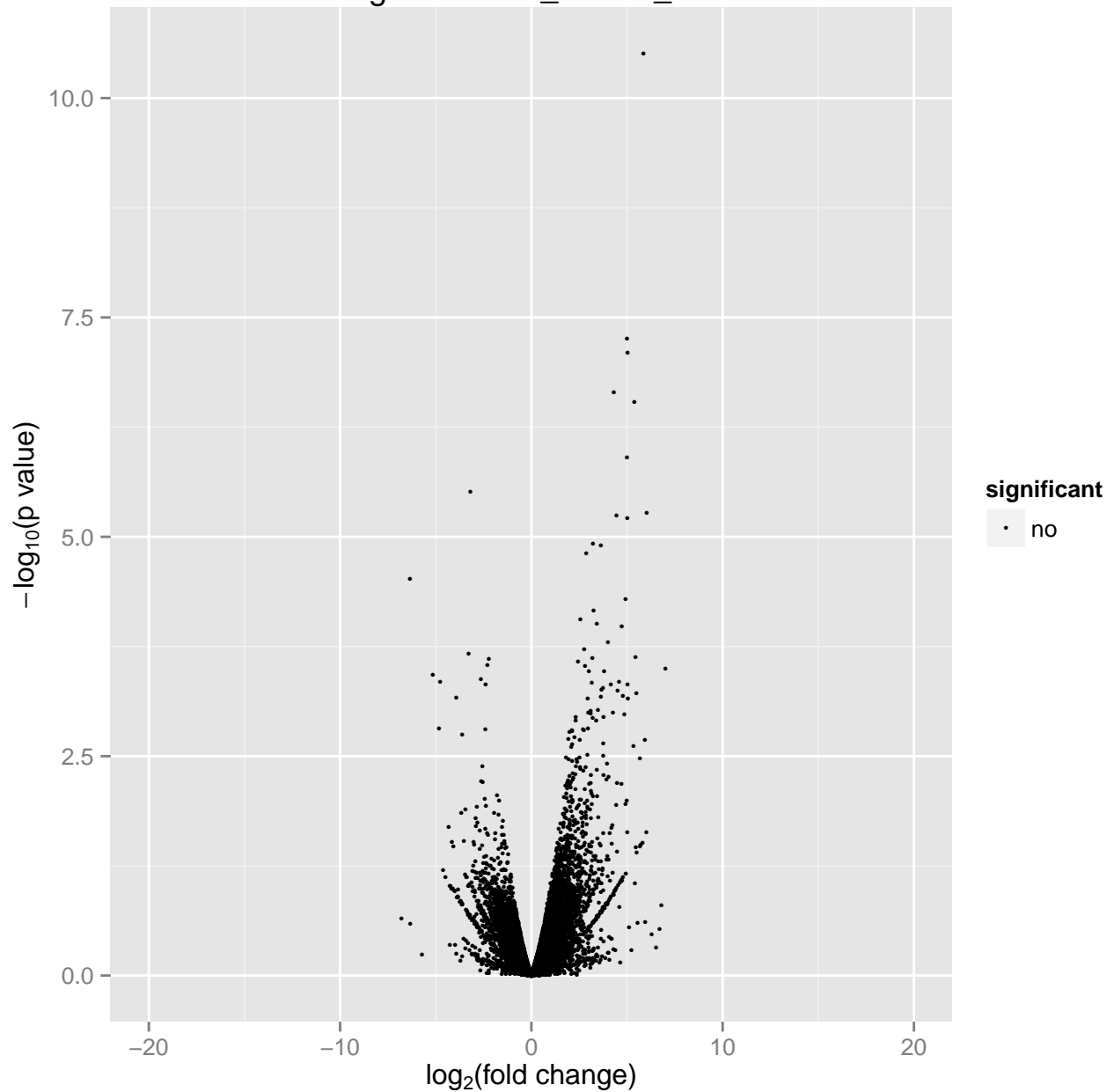
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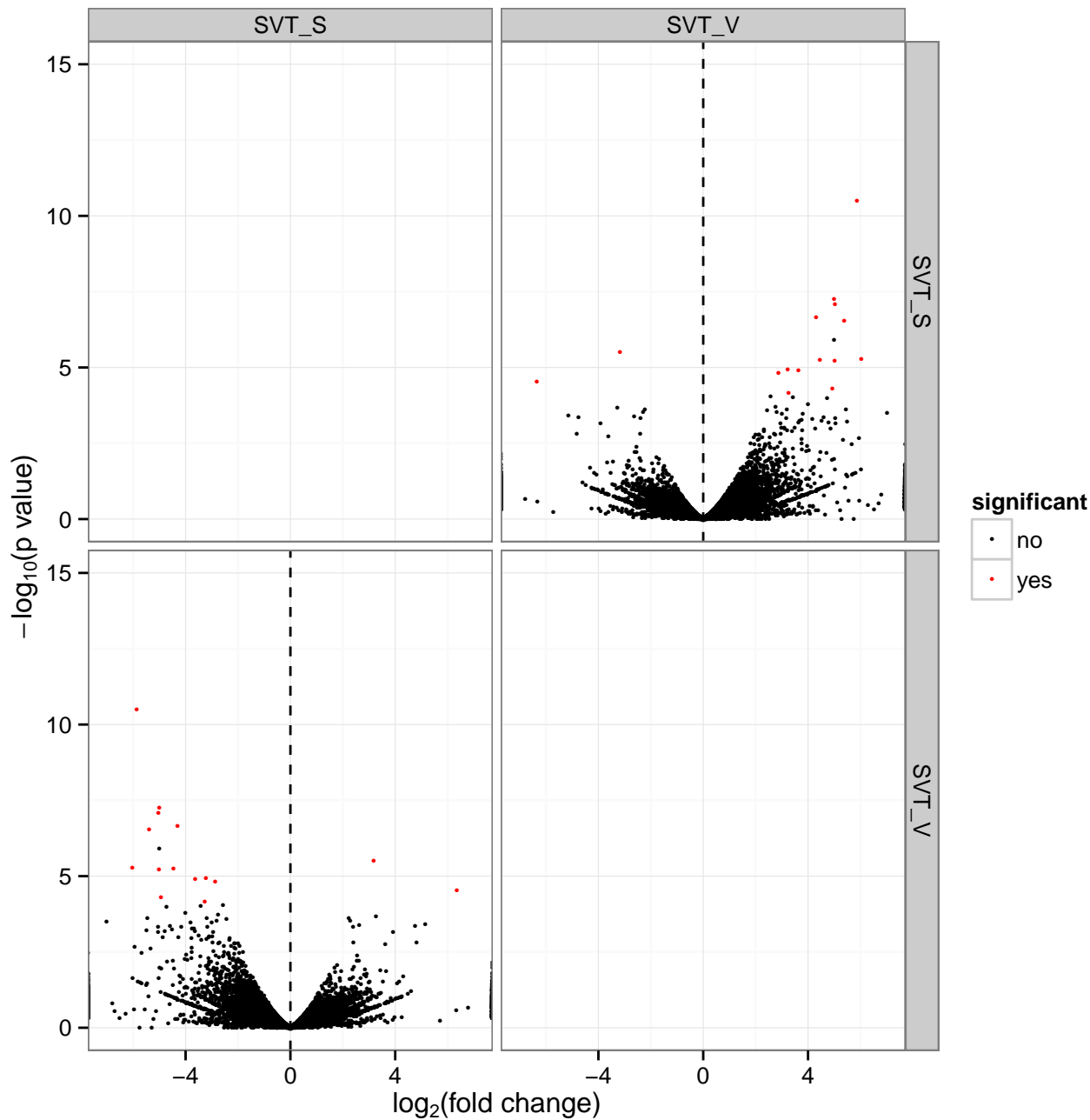
4

A

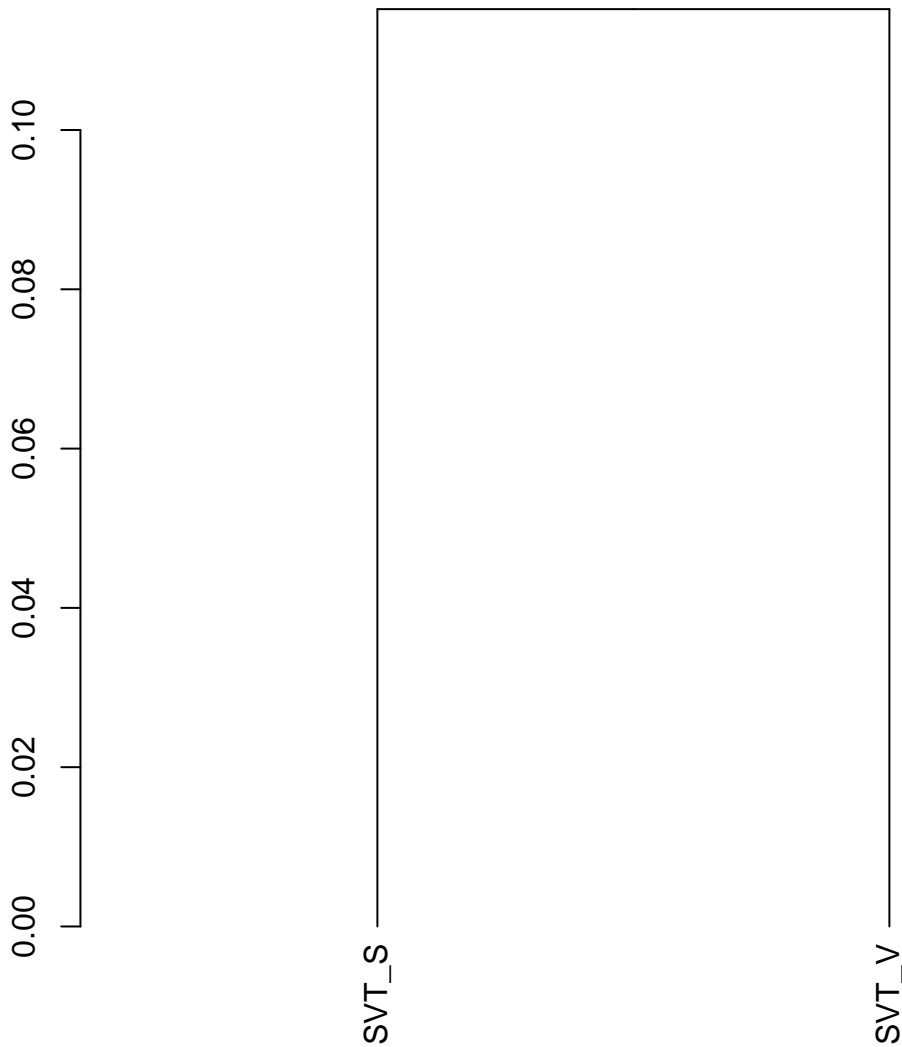


genes: SVT\_V/SVT\_S

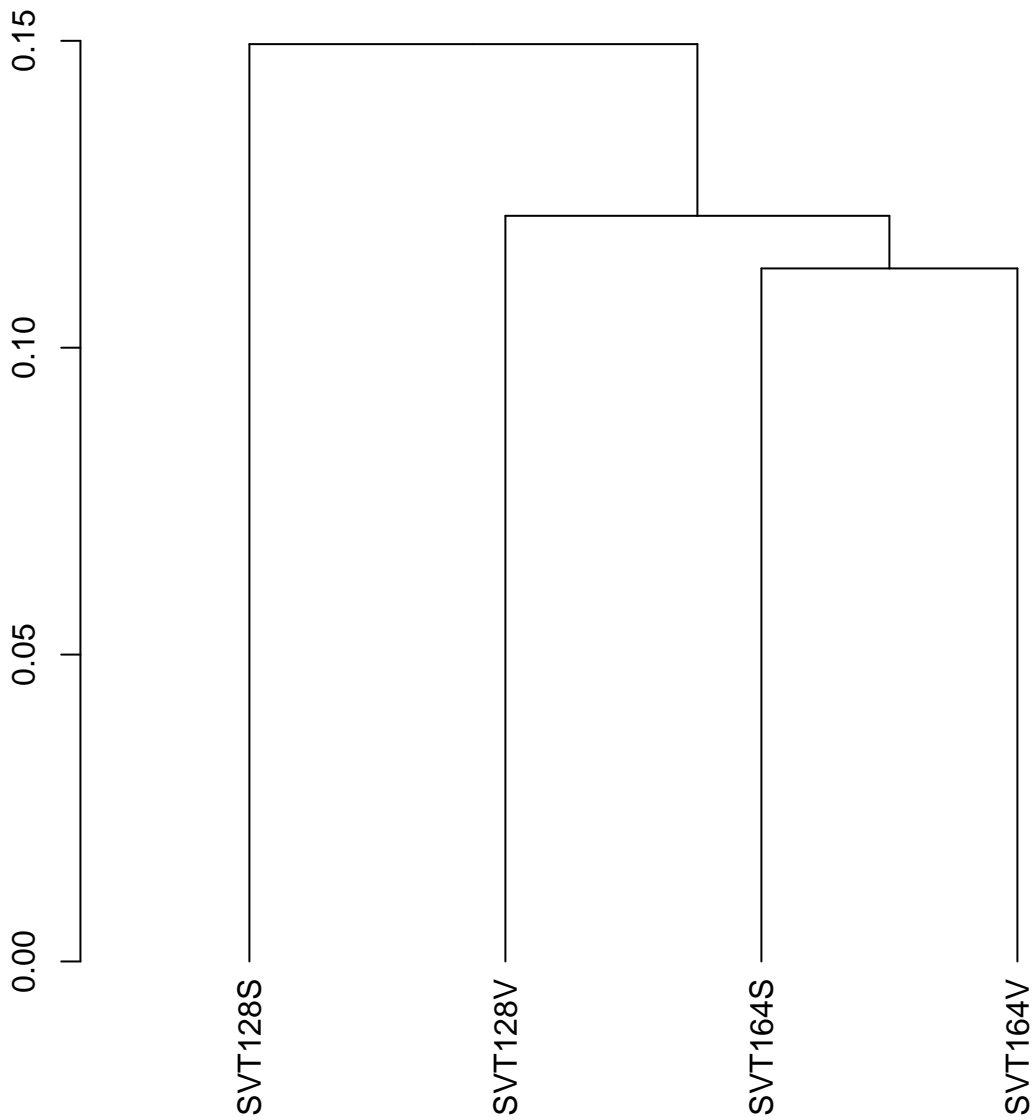


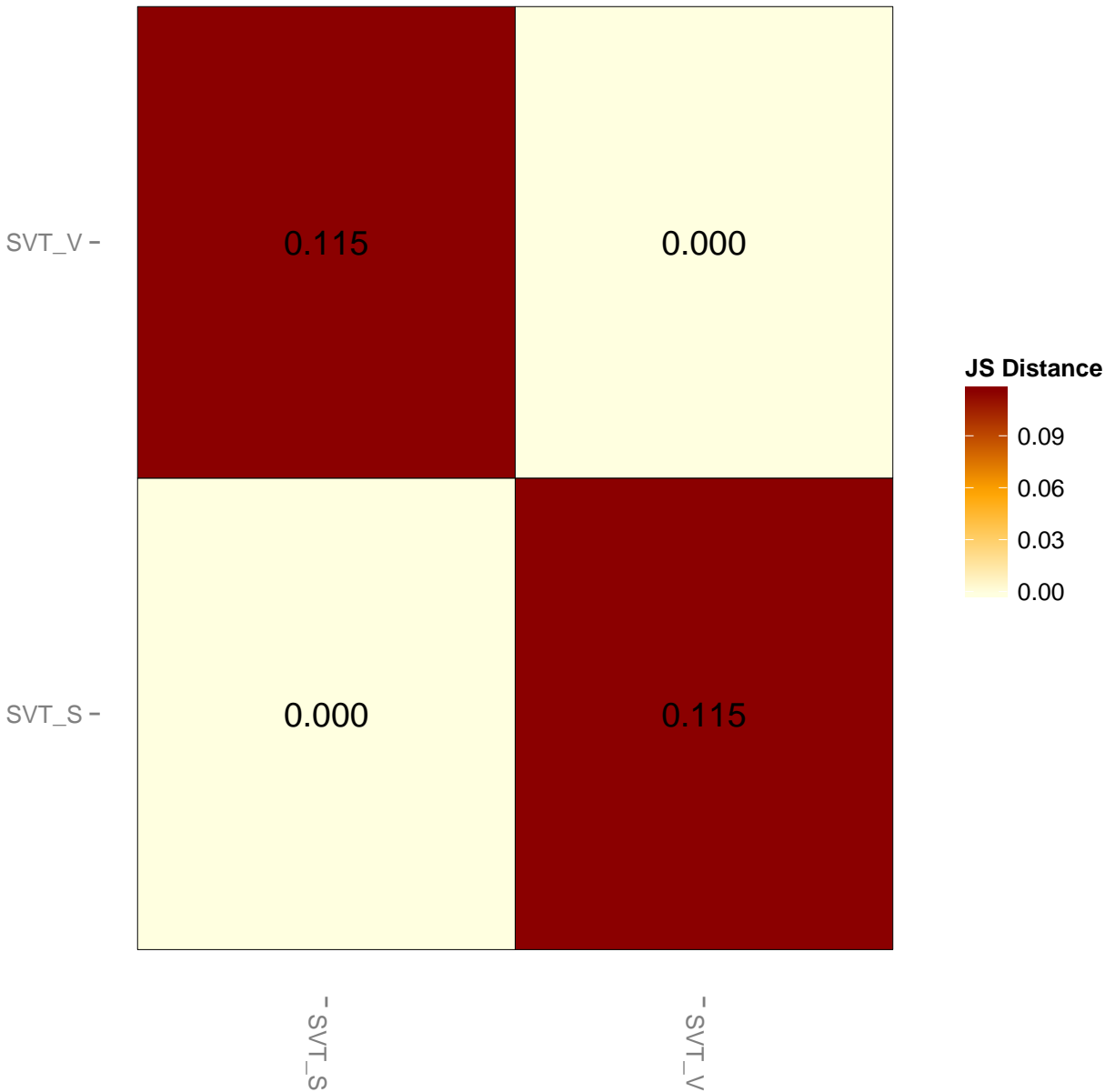


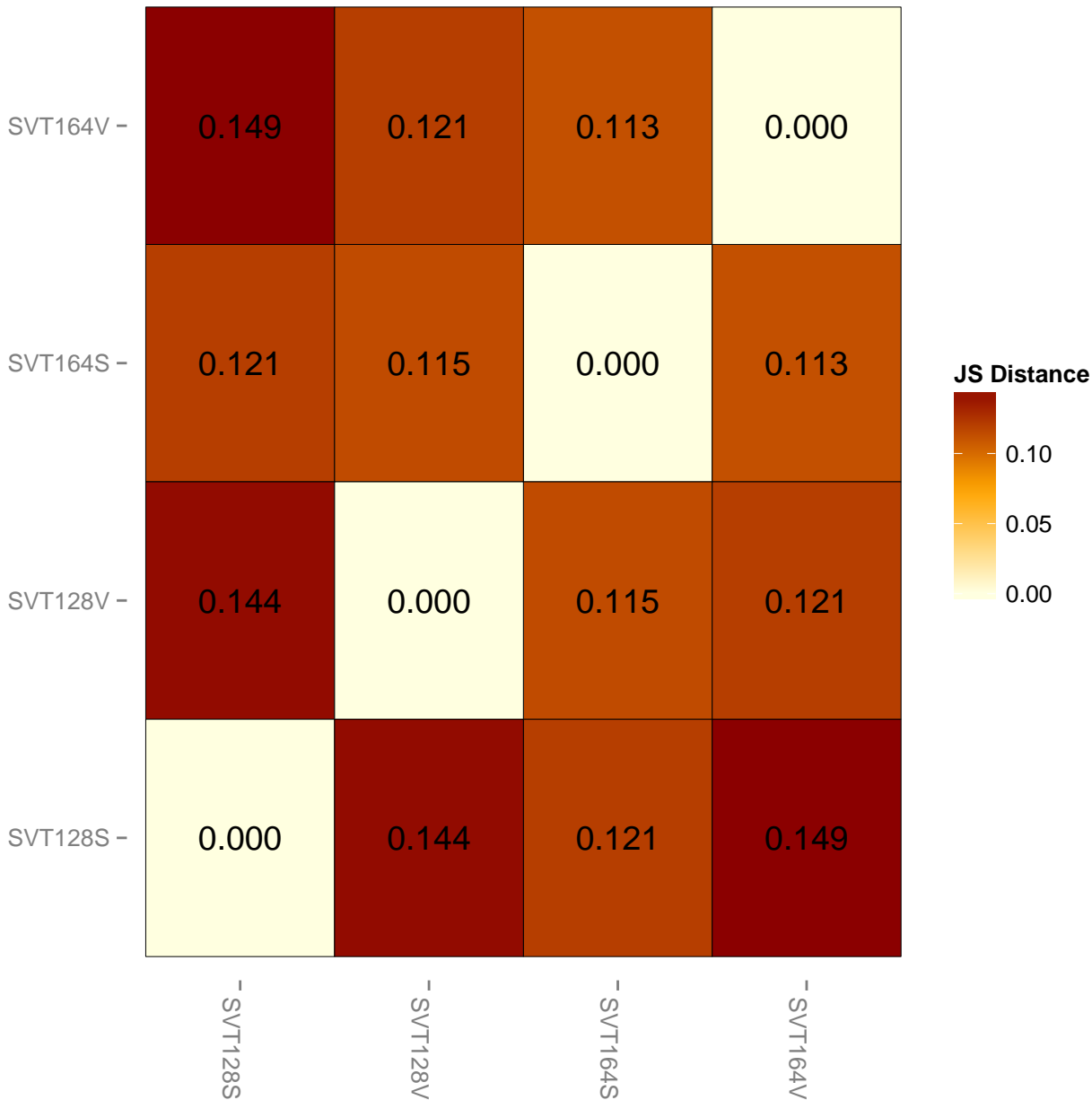
# All genes(cuff)

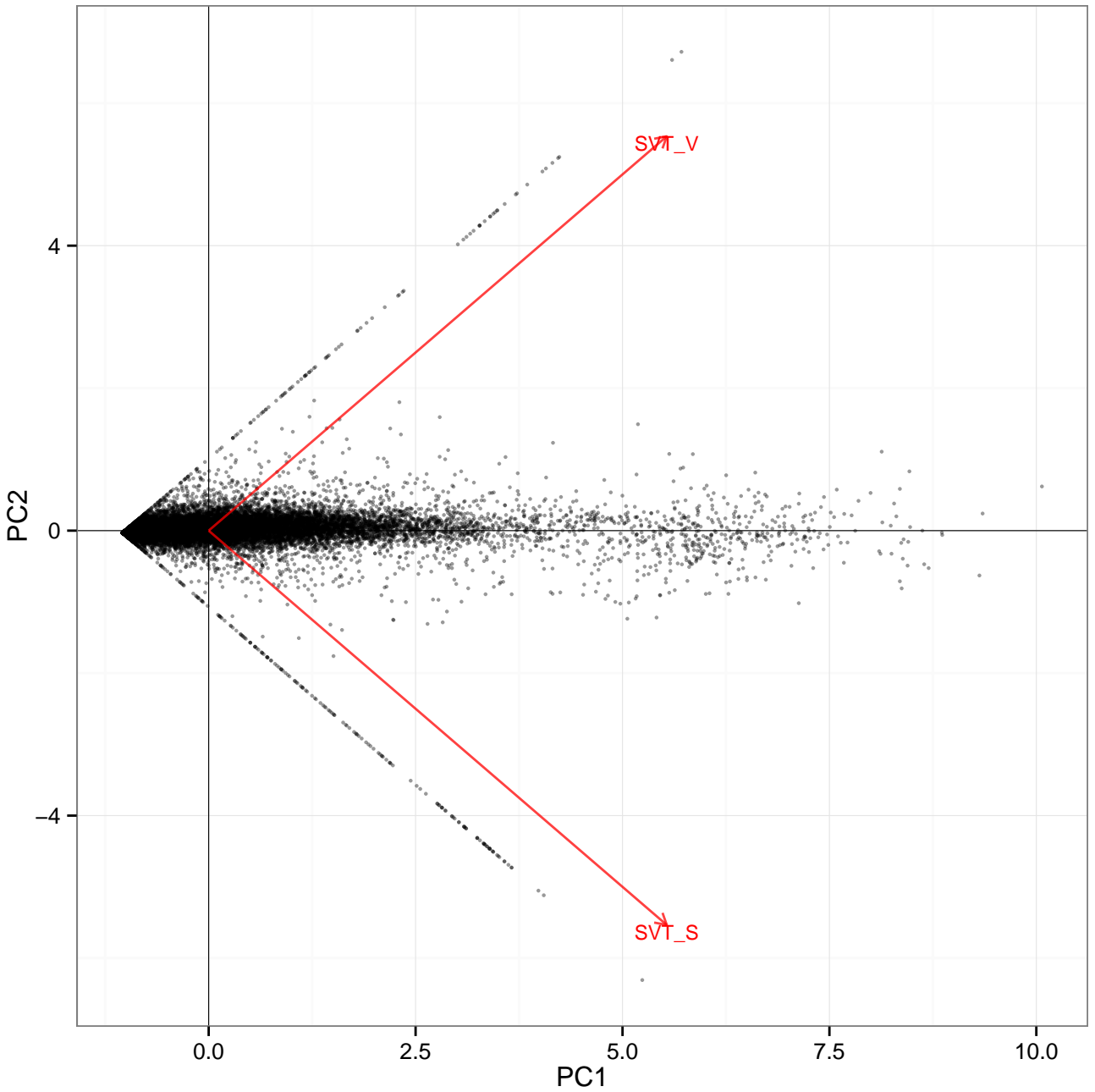


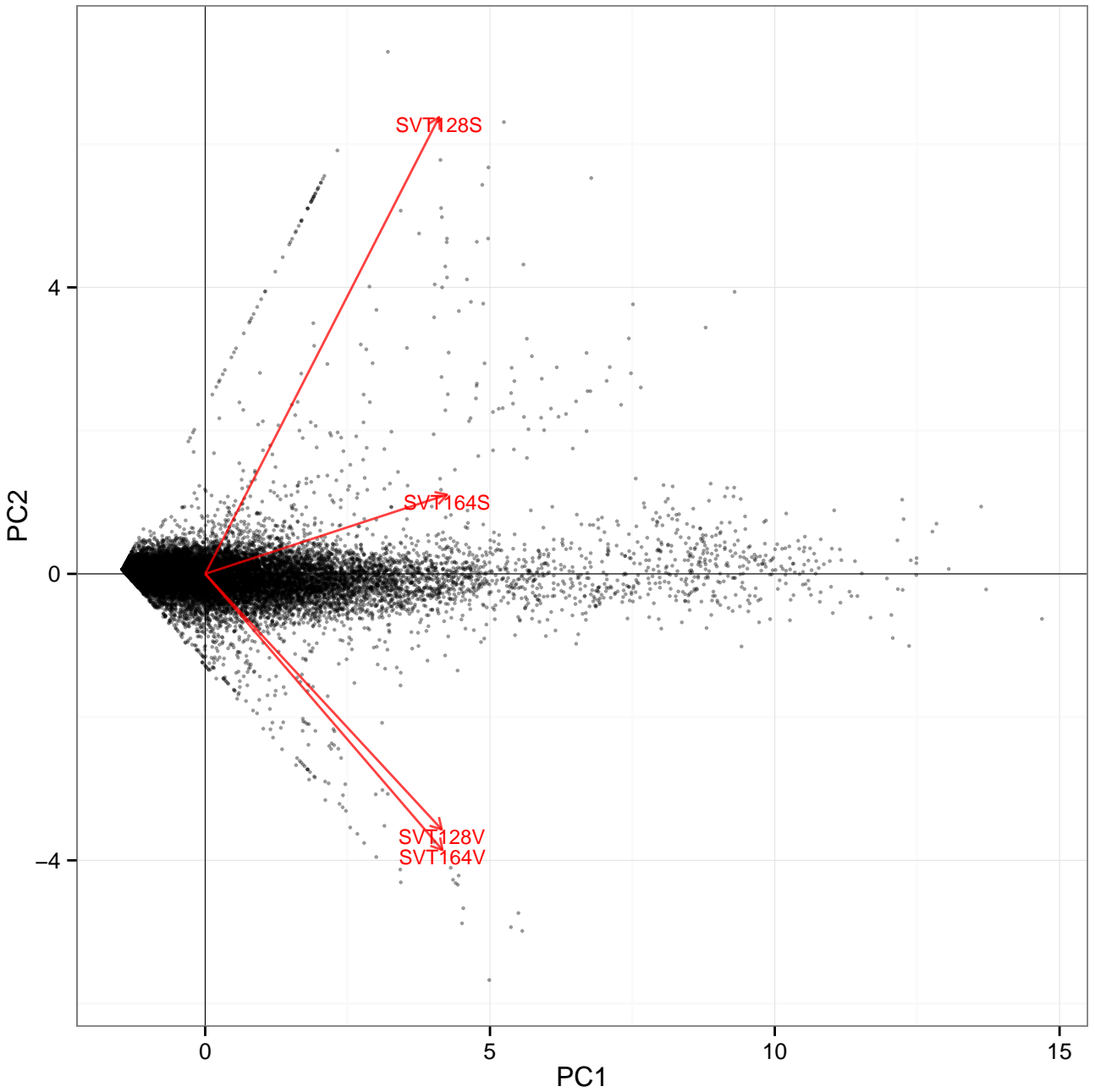
# All genes(cuff)

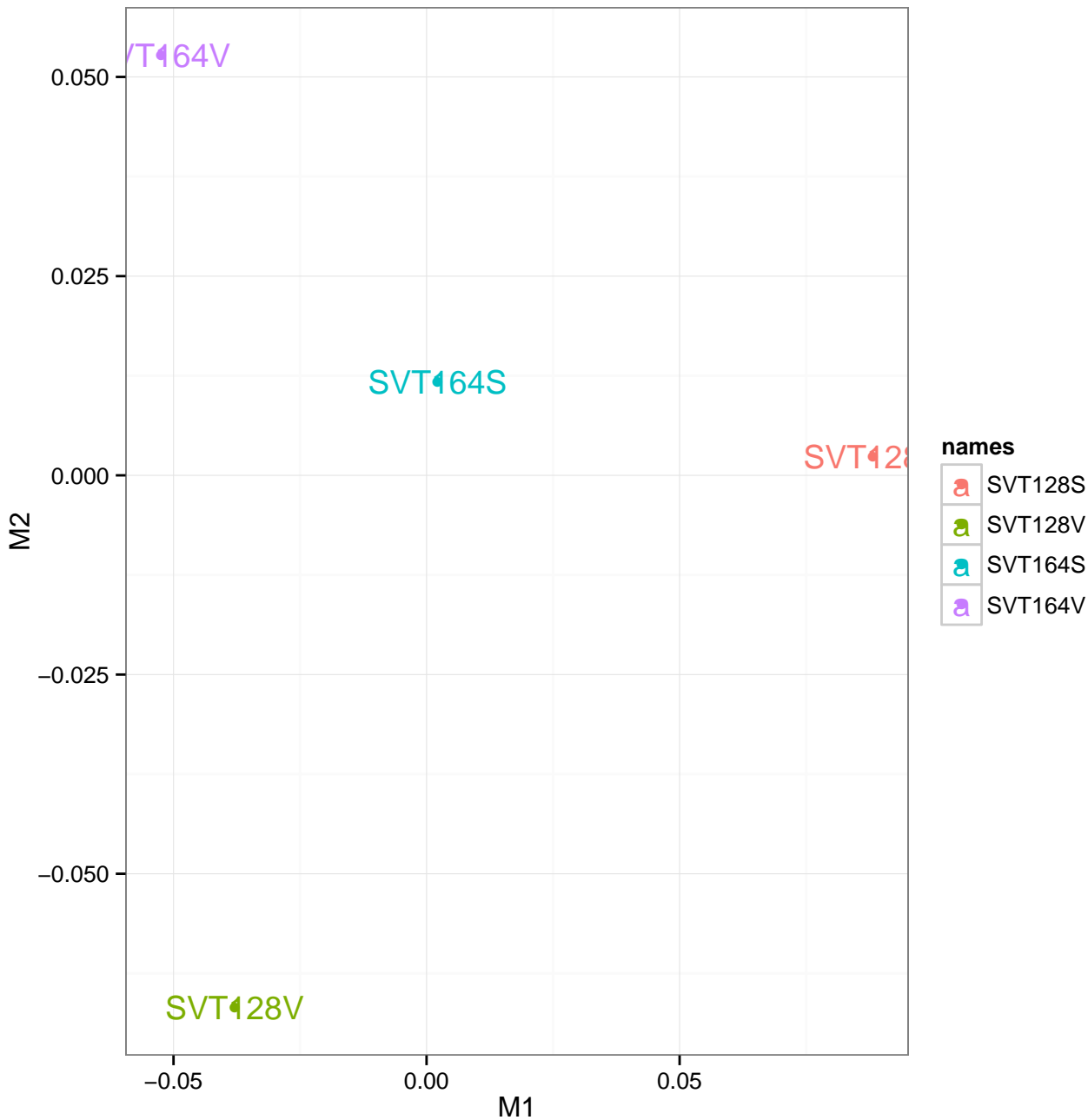




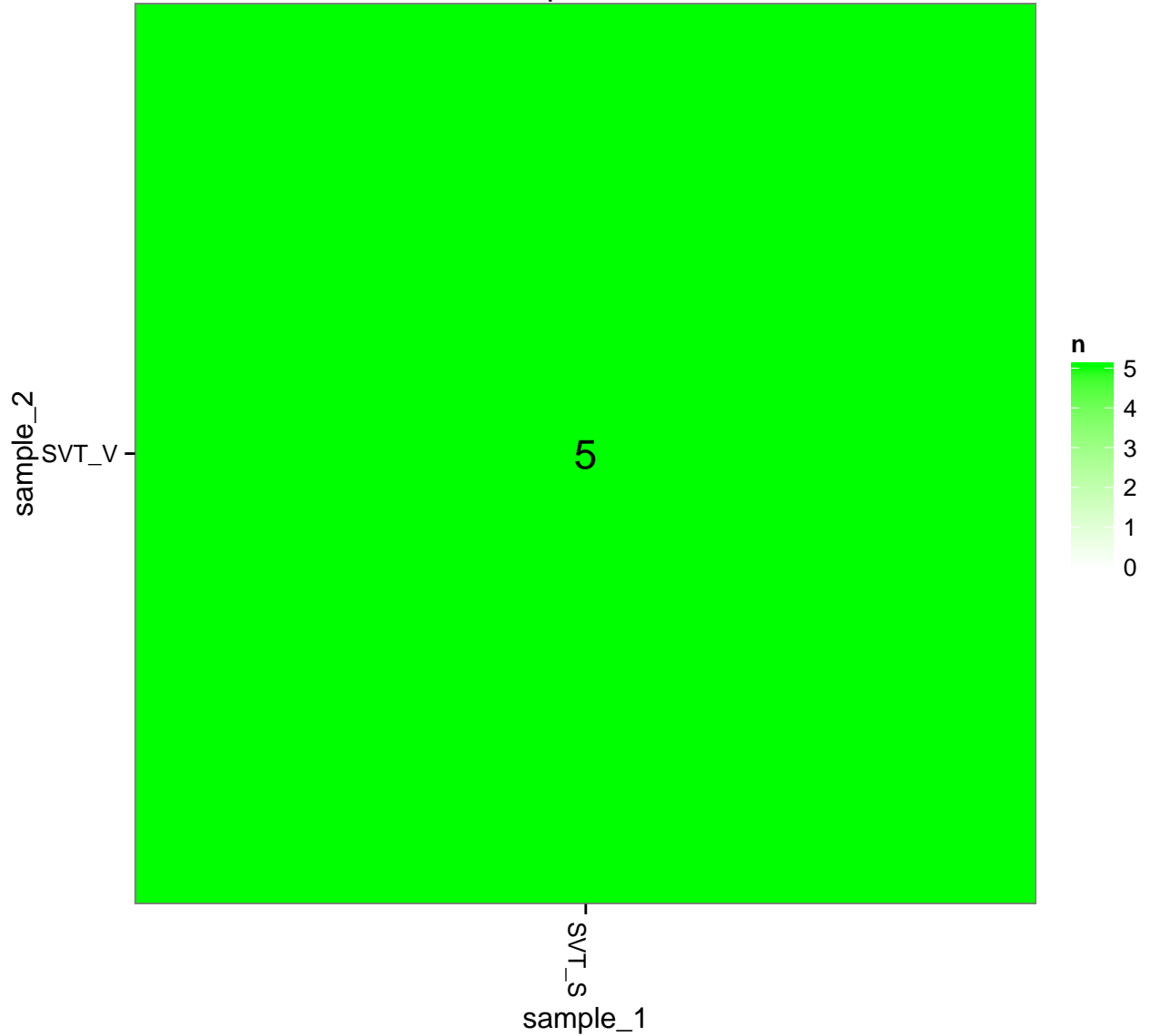




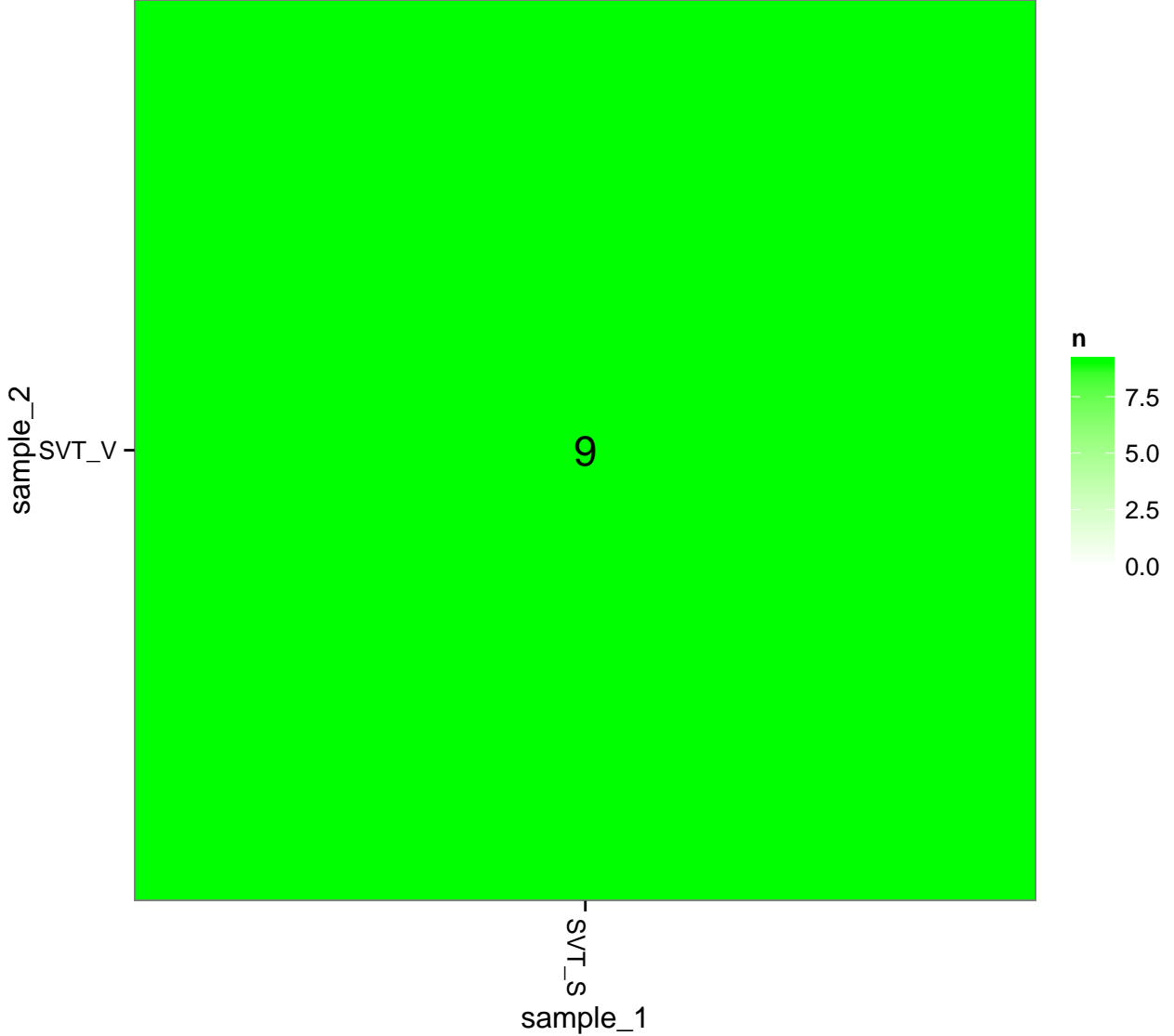




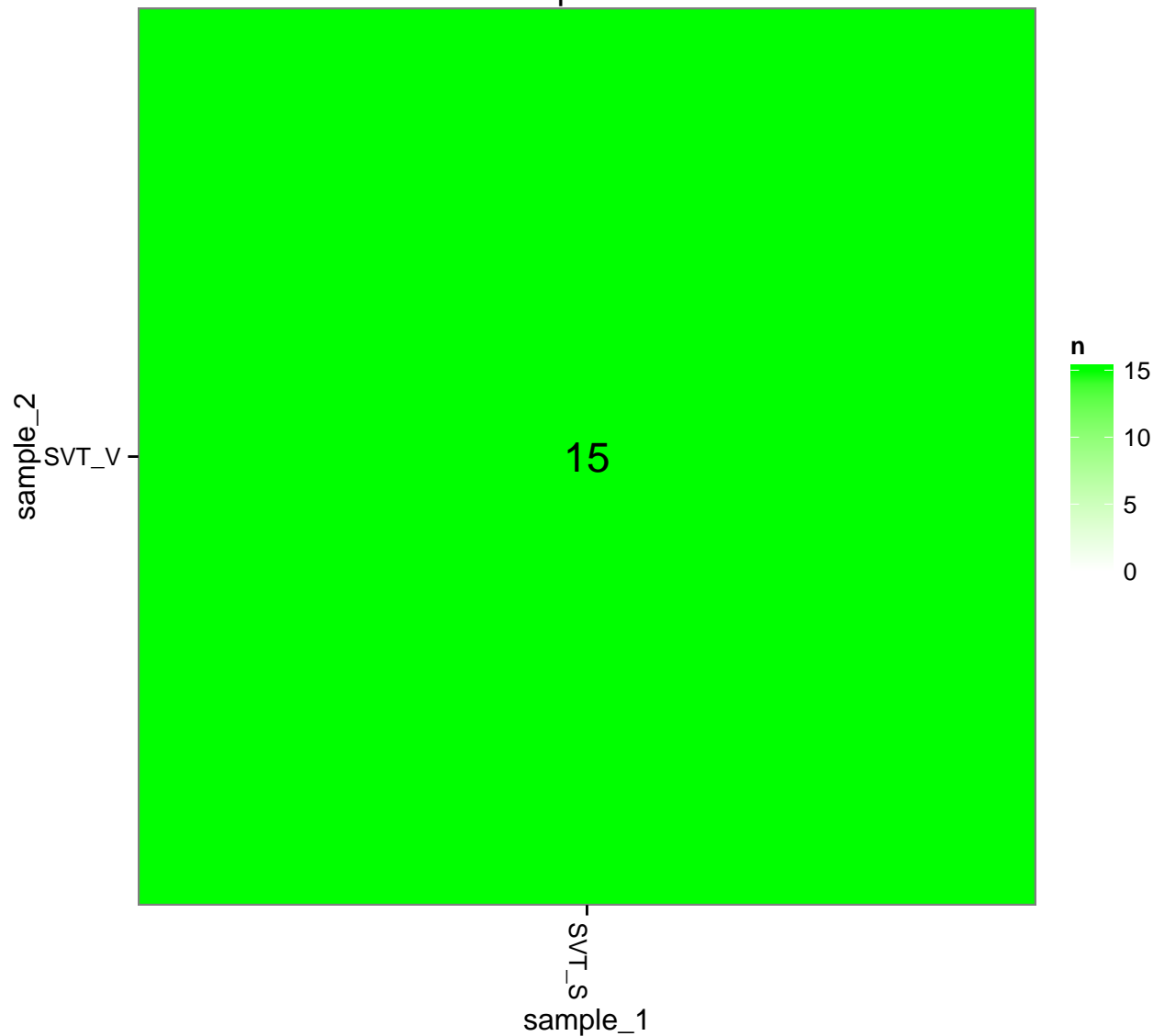
Significant genes  
at alpha 0.001

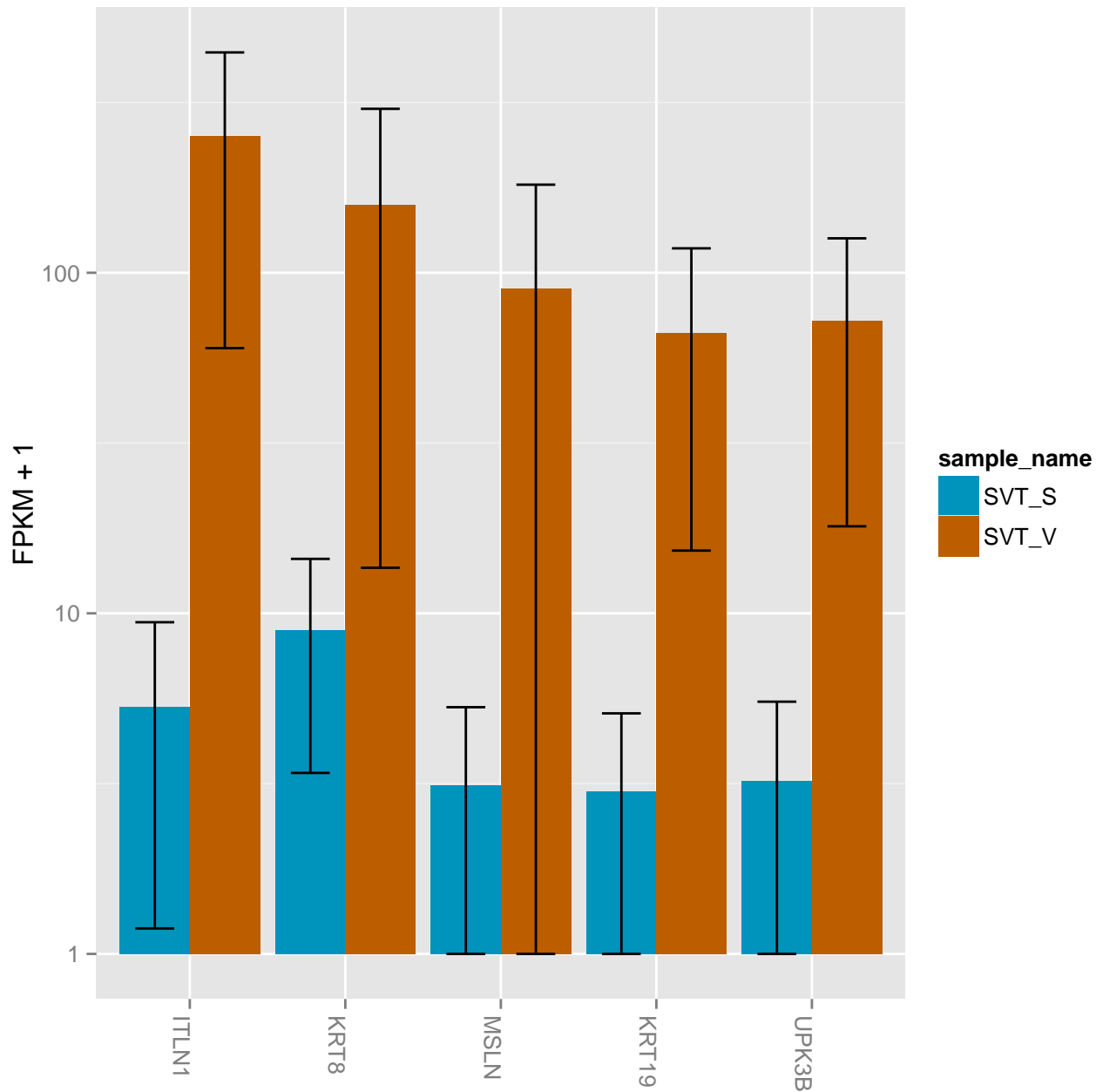


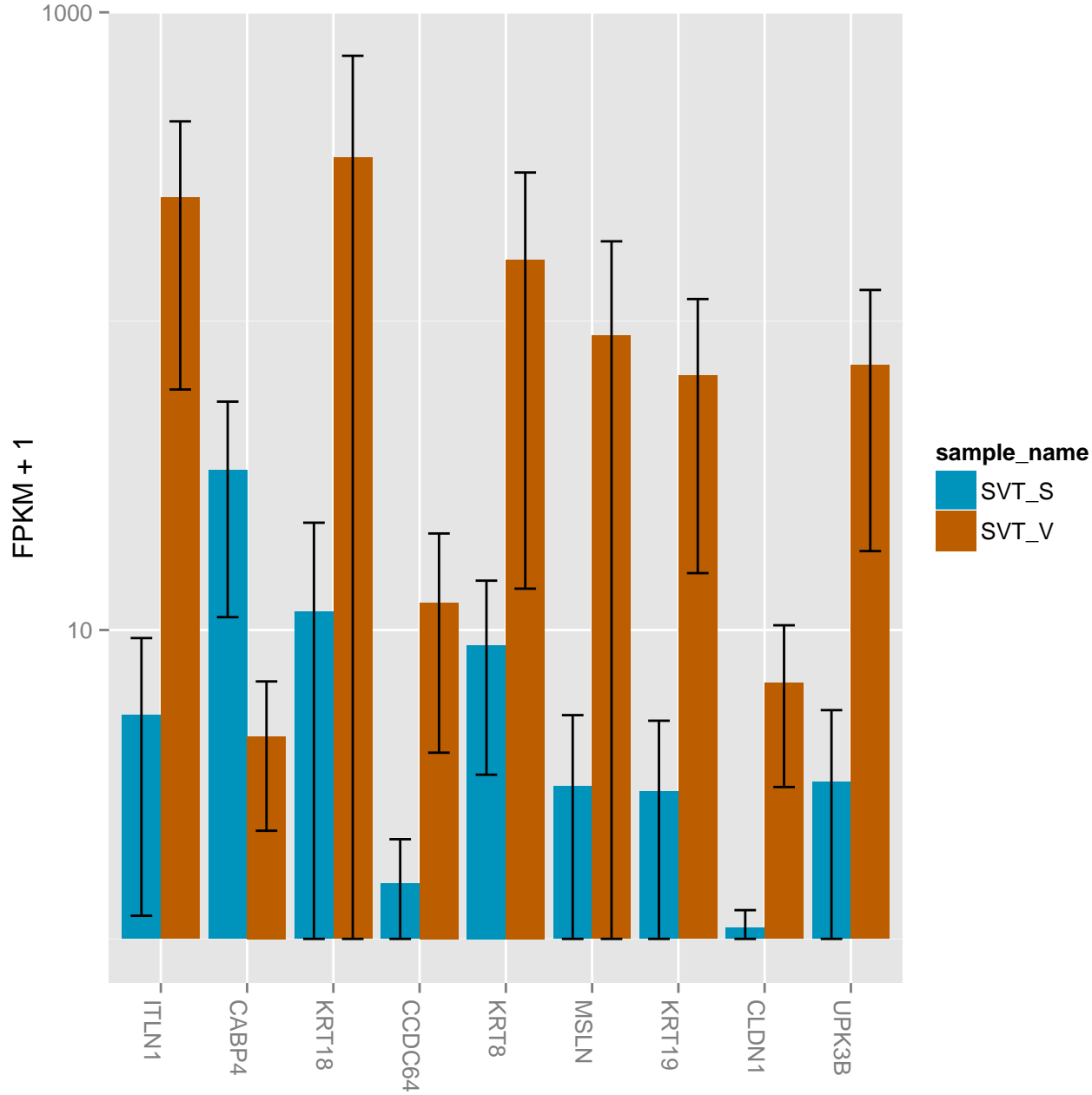
# Significant genes at alpha 0.01

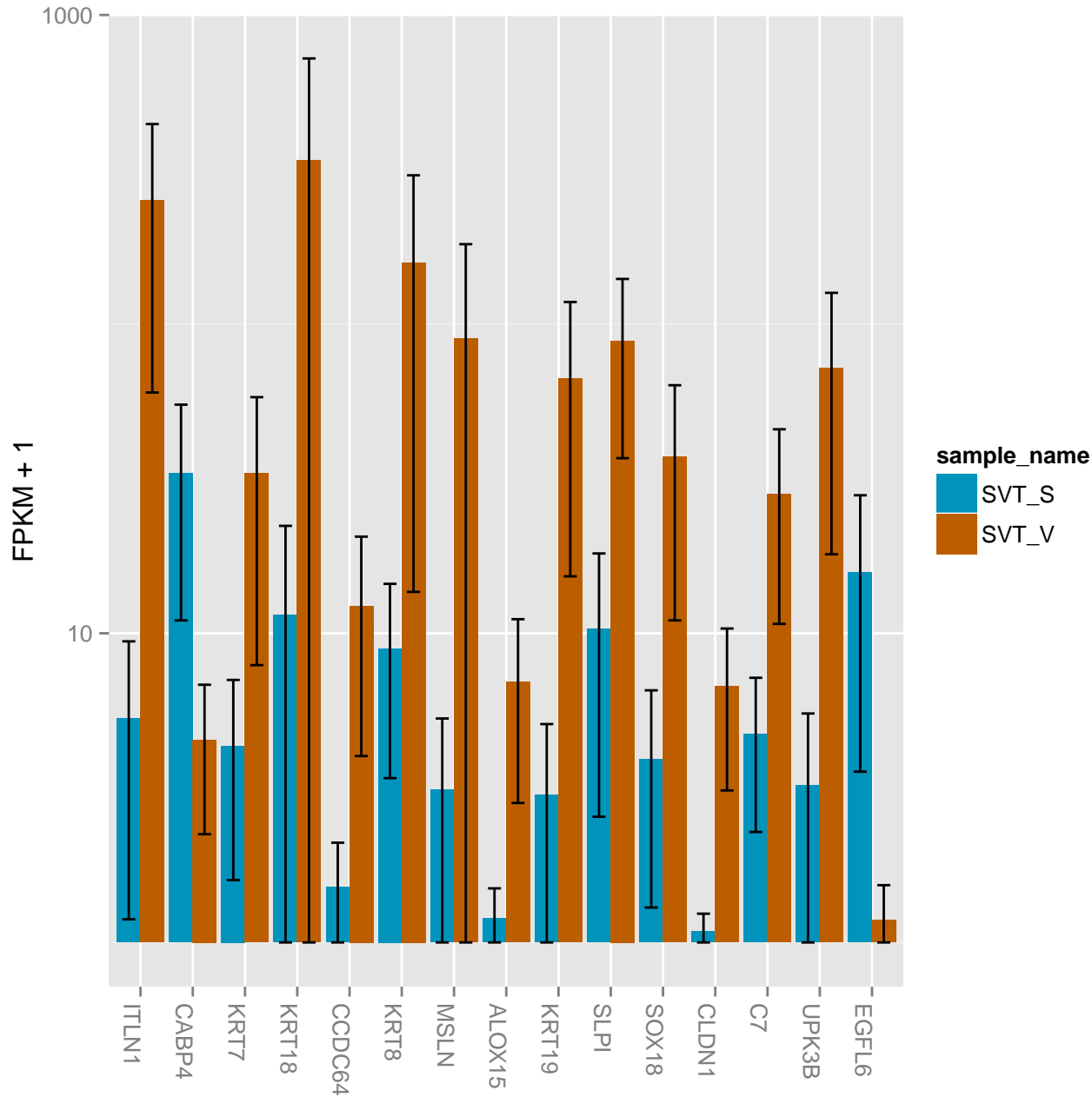


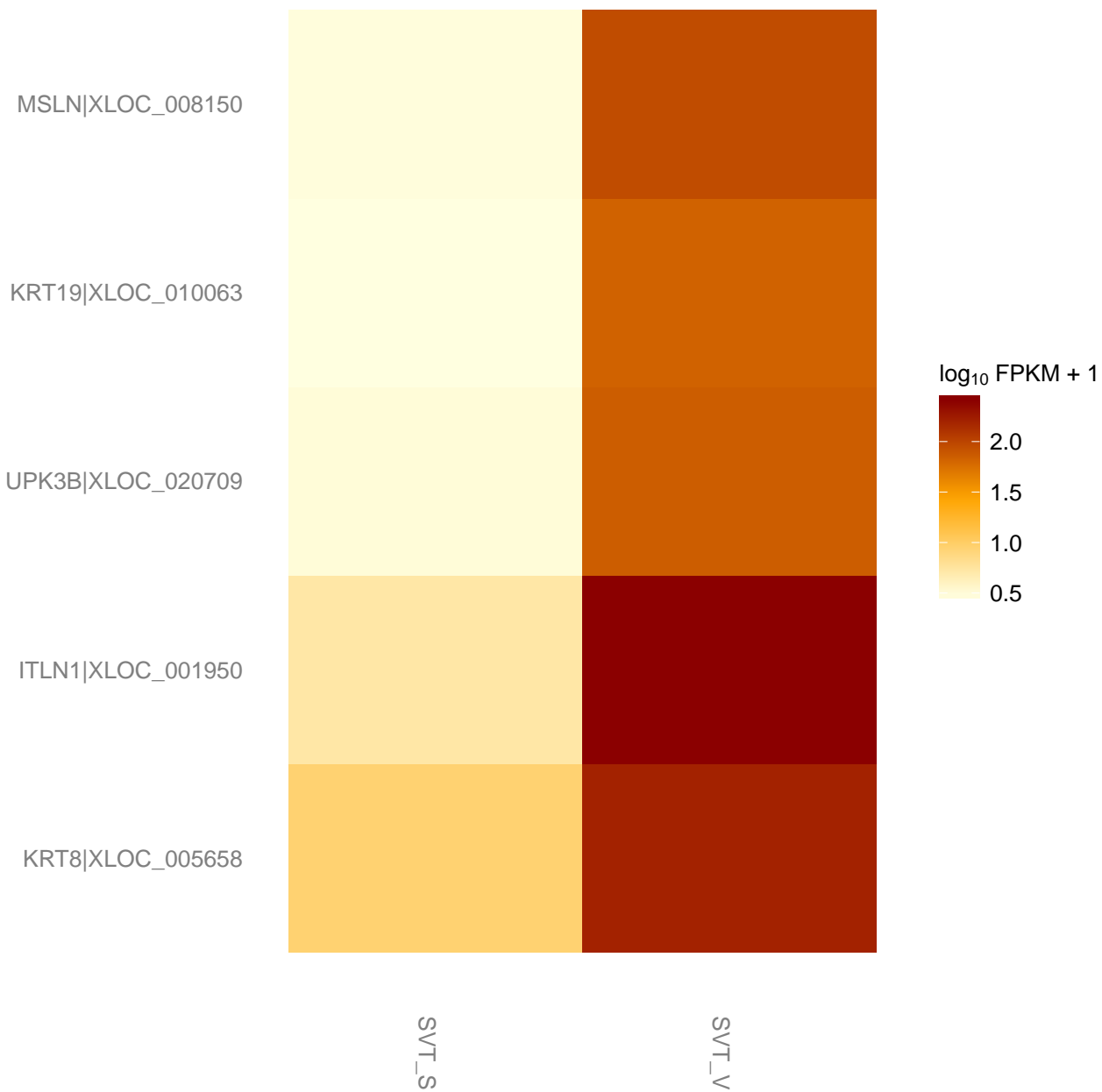
Significant genes  
at alpha 0.05

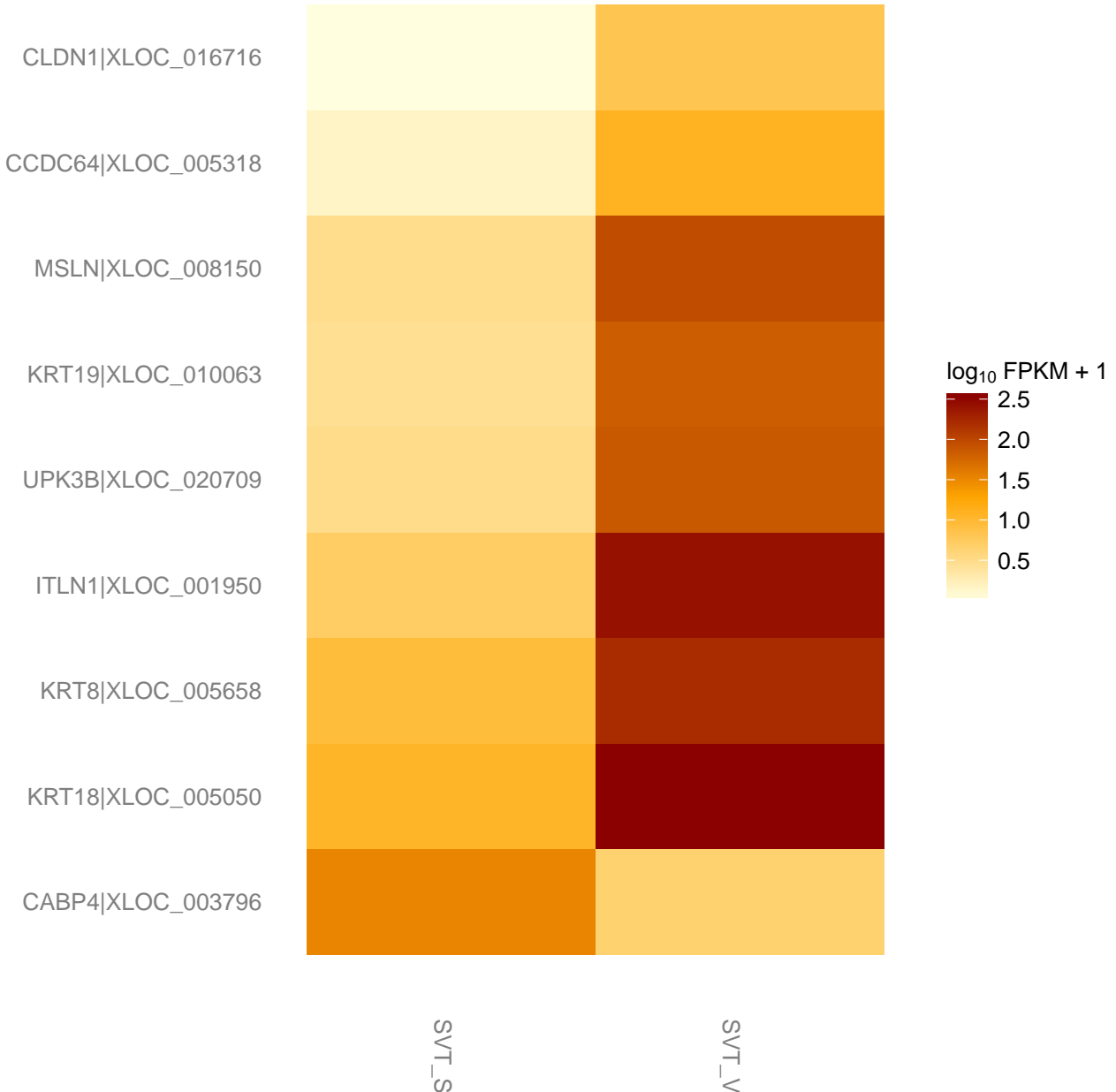


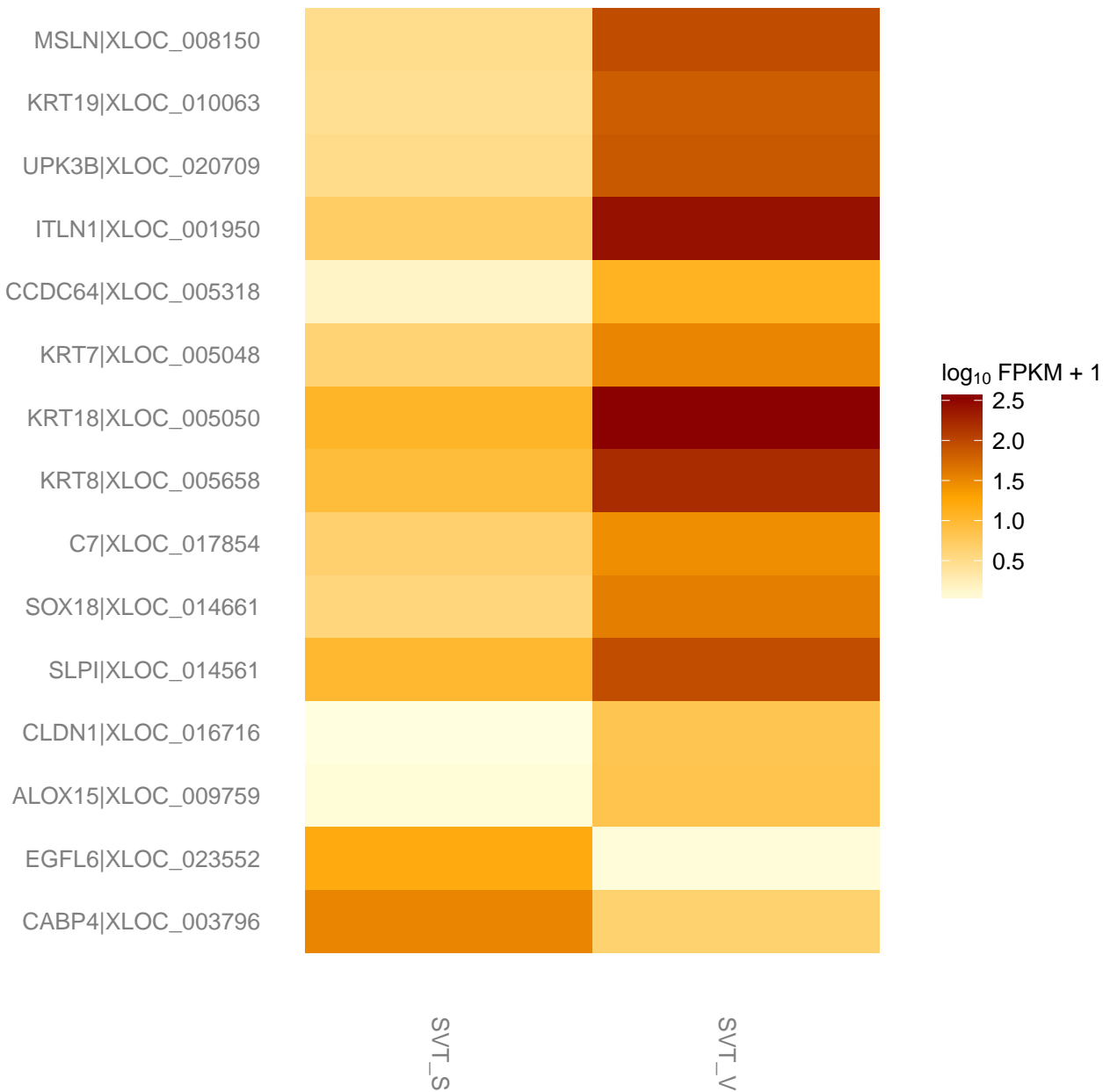












SVT\_V FPKM + 0

100

SVT\_S FPKM + 0



