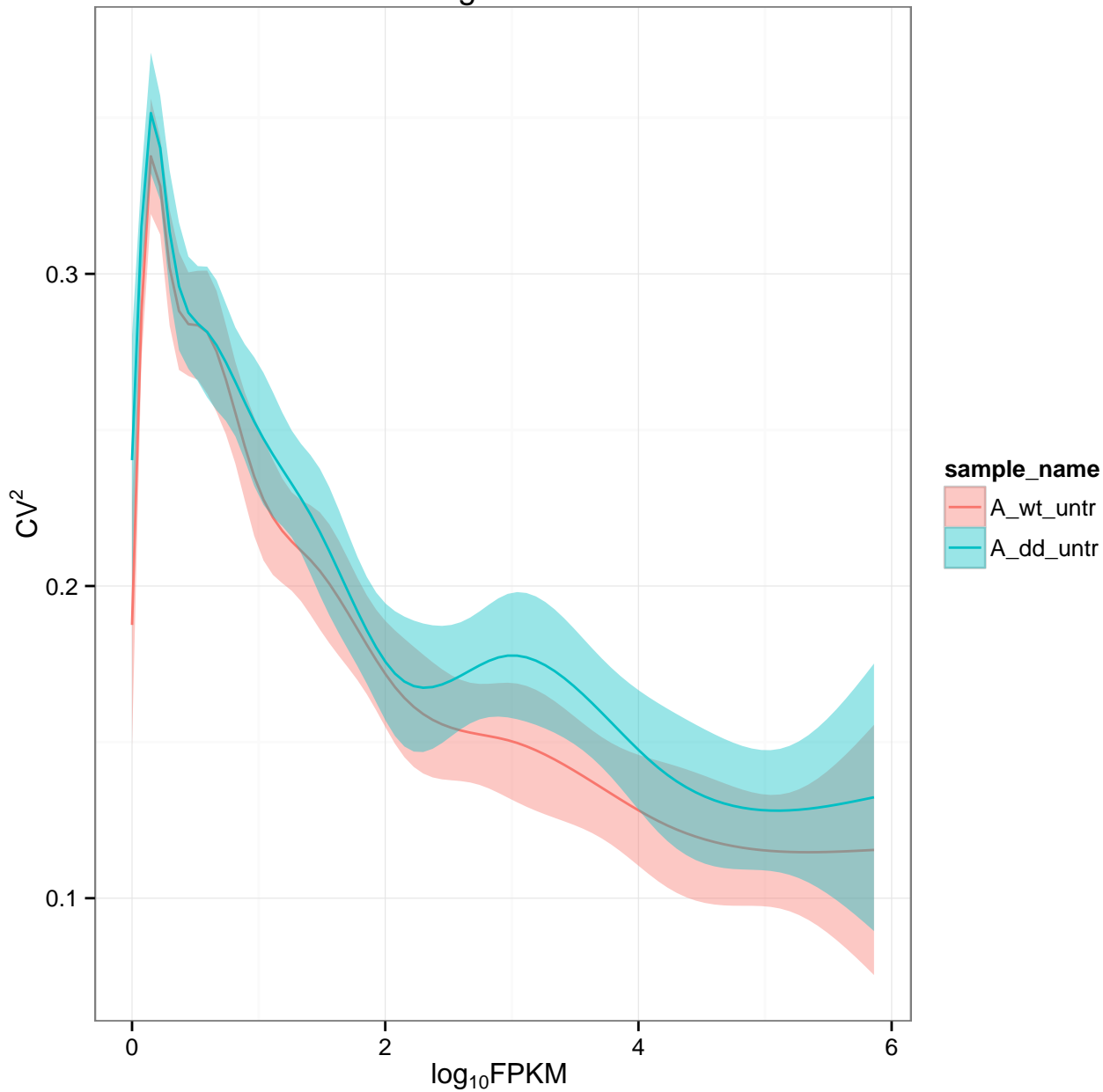
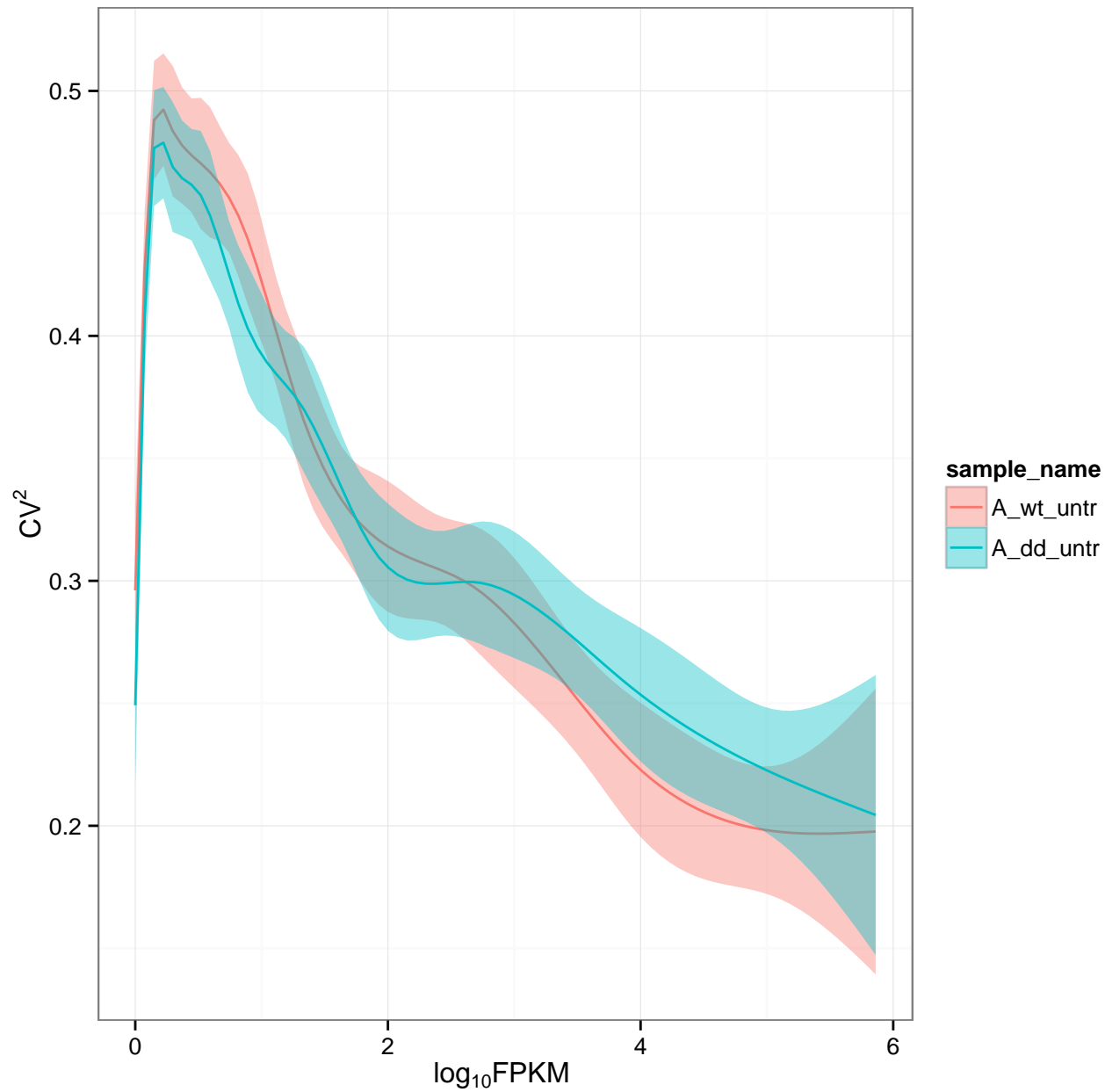


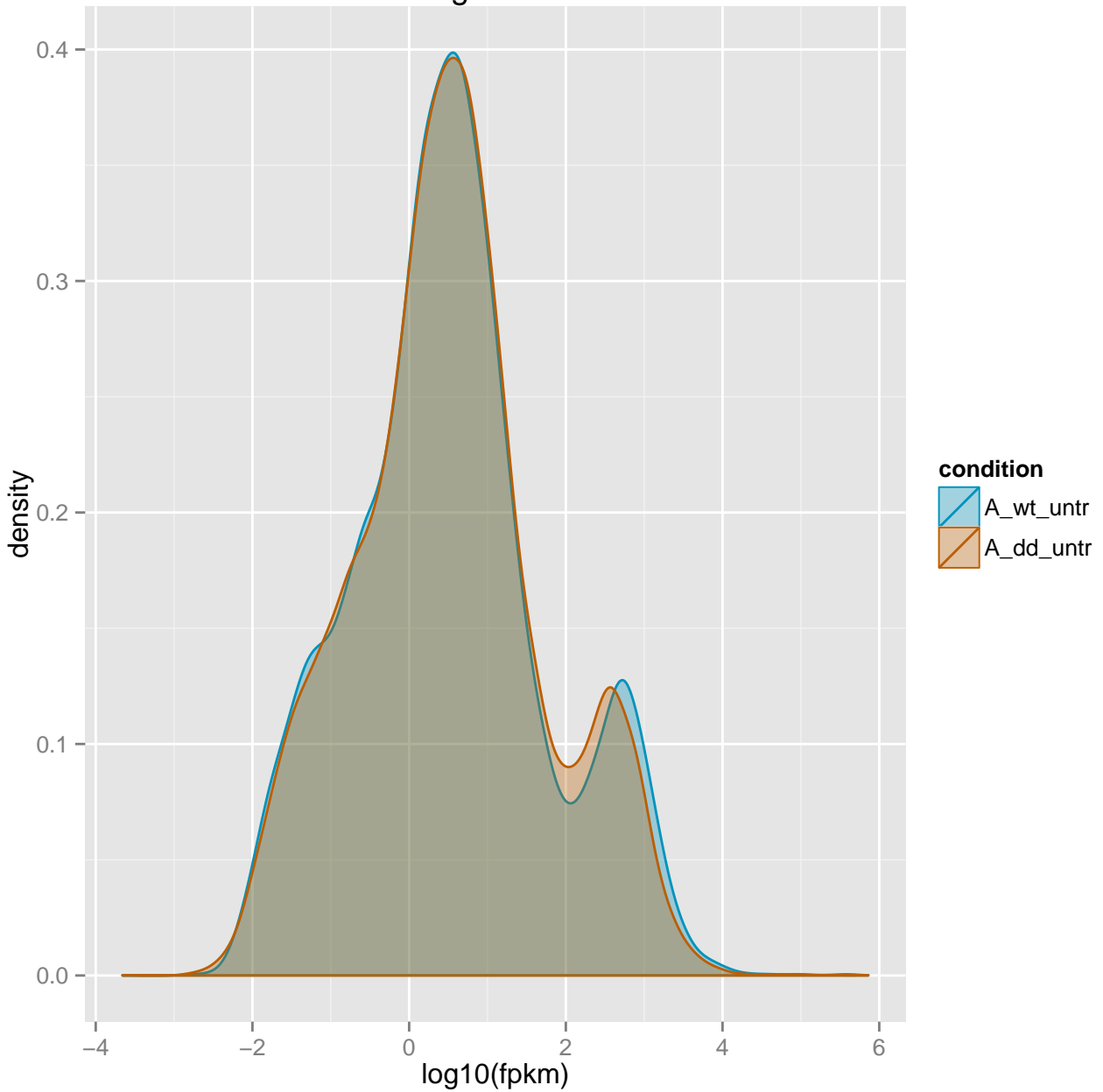
# genes



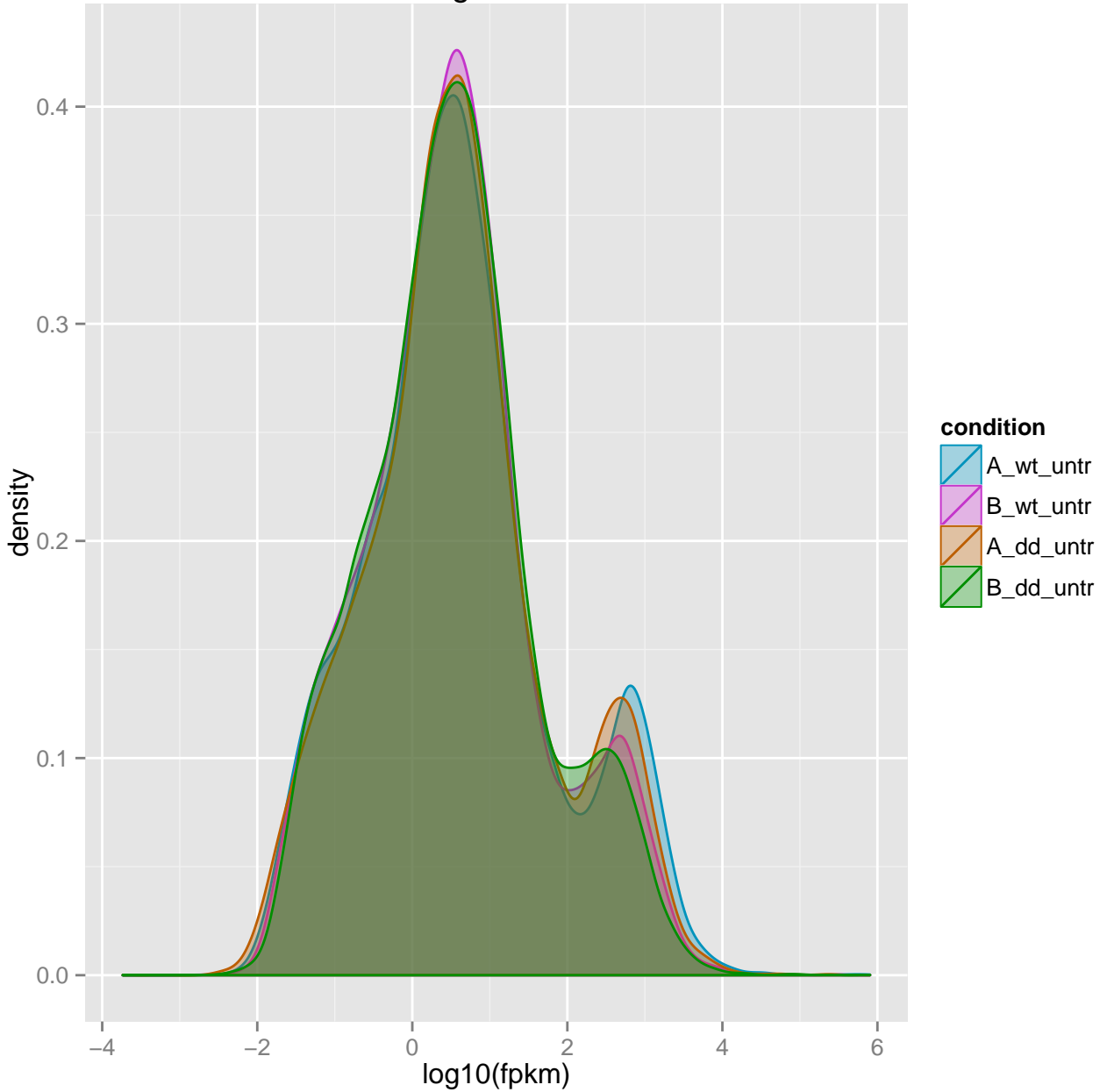
# isoforms

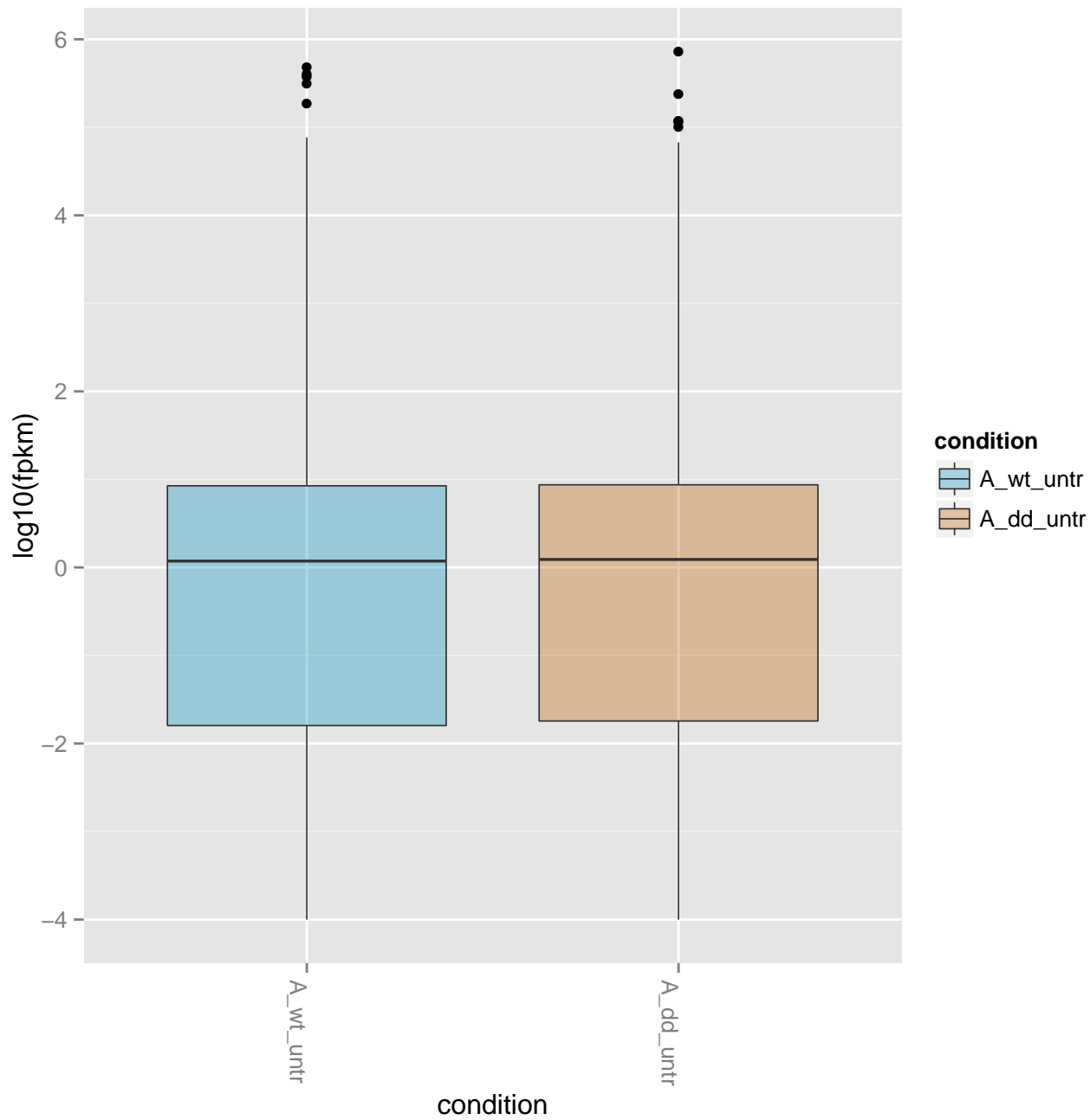


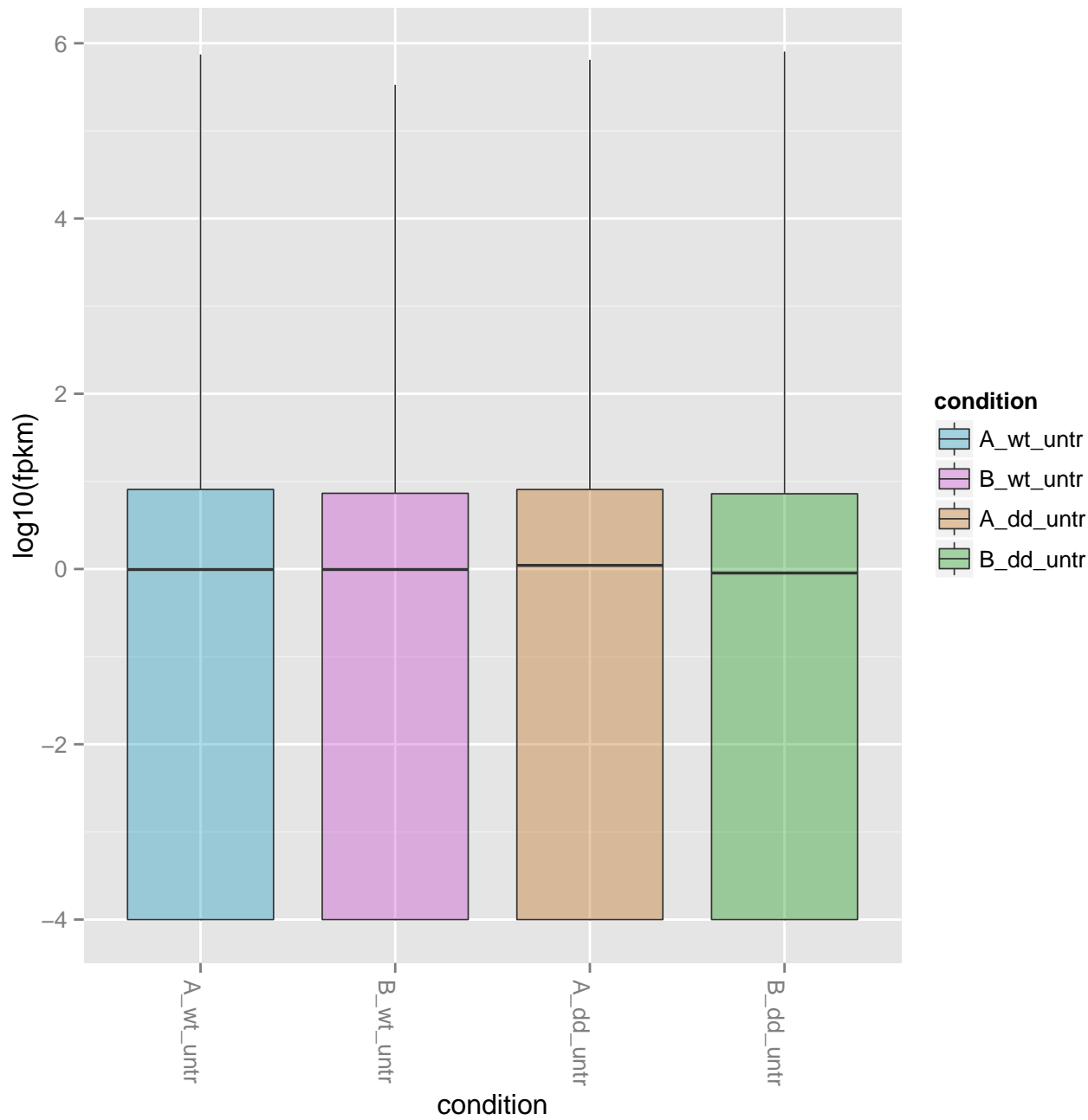
# genes

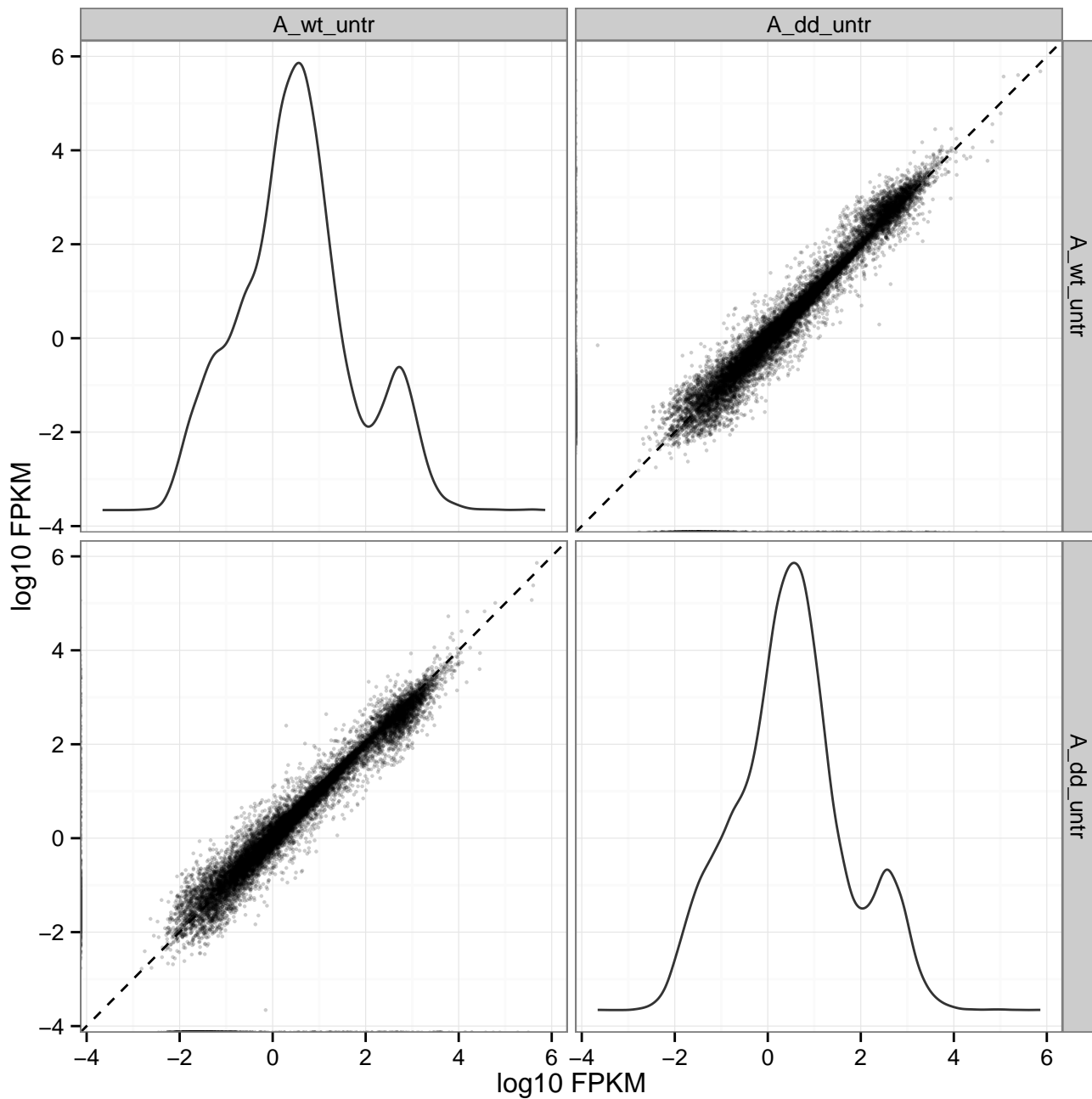


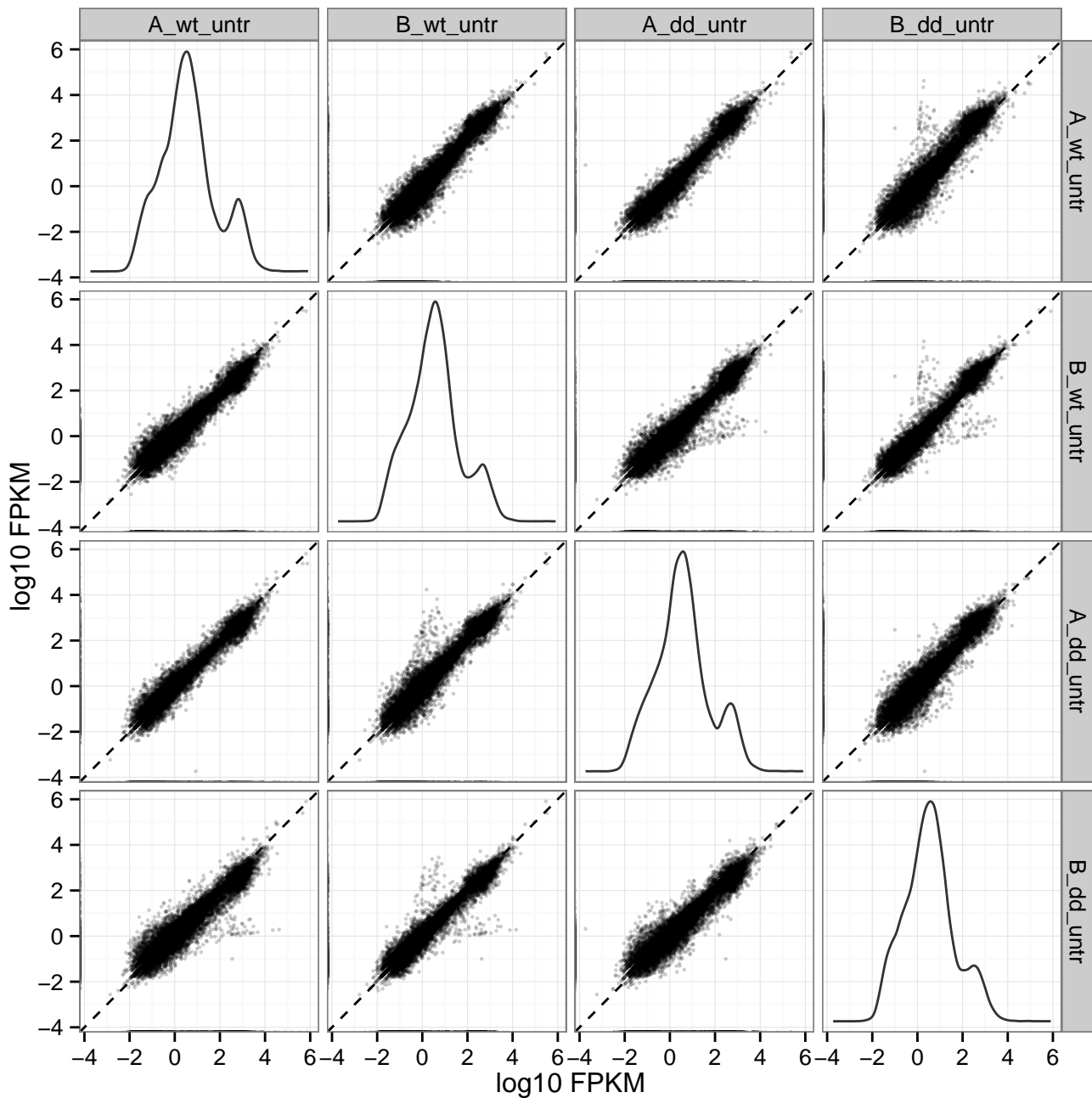
# genes



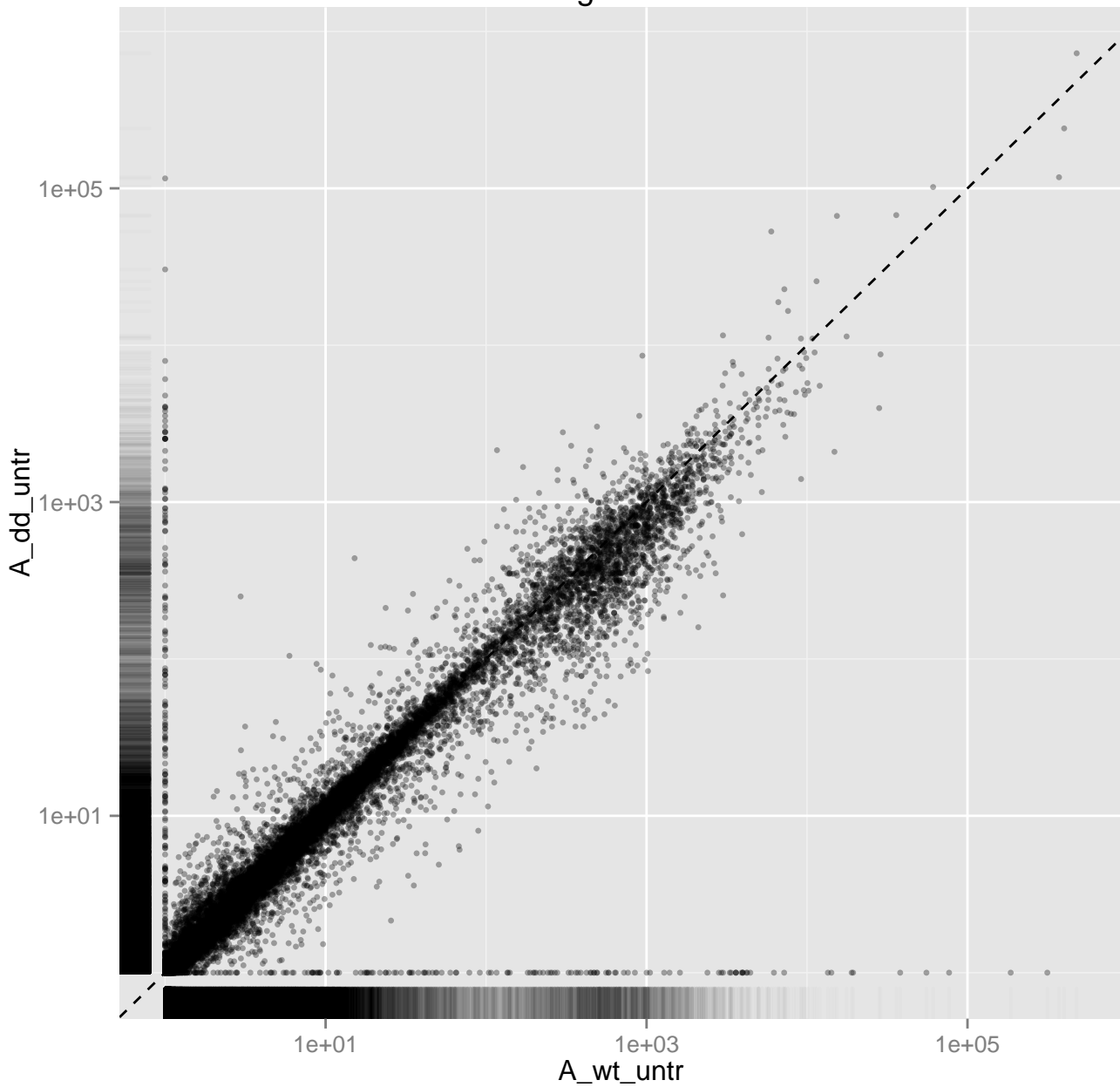




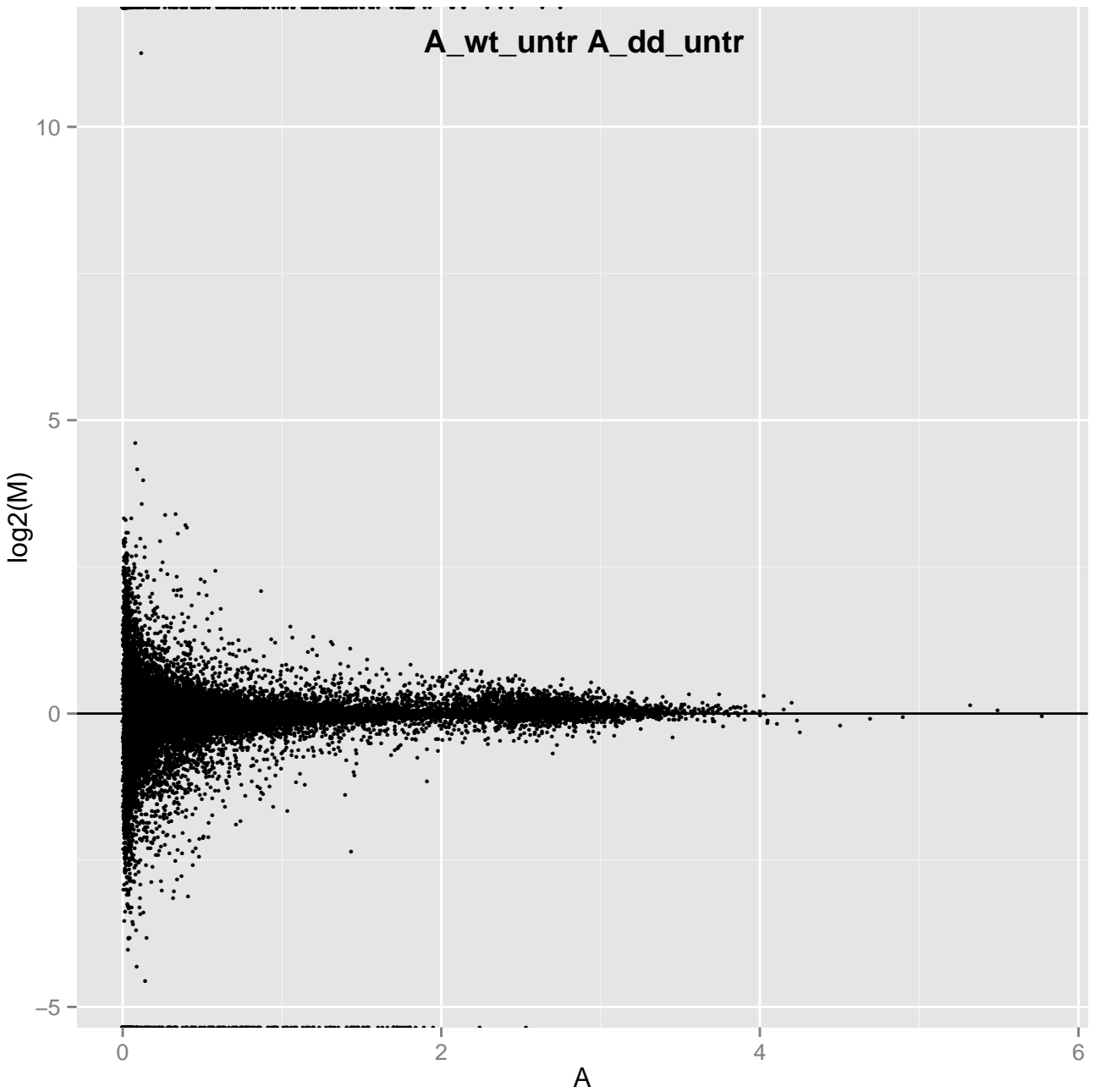




# genes









normalized count values A\_wt\_untr A\_dd\_untr

$\log_2(M)$

8

4

0

0

1

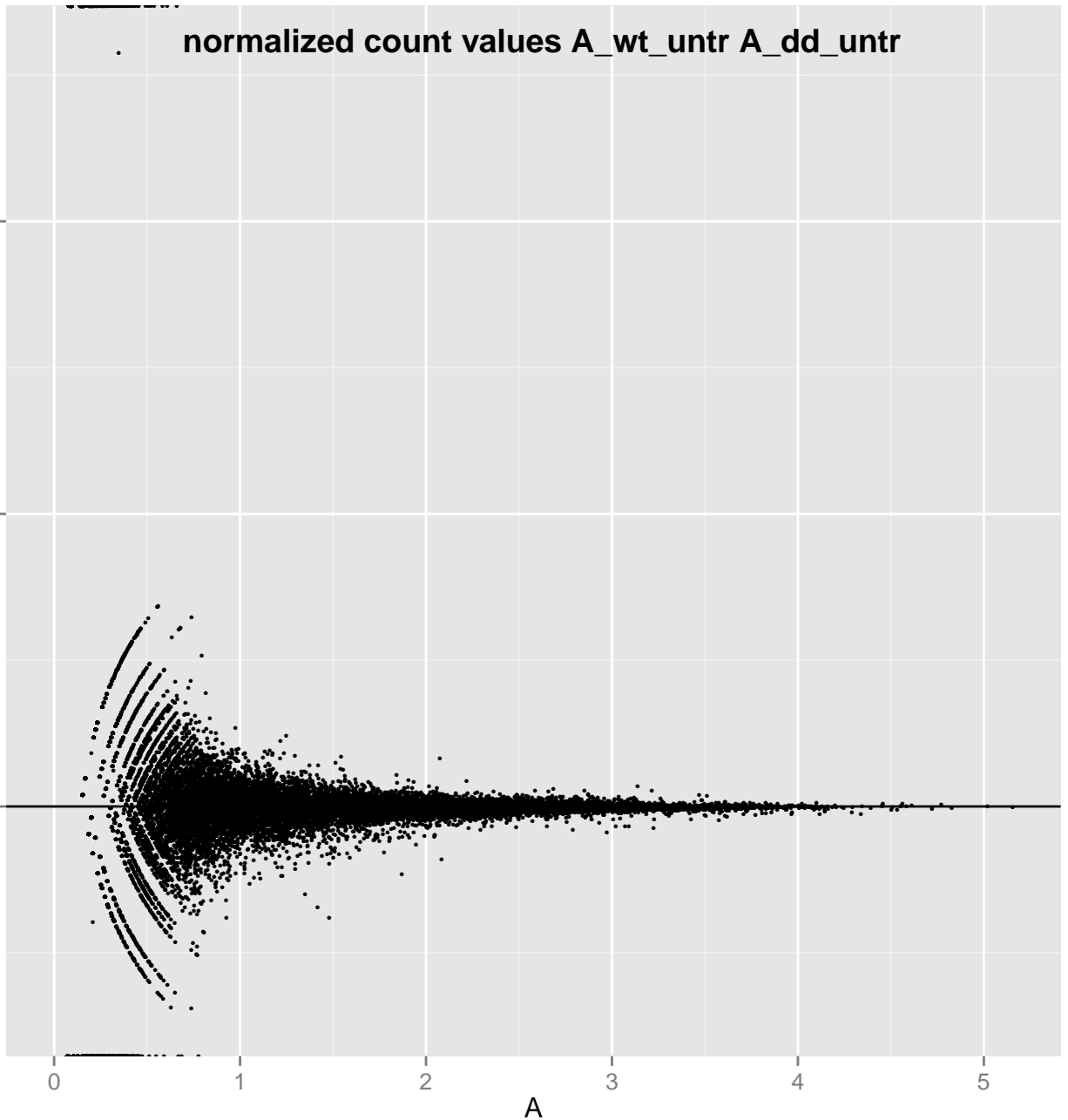
2

3

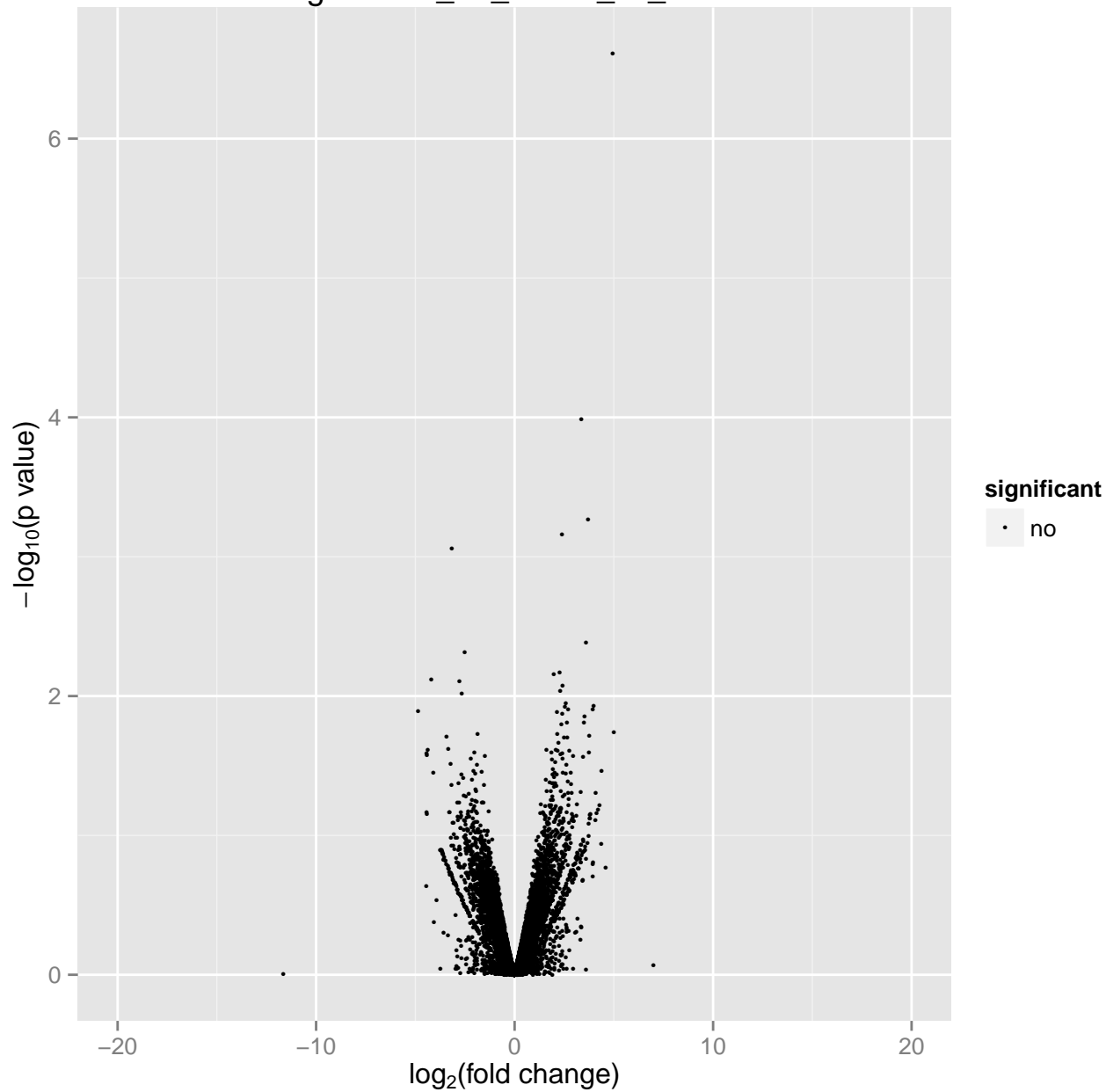
4

5

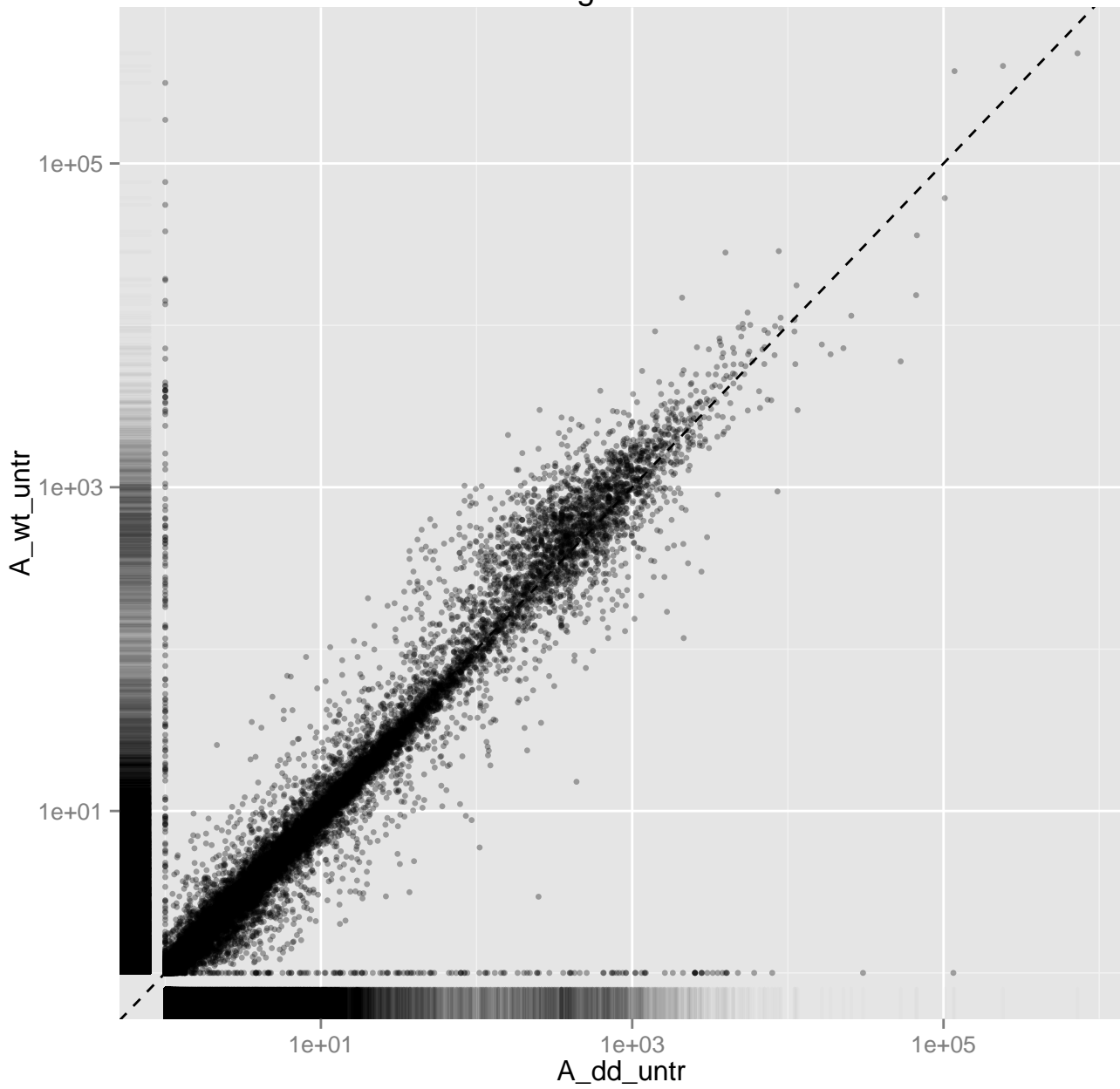
A



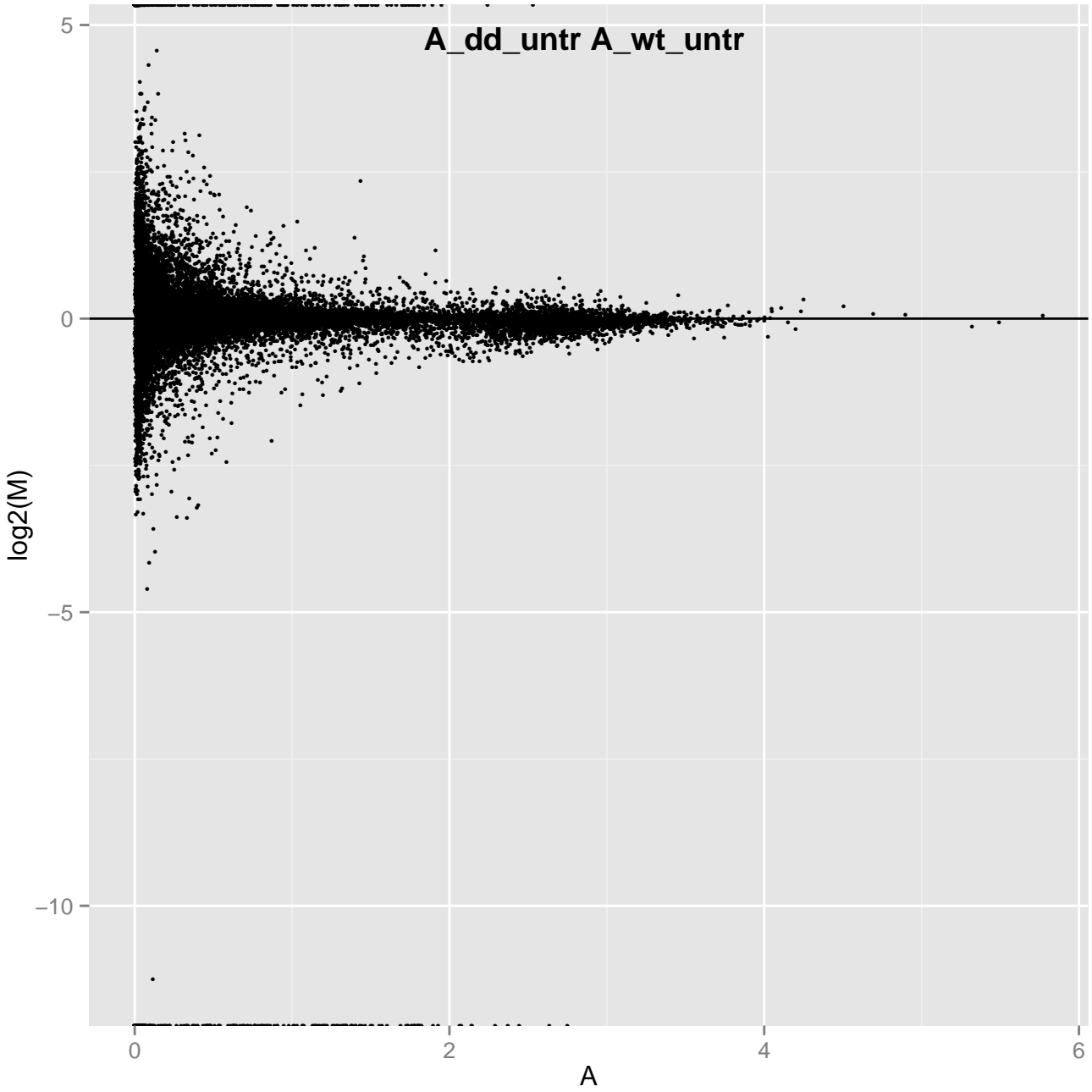
genes: A\_dd\_untr/A\_wt\_untr



genes









normalized count values A\_dd\_untr A\_wt\_untr

$\log_2(M)$

0

-4

-8

0

1

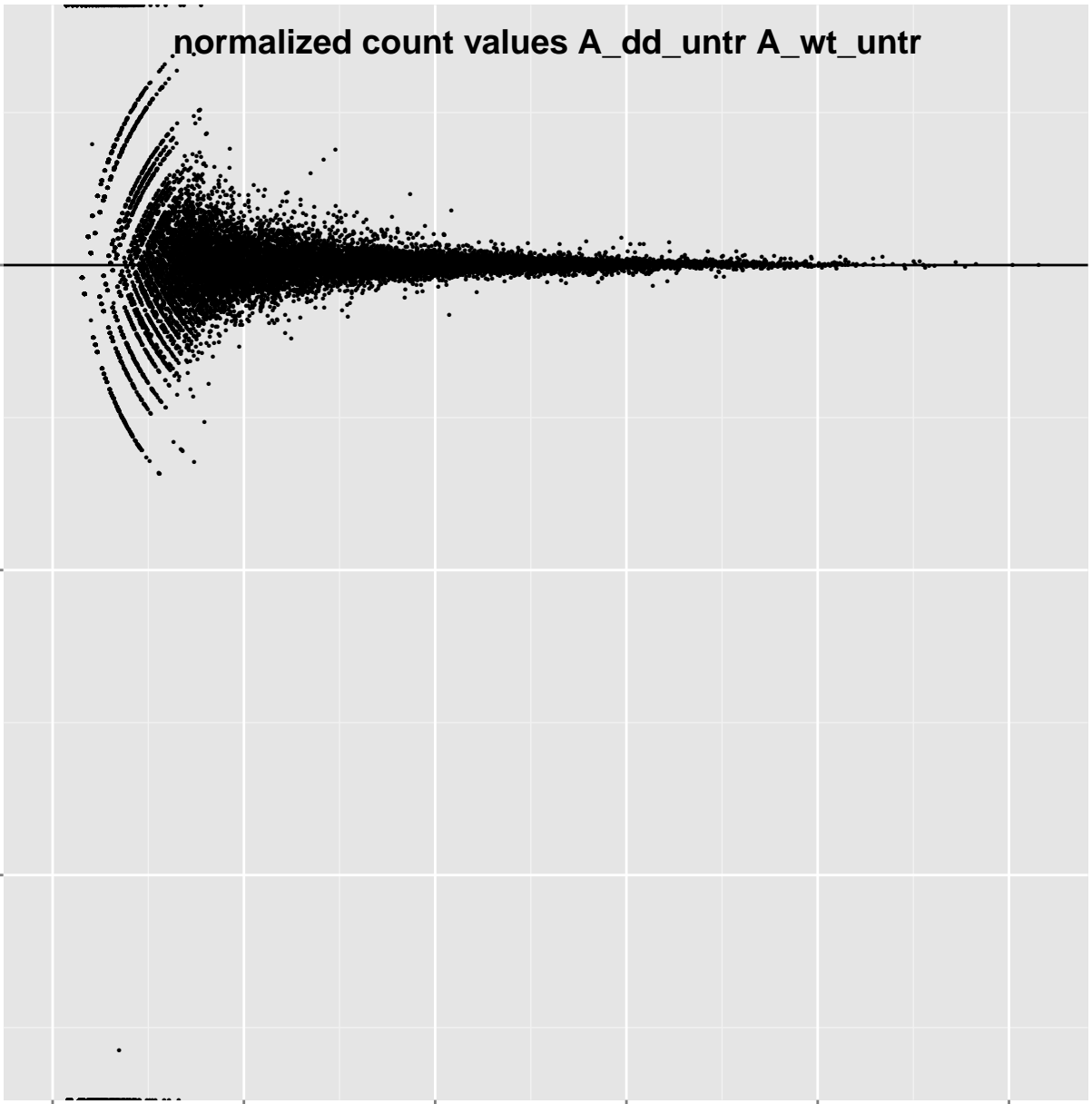
2

3

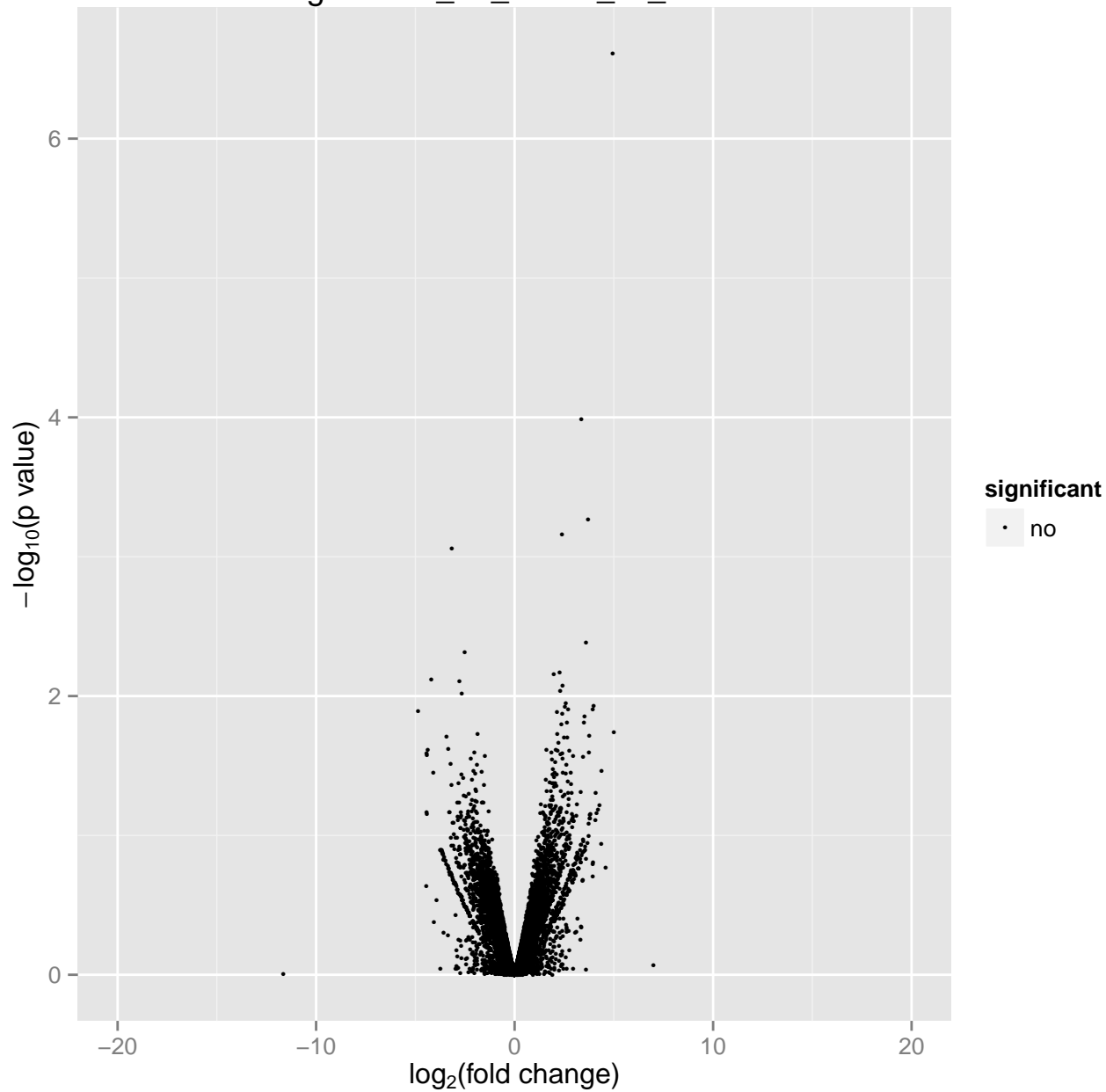
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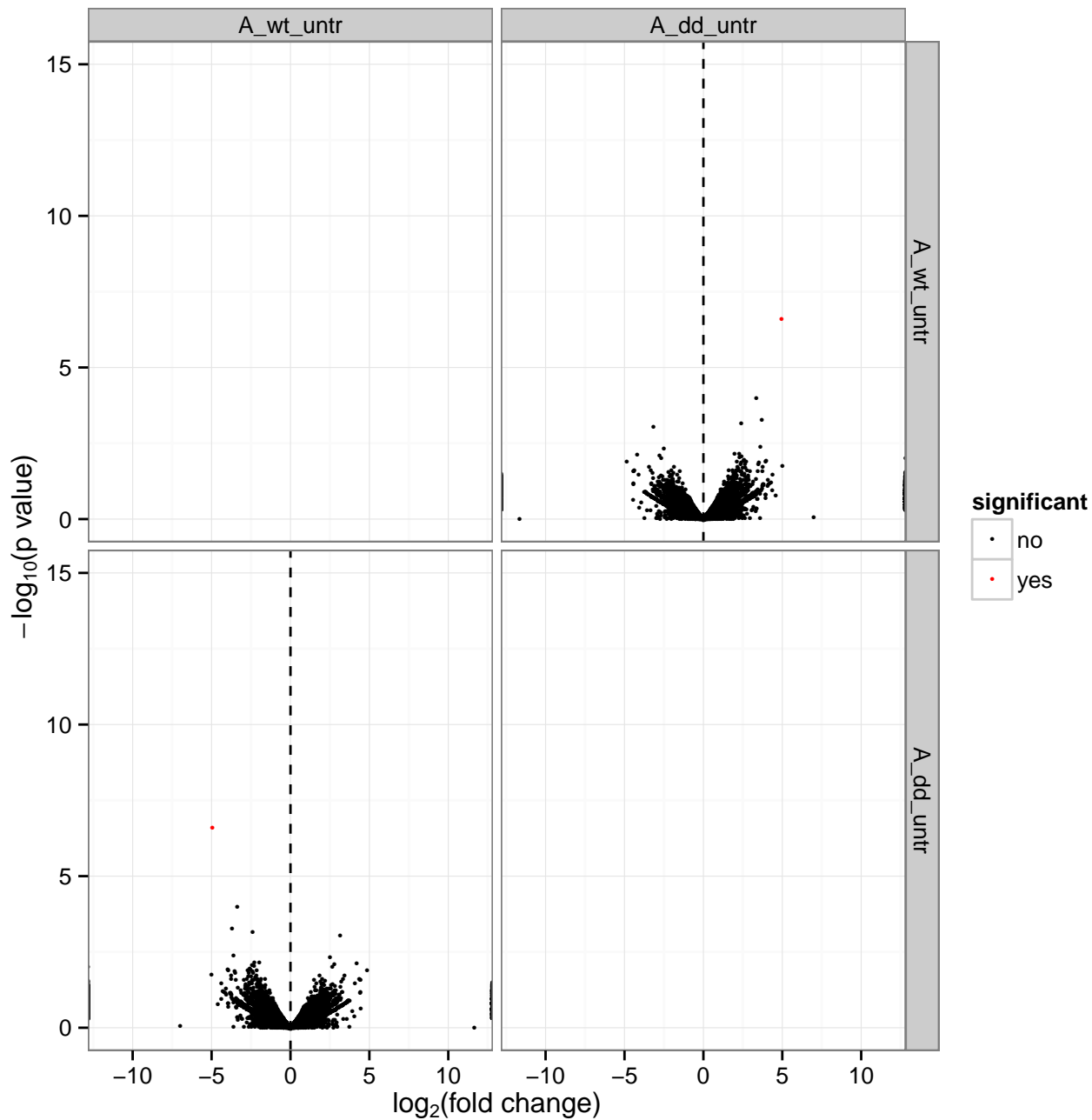
5

A

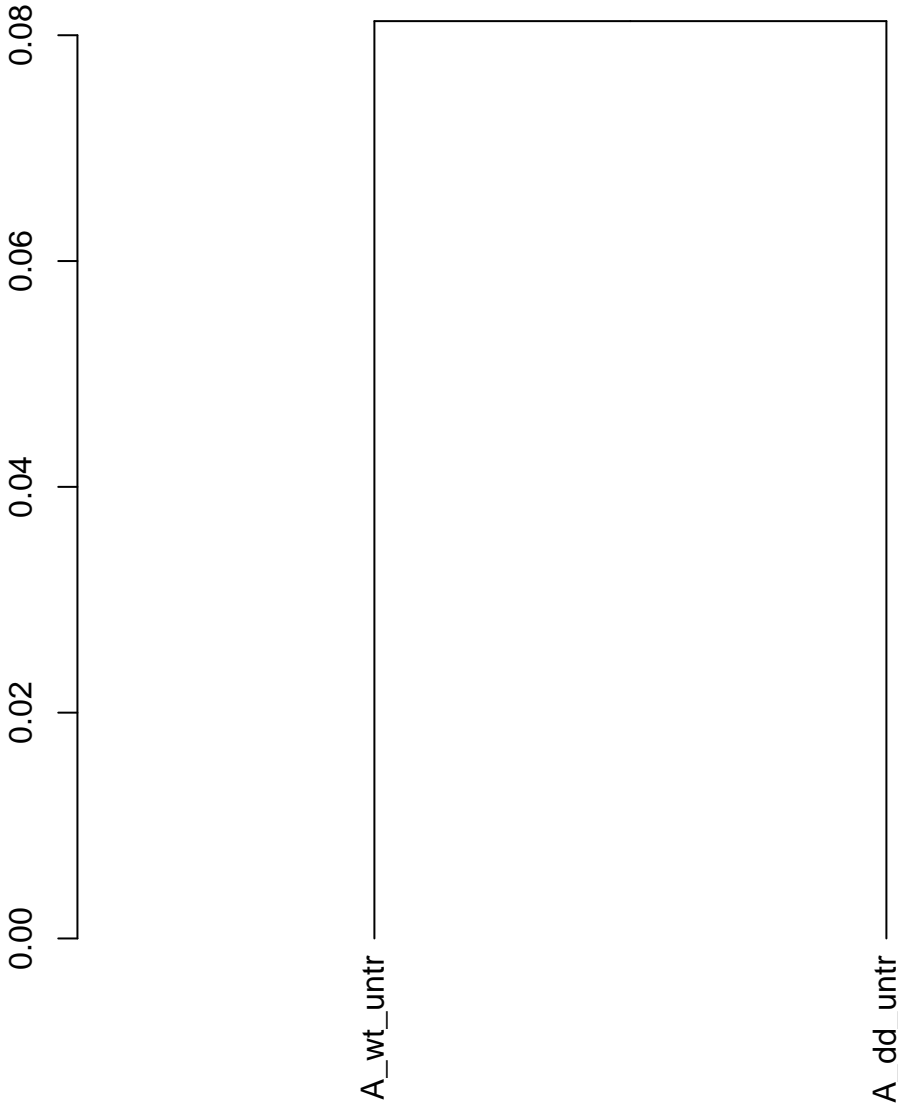


genes: A\_dd\_untr/A\_wt\_untr

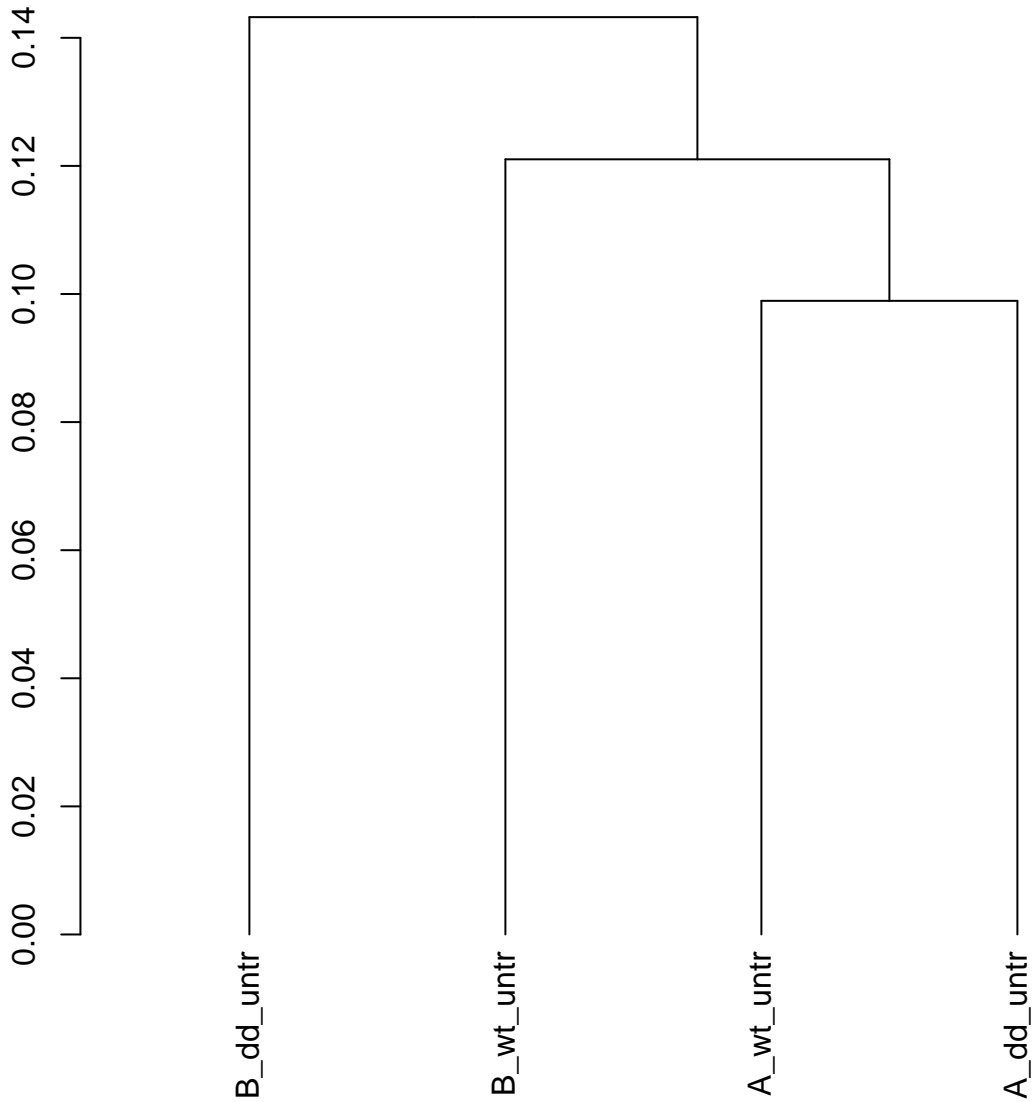


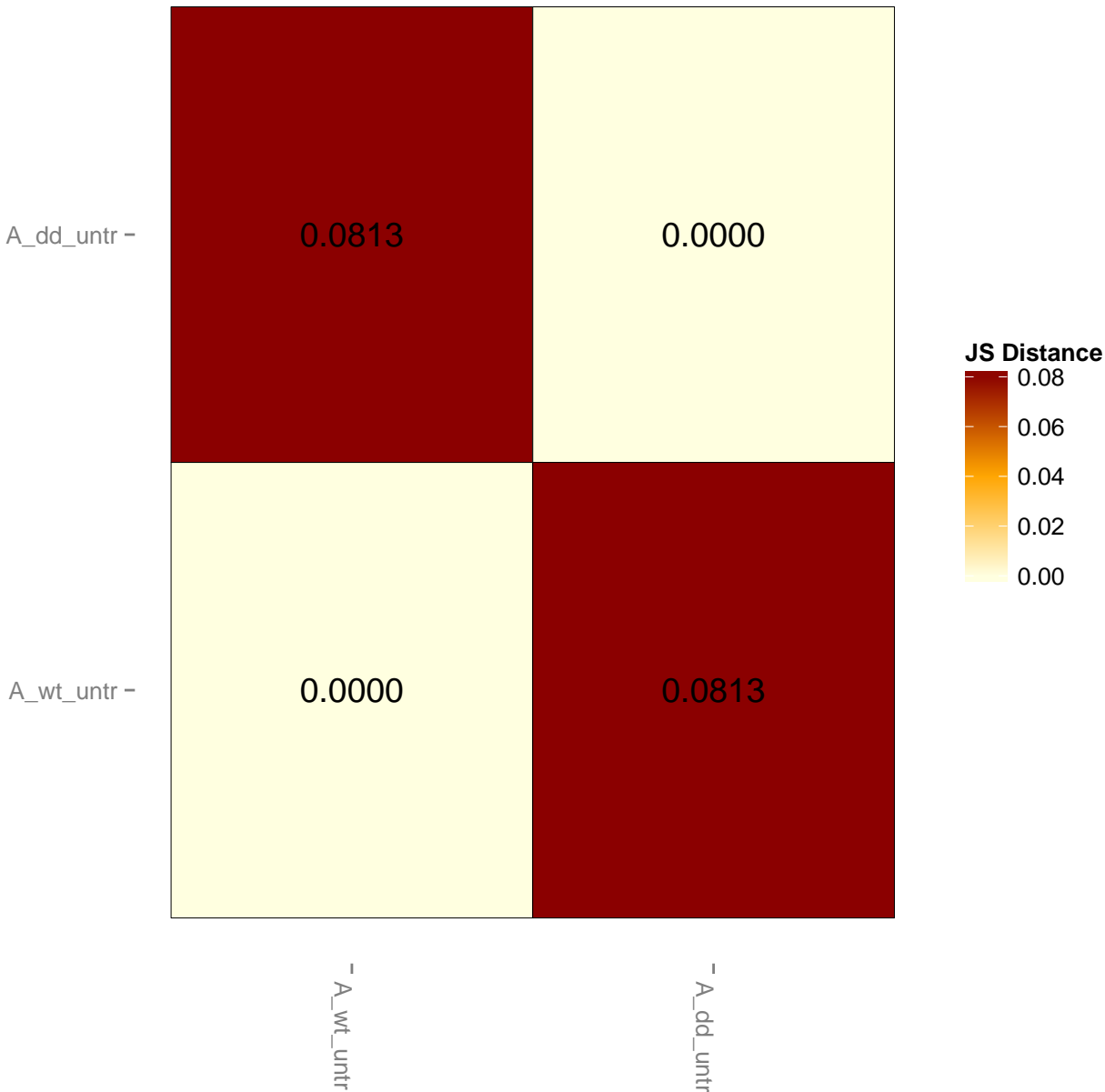


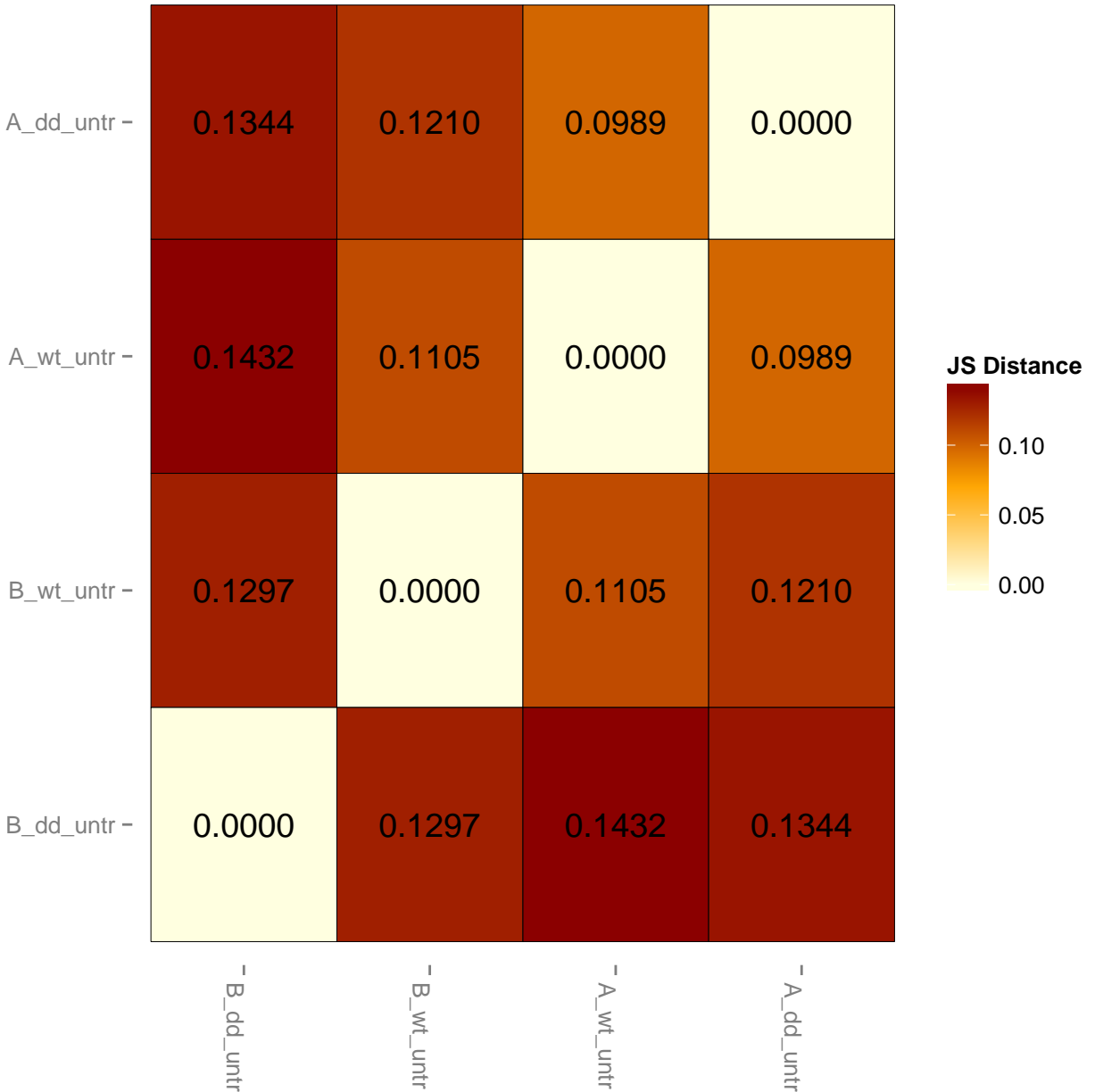
# All genes(cuff)

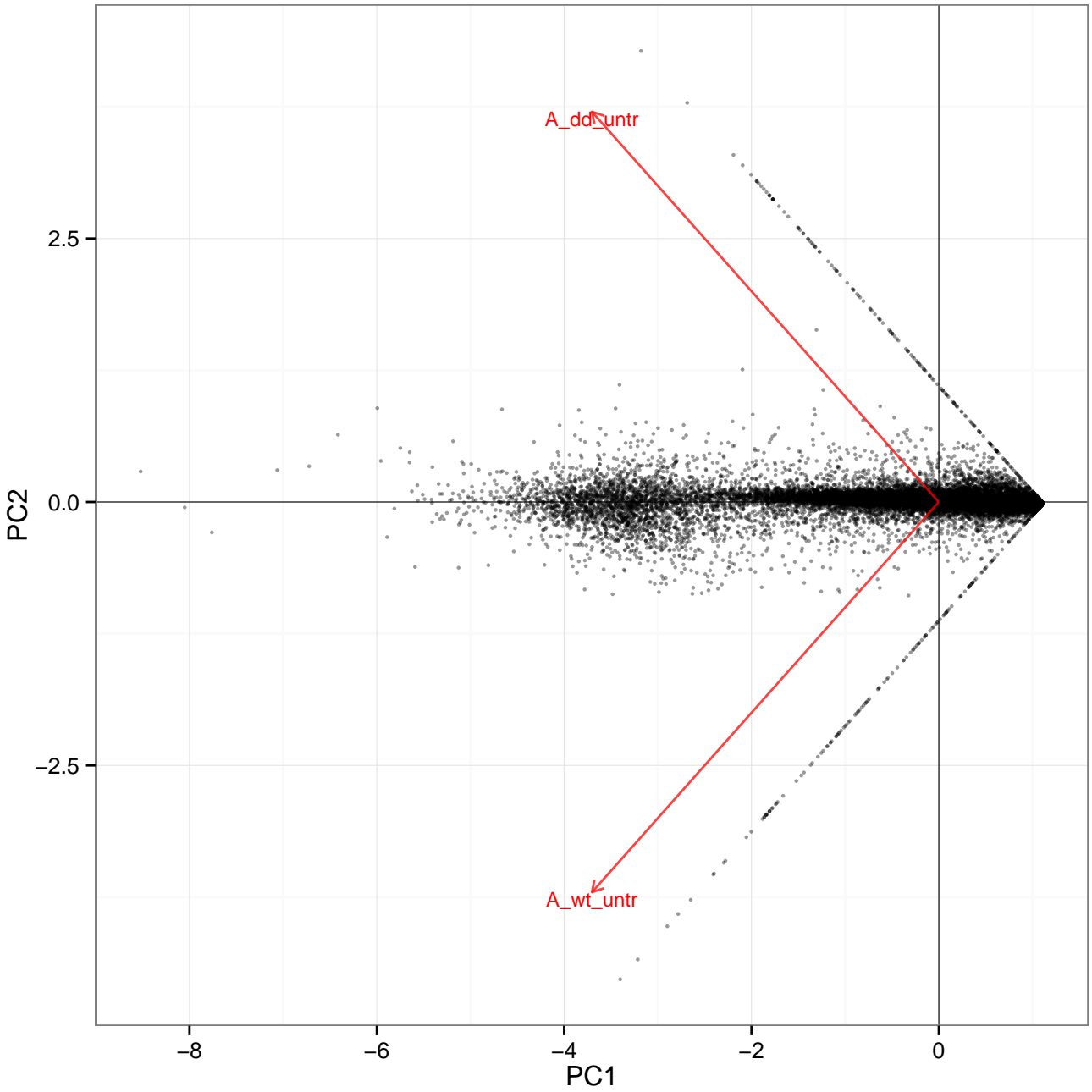


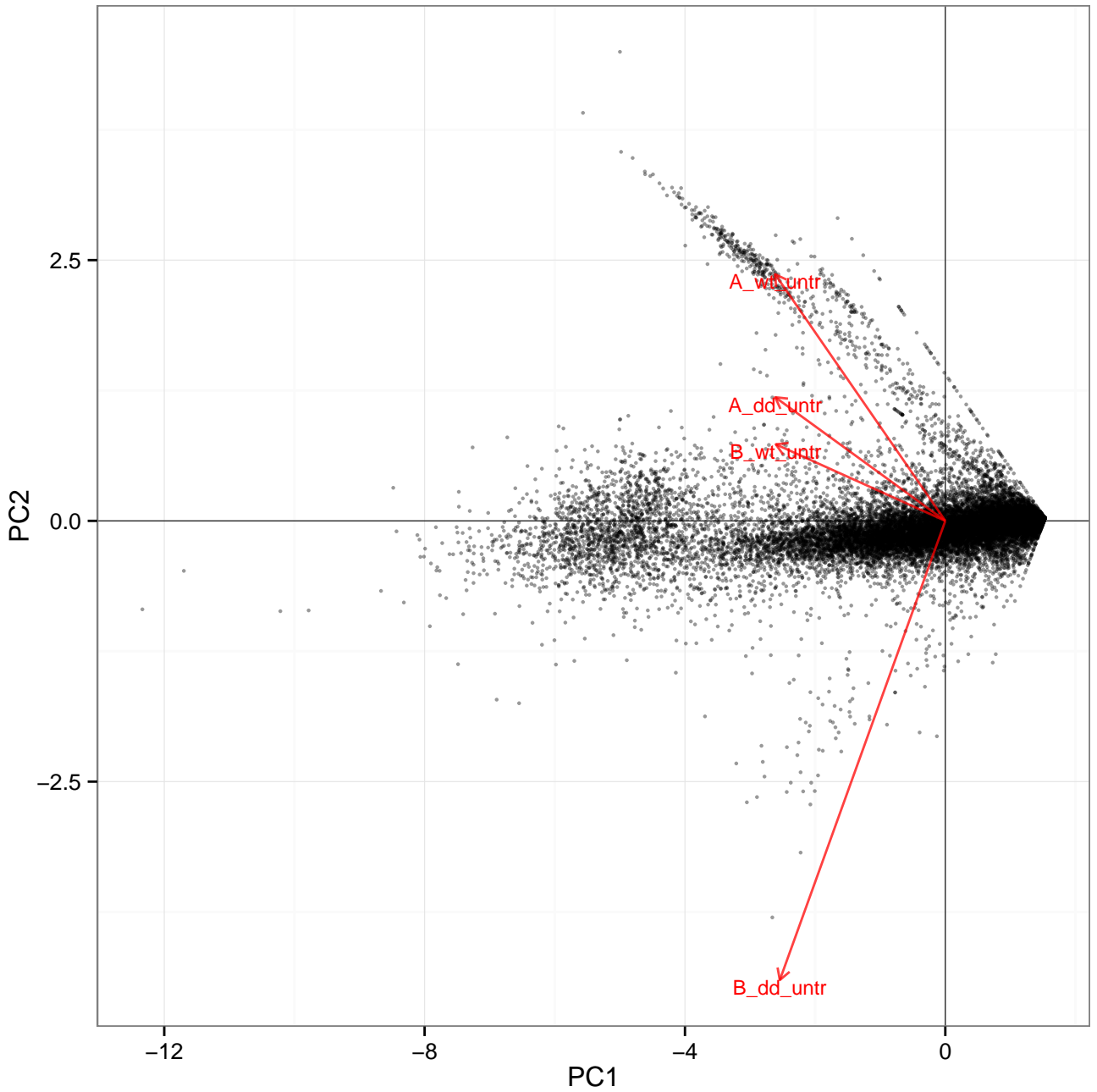
# All genes(cuff)

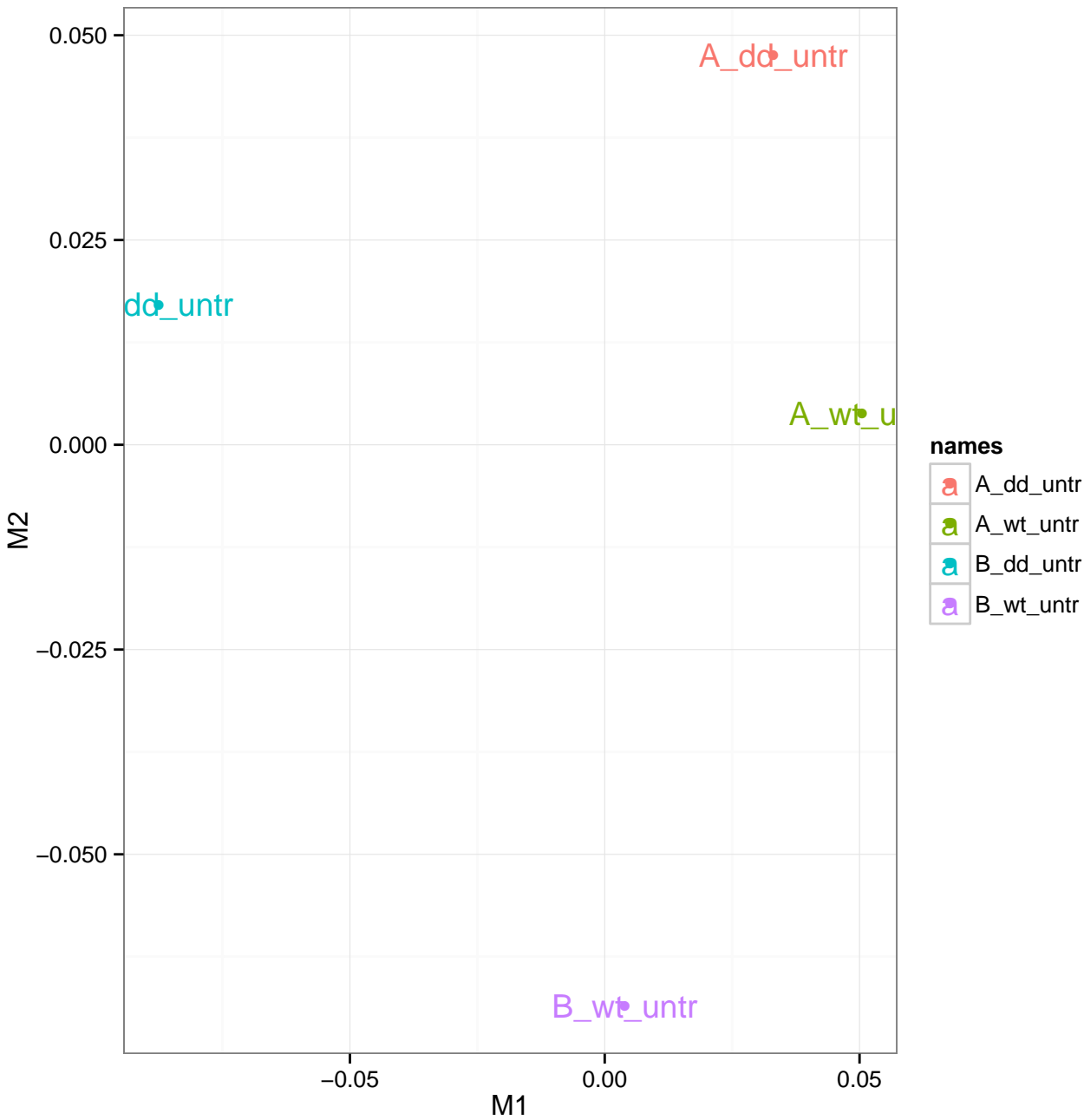


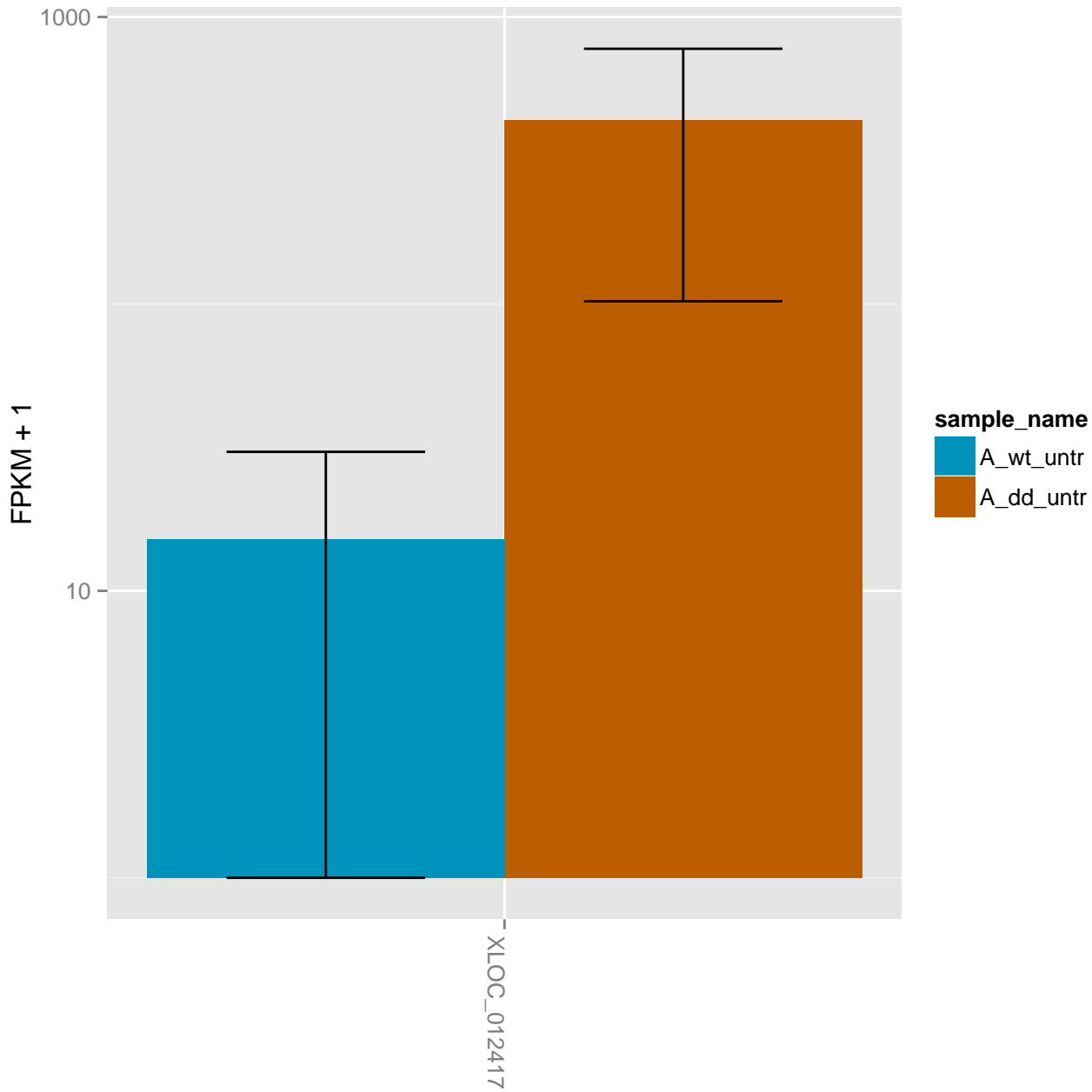


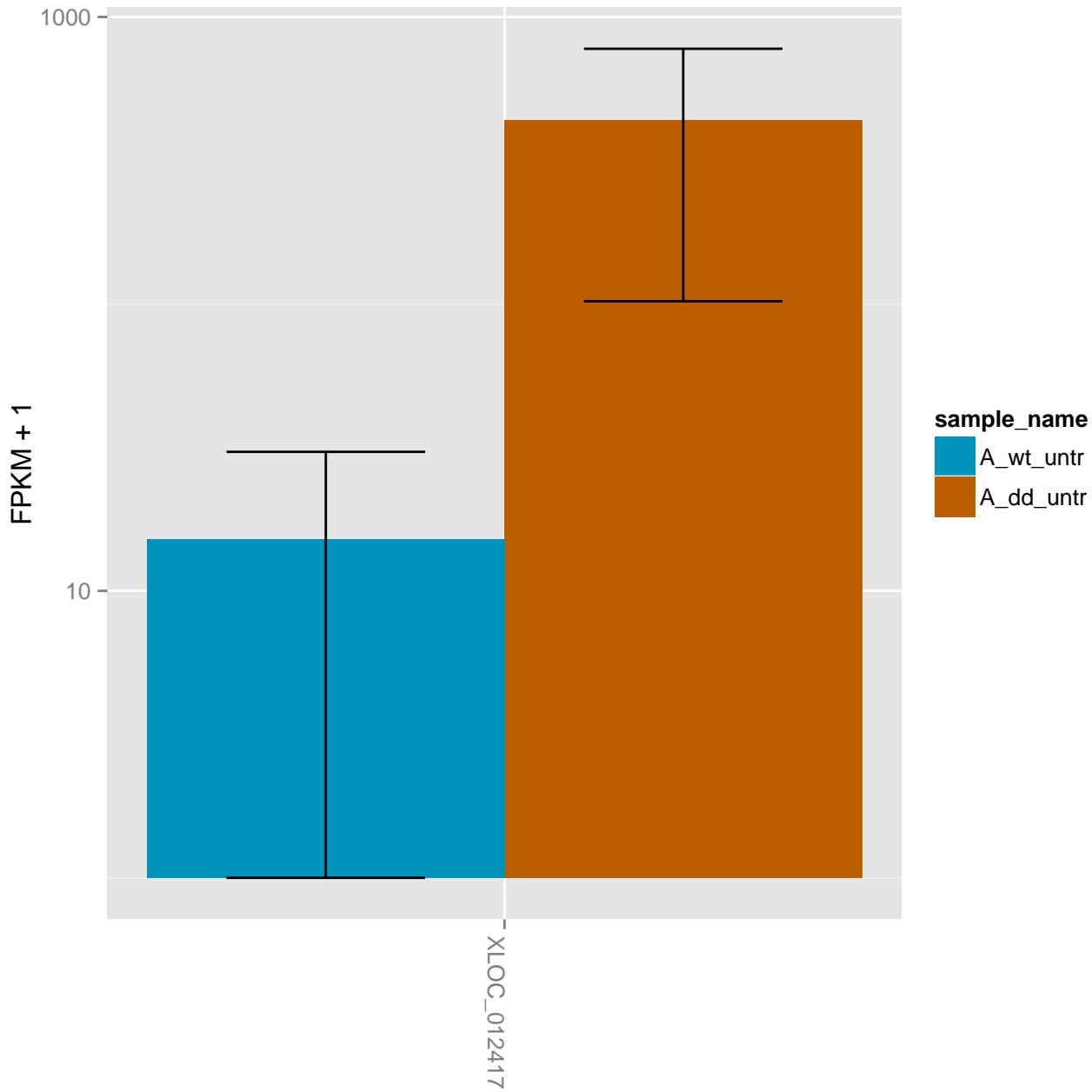












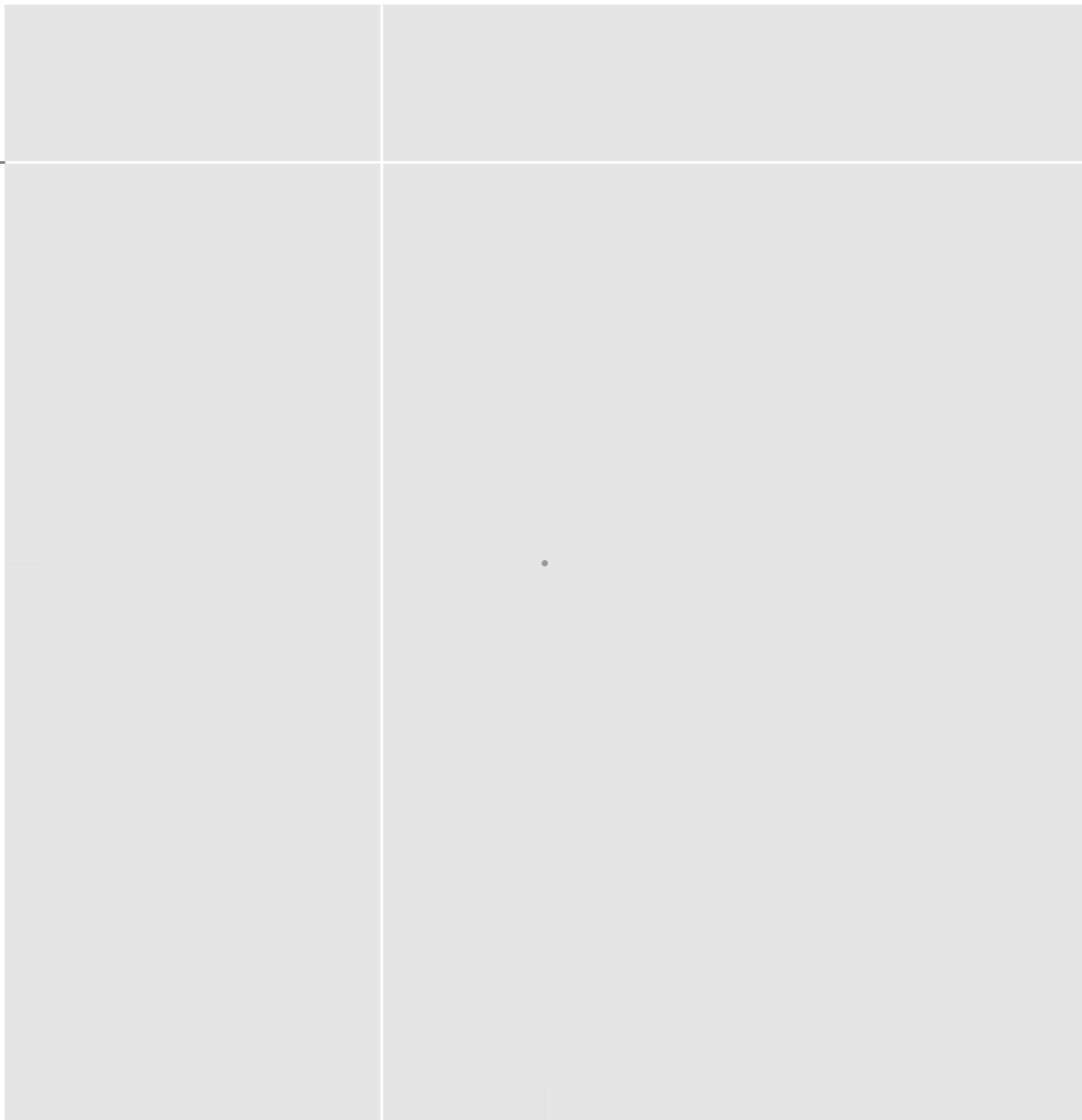
A\_dd\_untr FPKM + 0

1000

10

A\_wt\_untr FPKM + 0

.



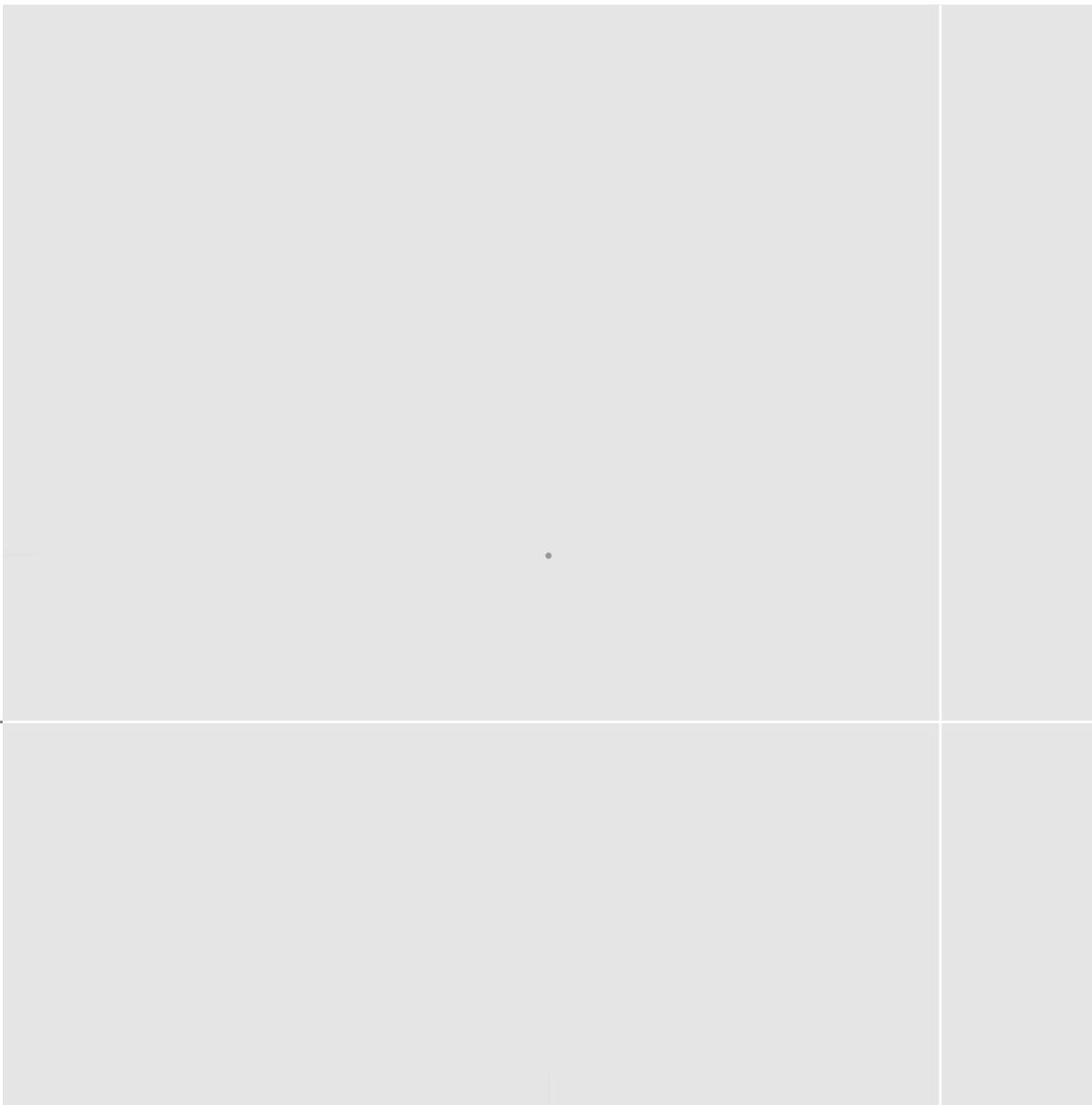
A\_wt\_untr FPKM + 0

10

A\_dd\_untr FPKM + 0

1000

.



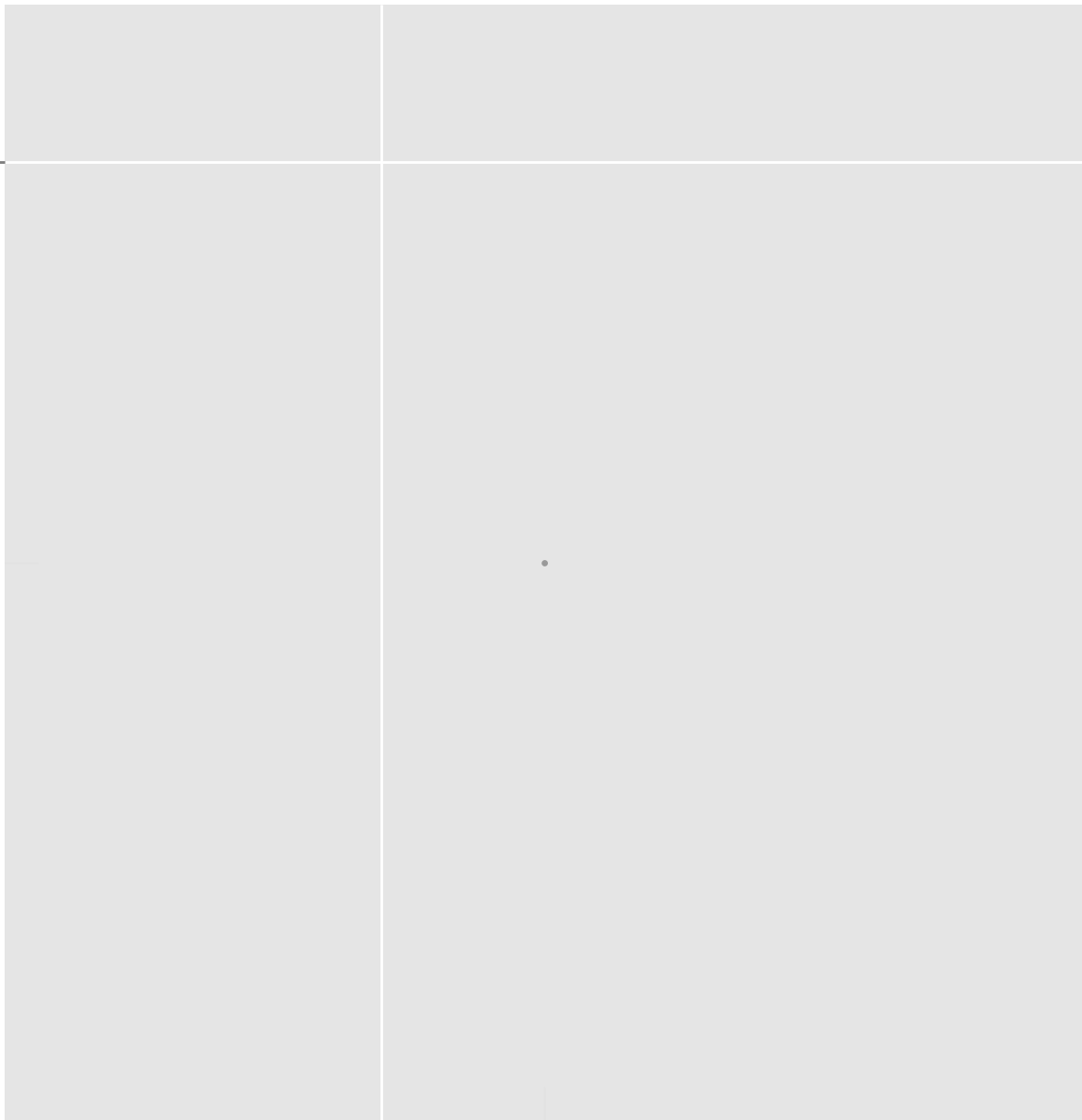
A\_dd\_untr FPKM + 0

1000

10

A\_wt\_untr FPKM + 0

.



A\_wt\_untr FPKM + 0

10

A\_dd\_untr FPKM + 0

1000

