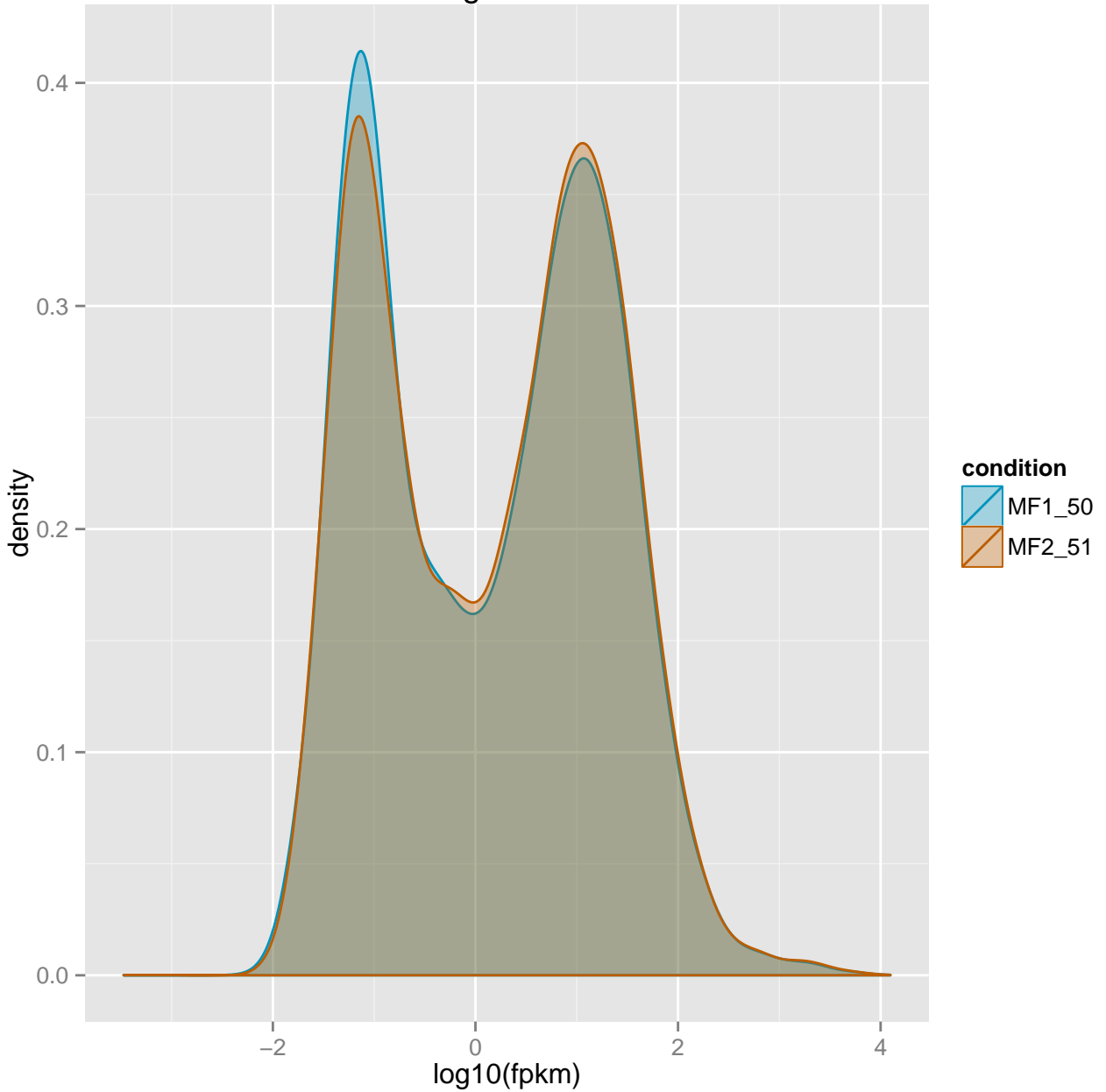
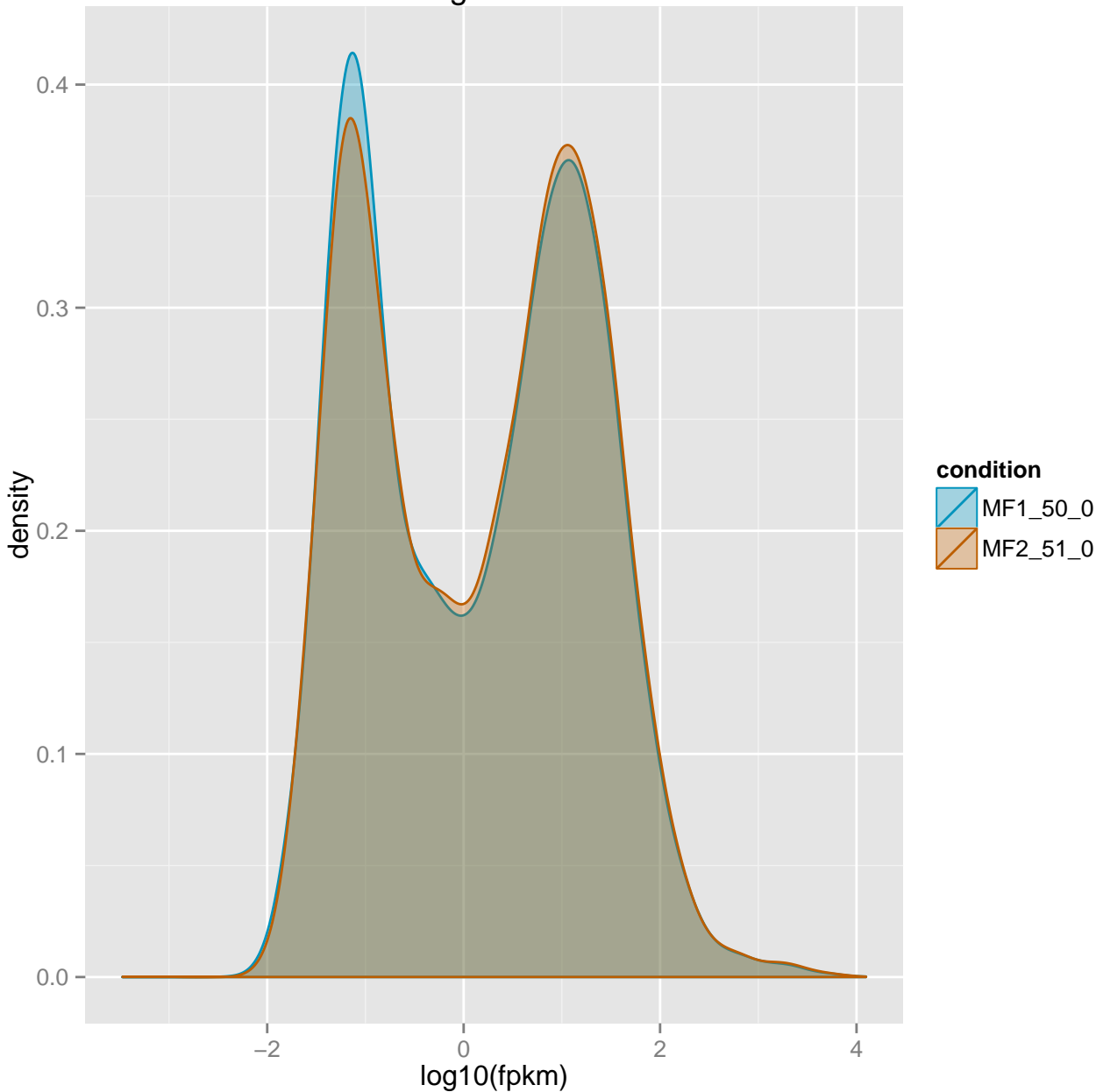
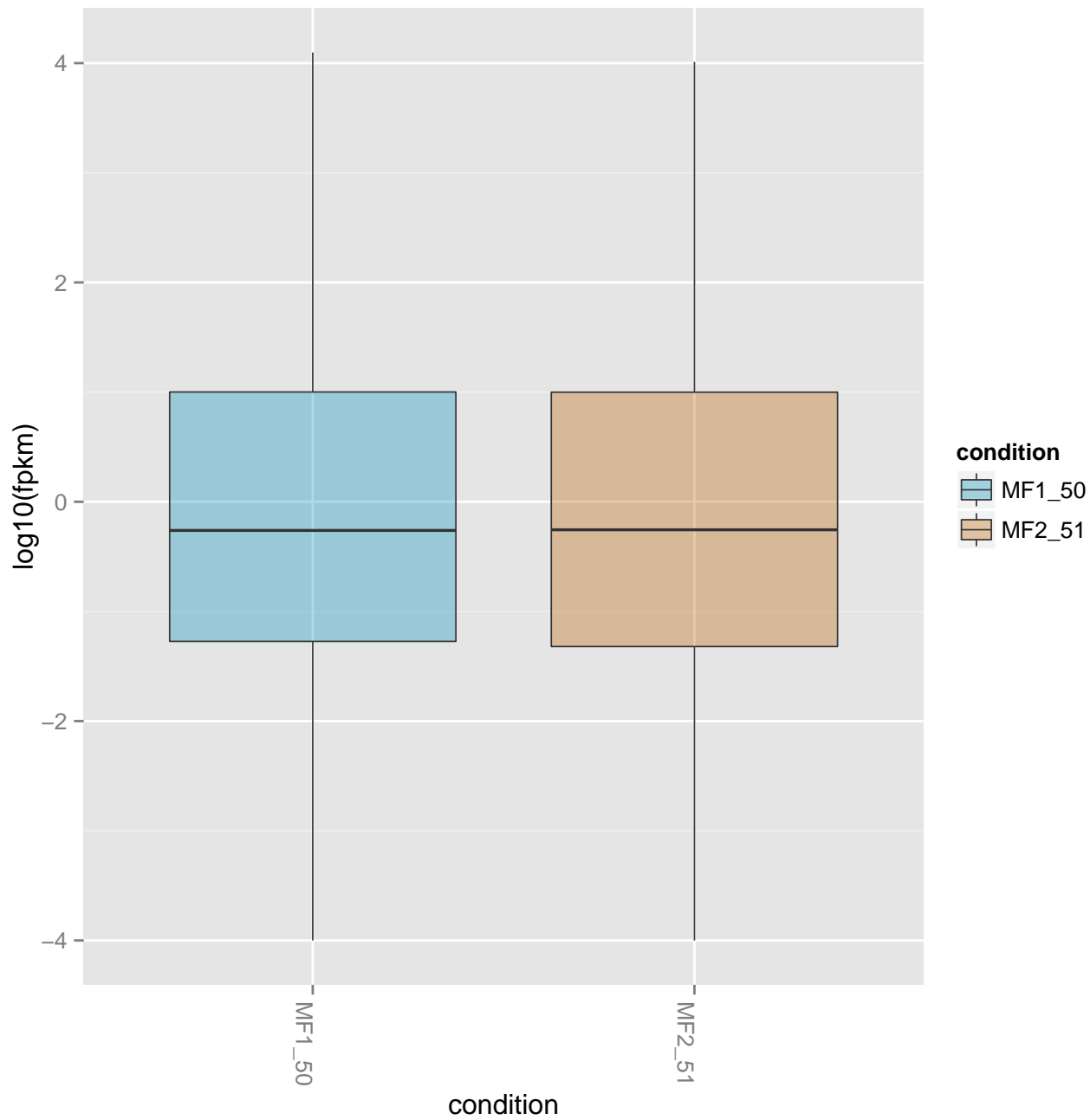


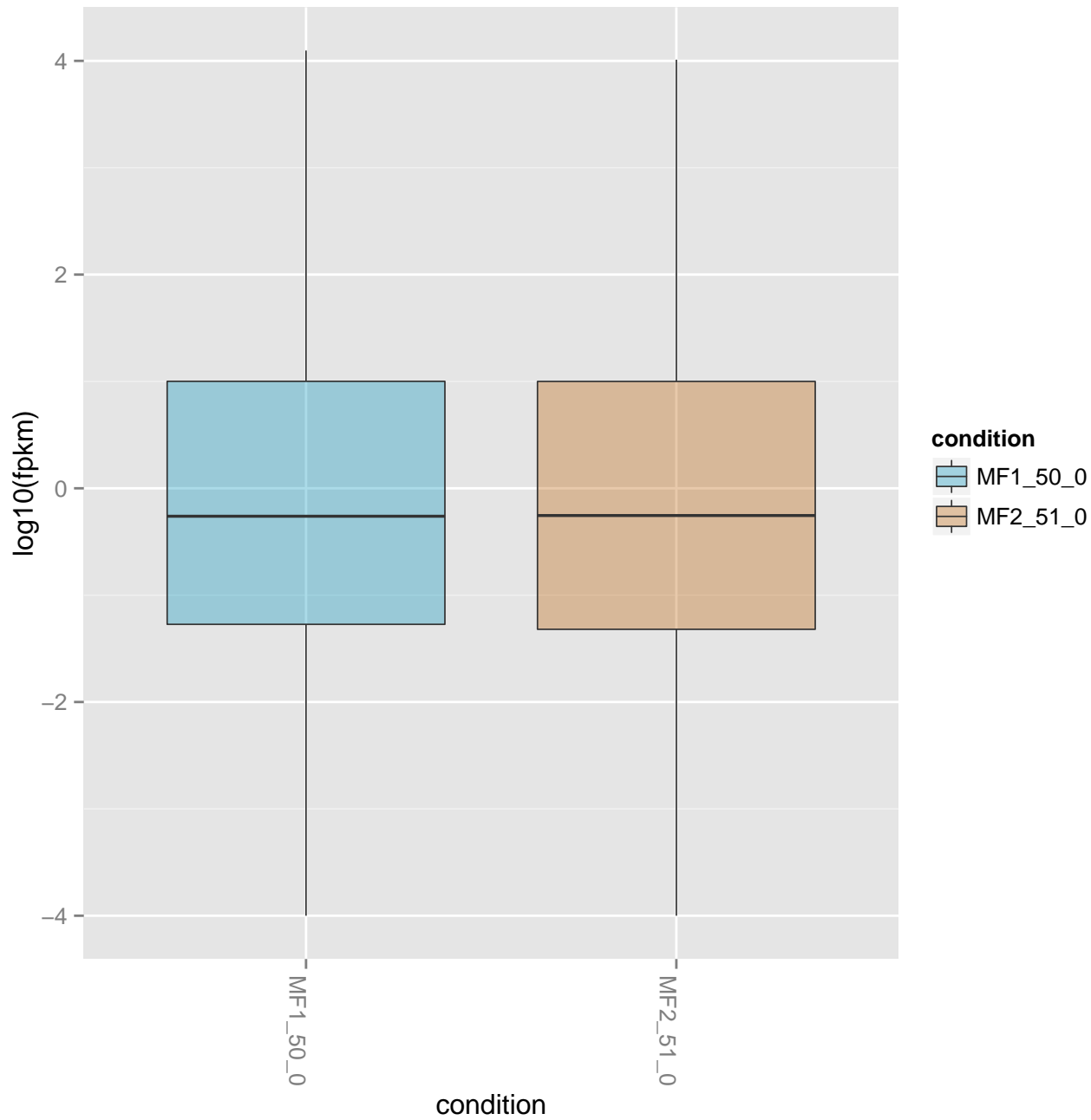
genes

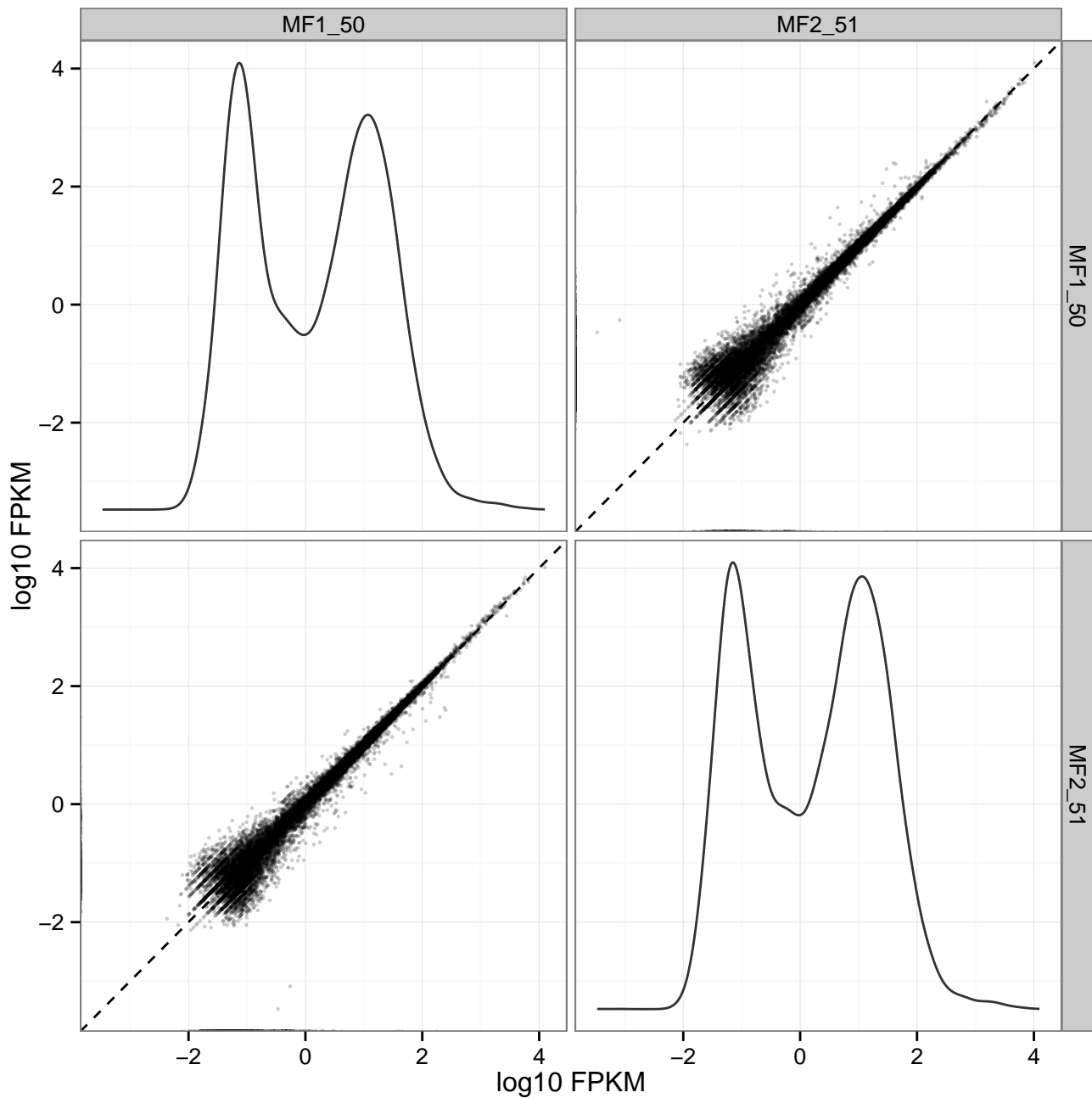


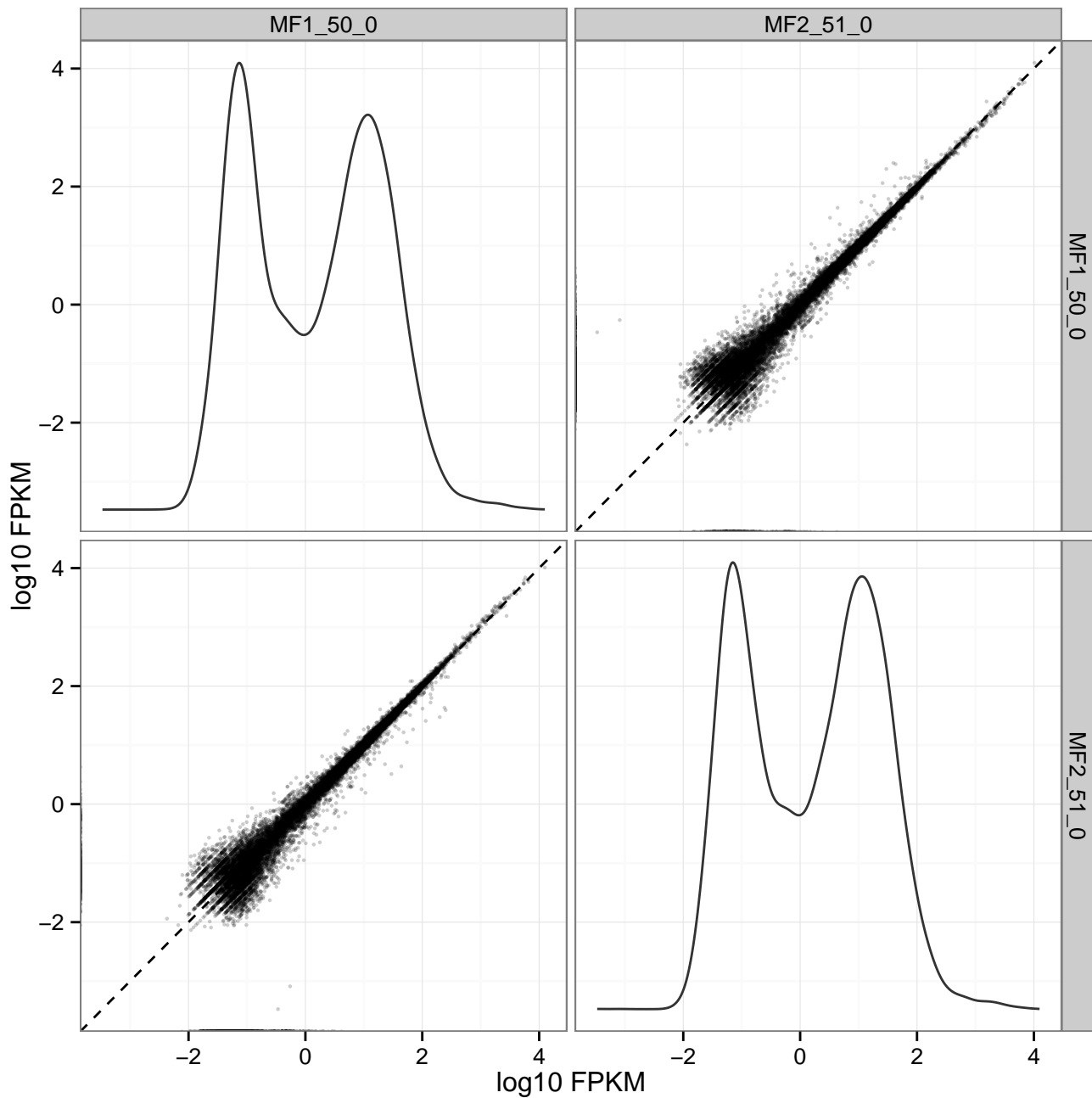
genes



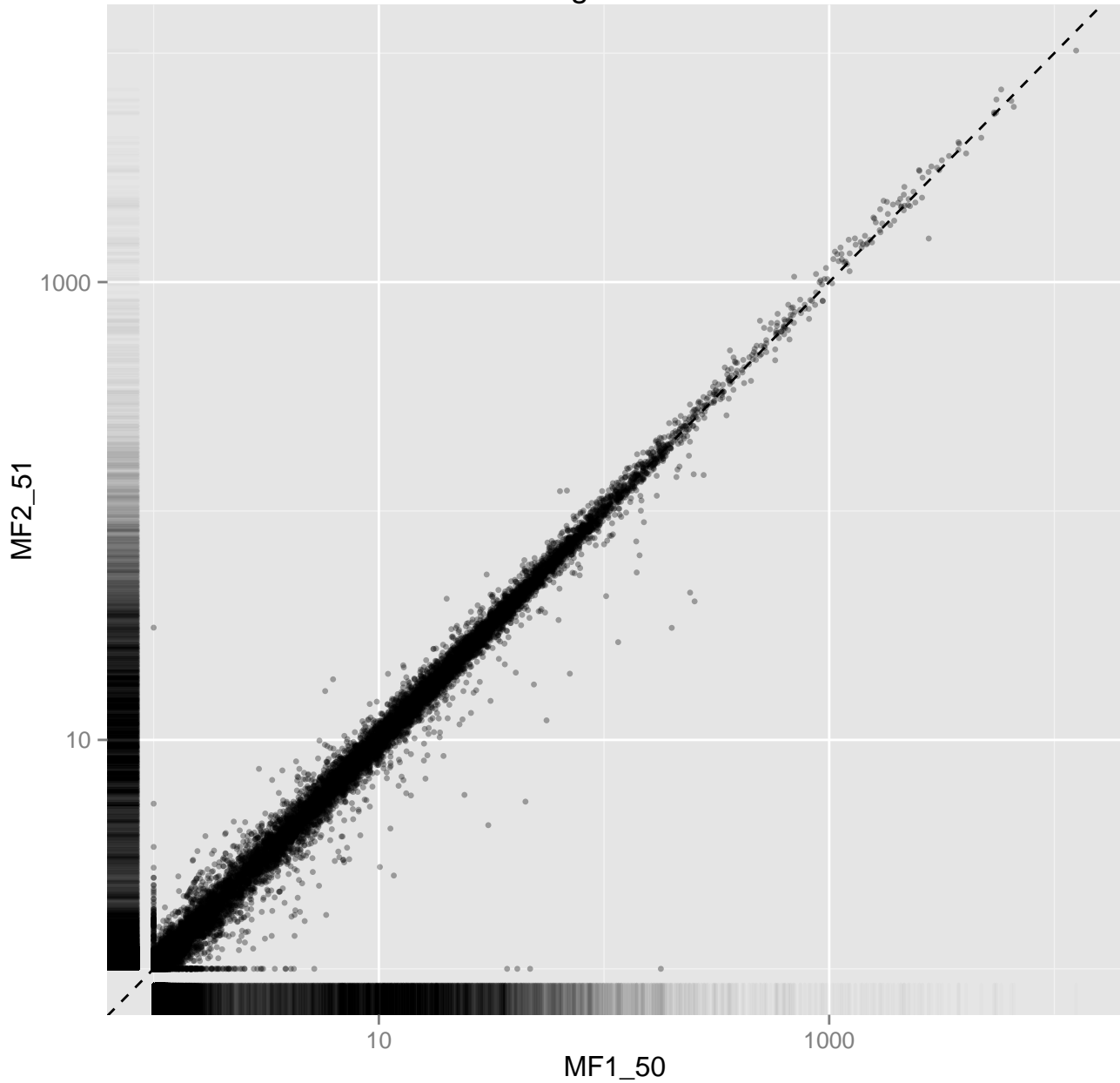








genes



MF1_50 MF2_51

$\log_2(M)$

10

5

0

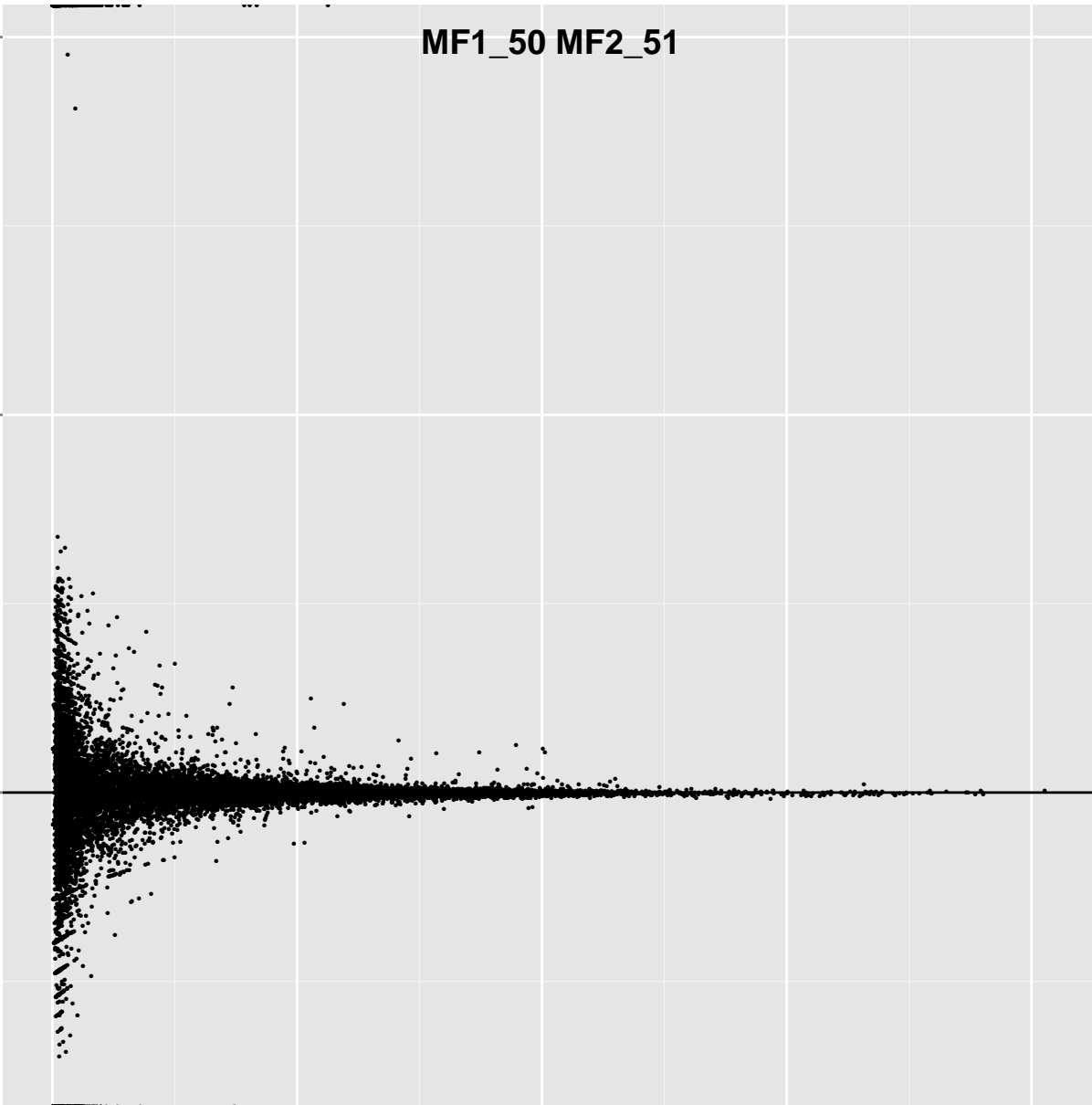
0

1

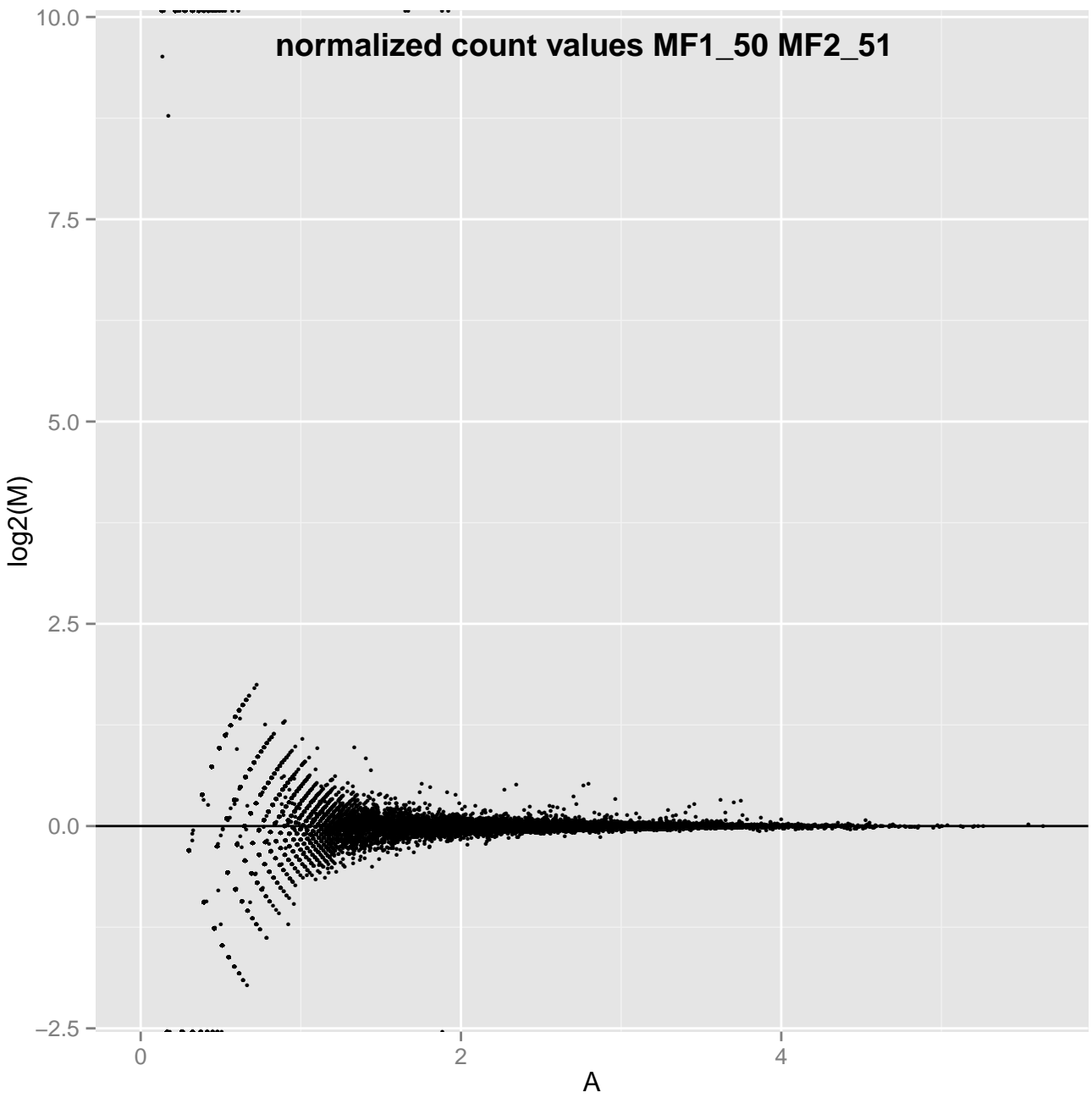
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3

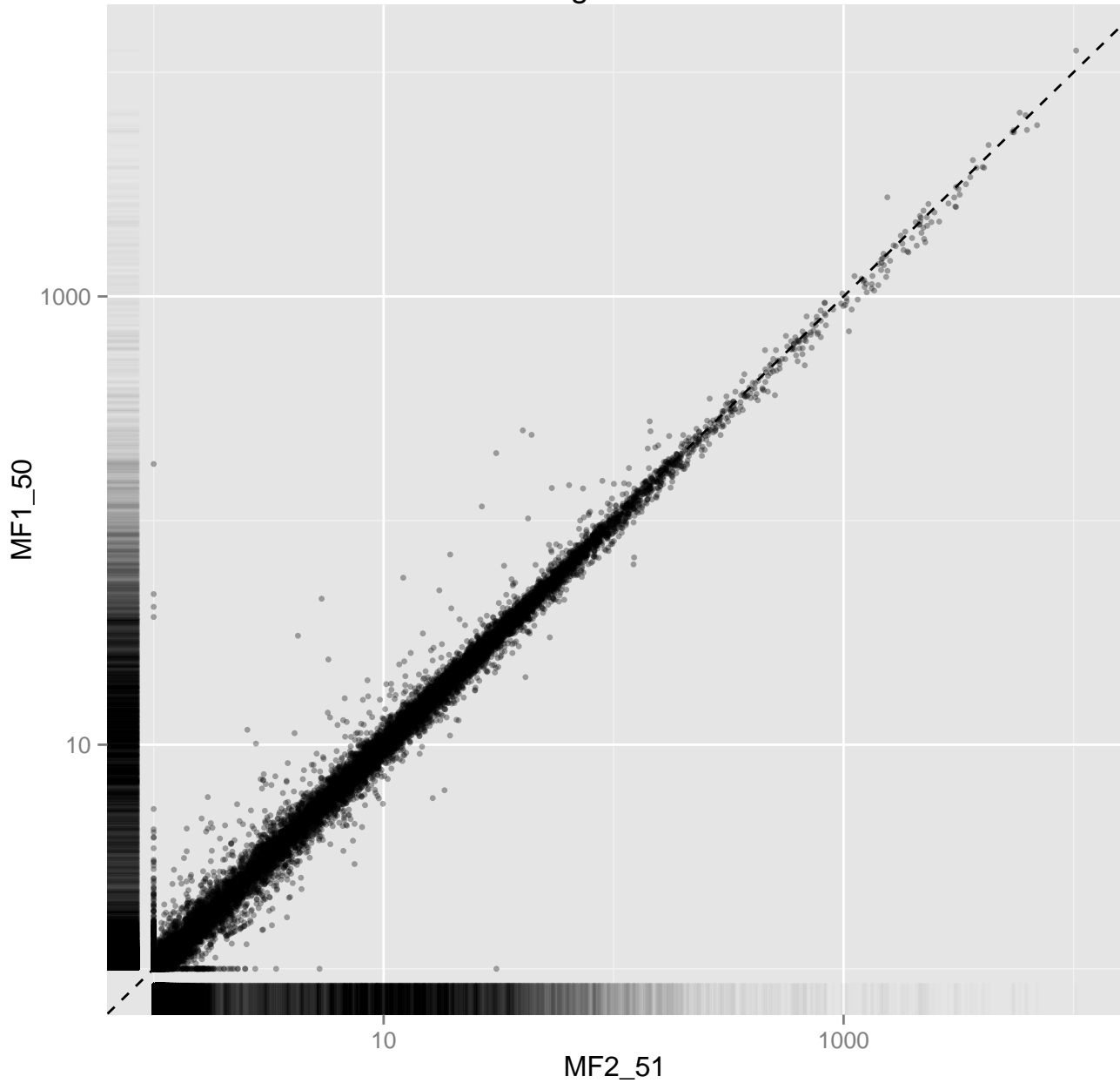
4



normalized count values MF1_50 MF2_51



genes



MF2_51 MF1_50

$\log_2(M)$

0

-5

-10

0

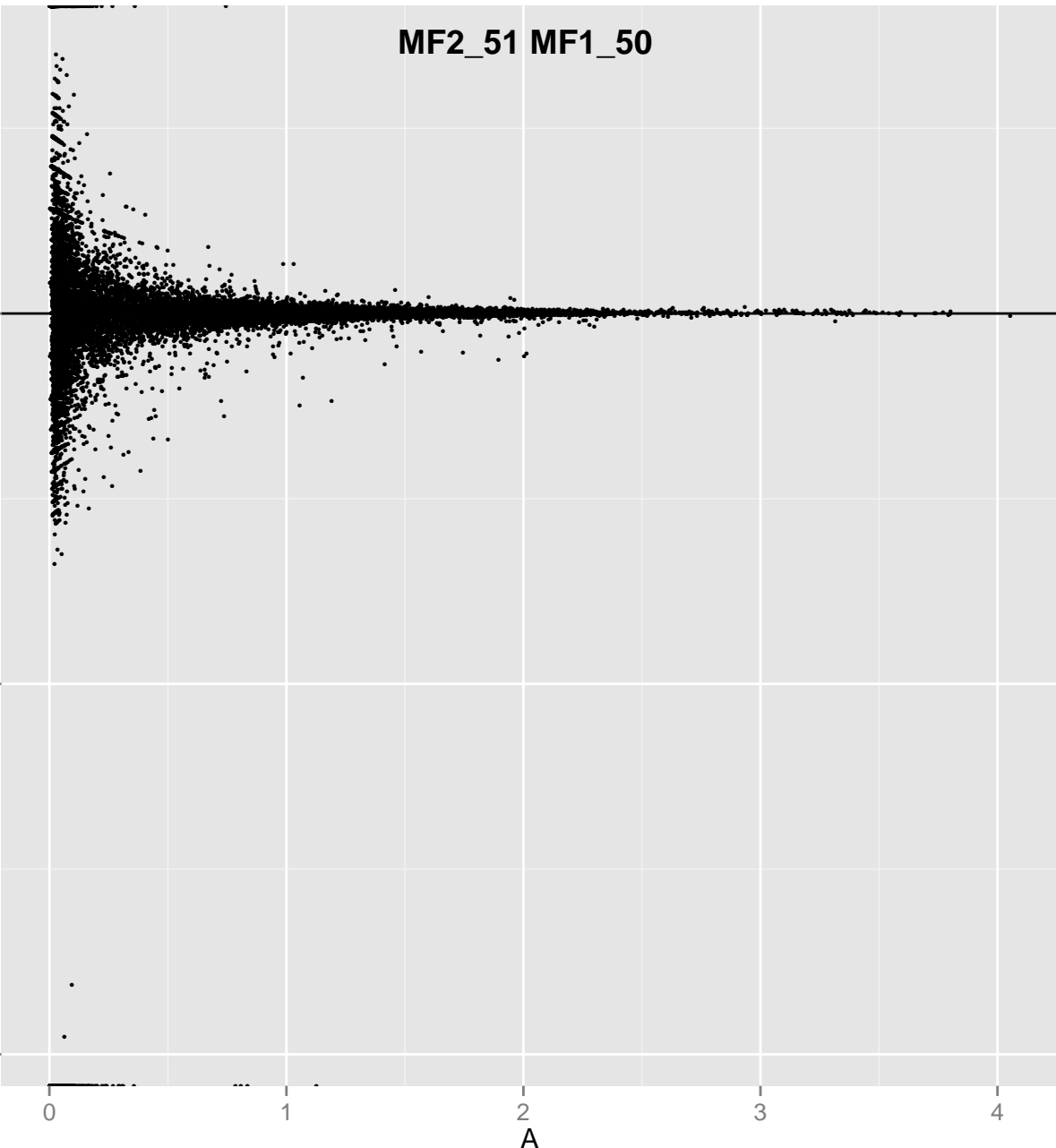
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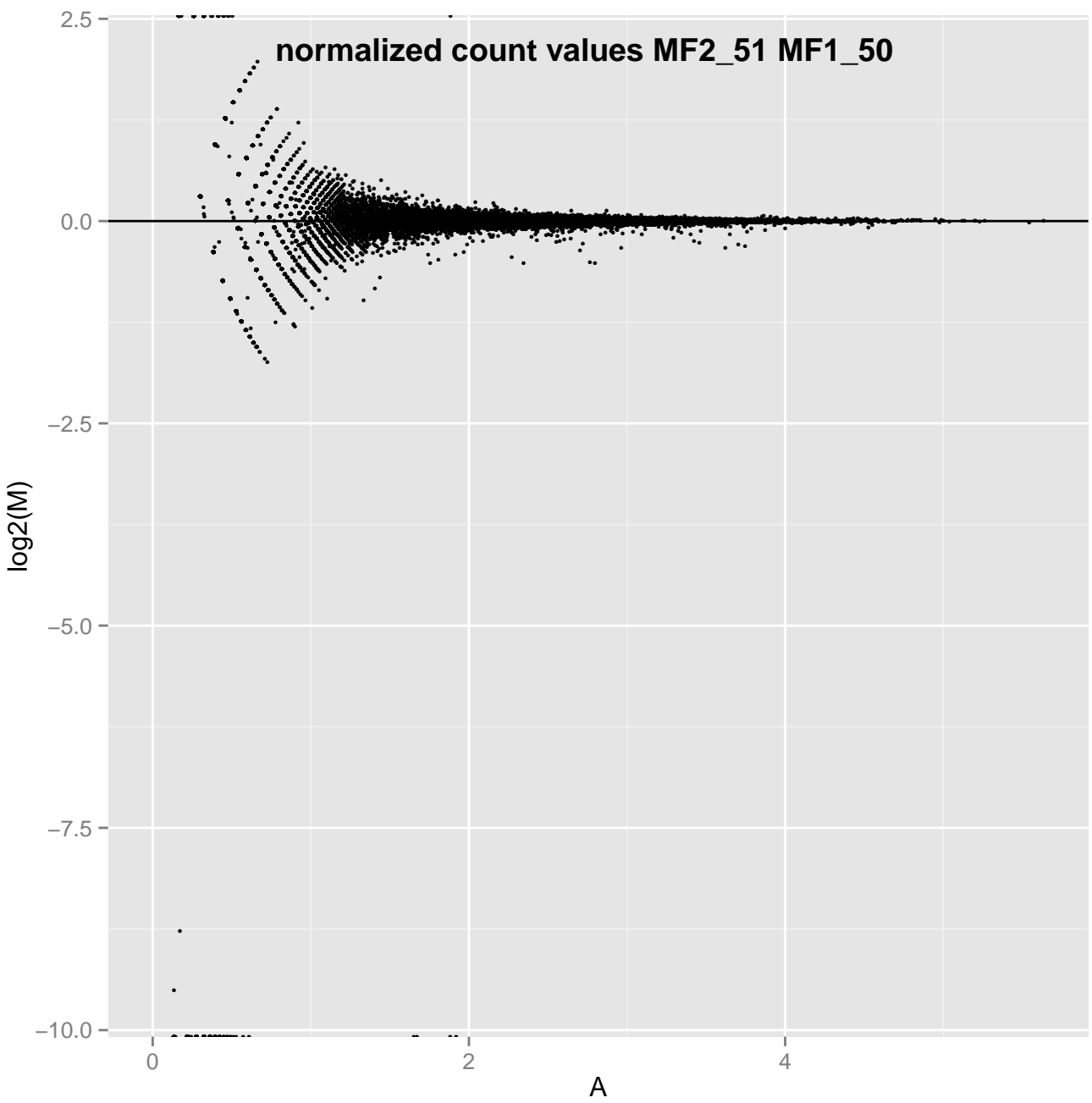
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2

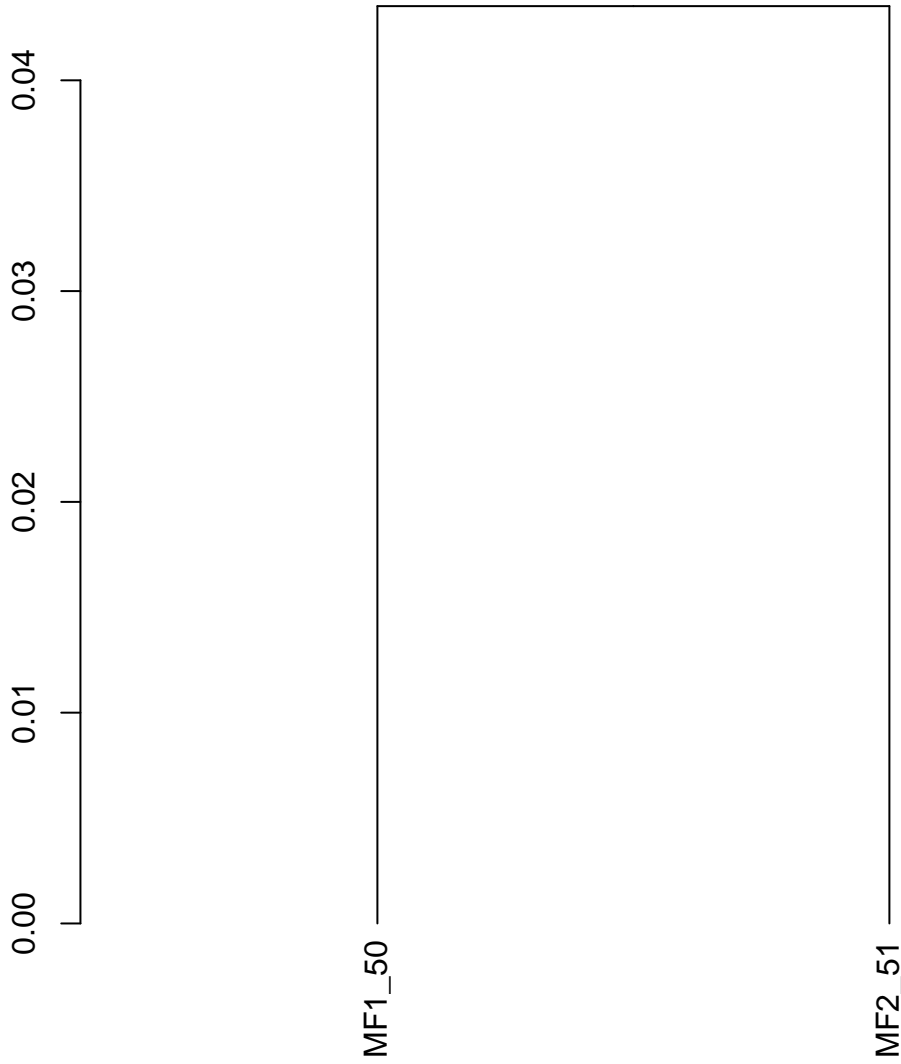
3

4





All genes(cuff)



All genes(cuff)

