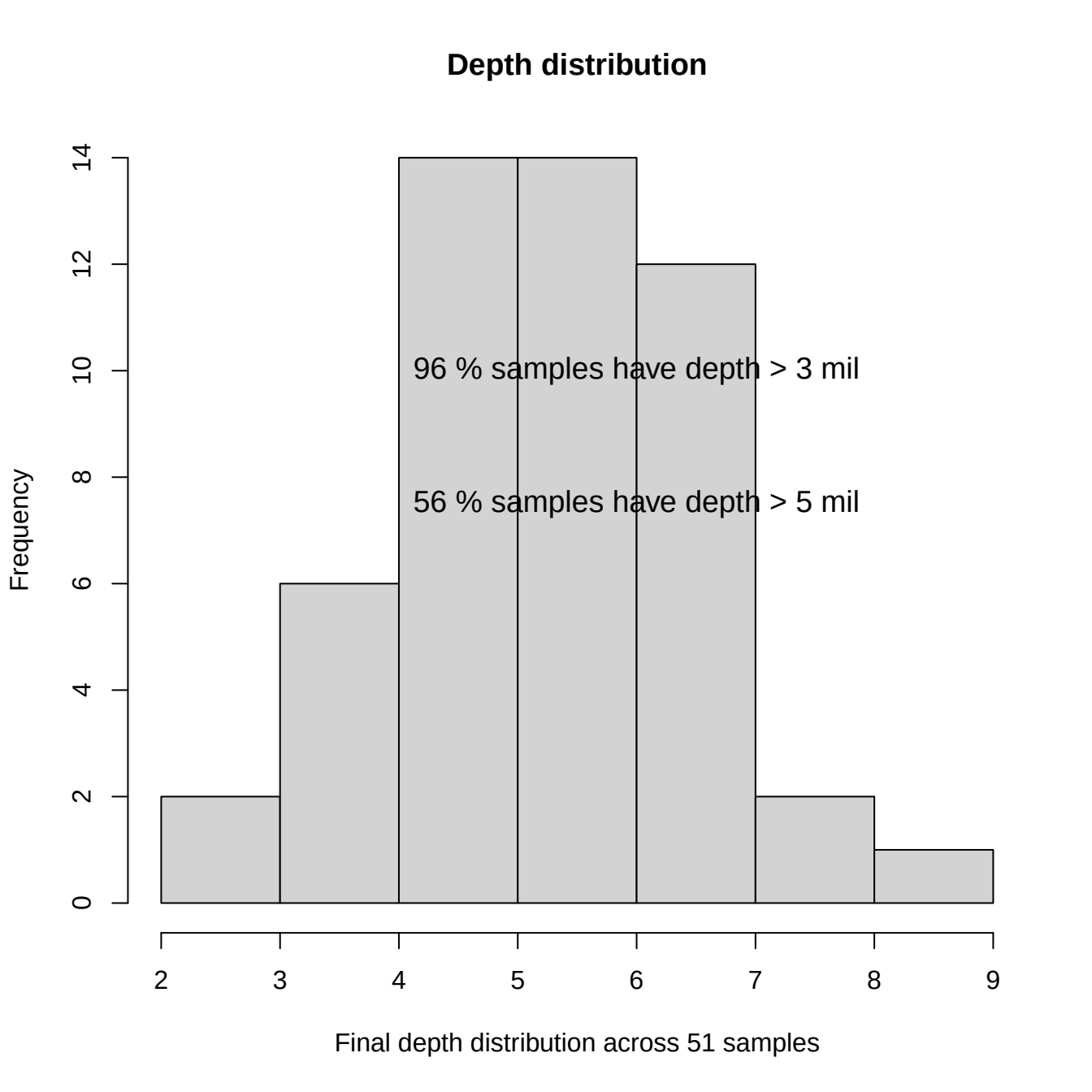
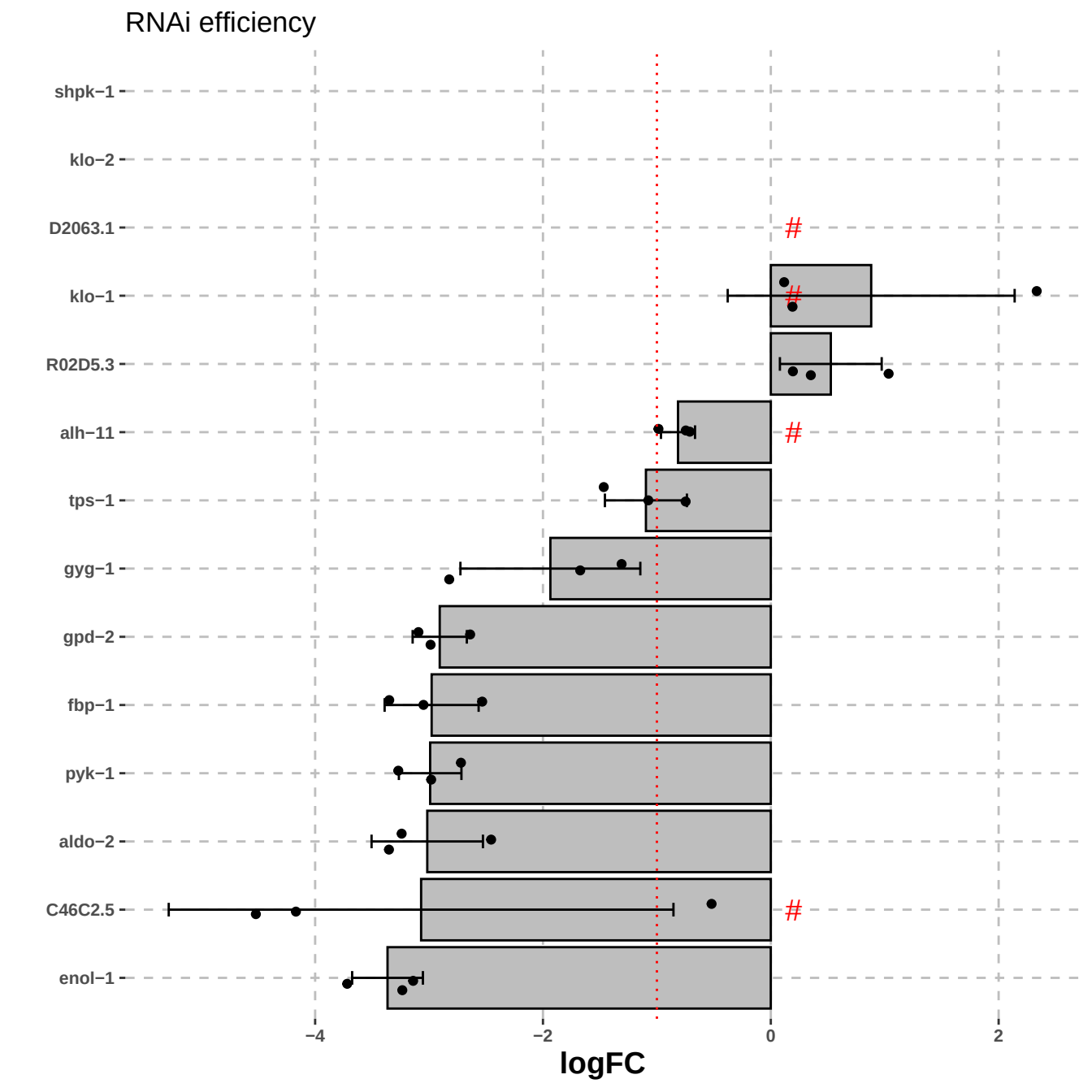
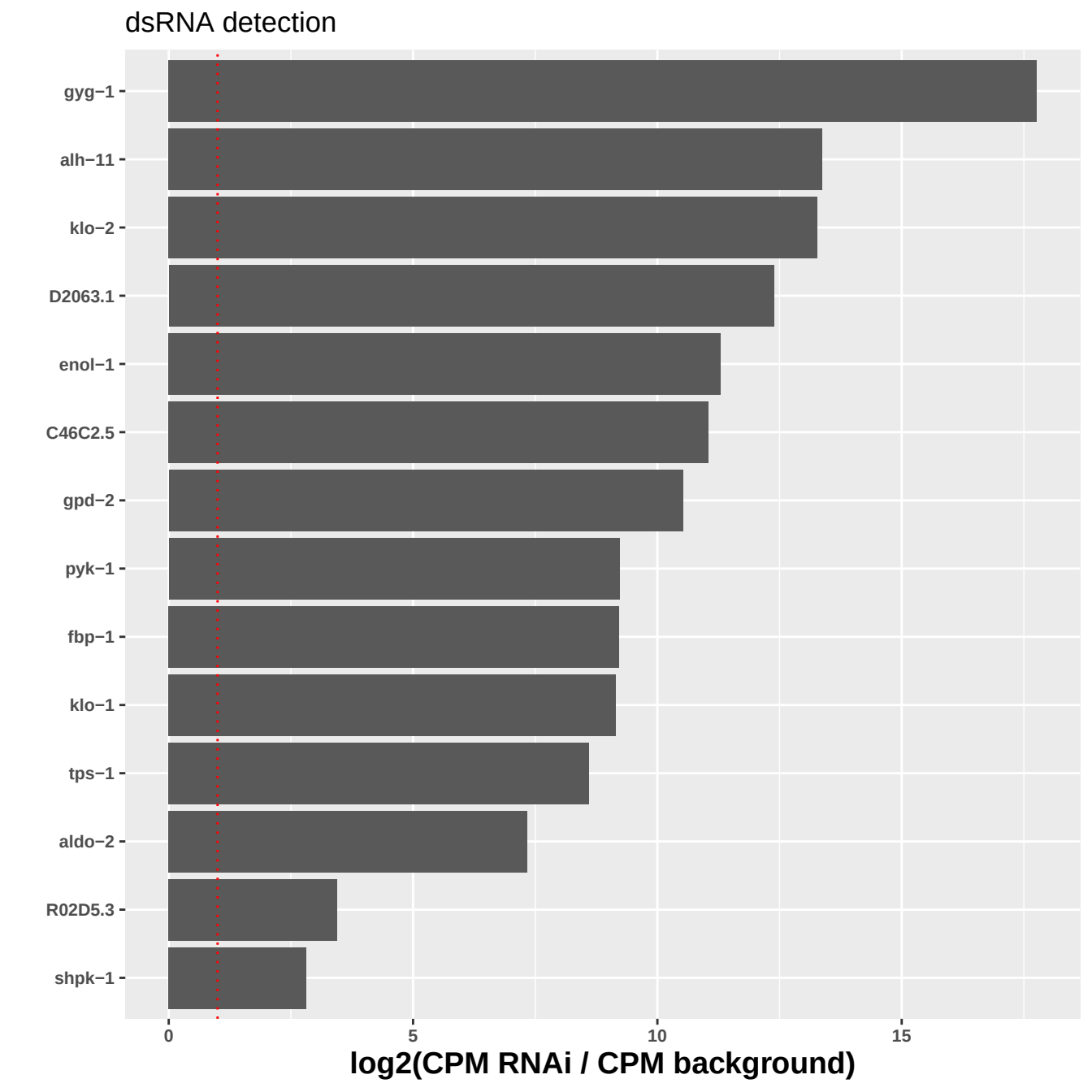
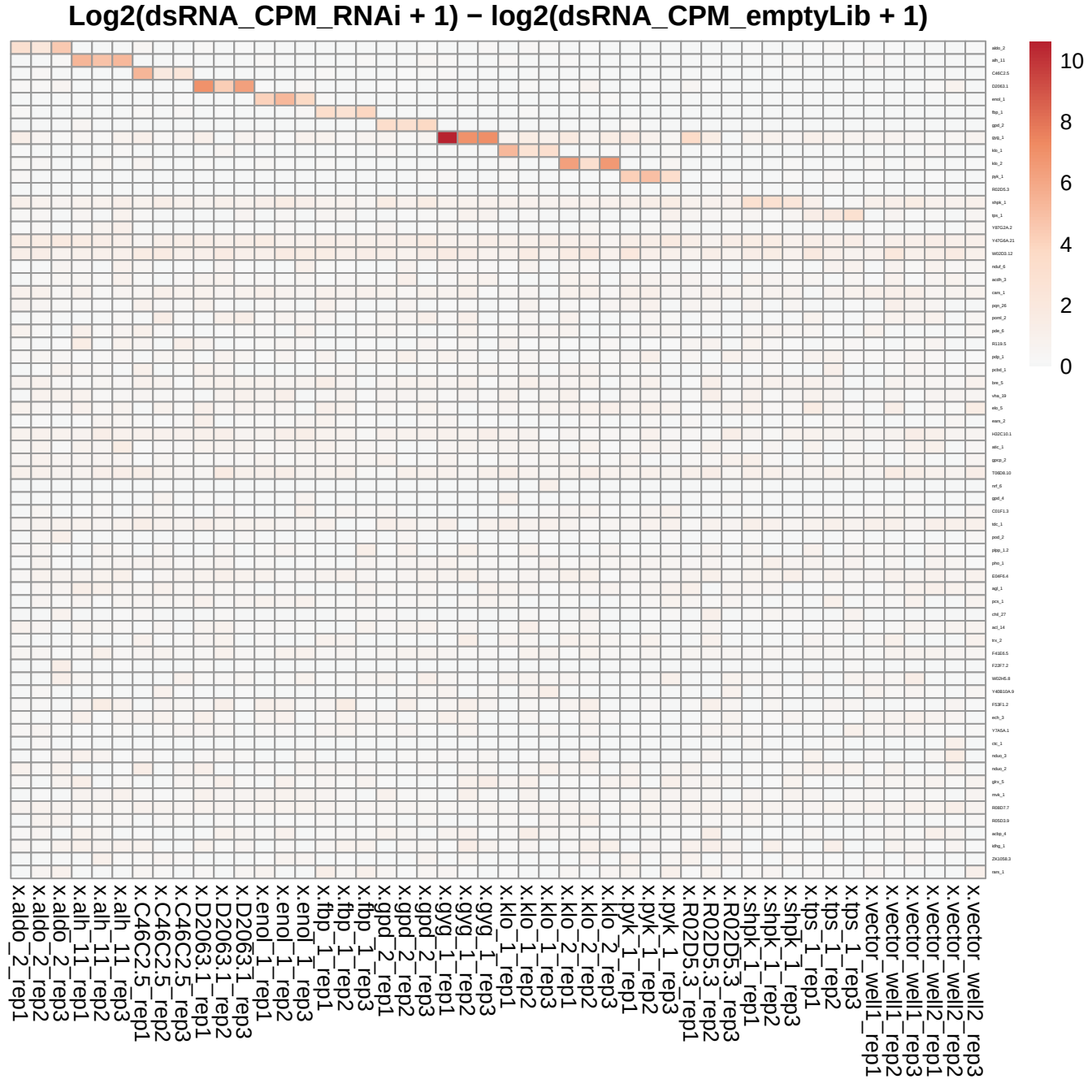

Depth distribution
Frequency
0
2
4
6
8
10
12
14
96 % samples have depth > 3 mil
56 % samples have depth > 5 mil
2
3
4
5
6
7
8
9
Final depth distribution across 51 samples


RNAi efficiency
shpk−1
klo−2
D2063.1
klo−1
R02D5.3
alh−11
tps−1
gyg−1
gpd−2
fbp−1
pyk−1
aldo−2
C46C2.5
enol−1
#
#
#
#
−4
−2
0
2
logFC


dsRNA detection
gyg−1
alh−11
klo−2
D2063.1
enol−1
C46C2.5
gpd−2
pyk−1
fbp−1
klo−1
tps−1
aldo−2
R02D5.3
shpk−1
0
5
10
15
log2(CPM RNAi / CPM background)


Log2(dsRNA_CPM_RNAi + 1) − log2(dsRNA_CPM_emptyLib + 1)
10
8
6
4
2
0
x.aldo_2_rep1
x.aldo_2_rep2
x.aldo_2_rep3
x.alh_11_rep1
x.alh_11_rep2
x.alh_11_rep3
x.C46C2.5_rep1
x.C46C2.5_rep2
x.C46C2.5_rep3
x.D2063.1_rep1
x.D2063.1_rep2
x.D2063.1_rep3
x.enol_1_rep1
x.enol_1_rep2
x.enol_1_rep3
x.fbp_1_rep1
x.fbp_1_rep2
x.fbp_1_rep3
x.gpd_2_rep1
x.gpd_2_rep2
x.gpd_2_rep3
x.gyg_1_rep1
x.gyg_1_rep2
x.gyg_1_rep3
x.klo_1_rep1
x.klo_1_rep2
x.klo_1_rep3
x.klo_2_rep1
x.klo_2_rep2
x.klo_2_rep3
x.pyk_1_rep1
x.pyk_1_rep2
x.pyk_1_rep3
x.R02D5.3_rep1
x.R02D5.3_rep2
x.R02D5.3_rep3
x.shpk_1_rep1
x.shpk_1_rep2
x.shpk_1_rep3
x.tps_1_rep1
x.tps_1_rep2
x.tps_1_rep3
x.vector_well1_rep1
x.vector_well1_rep2
x.vector_well1_rep3
x.vector_well2_rep1
x.vector_well2_rep2
x.vector_well2_rep3