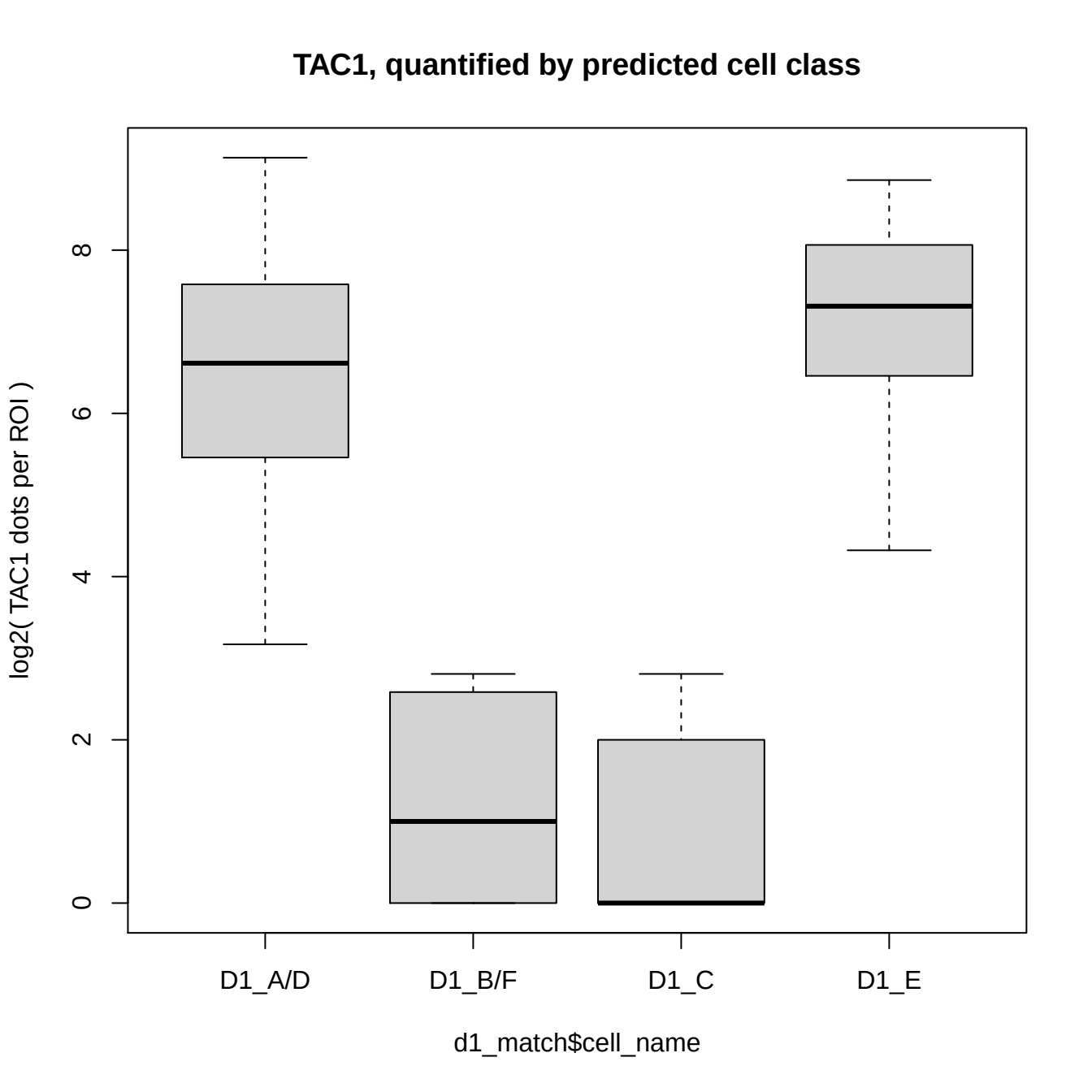
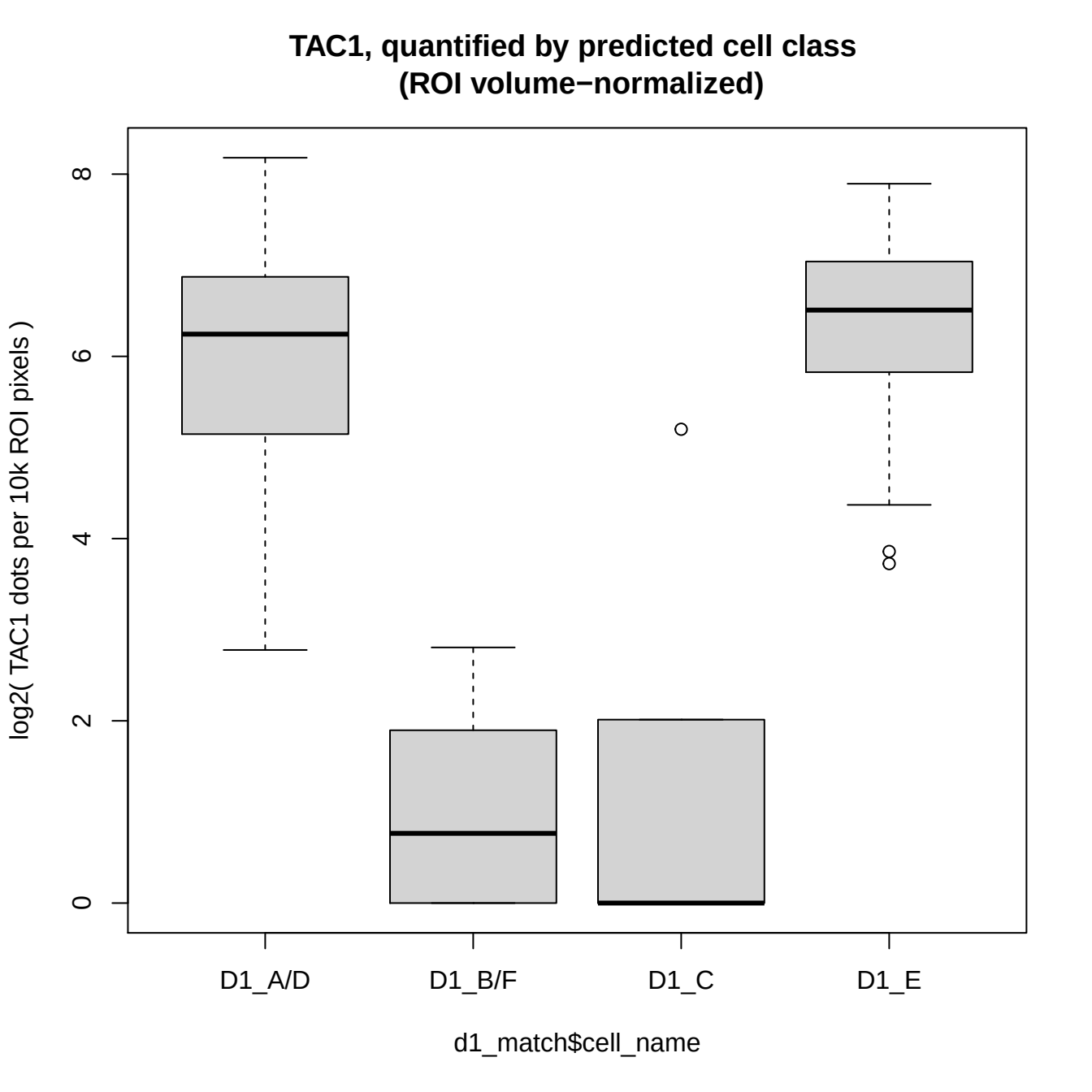
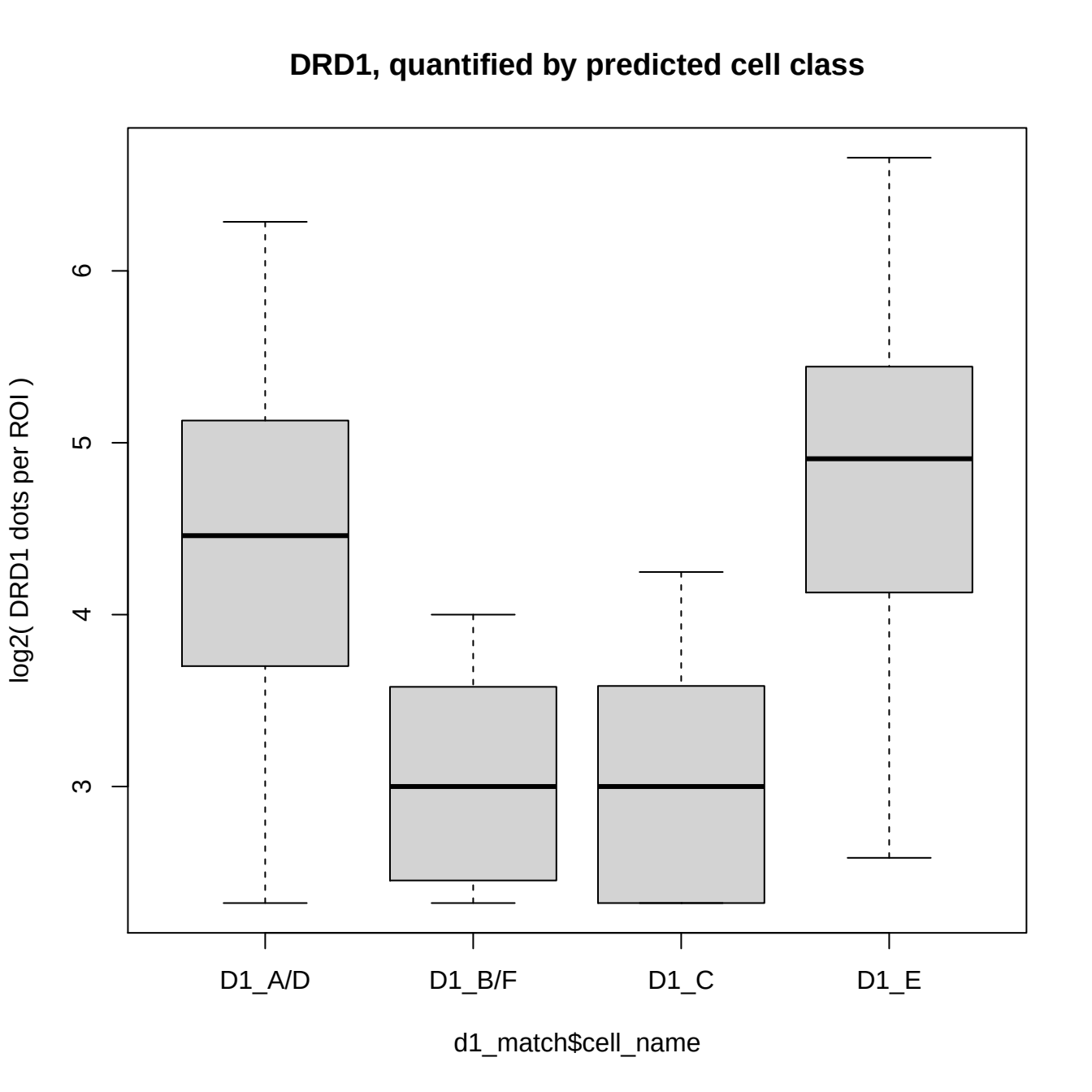
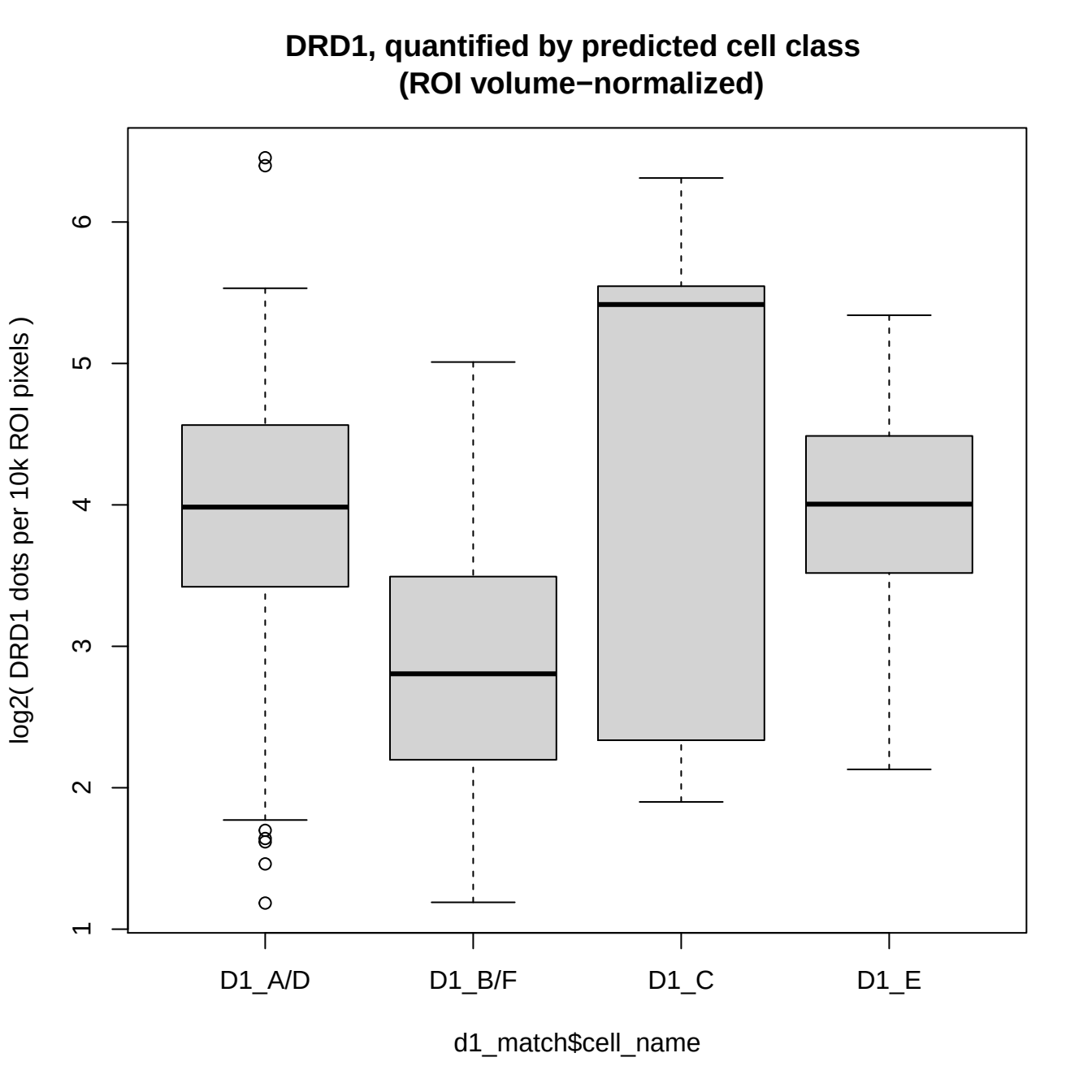
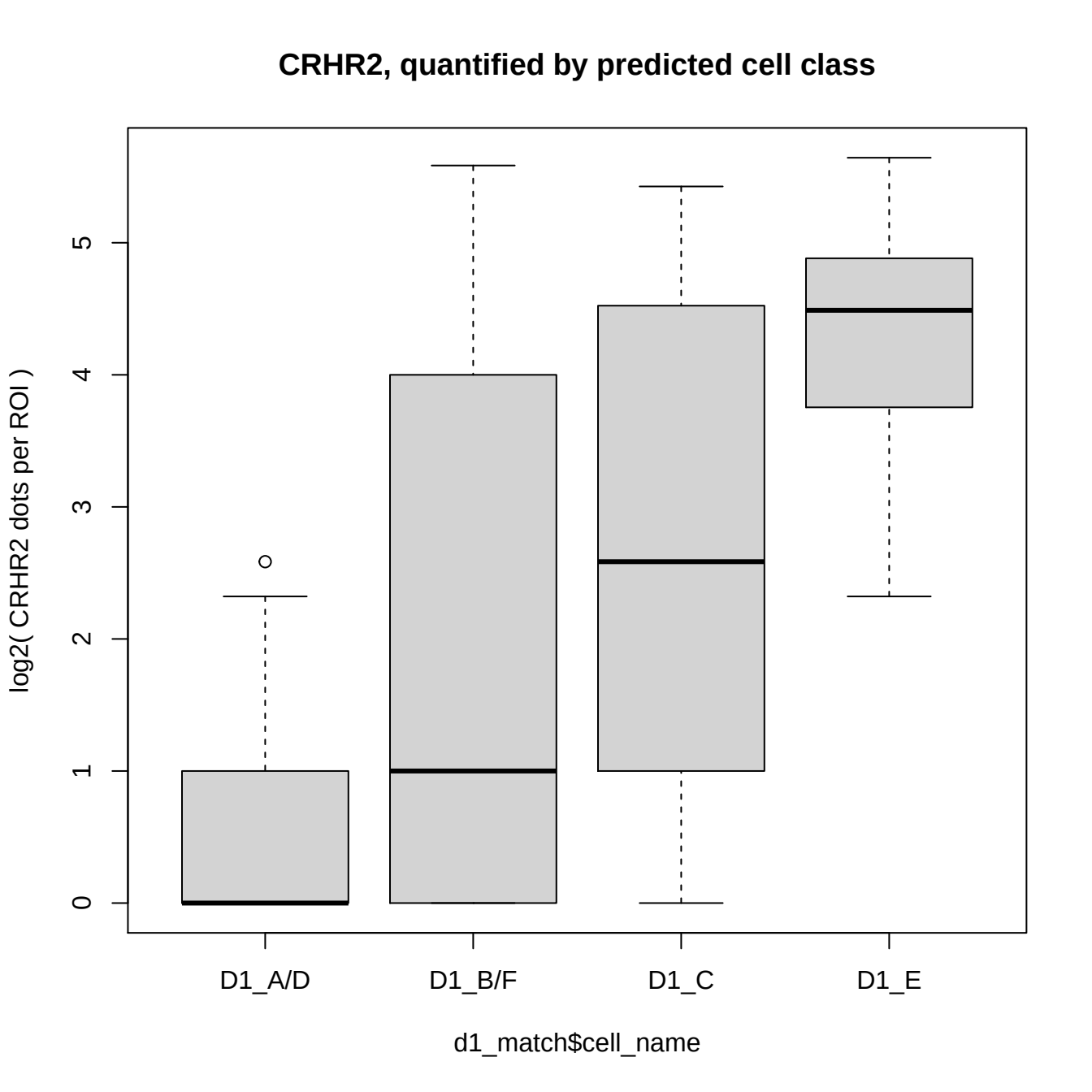
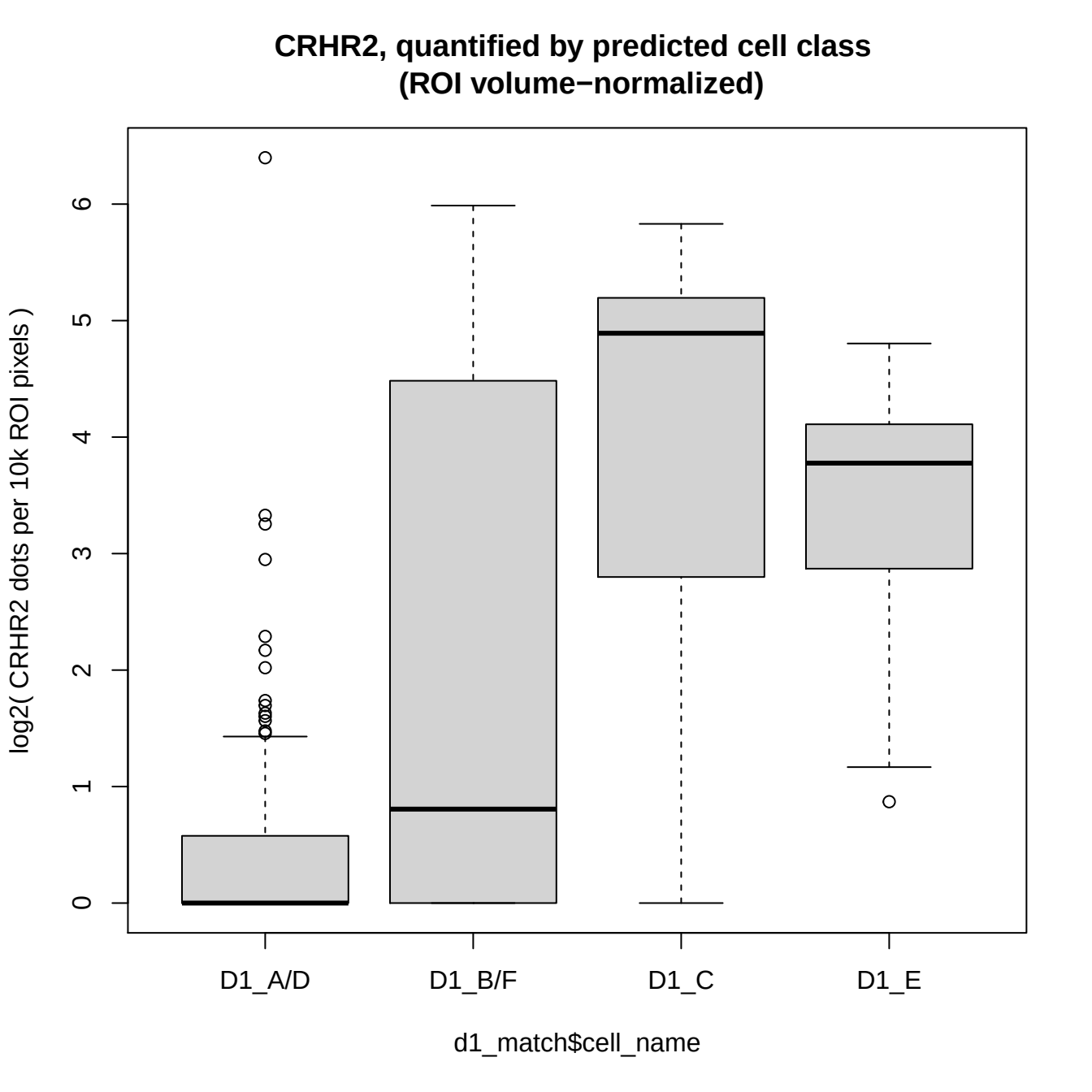

TAC1, quantified by predicted cell class
log2(TAC1 dots per ROI)
0
2
4
6
8
D1_A/D
D1_B/F
D1_C
D1_E
d1_match$cell_name

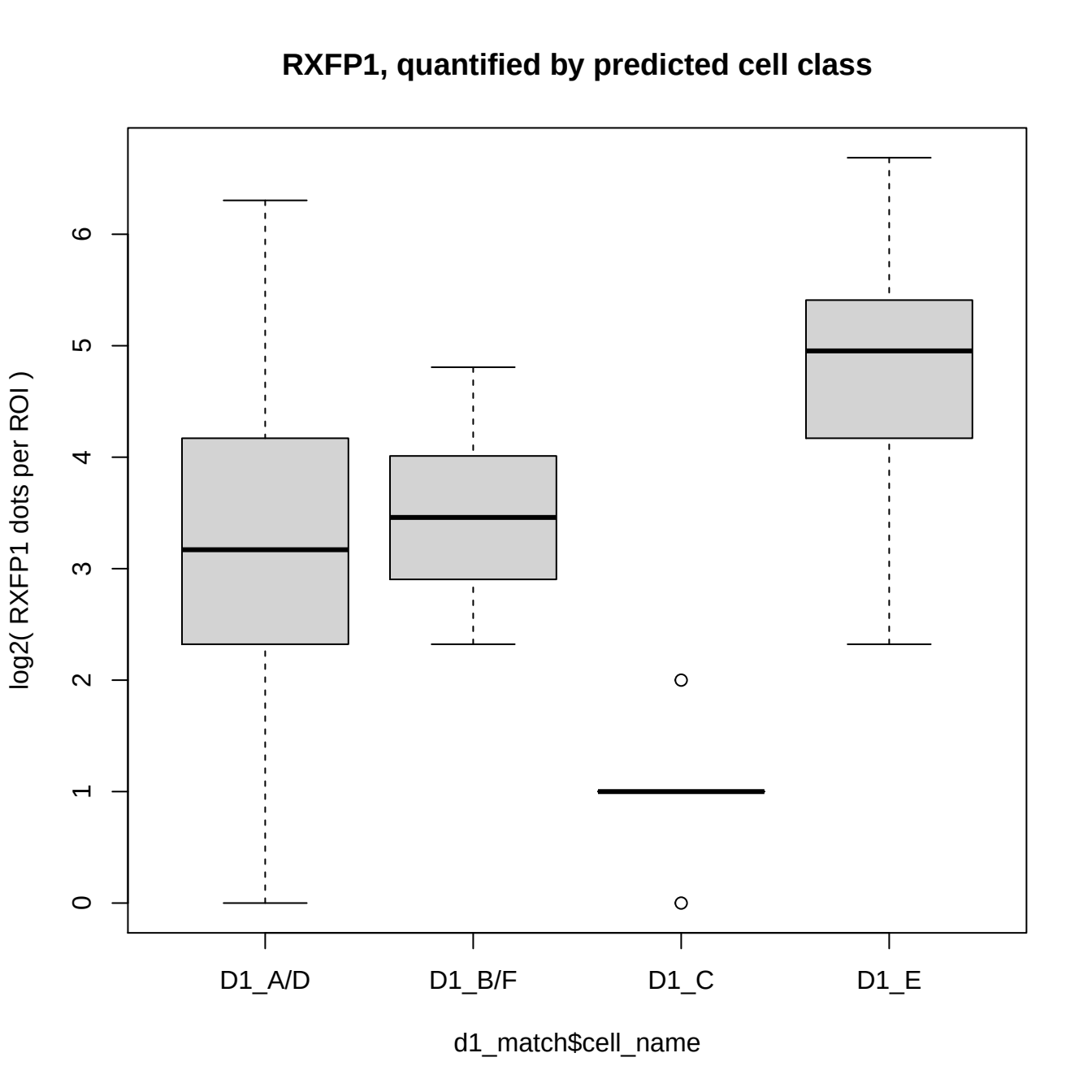

TAC1, quantified by predicted cell class
(ROI volume−normalized)
log2(TAC1 dots per 10k ROI pixels)
0
2
4
6
8
D1_A/D
D1_B/F
D1_C
D1_E
d1_match$cell_name

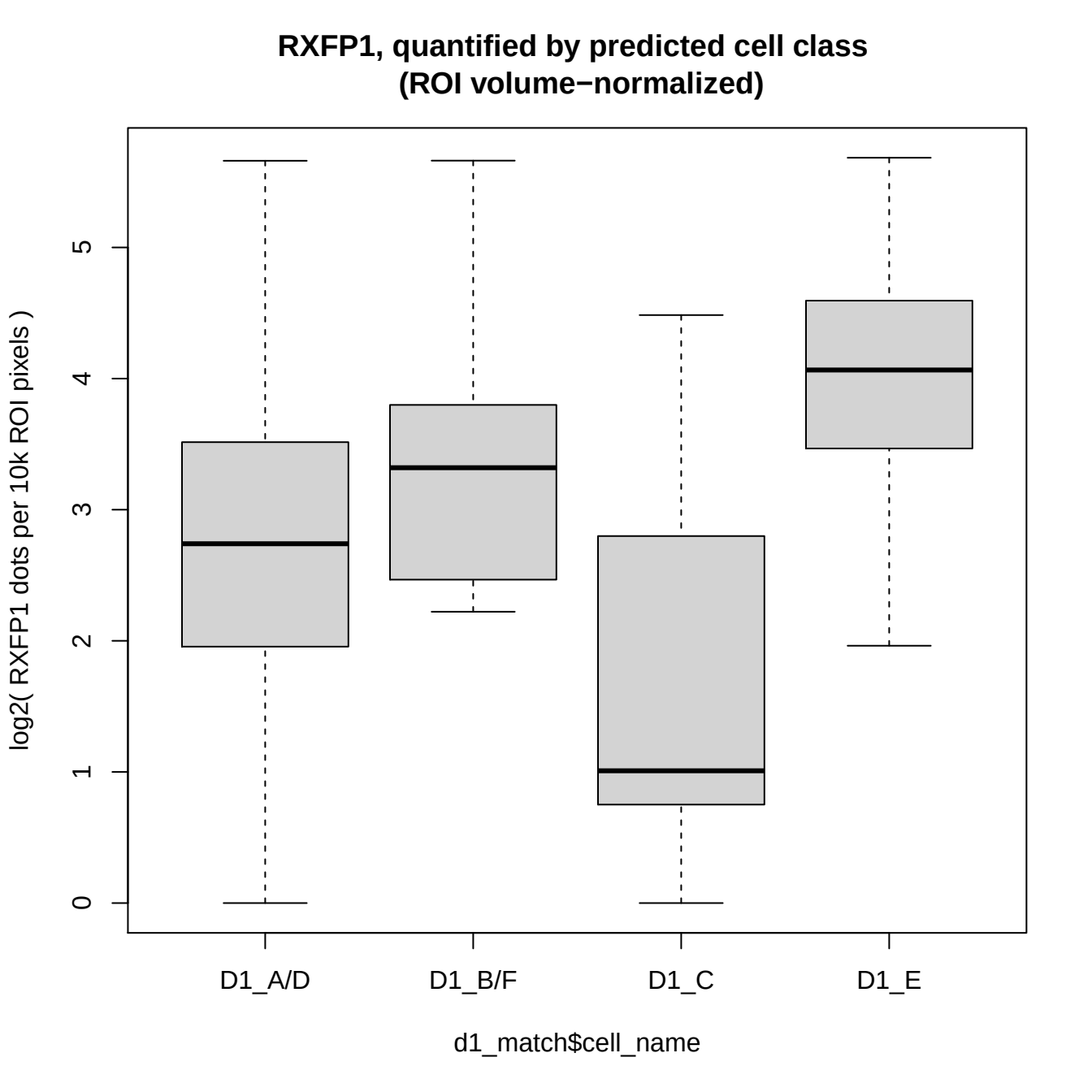

DRD1, quantified by predicted cell class
log2(DRD1 dots per ROI)
3
4
5
6
D1_A/D
D1_B/F
D1_C
D1_E
d1_match$cell_name


DRD1, quantified by predicted cell class
(ROI volume−normalized)
log2(DRD1 dots per 10k ROI pixels)
1
2
3
4
5
6
D1_A/D
D1_B/F
D1_C
D1_E
d1_match$cell_name


CRHR2, quantified by predicted cell class
log2(CRHR2 dots per ROI)
0
1
2
3
4
5
D1_A/D
D1_B/F
D1_C
D1_E
d1_match$cell_name


CRHR2, quantified by predicted cell class
(ROI volume−normalized)
log2(CRHR2 dots per 10k ROI pixels)
0
1
2
3
4
5
6
D1_A/D
D1_B/F
D1_C
D1_E
d1_match$cell_name


RXFP1, quantified by predicted cell class
log2(RXFP1 dots per ROI)
0
1
2
3
4
5
6
D1_A/D
D1_B/F
D1_C
D1_E
d1_match$cell_name


RXFP1, quantified by predicted cell class
(ROI volume−normalized)
log2(RXFP1 dots per 10k ROI pixels)
0
1
2
3
4
5
D1_A/D
D1_B/F
D1_C
D1_E
d1_match$cell_name