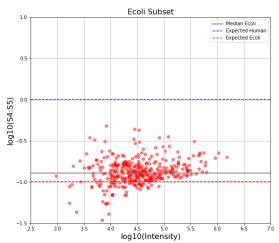
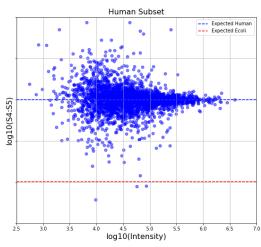
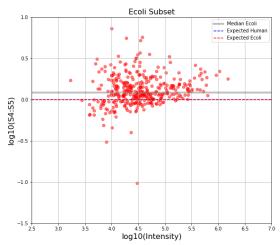
```
# workings for human abundance in each sample
human_light_s5 = 66.8
human_light_s4 = 66.8
human_heavy = 60
human_total_s5 = human_light_s5 + human_heavy
human_total_s4 = human_light_s4 + human_heavy
# workings for ecoli abundance in each sample
ecoli_light_s5 = 66.8
ecoli_light_s4 = 6.8
ecoli_heavy = 6
ecoli_total_s5 = ecoli_light_s5 + ecoli_heavy
ecoli_total_s4 = ecoli_light_s4 + ecoli_heavy
# workings for expected ratios in each channel (human)
expected_human_light = human_light_s4/human_light_s5
expected_human_heavy = human_heavy/human_heavy
expected_human_total = human_total_s4/human_total_s5
# workings for expected ratios in each channel (ecoli)
expected_ecoli_light = ecoli_light_s4/ecoli_light_s5
expected_ecoli_heavy = ecoli_heavy/ecoli_heavy
expected_ecoli_total = ecoli_total_s4/ecoli_total_s5
```

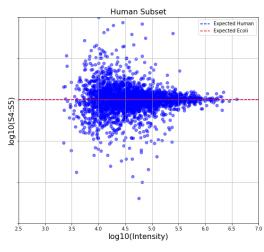
Light (no norm)



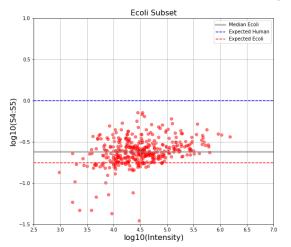


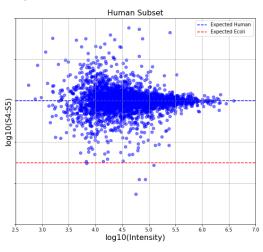
Heavy (no norm)



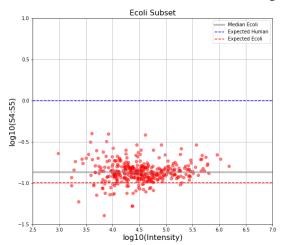


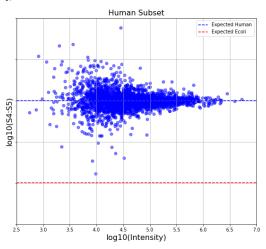
Total (no norm)



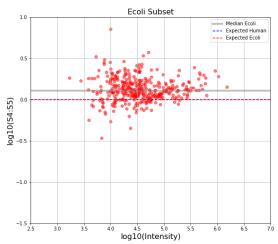


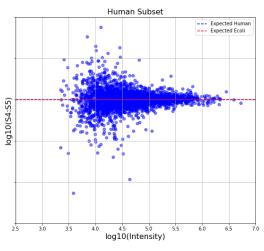
Light (Ifq)



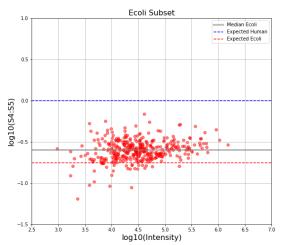


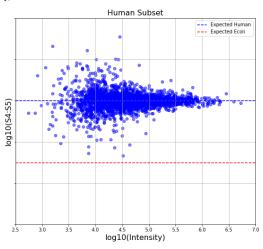
Heavy (Ifq)



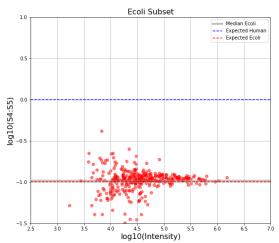


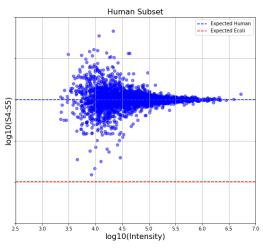
Total (Ifq)



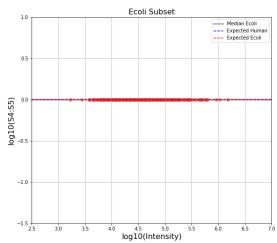


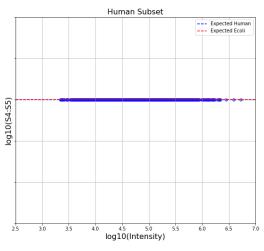
Light (href)





Heavy (href)





Total (href)

