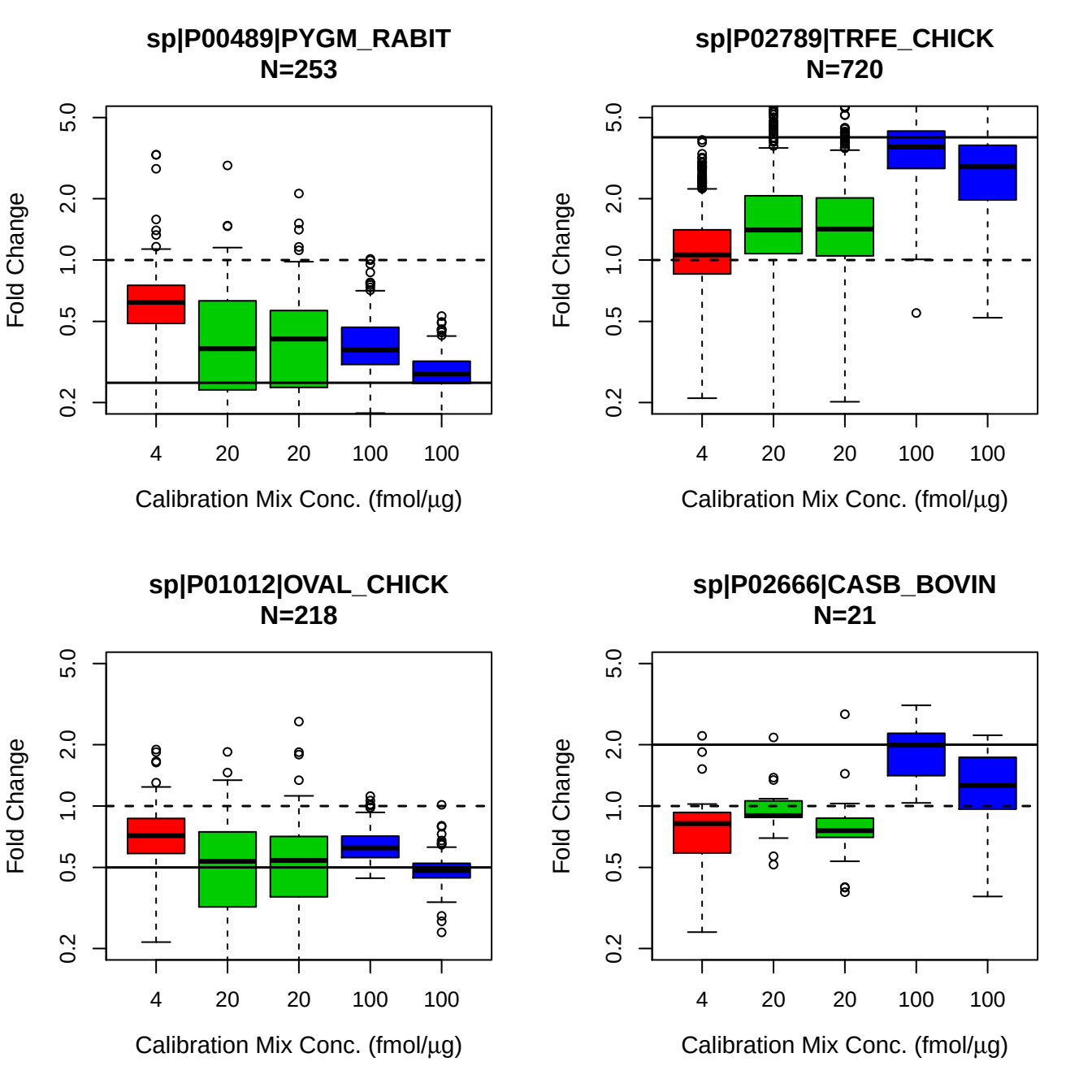
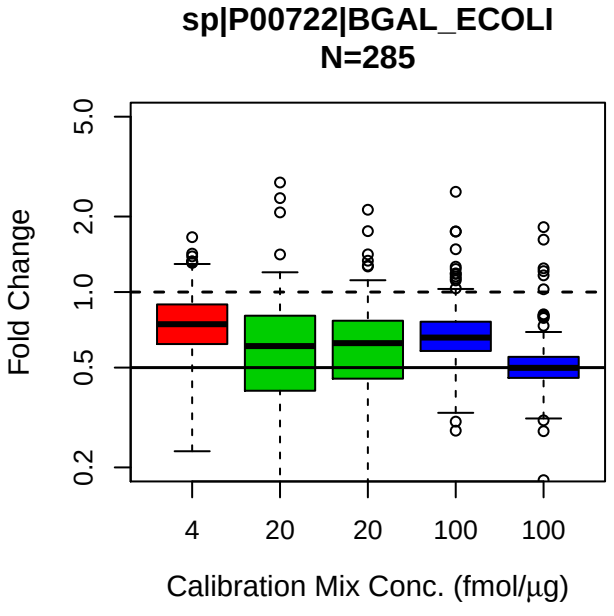
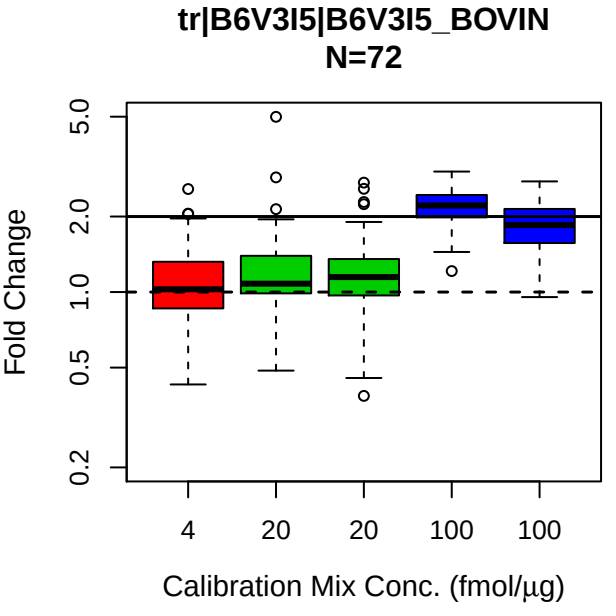
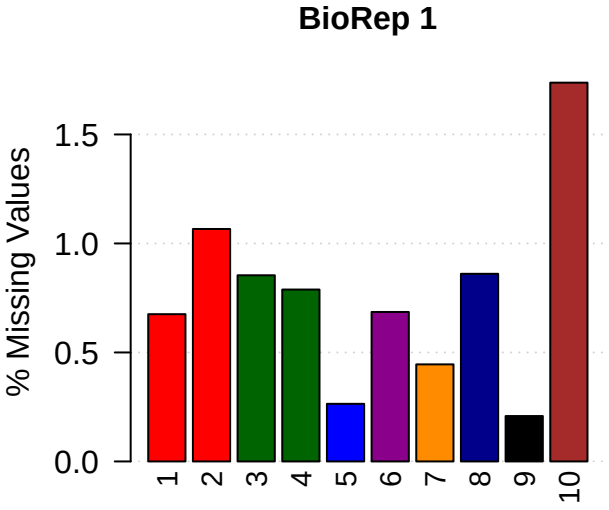
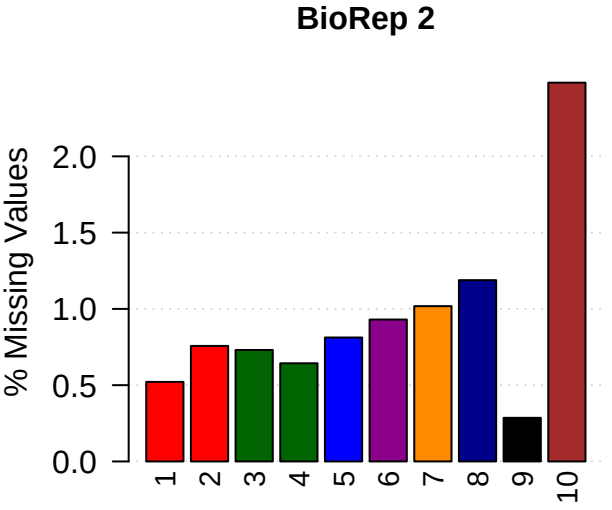
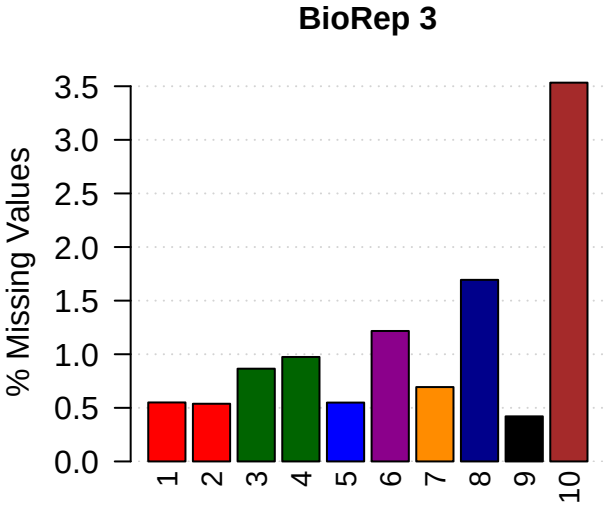

sp|P00489|PYGM_RABIT
N=253
sp|P02789|TRFE_CHICK
N=720
sp|P01012|OVAL_CHICK
N=218
sp|P02666|CASB_BOVIN
N=21
Fold Change
5.0
2.0
1.0
0.5
0.2
4
20
20
100
100
Calibration Mix Conc. (fmol/µg)

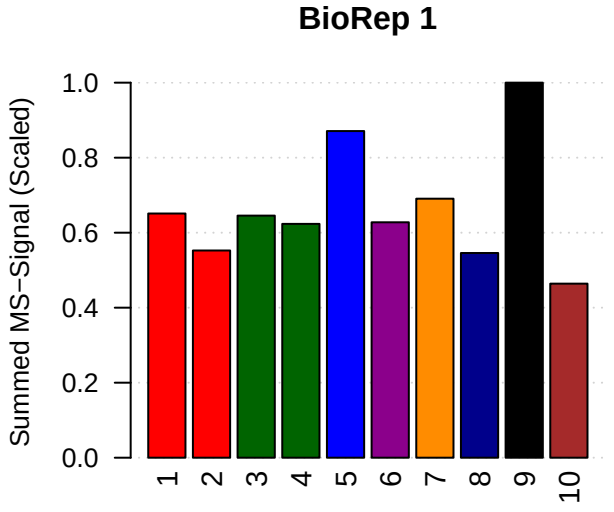

sp|P00722|BGAL_ECOLI
N=285
Fold Change
5.0
2.0
1.0
0.5
0.2
4
20
20
100
100
Calibration Mix Conc. (fmol/µg)

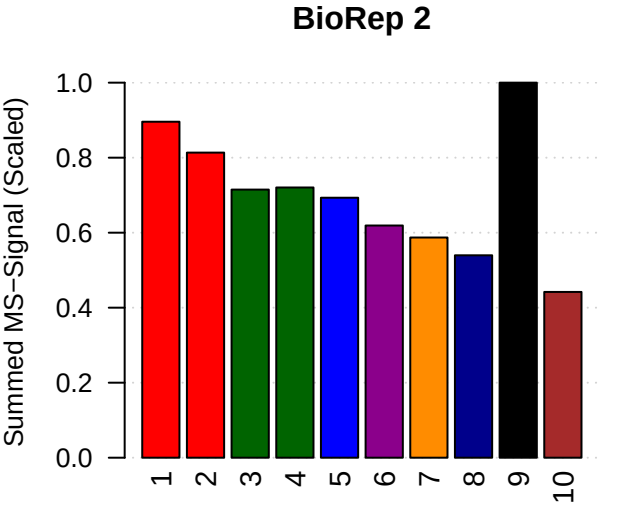

tr|B6V3I5|B6V3I5_BOVIN
N=72
Fold Change
5.0
2.0
1.0
0.5
0.2
4
20
20
100
100
Calibration Mix Conc. (fmol/µg)

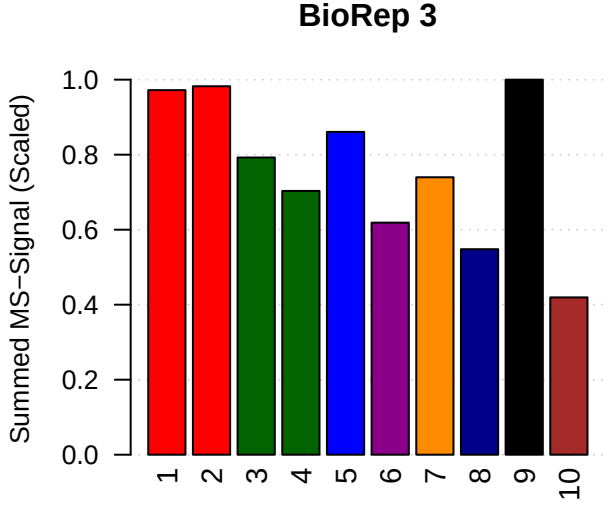

BioRep 1
% Missing Values
0.0
0.5
1.0
1.5
1
2
3
4
5
6
7
8
9
10

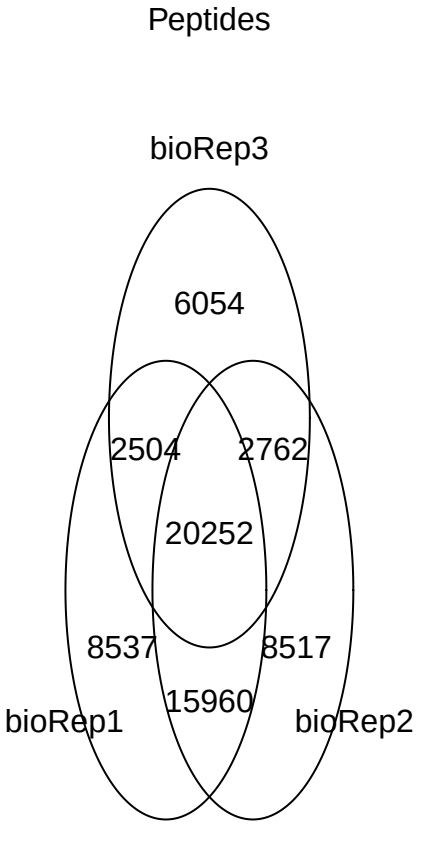

BioRep 2
% Missing Values
0.0
0.5
1.0
1.5
2.0
1
2
3
4
5
6
7
8
9
10

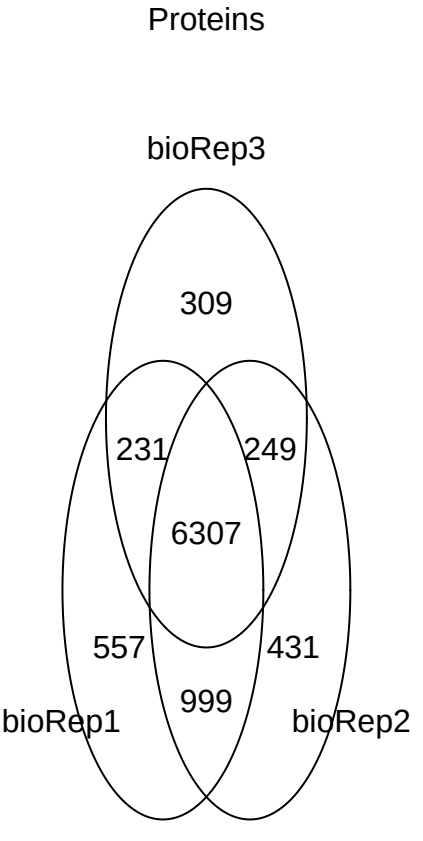

BioRep 3
% Missing Values
0.0
0.5
1.0
1.5
2.0
2.5
3.0
3.5
1
2
3
4
5
6
7
8
9
10

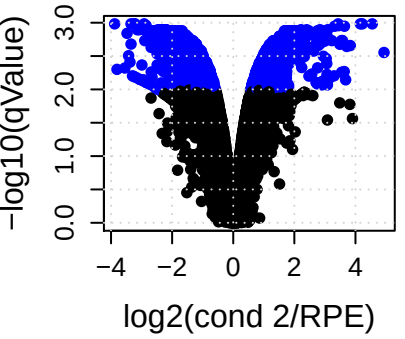

BioRep 1
Summed MS−Signal (Scaled)
0.0
0.2
0.4
0.6
0.8
1.0
1
2
3
4
5
6
7
8
9
10

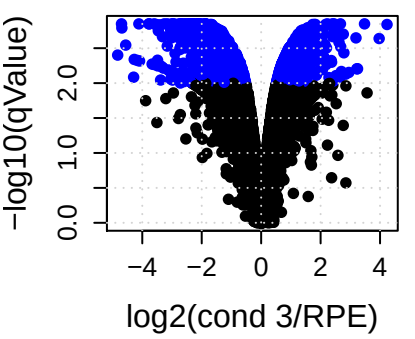

BioRep 2
Summed MS−Signal (Scaled)
0.0
0.2
0.4
0.6
0.8
1.0
1
2
3
4
5
6
7
8
9
10

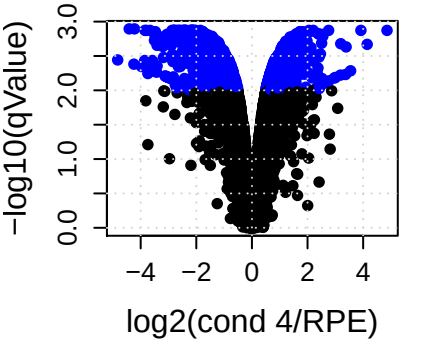

BioRep 3
Summed MS−Signal (Scaled)
0.0
0.2
0.4
0.6
0.8
1.0
1
2
3
4
5
6
7
8
9
10

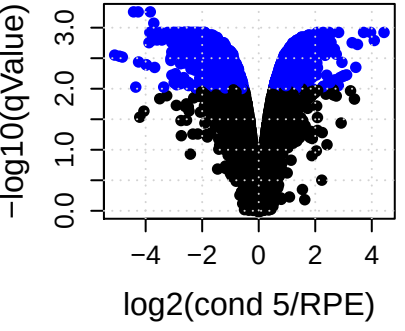

Peptides
bioRep3
6054
2504
2762
20252
8537
8517
15960
bioRep1
bioRep2

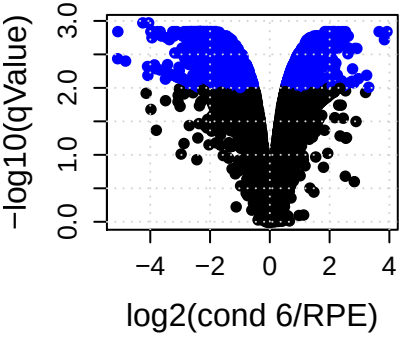

Proteins
bioRep3
309
231
249
6307
557
431
999
bioRep1
bioRep2

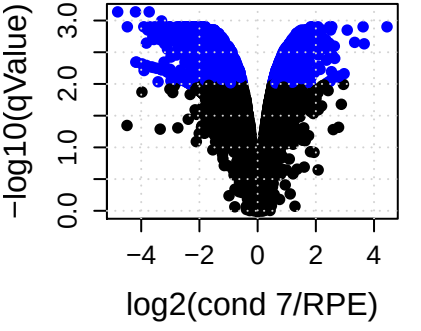

−log10(qValue)
0.0
1.0
2.0
3.0
−4
−2
0
2
4
log2(cond 2/RPE)

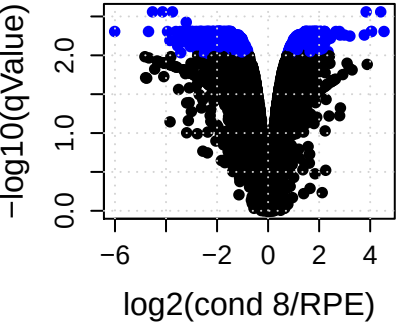

−log10(qValue)
0.0
1.0
2.0
−4
−2
0
2
4
log2(cond 3/RPE)

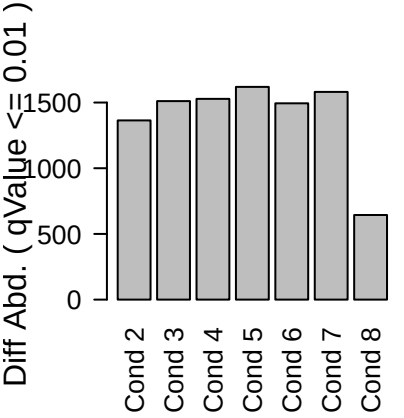

−log10(qValue)
0.0
1.0
2.0
3.0
−4
−2
0
2
4
log2(cond 4/RPE)


−log10(qValue)
0.0
1.0
2.0
3.0
−4
−2
0
2
4
log2(cond 5/RPE)


−log10(qValue)
0.0
1.0
2.0
3.0
−4
−2
0
2
4
log2(cond 6/RPE)


−log10(qValue)
0.0
1.0
2.0
3.0
−4
−2
0
2
4
log2(cond 7/RPE)


−log10(qValue)
0.0
1.0
2.0
−6
−2
0
2
4
log2(cond 8/RPE)


Diff Abd. (qValue <= 0.01)
0
500
1000
1500
Cond 2
Cond 3
Cond 4
Cond 5
Cond 6
Cond 7
Cond 8