

Aggregatilineales_Genus

FDR: 1.539e-19
Coefficient: 6.97e-02
N: 236

75
50
25
0

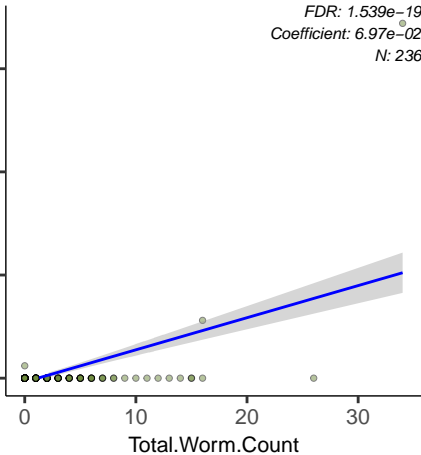
0

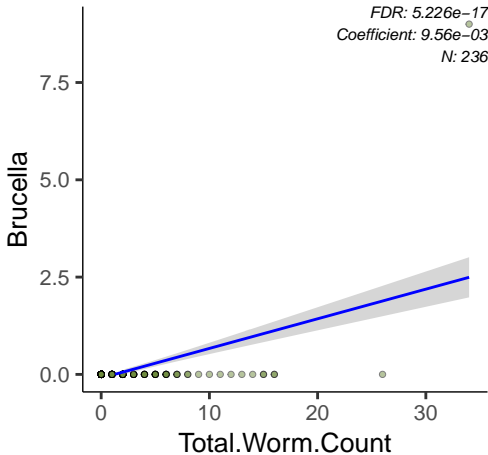
Total.Worm.Count

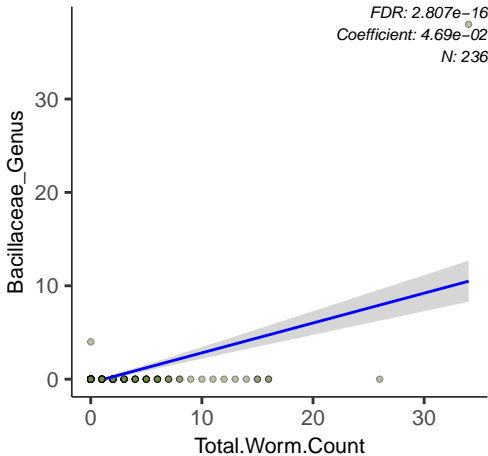
10

20

30







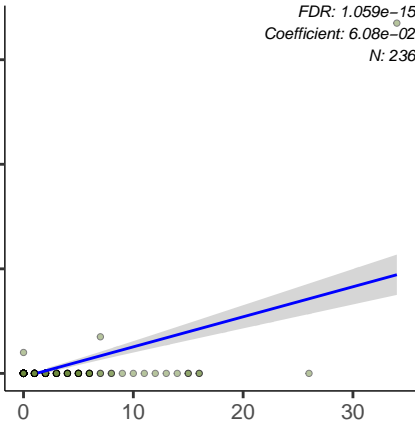
vadinHA49_Genus

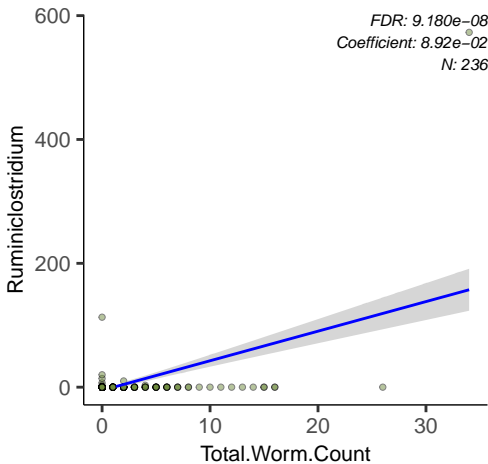
FDR: 1.059e-15
Coefficient: 6.08e-02
N: 236

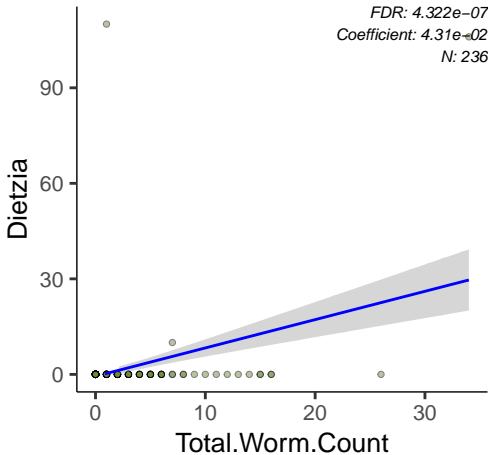
60
40
20
0

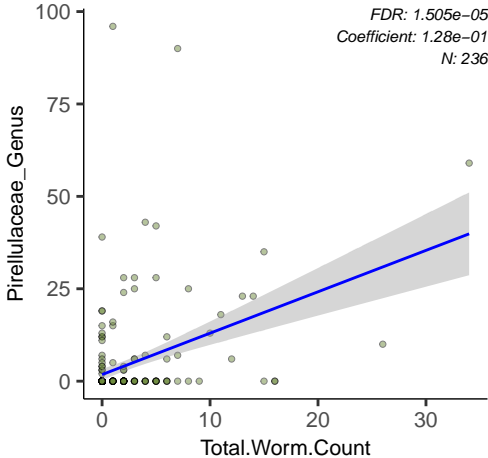
0 10 20 30

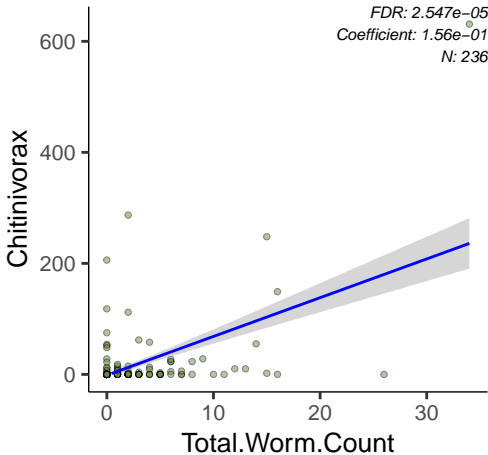
Total.Worm.Count

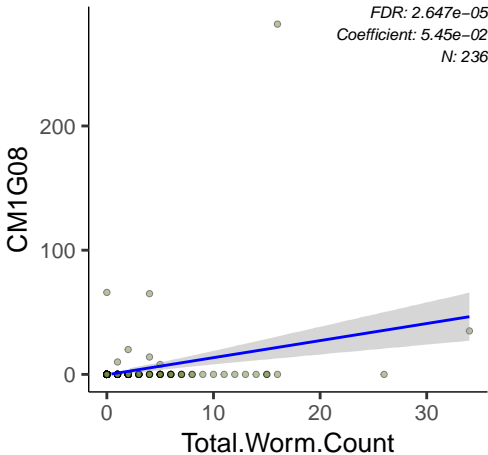


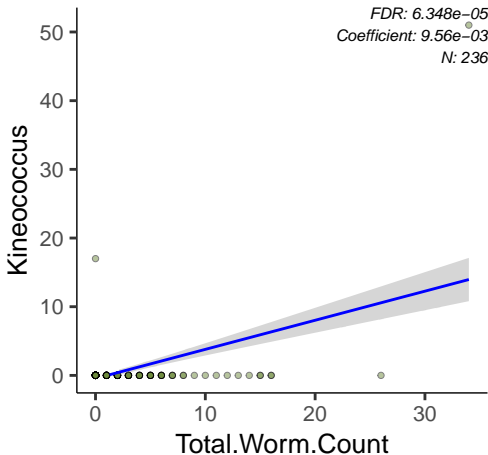


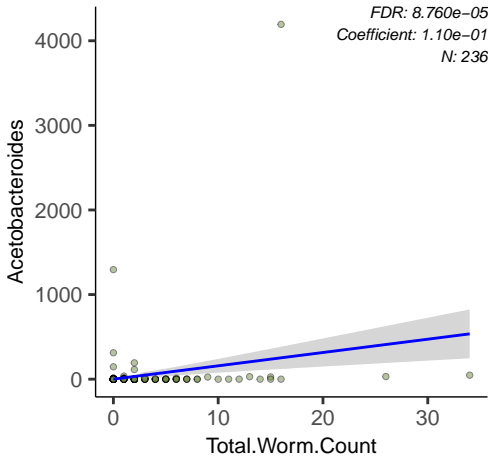


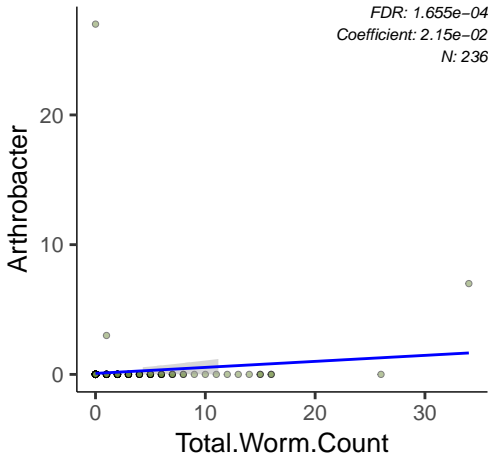


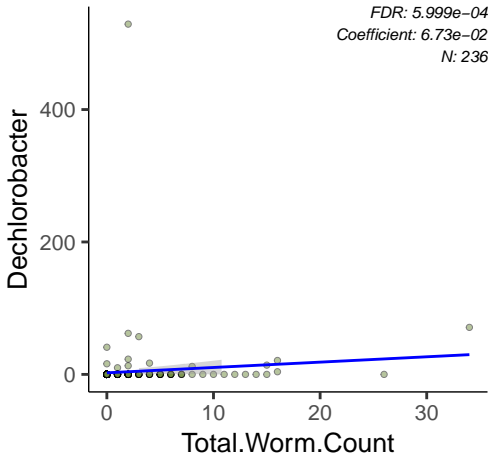


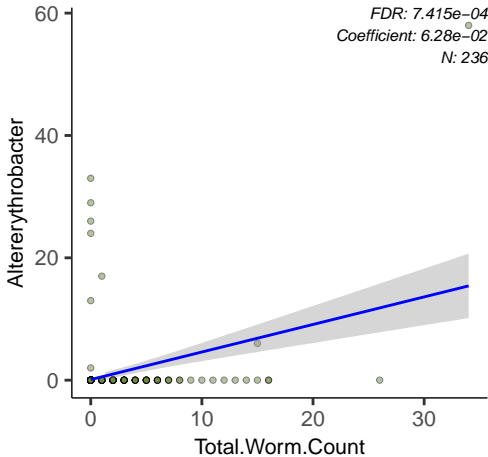












NS9_marine_group_Genus

FDR: 2.090e-03
Coefficient: 1.66e-01
N: 236

200

100

0

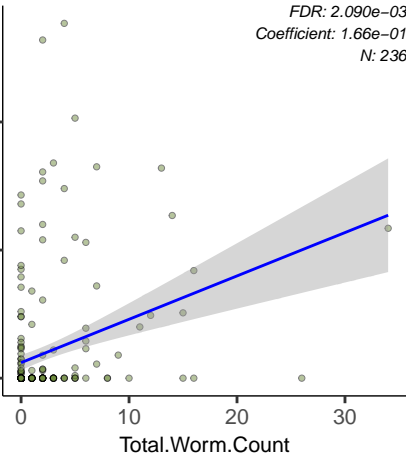
0

10

20

30

Total.Worm.Count



Spirosomataceae_Genus

100

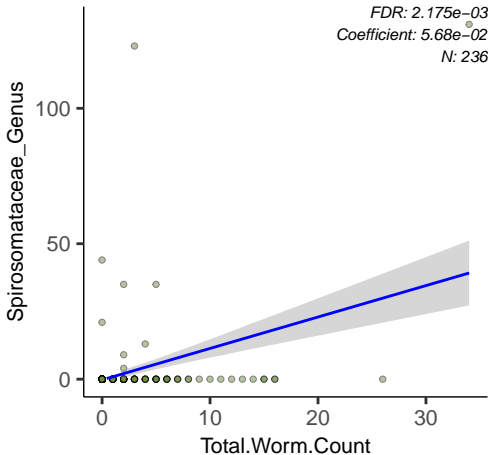
50

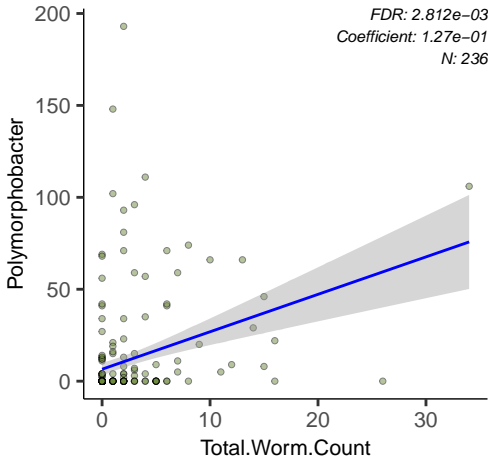
0

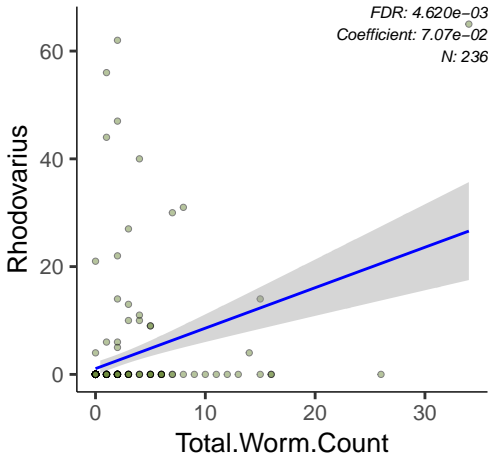
0

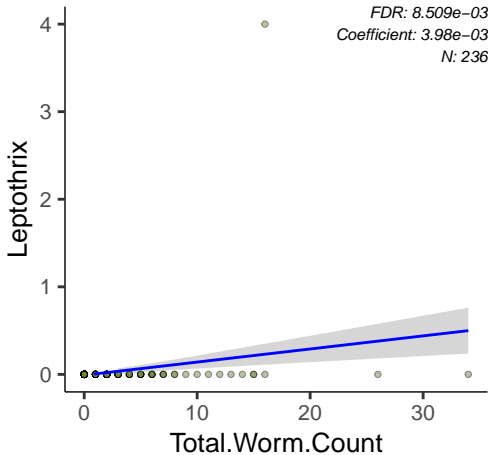
Total.Worm.Count

FDR: 2.175e-03
Coefficient: 5.68e-02
N: 236









Absconditabacteriales_SR1._Genus

FDR: 8.509e-03
Coefficient: 3.98e-03
N: 236

3
2
1
0

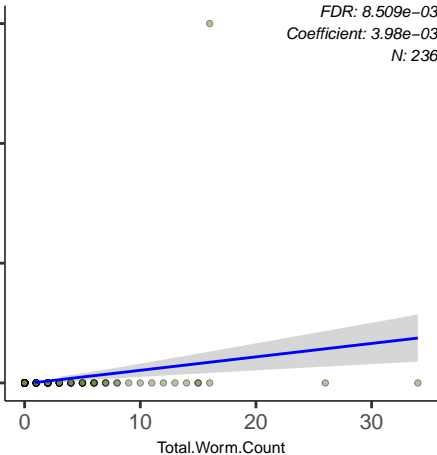
0

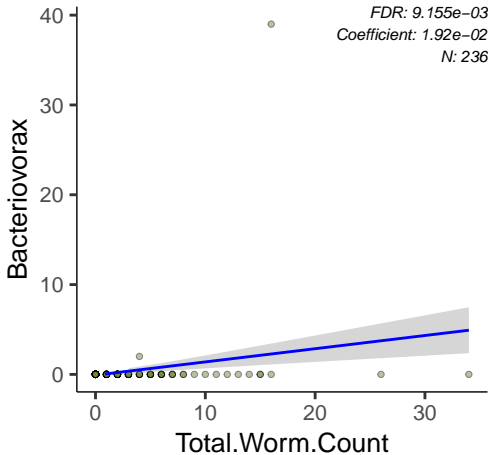
10

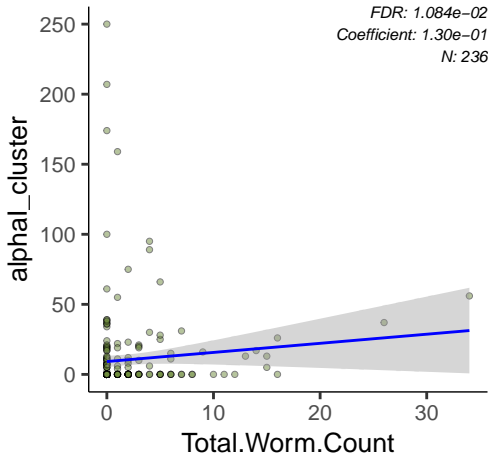
20

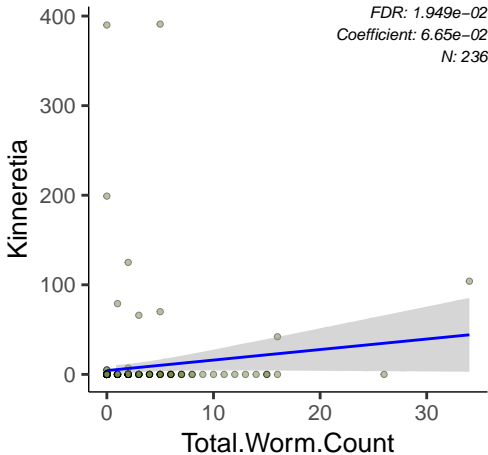
30

Total.Worm.Count









Candidatus_Alysiosphaera

FDR: 1.989e-02
Coefficient: 1.01e-01
N: 236

1500

1000

500

0

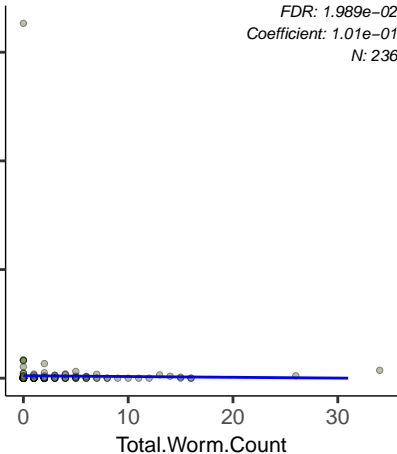
0

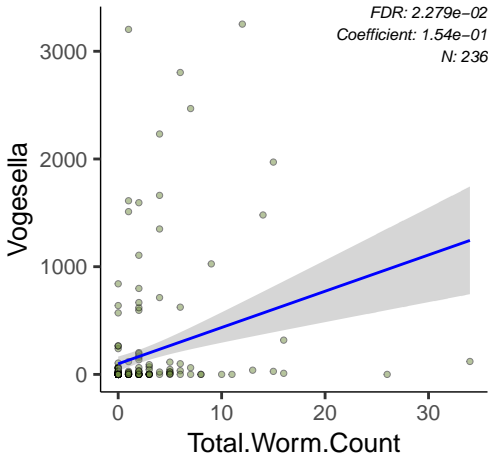
10

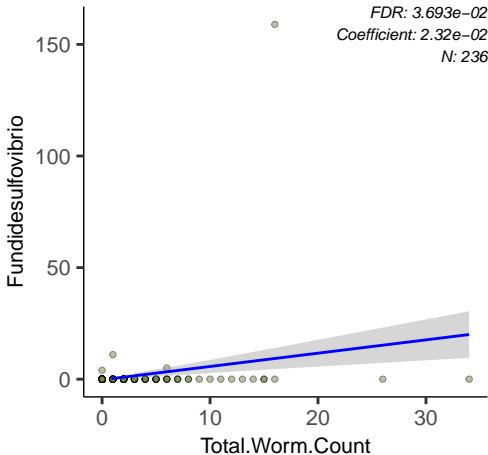
20

30

Total.Worm.Count







X0319.6G20_Genus

FDR: 3.806e-02
Coefficient: 1.22e-01
N: 236

150

100

50

0

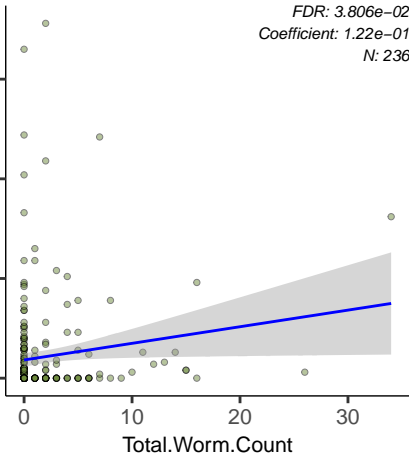
0

10

20

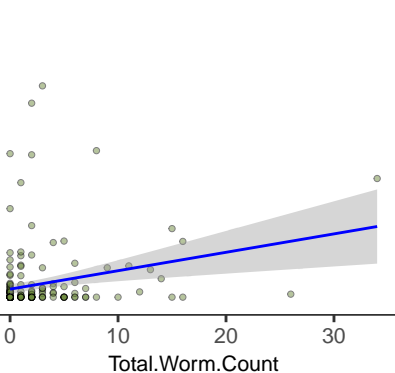
30

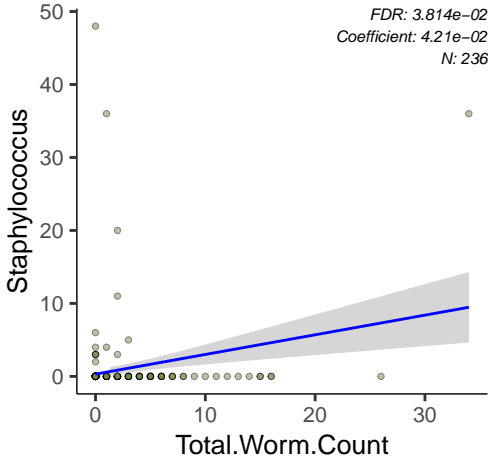
Total.Worm.Count

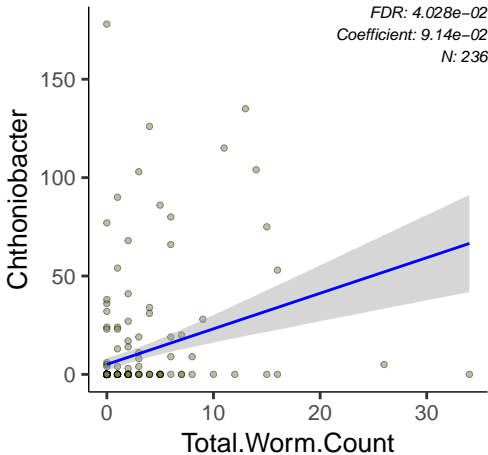


Rhodocyclaceae_Genus

FDR: 3.806e-02
Coefficient: 1.05e-01
N: 236







Microscillaceae_Genus

FDR: 4.337e-02
Coefficient: 1.13e-01
N: 236

750
500
250
0

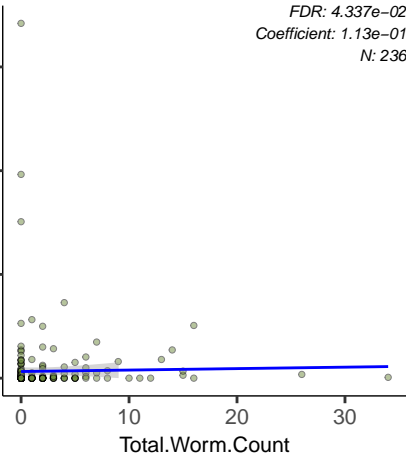
0

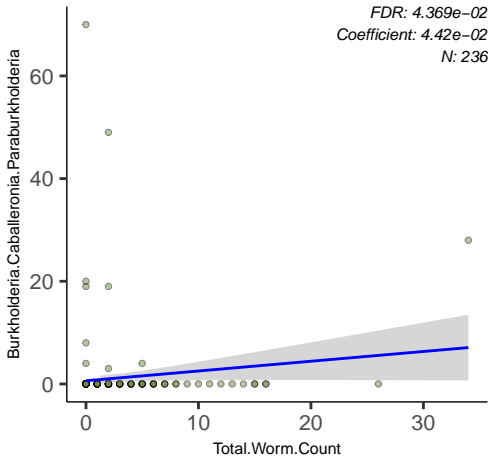
Total.Worm.Count

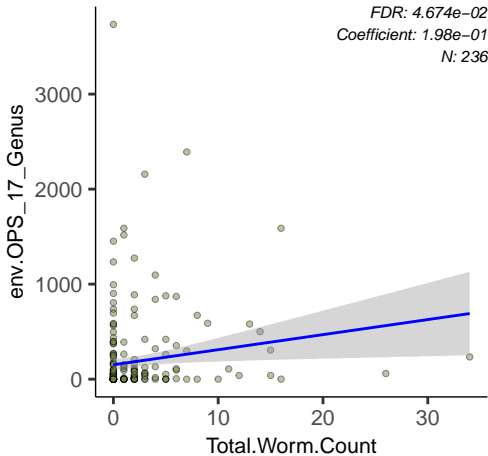
10

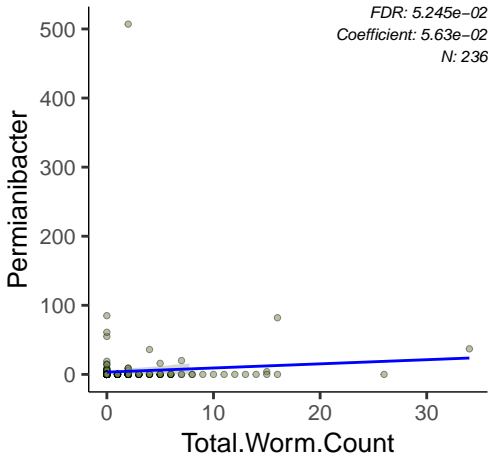
20

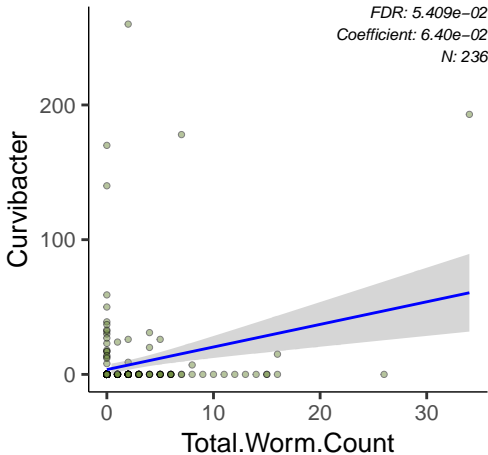
30











Chitinibacter

FDR: 5.478e-02
Coefficient: 8.76e-02
N: 236

150

100

50

0

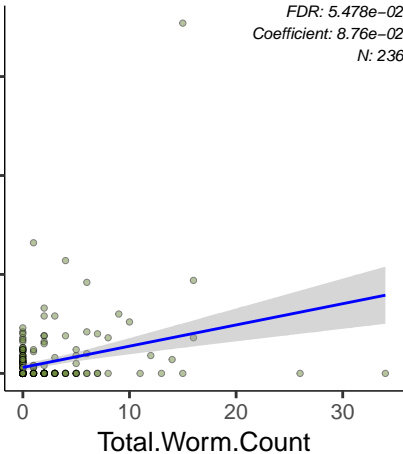
0

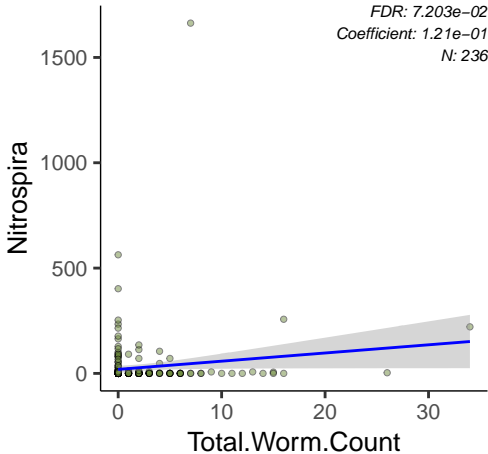
10

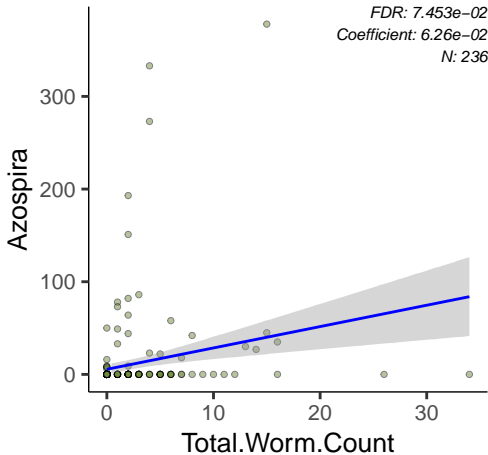
20

30

Total.Worm.Count

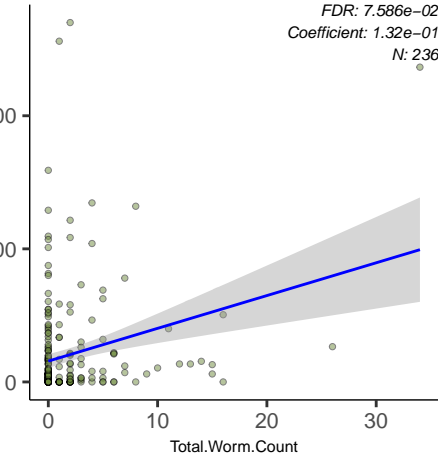


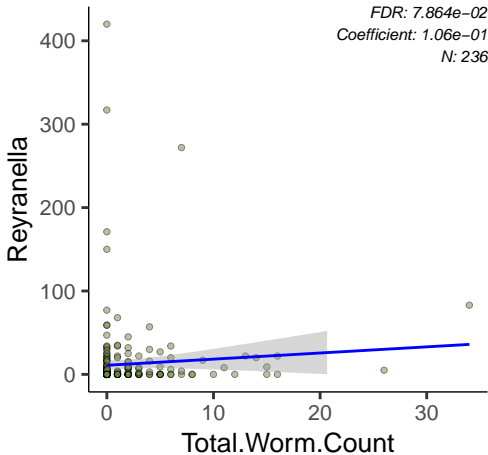


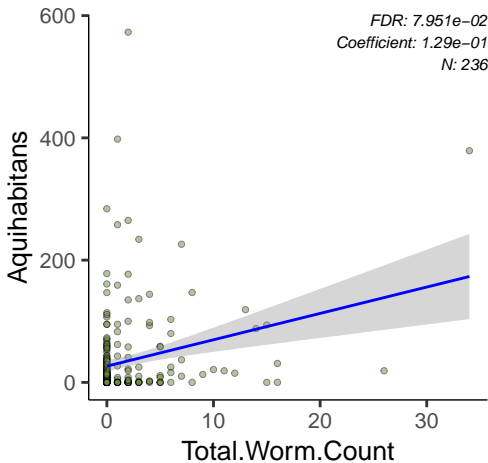


Pseudobdellovibrionaceae_Genus

FDR: 7.586e-02
Coefficient: 1.32e-01
N: 236







Microtrichaceae_Genus

300
200
100
0

FDR: 8.172e-02
Coefficient: 5.34e-02
N: 236

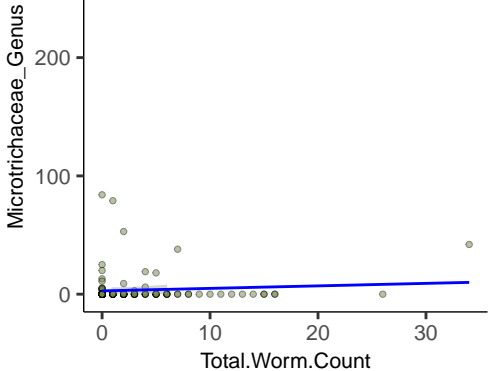
0

Total.Worm.Count

10

20

30



Mycobacterium

FDR: 1.022e-01
Coefficient: 1.33e-01
N: 236

20000

15000

10000

5000

0

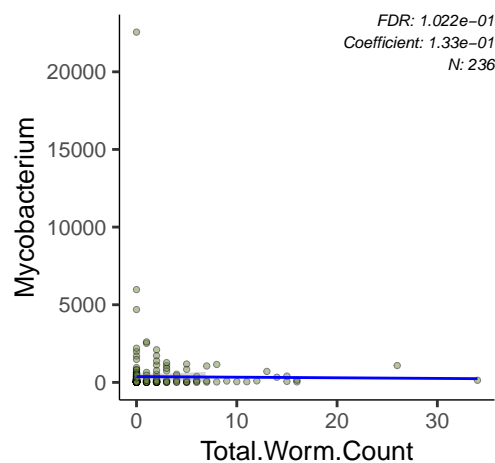
0

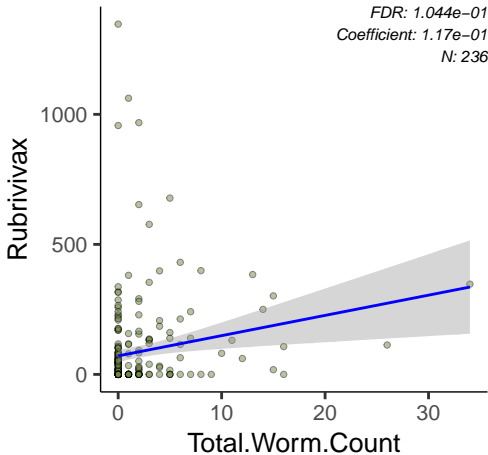
10

20

30

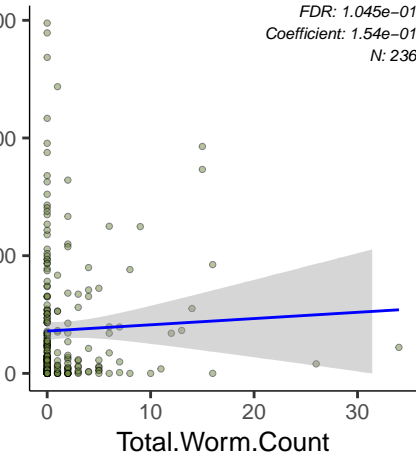
Total.Worm.Count



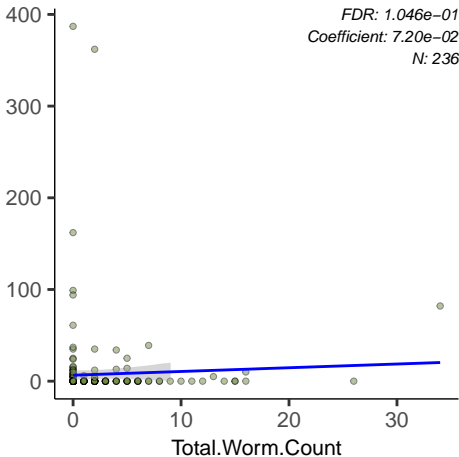


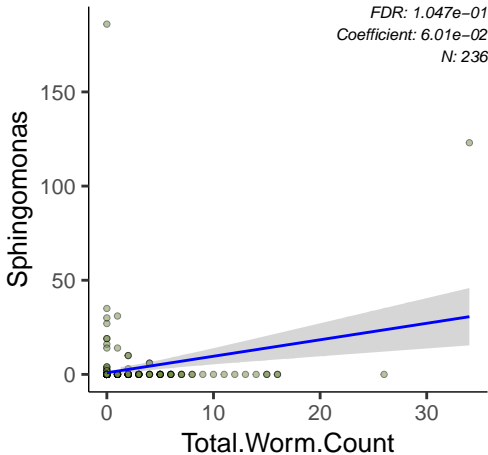
Dechloromonas

FDR: 1.045e-01
Coefficient: 1.54e-01
N: 236



Kapabacteriales_Genus





Sporomusaceae_Genus

FDR: 1.069e-01
Coefficient: 9.57e-02
N: 236

1000

500

0

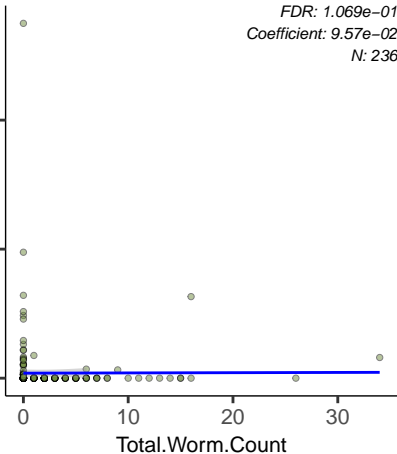
0

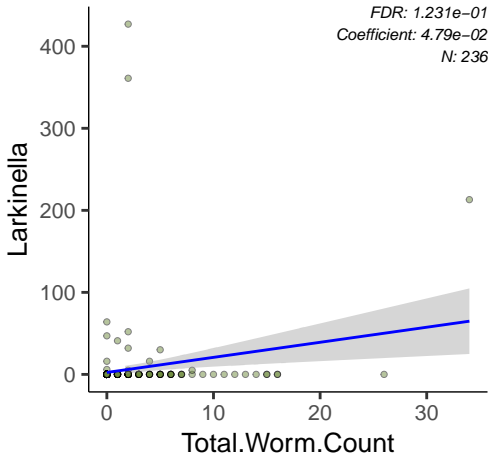
Total.Worm.Count

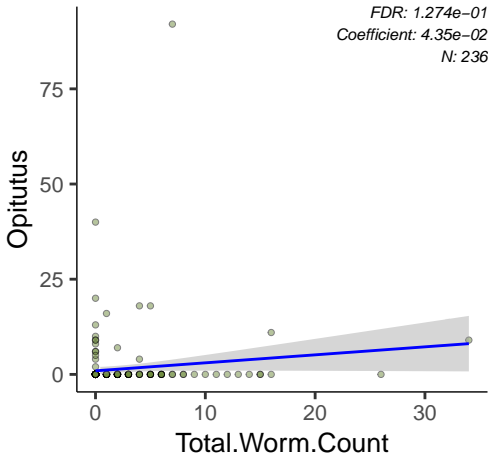
10

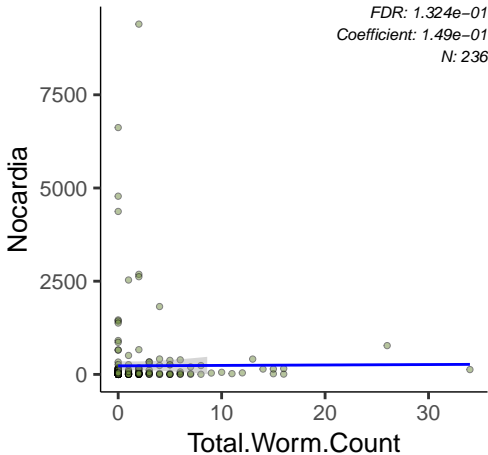
20

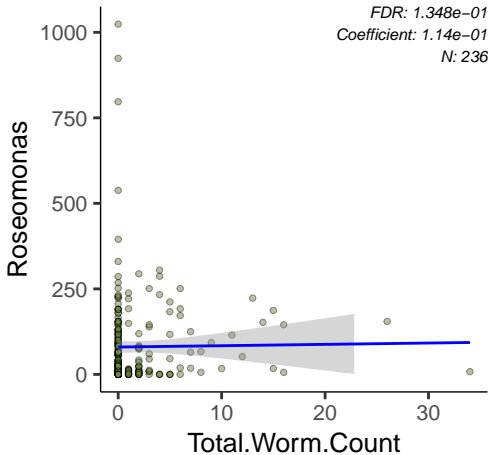
30

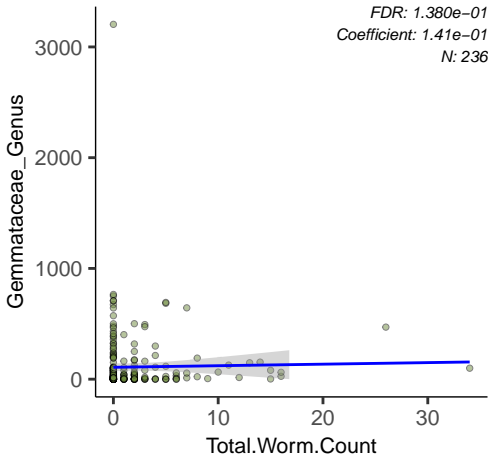


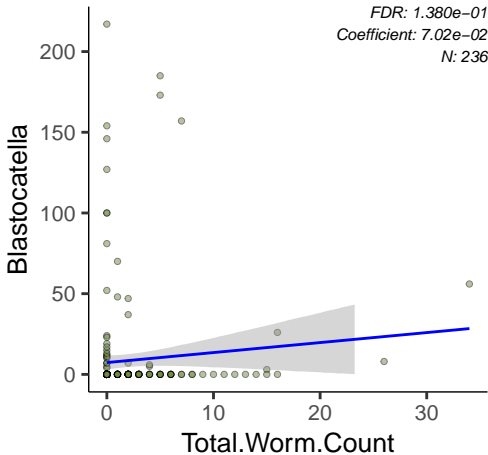




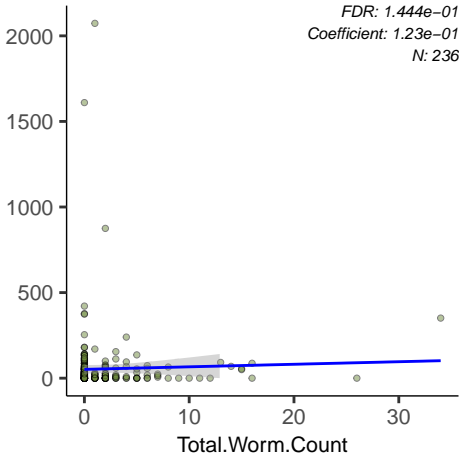


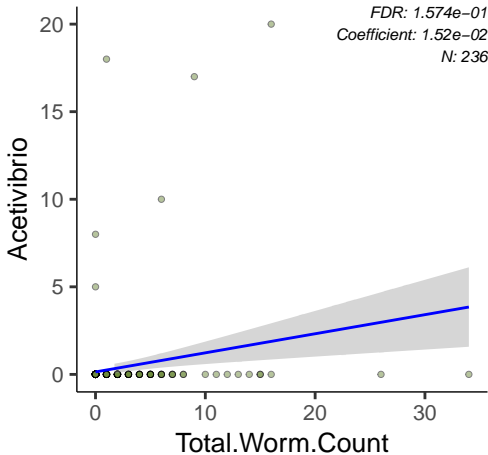






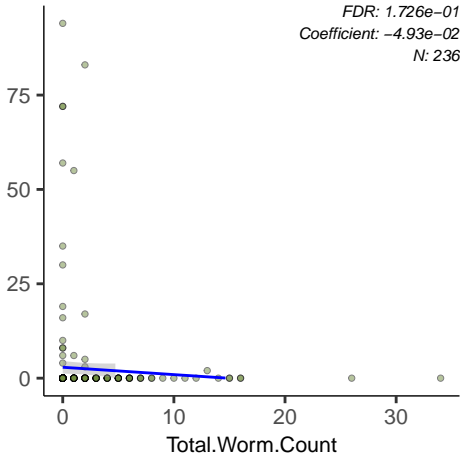
Rhizobiaceae_Genus

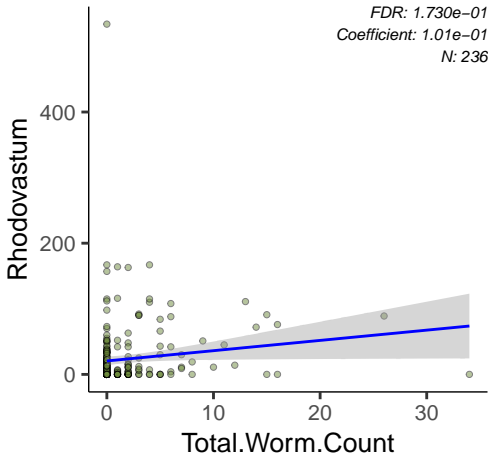


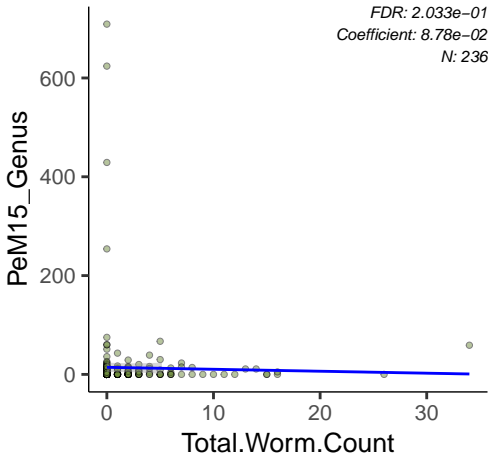


Unknown_Family_Genus

FDR: 1.726e-01
Coefficient: -4.93e-02
N: 236







Vampirovibrionaceae_Genus

FDR: 2.037e-01
Coefficient: 1.13e-01
N: 236

2000

1000

0

0

10

20

30

Total.Worm.Count

