

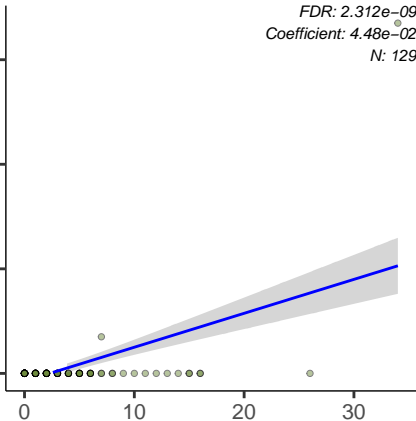
vadinHA49_Genus

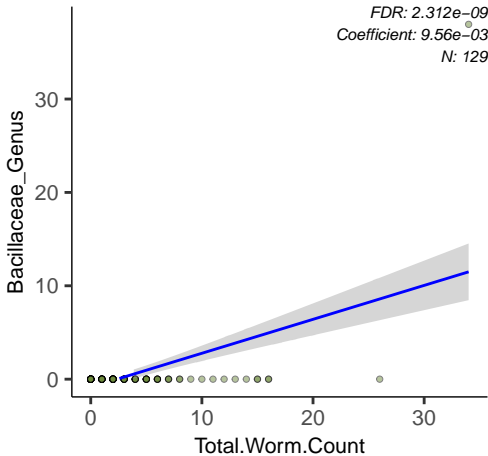
FDR: 2.312e-09
Coefficient: 4.48e-02
N: 129

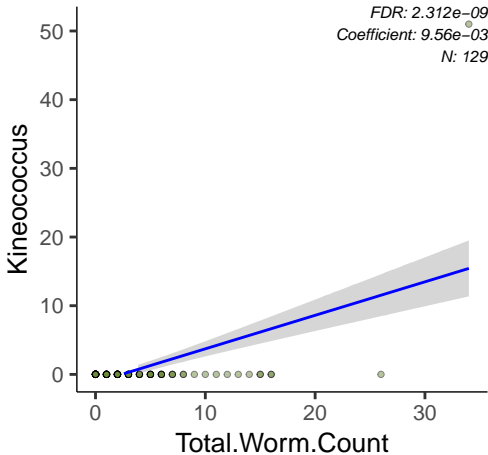
60
40
20
0

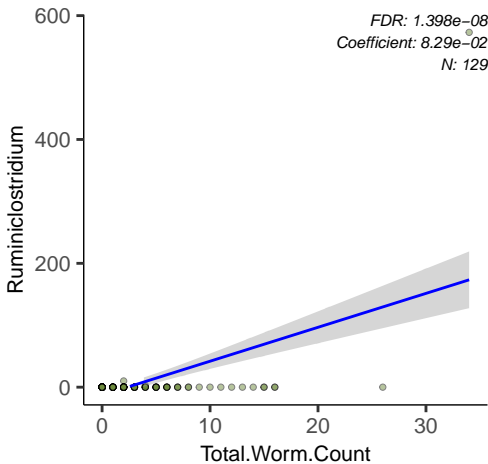
0 10 20 30

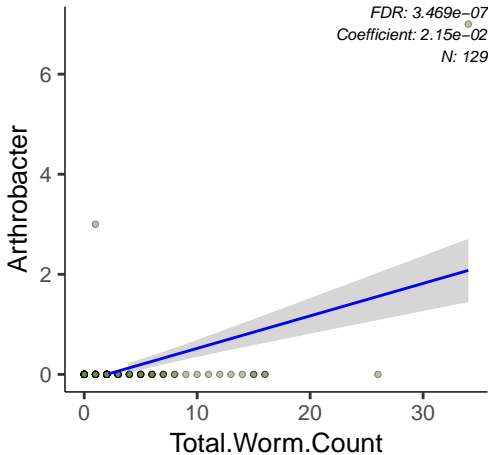
Total.Worm.Count

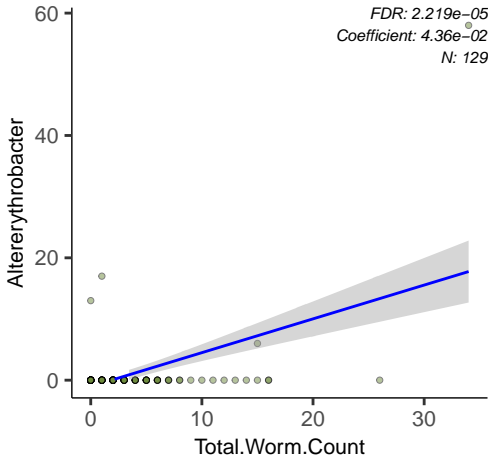


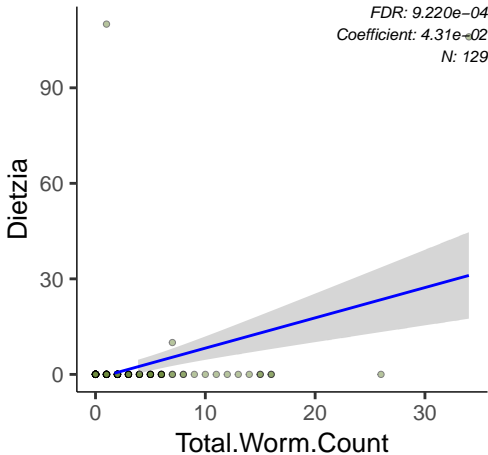


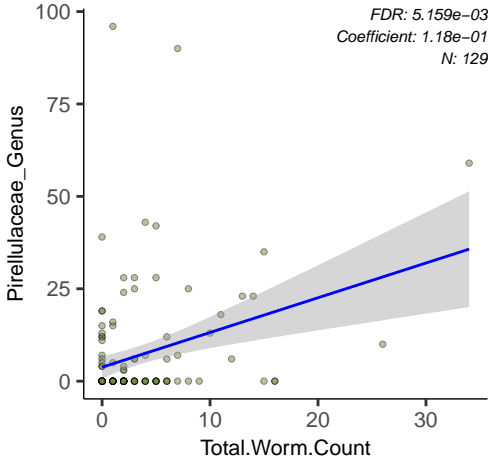


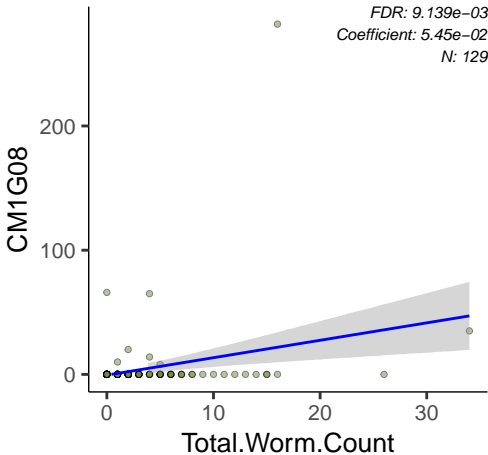


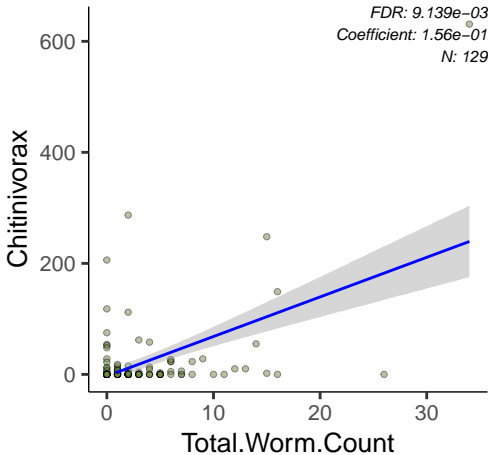


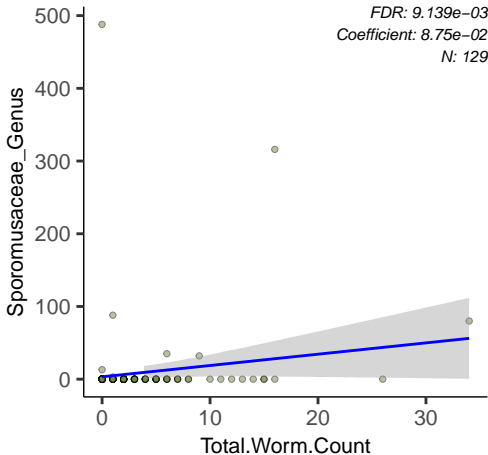


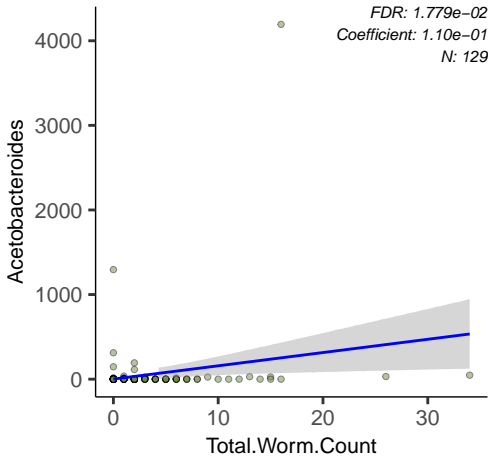


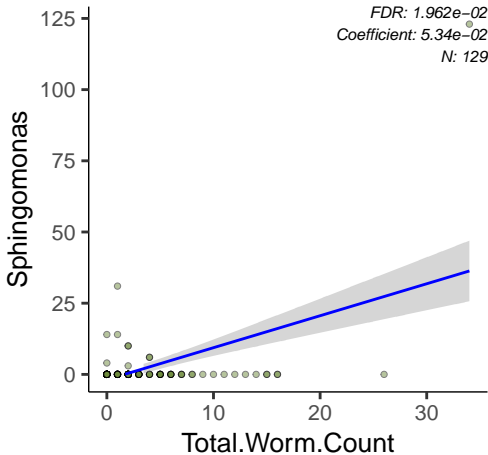


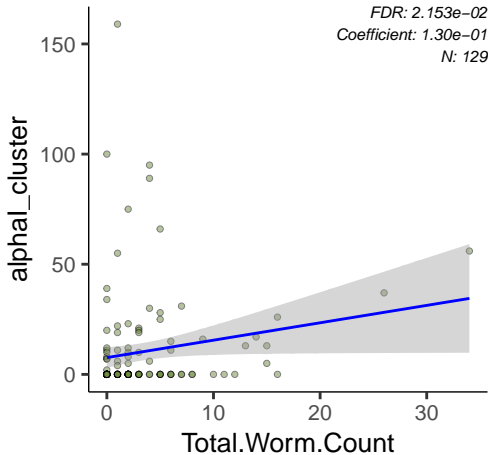






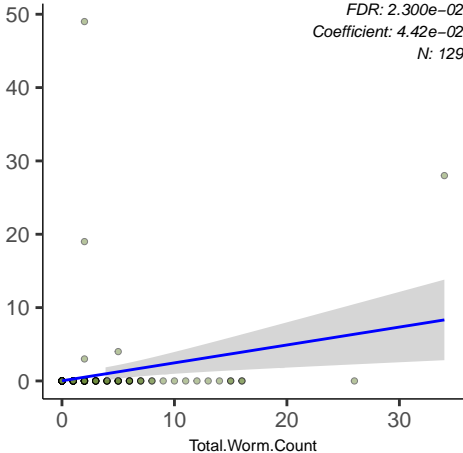






Burkholderia.Caballeronia.Paraburkholderia

FDR: 2.300e-02
Coefficient: 4.42e-02
N: 129



Candidatus_Alysiosphaera

FDR: 2.376e-02
Coefficient: 1.01e-01
N: 129

60

40

20

0

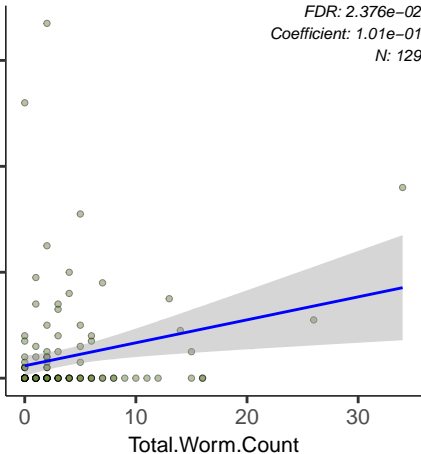
0

10

20

30

Total.Worm.Count



NS9_marine_group_Genus

FDR: 2.766e-02
Coefficient: 1.66e-01
N: 129

200

100

0

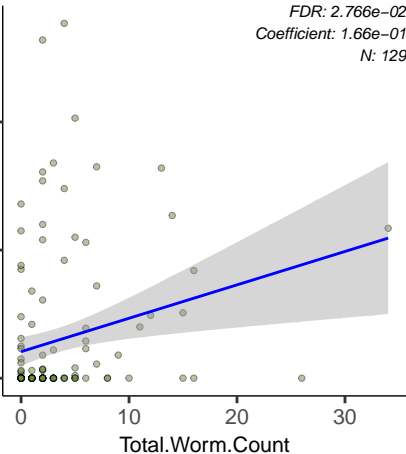
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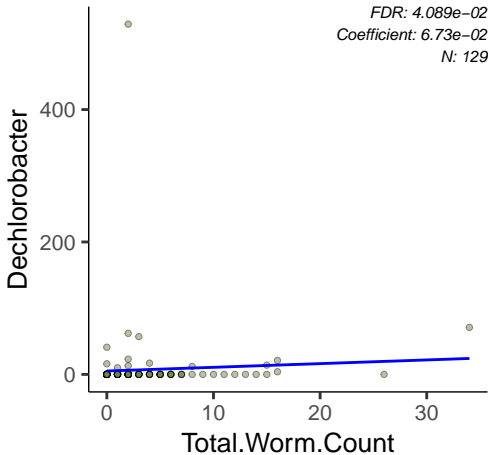
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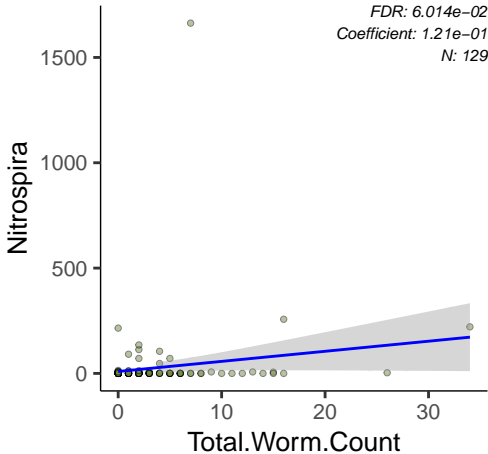
20

30

Total.Worm.Count

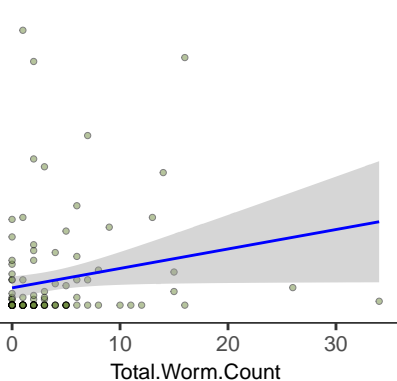


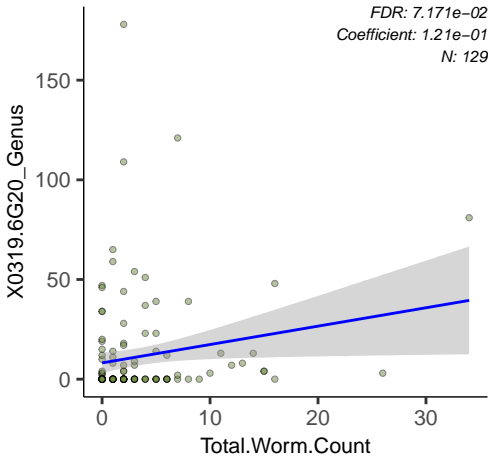


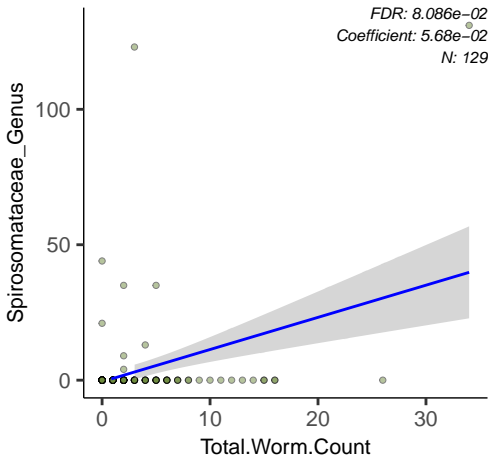


Microscillaceae_Genus

FDR: 6.535e-02
Coefficient: 1.13e-01
N: 129

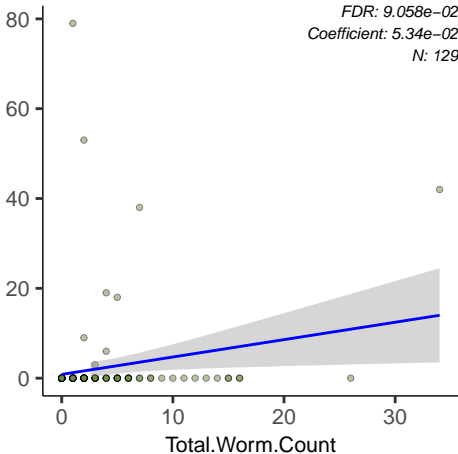


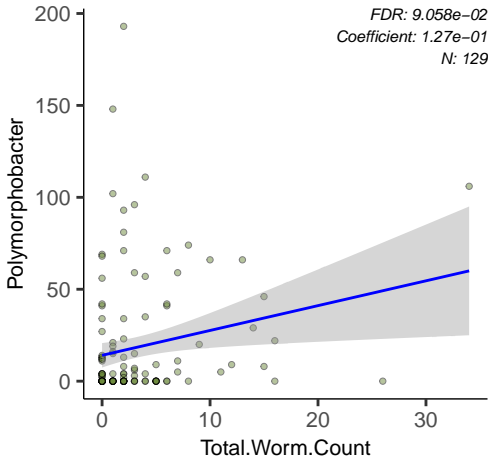


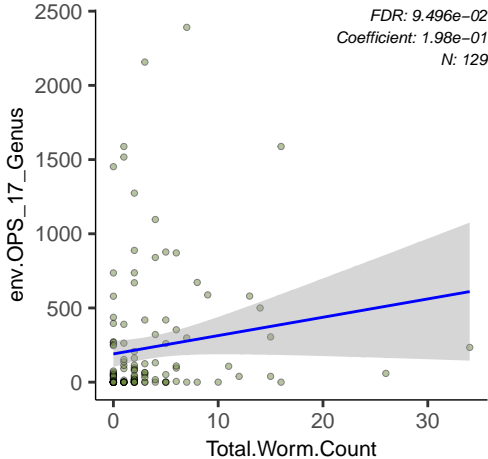


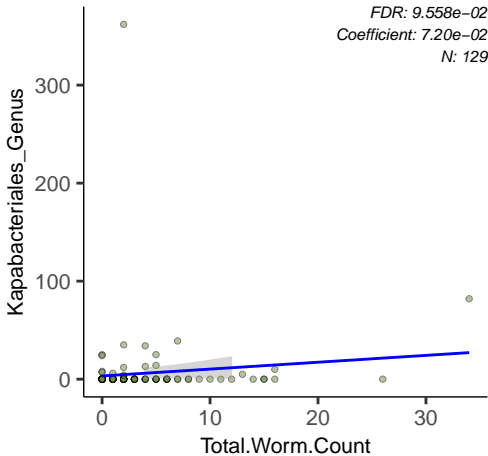
Microtrichaceae_Genus

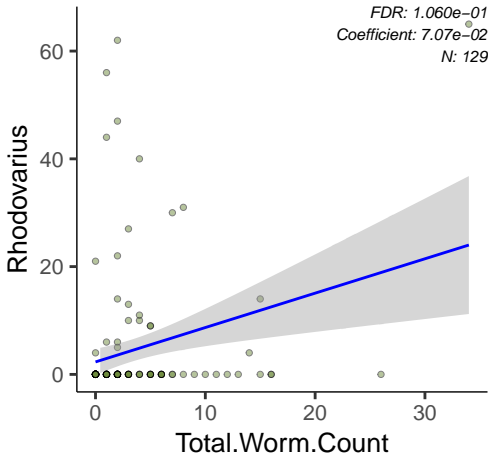
FDR: 9.058e-02
Coefficient: 5.34e-02
N: 129

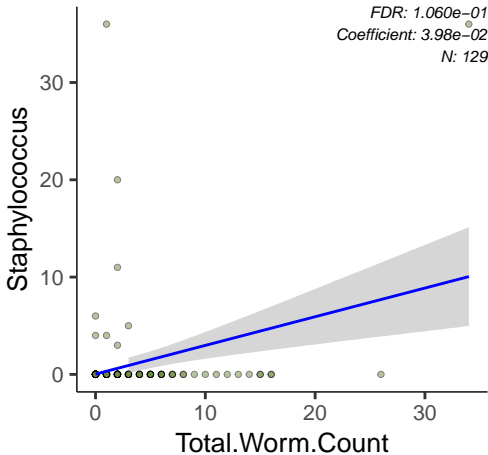


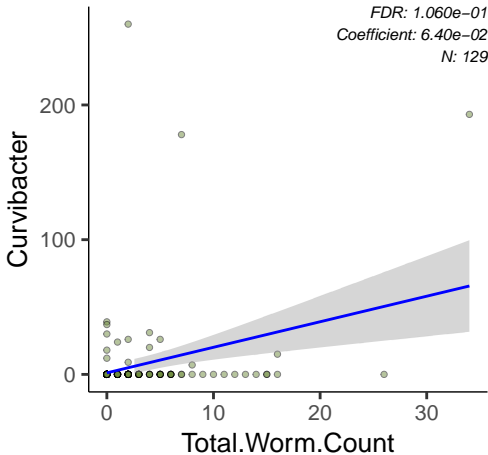


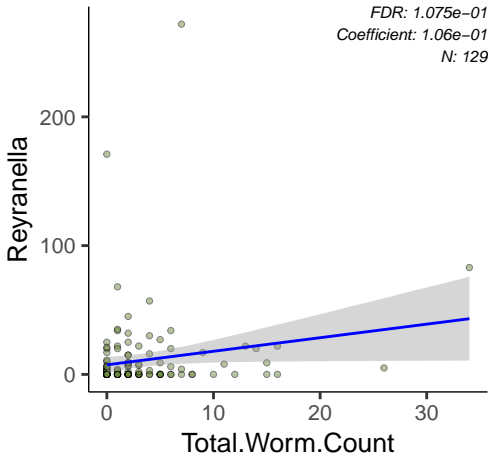






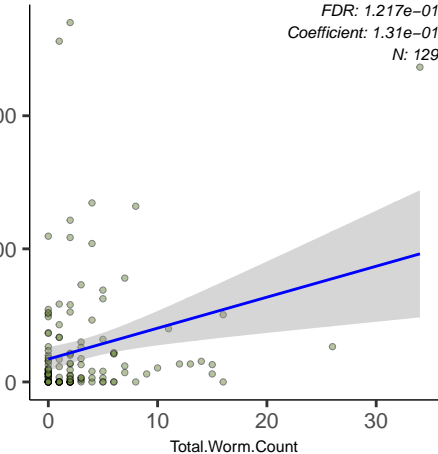






Pseudobdellovibrionaceae_Genus

FDR: 1.217e-01
Coefficient: 1.31e-01
N: 129



Chitinibacter

FDR: 1.217e-01
Coefficient: 7.97e-02
N: 129

150

100

50

0

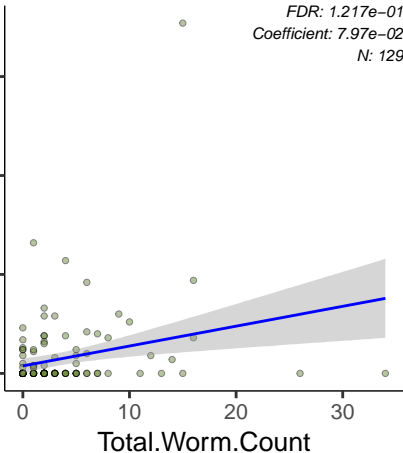
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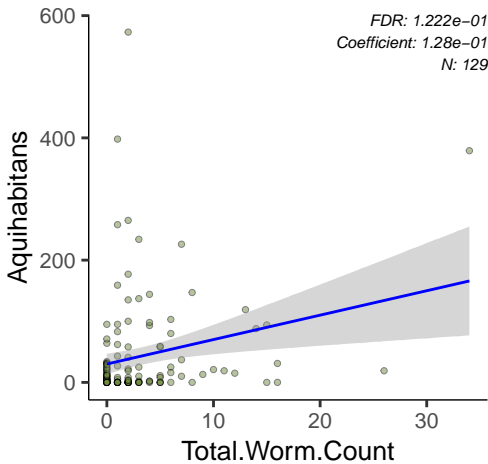
10

20

30

Total.Worm.Count





Candidatus_Nomurabacteria_Genus

FDR: 1.222e-01
Coefficient: 3.98e-03
N: 129

5
4
3
2
1
0

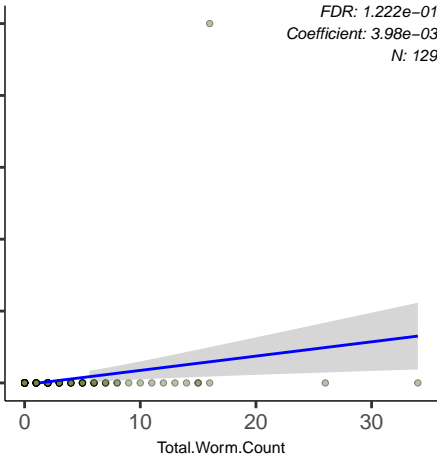
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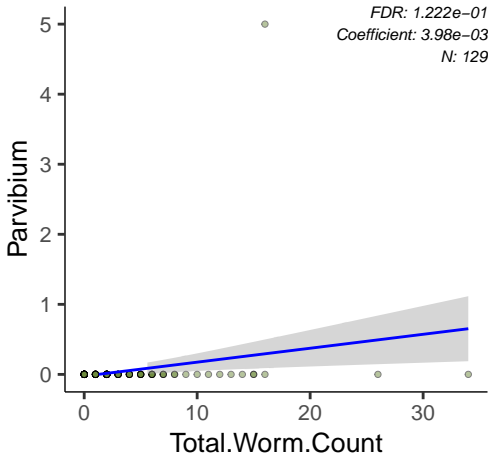
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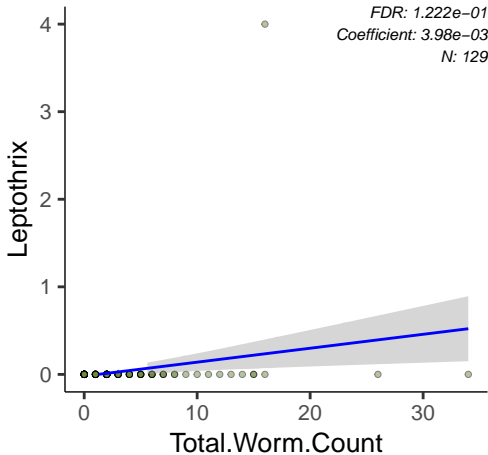
20

30

Total.Worm.Count







Absconditabacteriales_SR1._Genus

FDR: 1.222e-01
Coefficient: 3.98e-03
N: 129

3
2
1
0

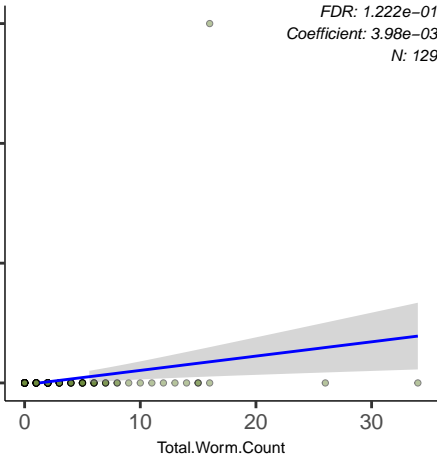
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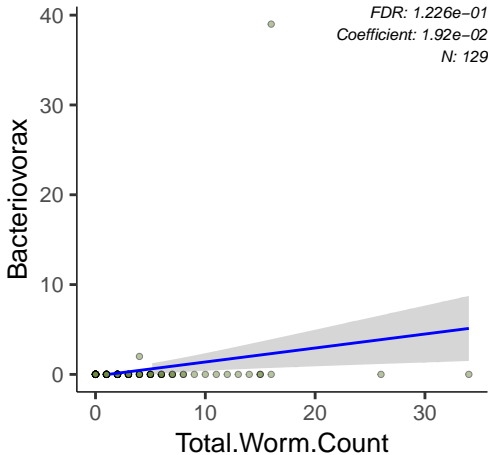
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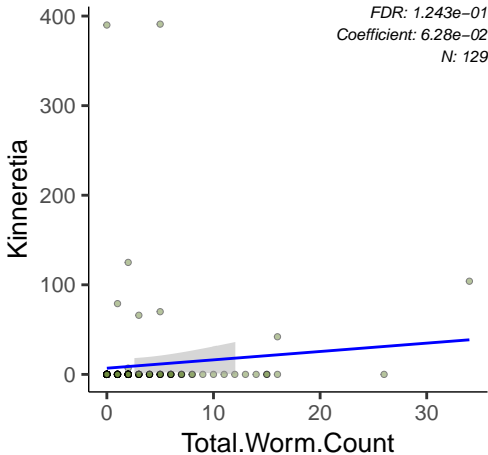
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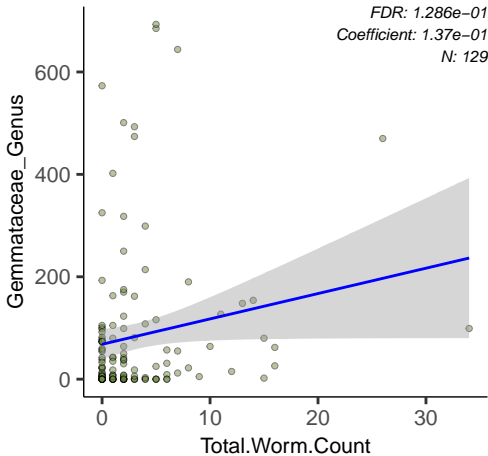
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Total.Worm.Count









PeM15_Genus

FDR: 1.286e-01
Coefficient: 8.78e-02
N: 129

60

40

20

0

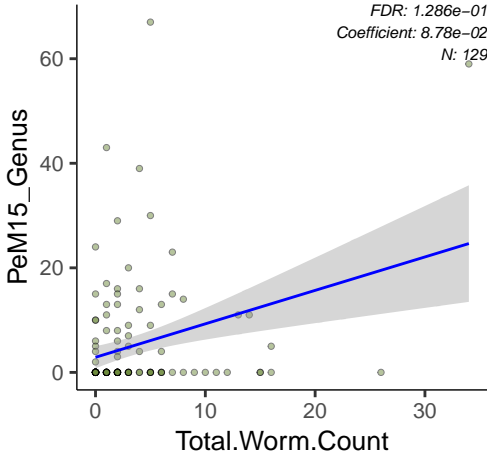
Total.Worm.Count

0

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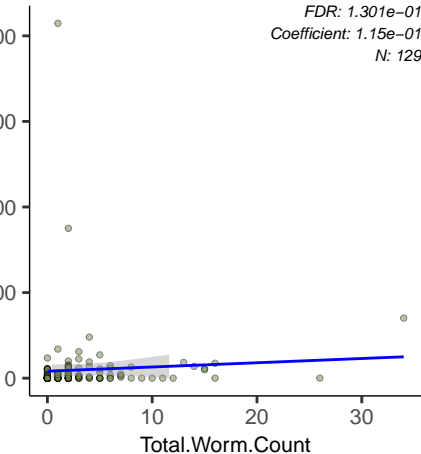
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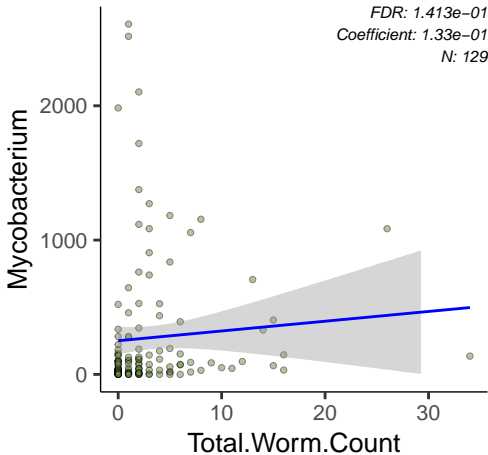
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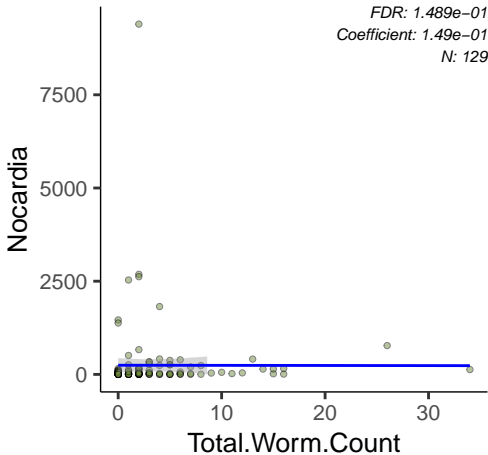


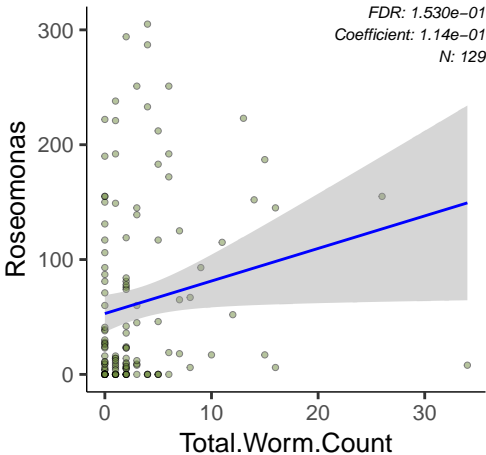
Rhizobiaceae_Genus

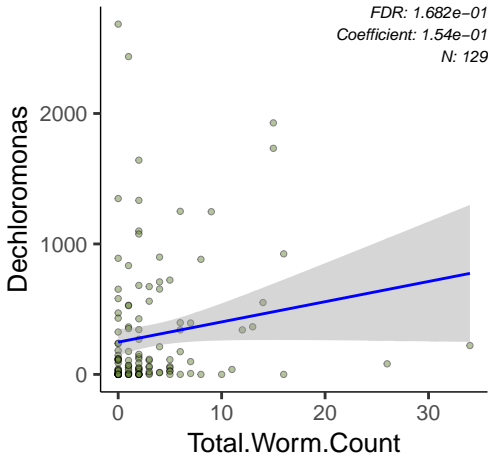
FDR: 1.301e-01
Coefficient: 1.15e-01
N: 129

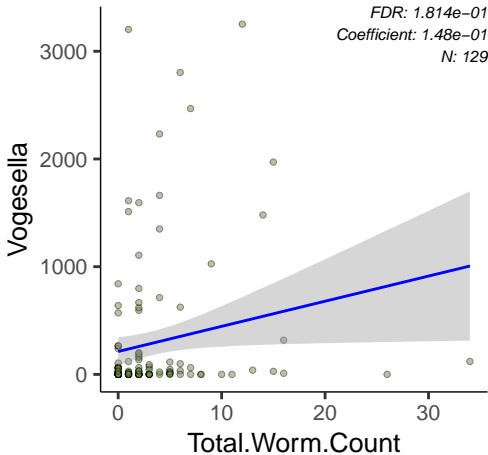


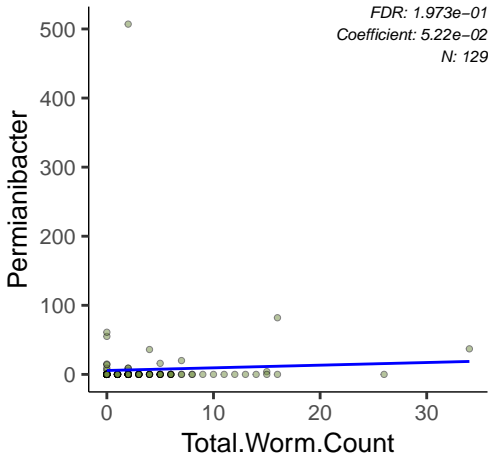


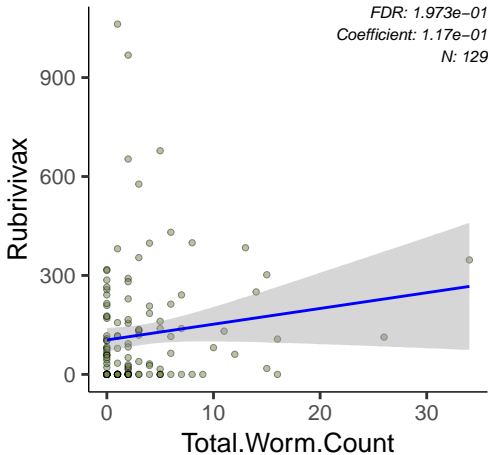


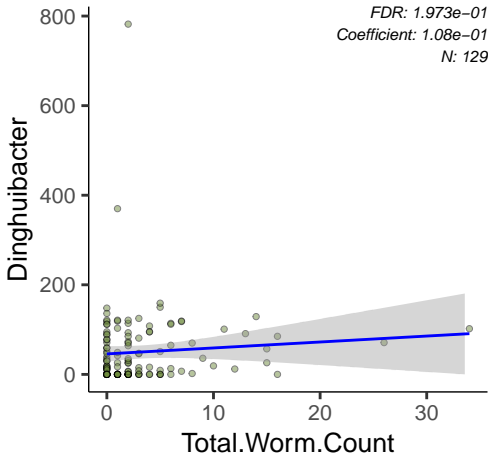


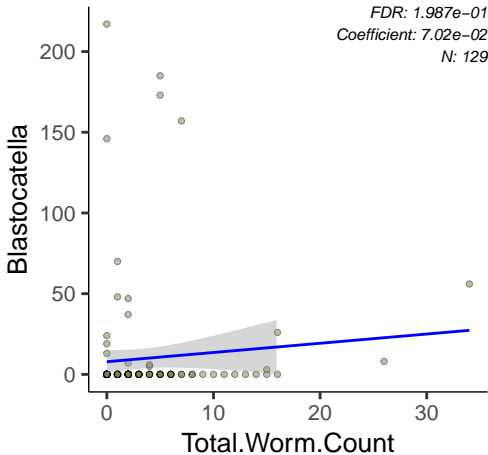


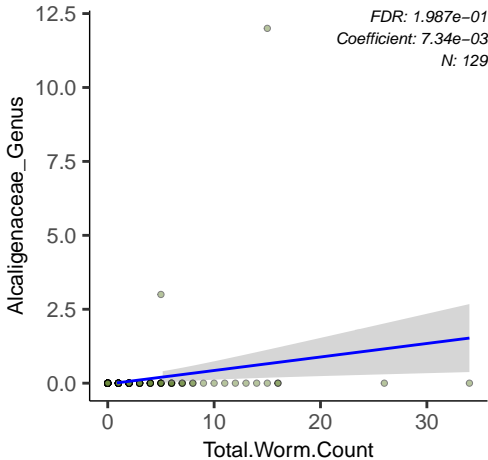






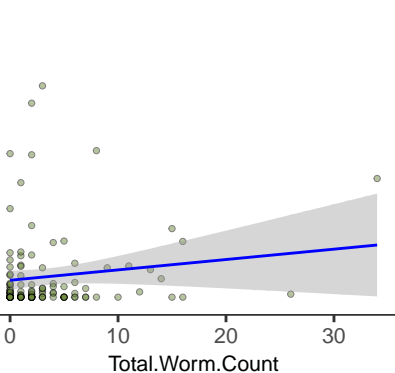


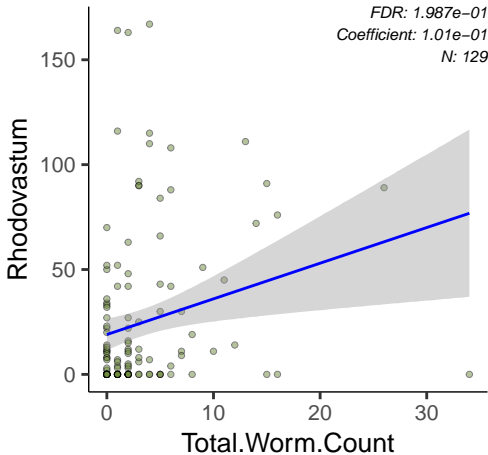


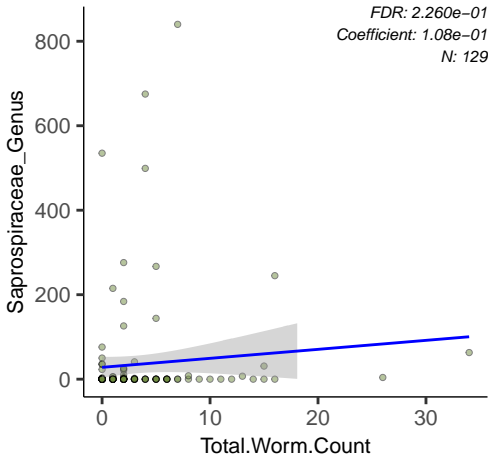


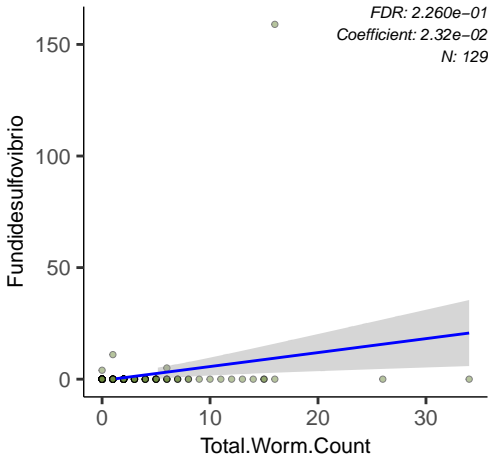
Rhodocyclaceae_Genus

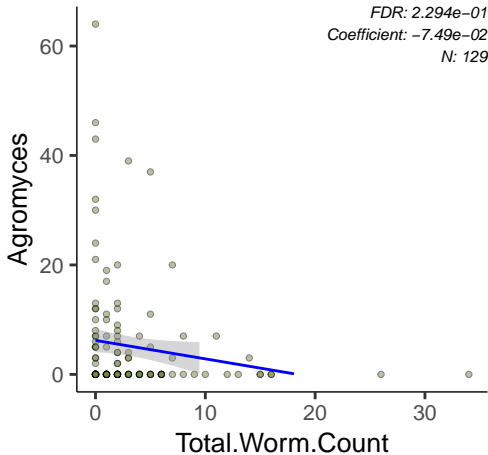
FDR: 1.987e-01
Coefficient: 1.02e-01
N: 129











Bradyrhizobium

FDR: 2.294e-01
Coefficient: 8.98e-02
N: 129

600

400

200

0

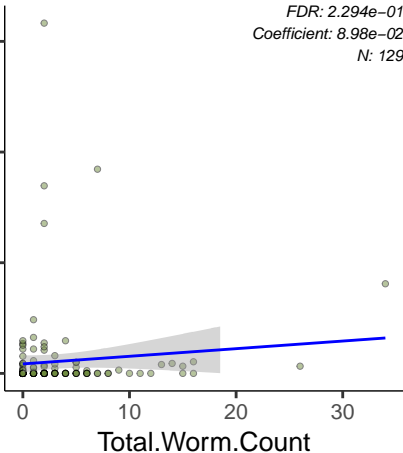
0

10

20

30

Total.Worm.Count



Chthoniobacter

FDR: 2.324e-01
Coefficient: 9.14e-02
N: 129

150

100

50

0

0

Total.Worm.Count

20

30

