

Atopobiaceae_sp.____HG3A.0704

Atopobiaceae_sp.____HG3A.0704

0.6

0.4

0.2

0.0

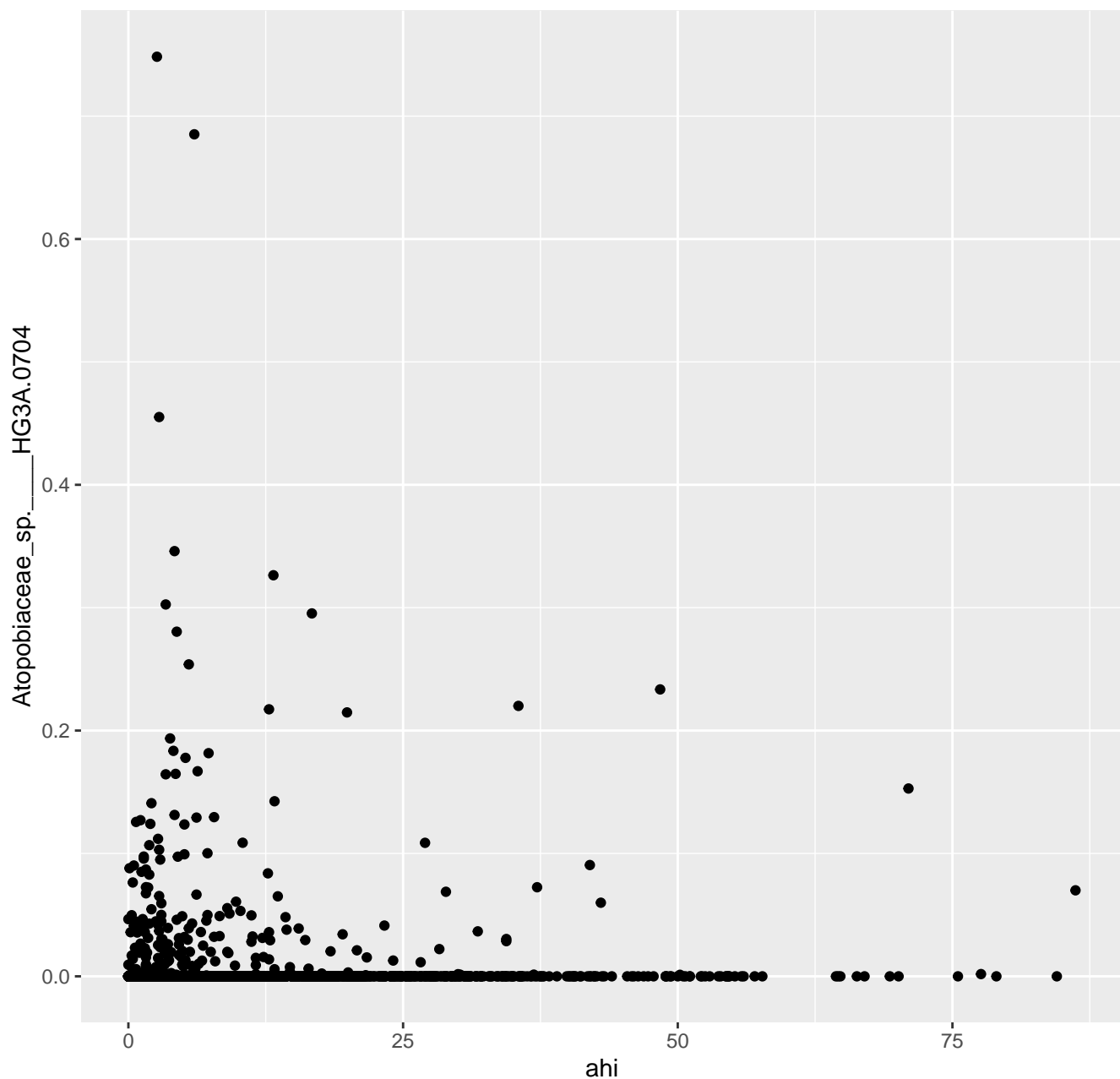
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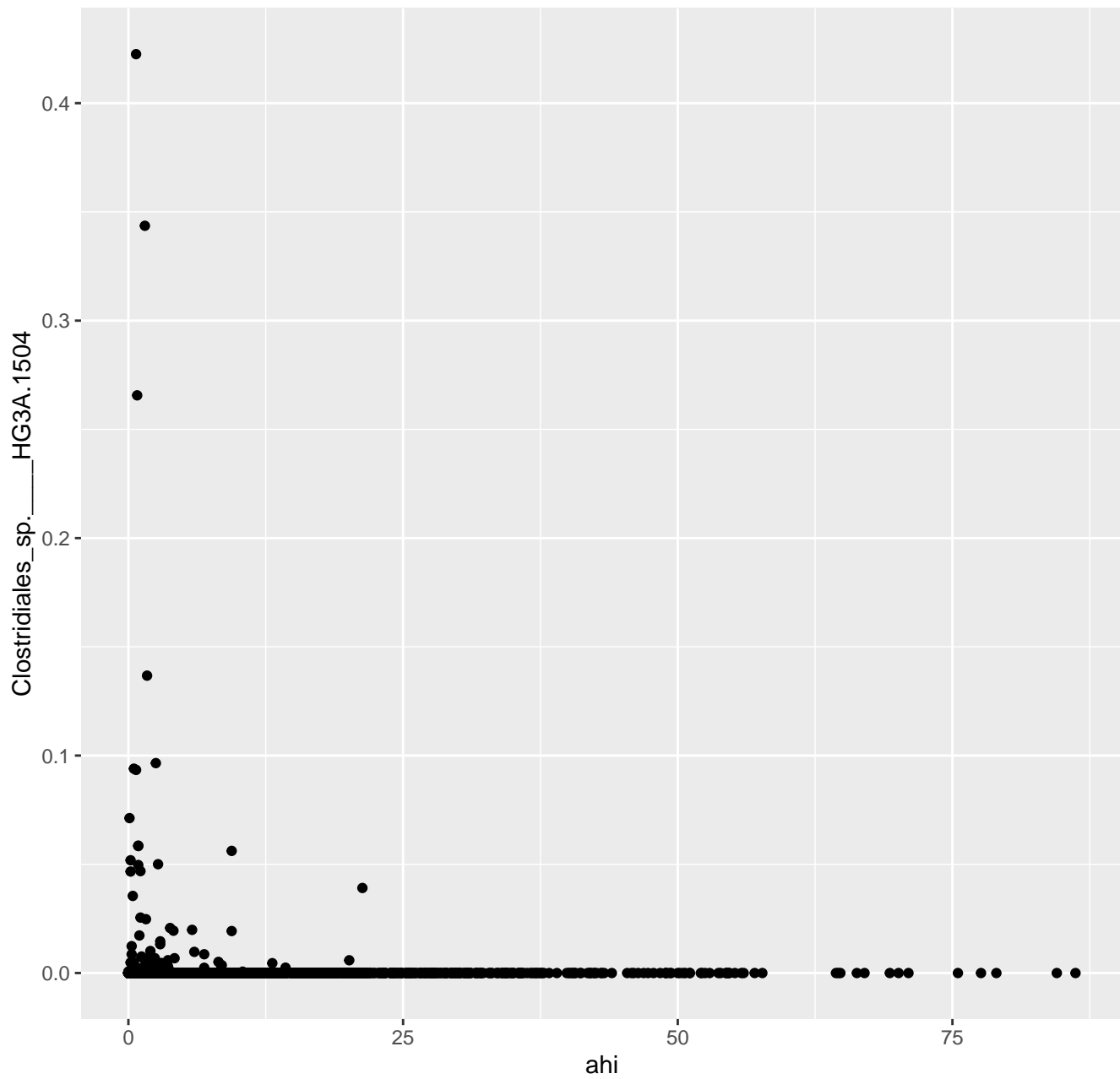
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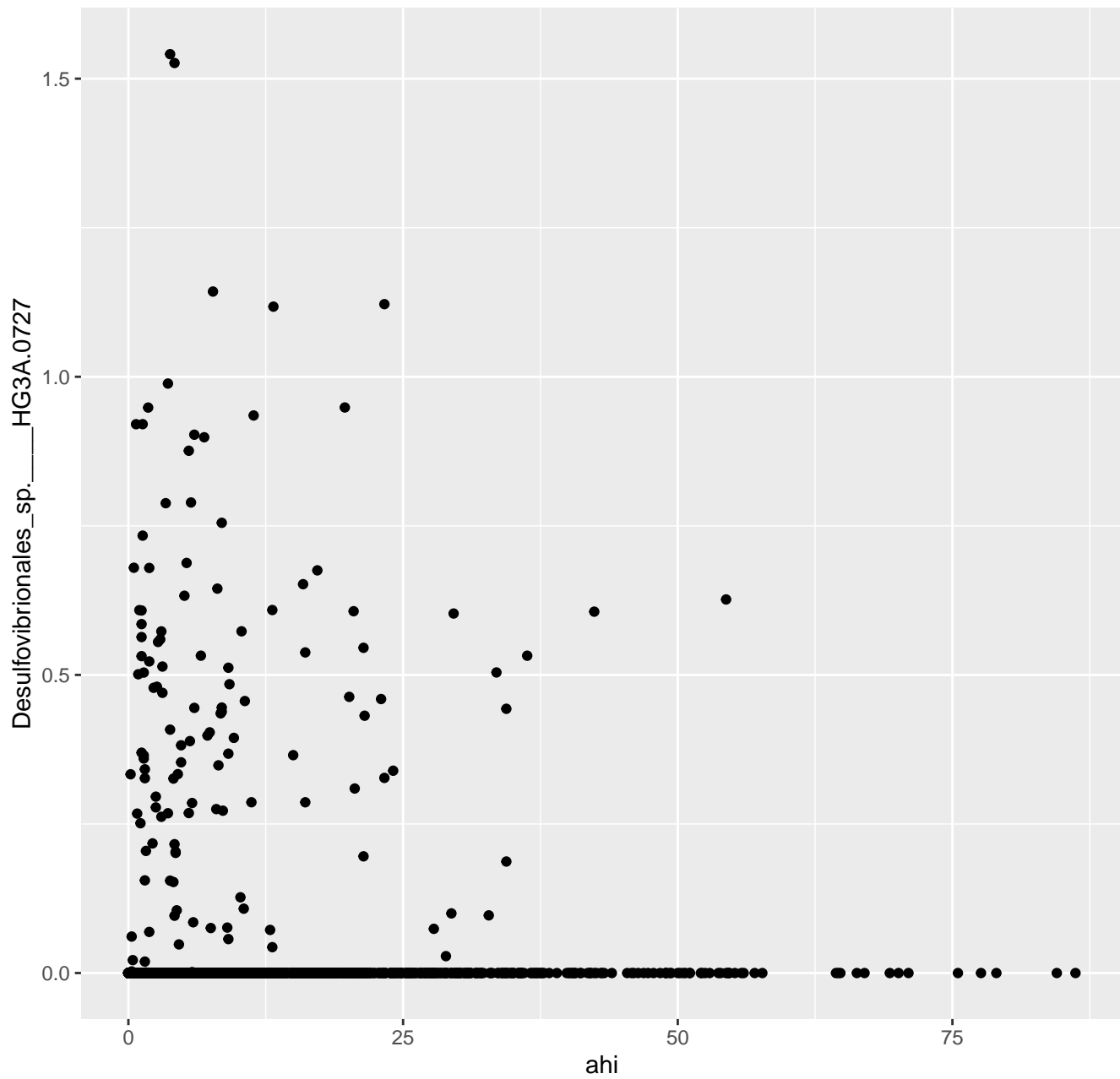
ahi



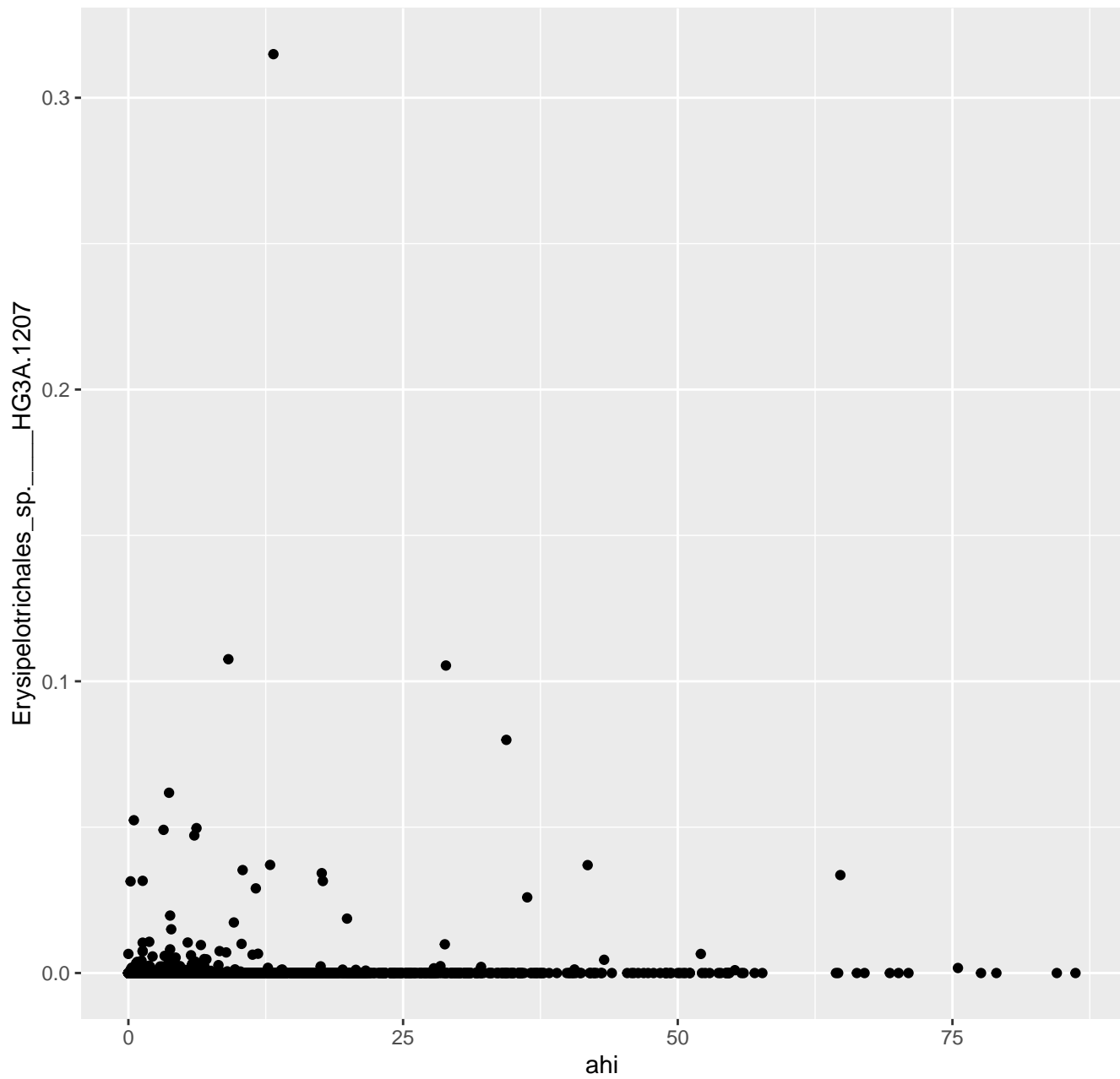
Clostridiales_sp.____HG3A.1504



Desulfovibrionales_sp.____HG3A.0727



Erysipelotrichales_sp.____HG3A.1207



Eubacteriales_sp.____HG3A.0829

Eubacteriales_sp.____HG3A.0829

2.0

1.0

0.5

0.0

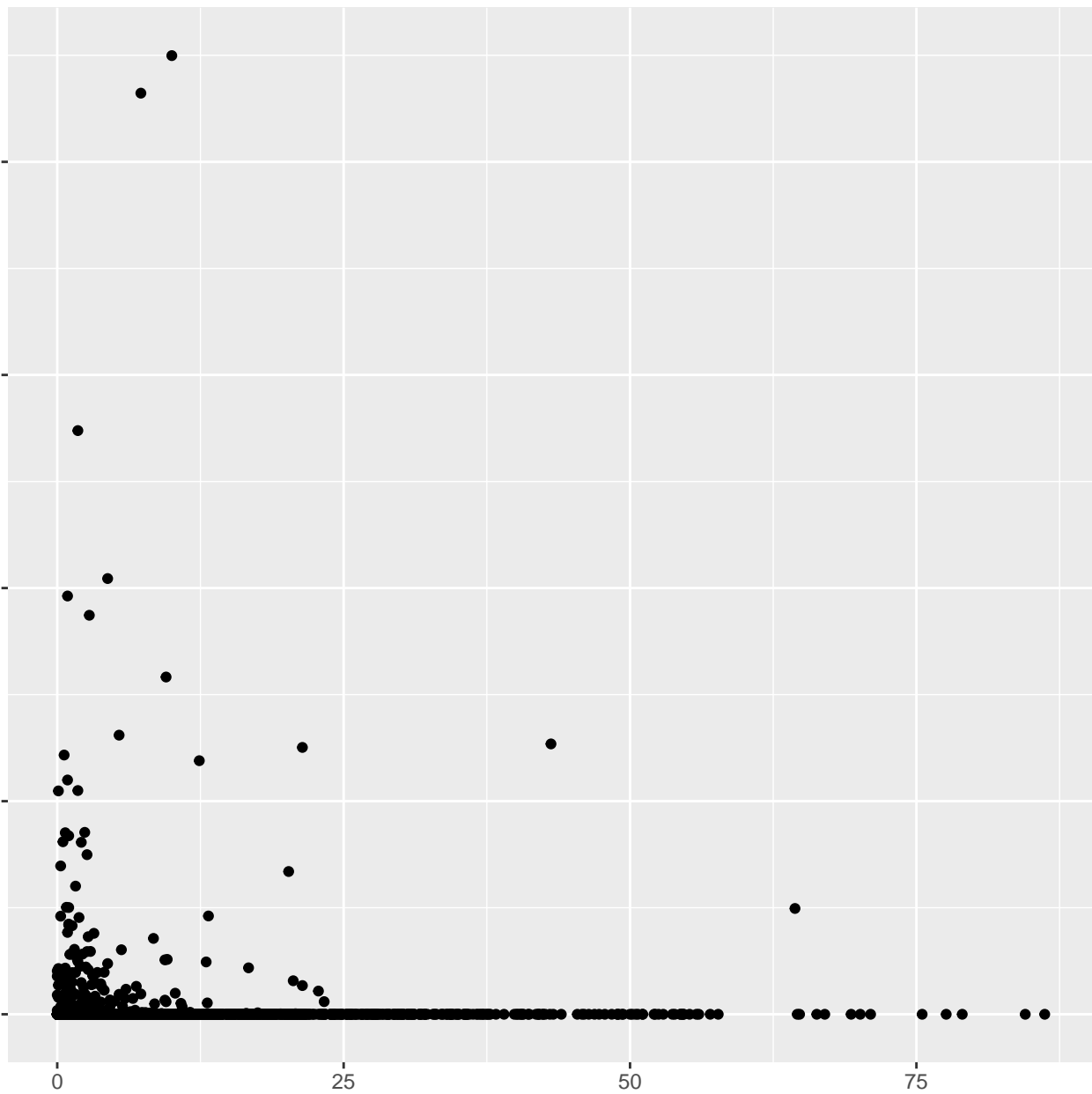
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ahi



Eubacteriales_sp.____HG3A.0964

Eubacteriales_sp.____HG3A.0964

0.3

0.2

0.1

0.0

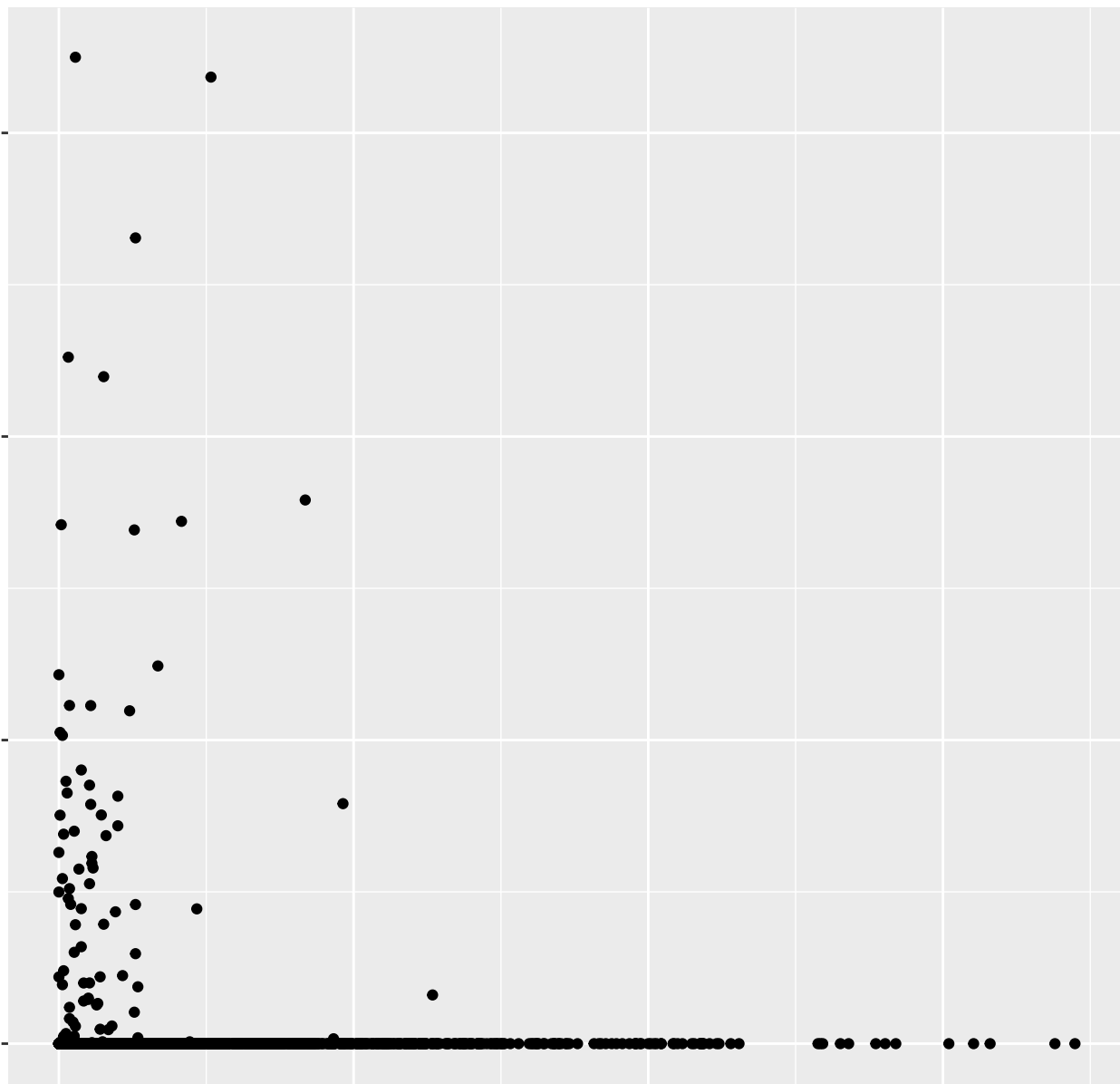
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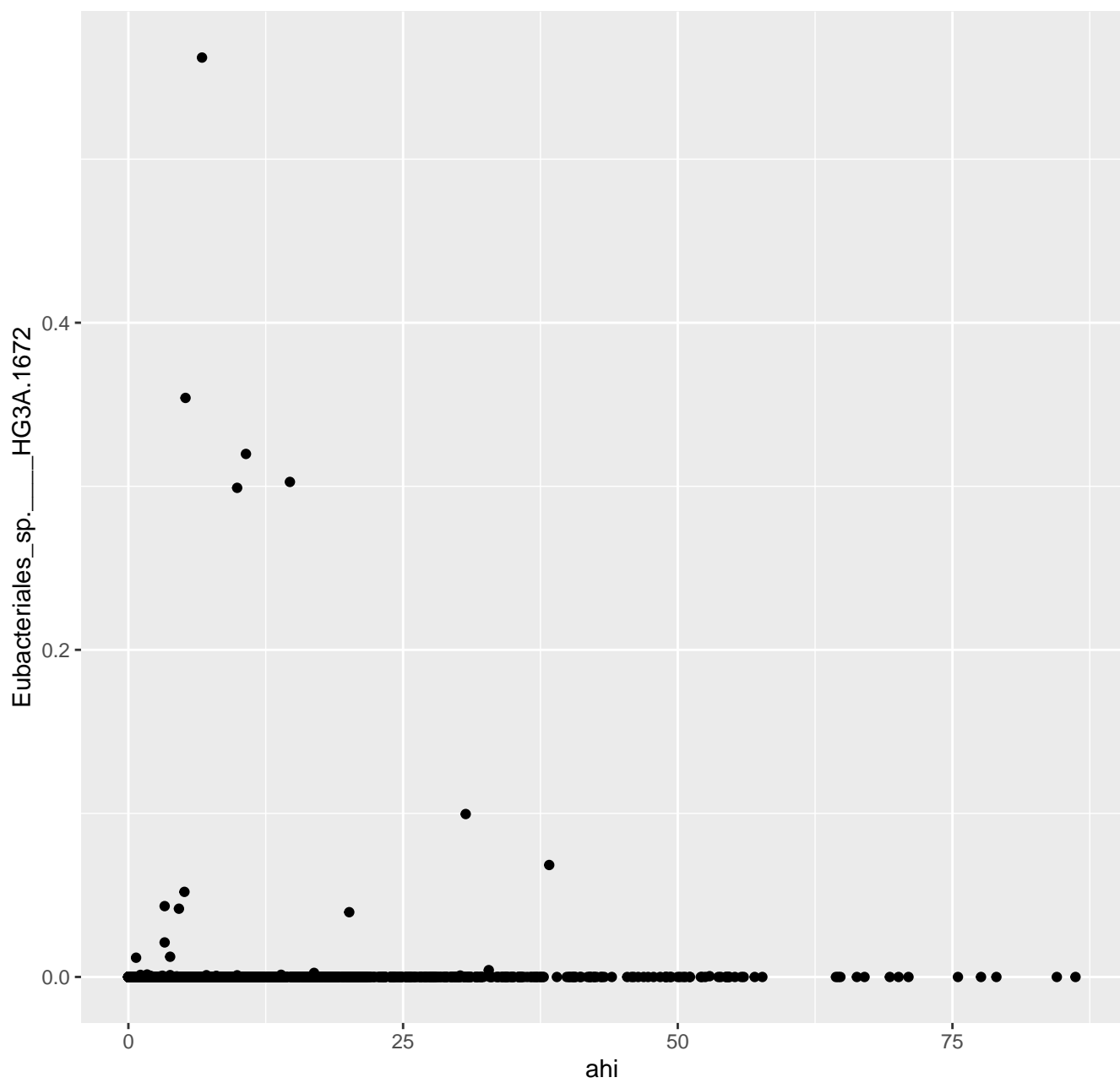
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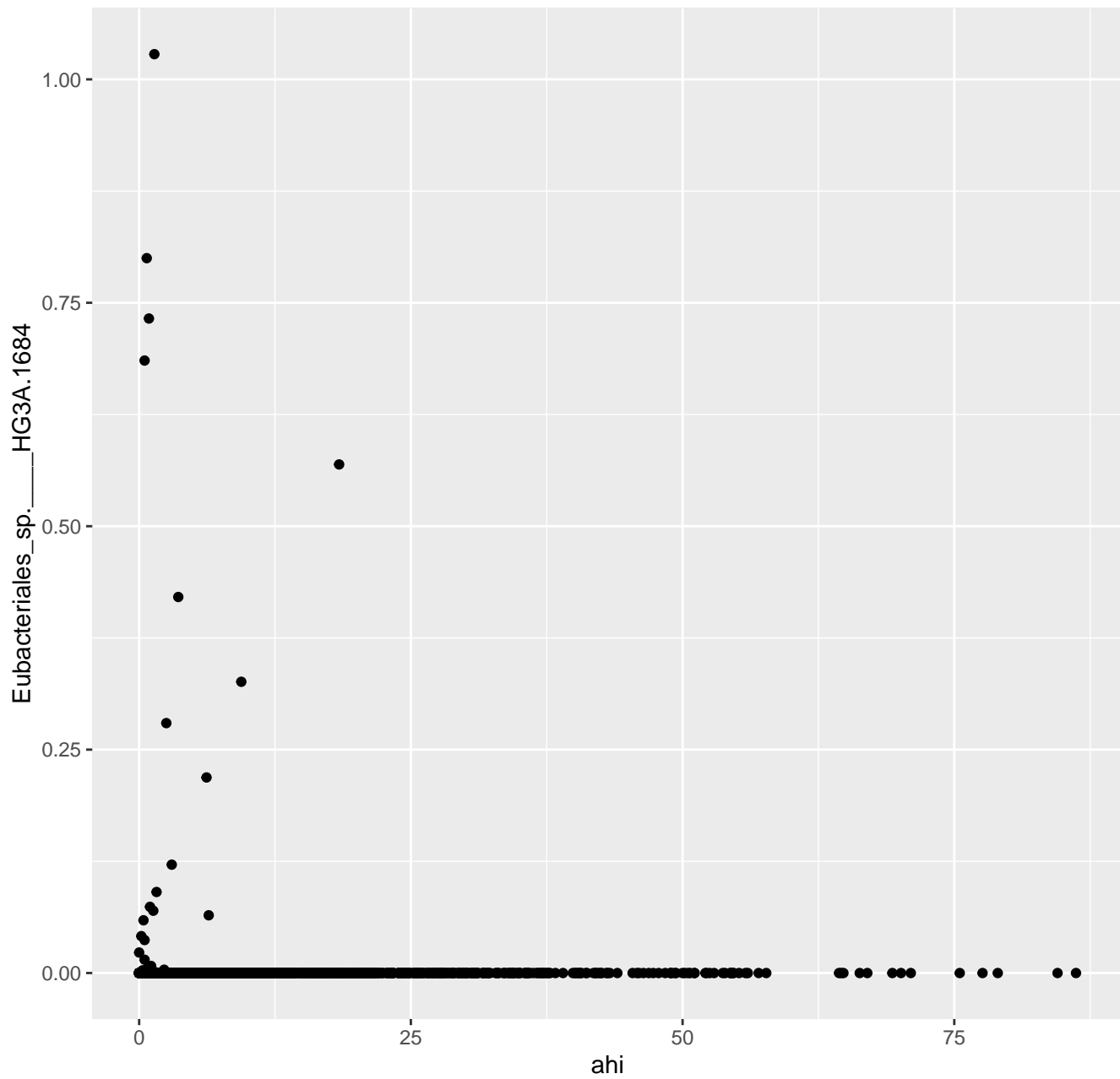
ahi



Eubacteriales_sp.____HG3A.1672



Eubacteriales_sp.____HG3A.1684



Firmicutes_sp.____HG3A.0398

Firmicutes_sp.____HG3A.0398

1.5

1.0

0.5

0.0

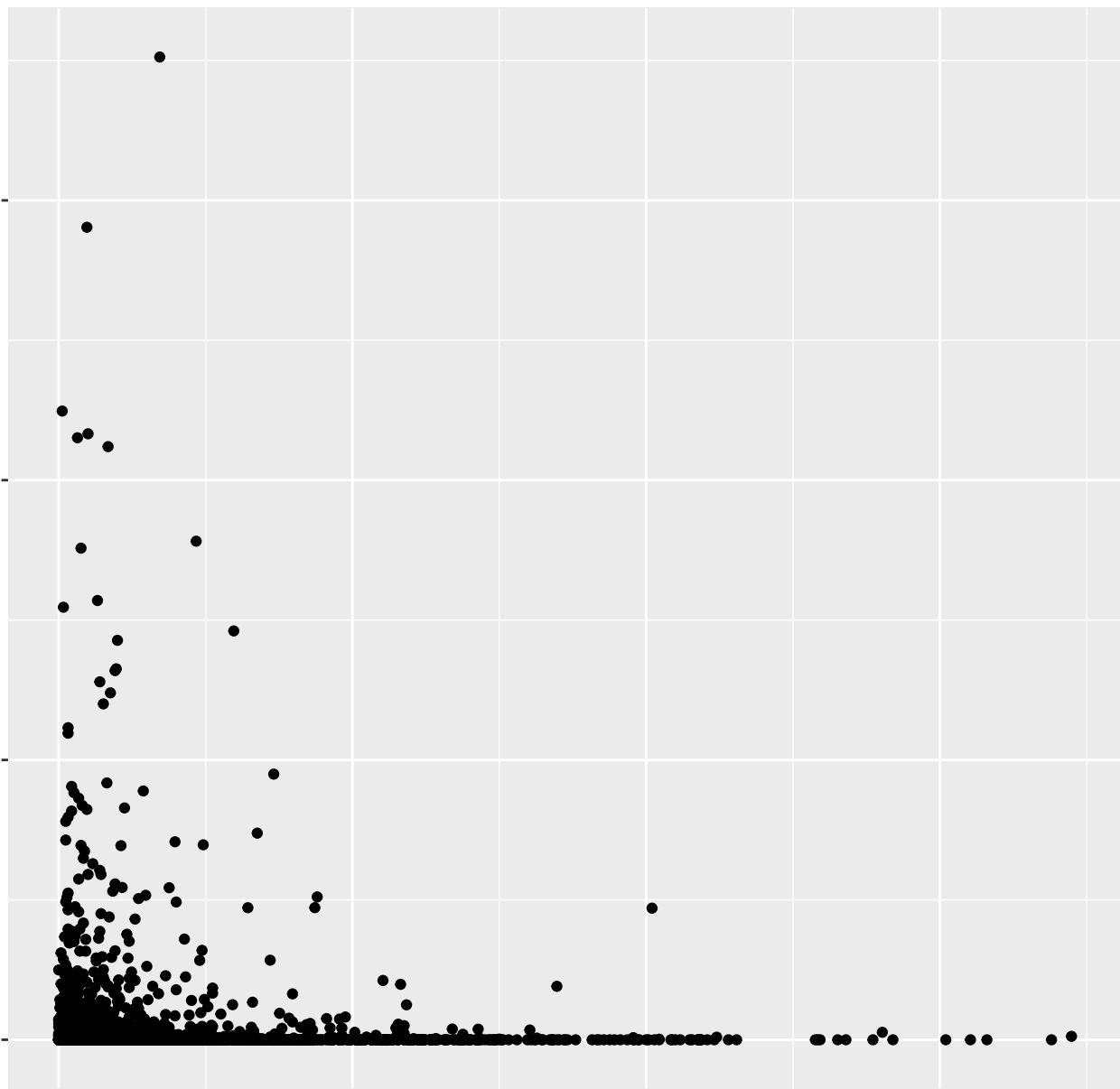
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ahi



Firmicutes_sp.____HG3A.0681

Firmicutes_sp.____HG3A.0681

