



ADDITIONAL MATERIALS

Research Report 237

Early-Life Air Pollution Exposure Is Associated with the Infant Gut Microbiome and Fecal Metabolome in the First Two Years of Life

Tanya L. Alderete et al.

Supplemental File 11

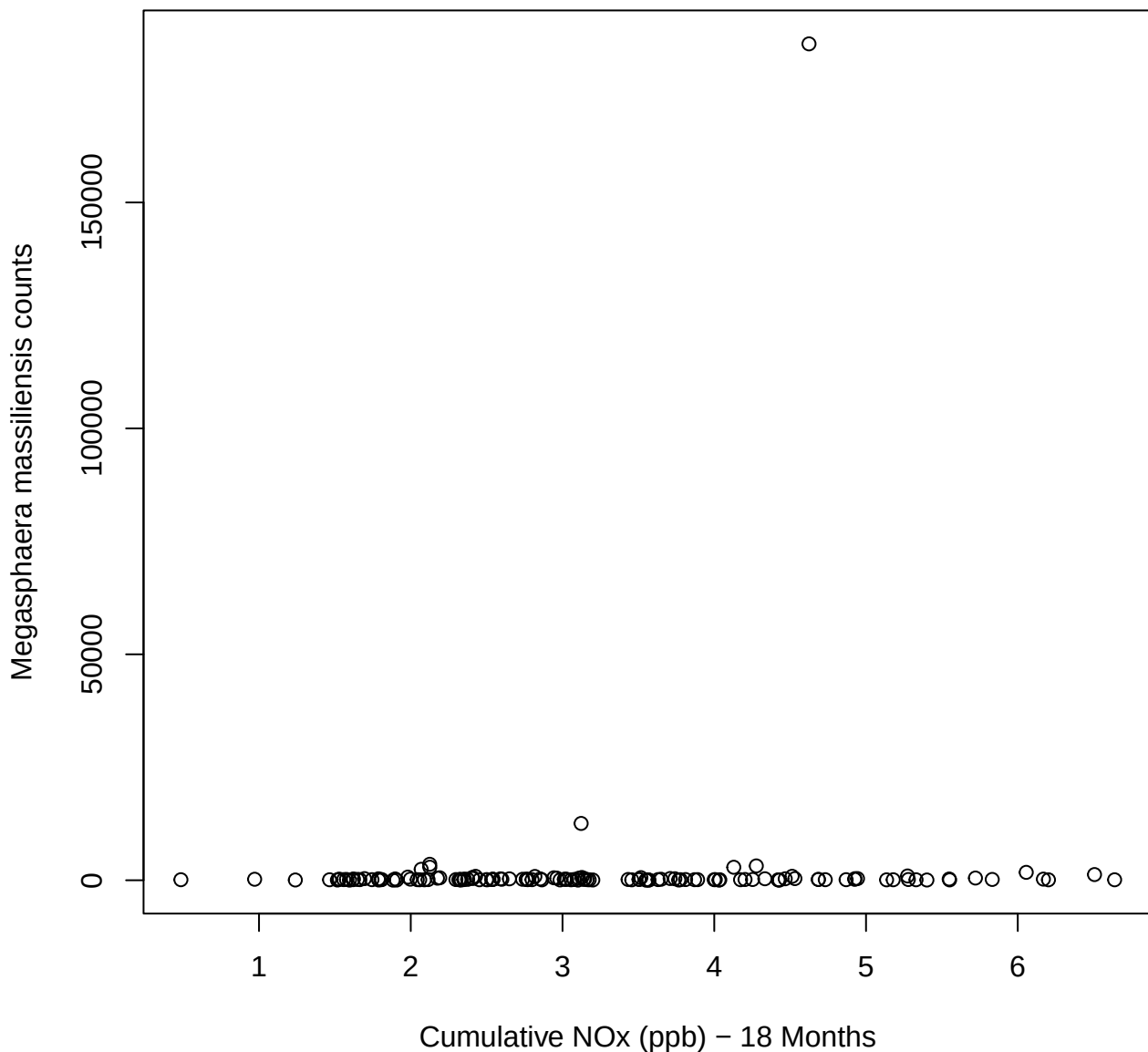
The Additional Materials were reviewed by the HEI Review Committee. They have not been fully edited or formatted by HEI.

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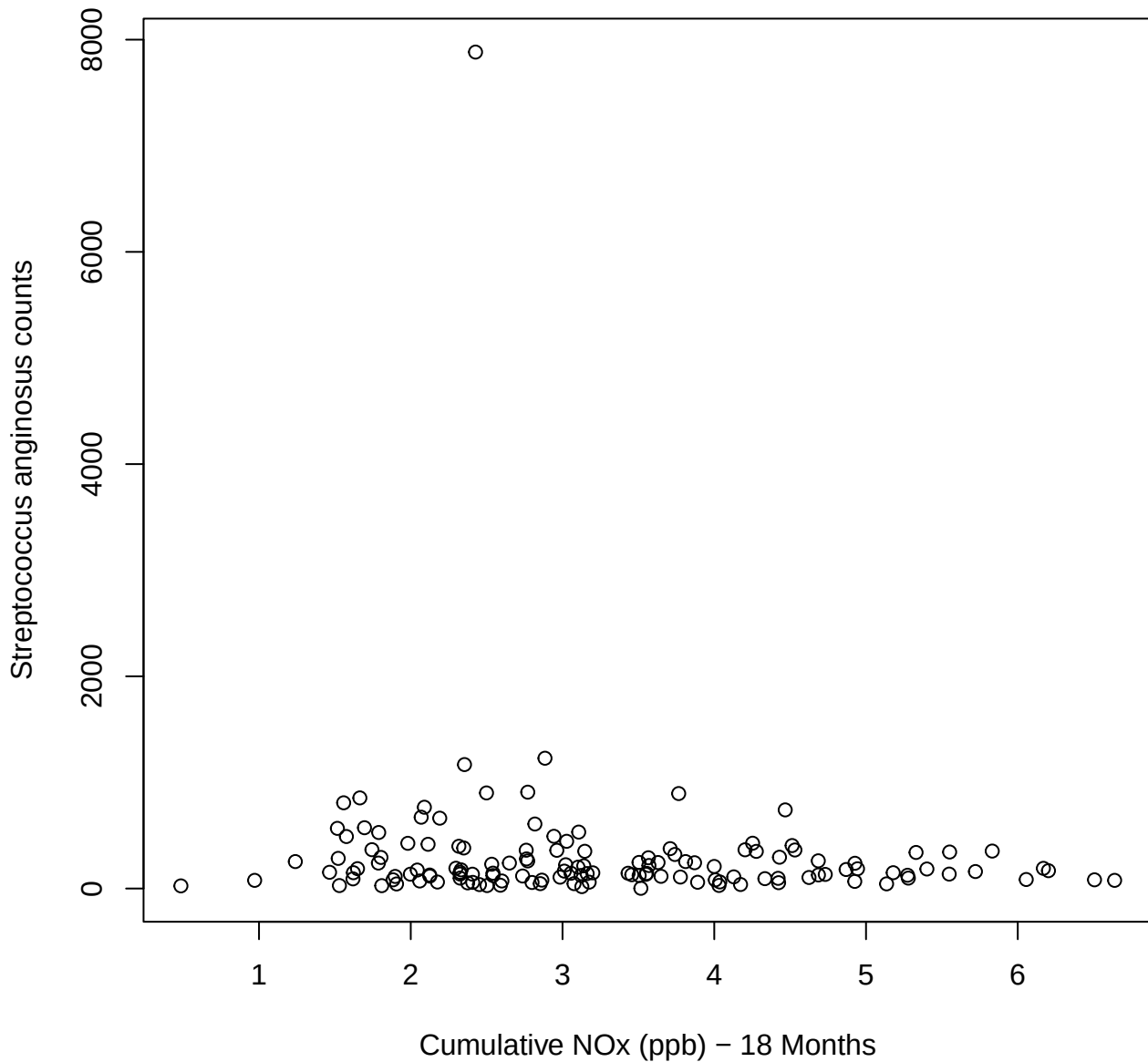
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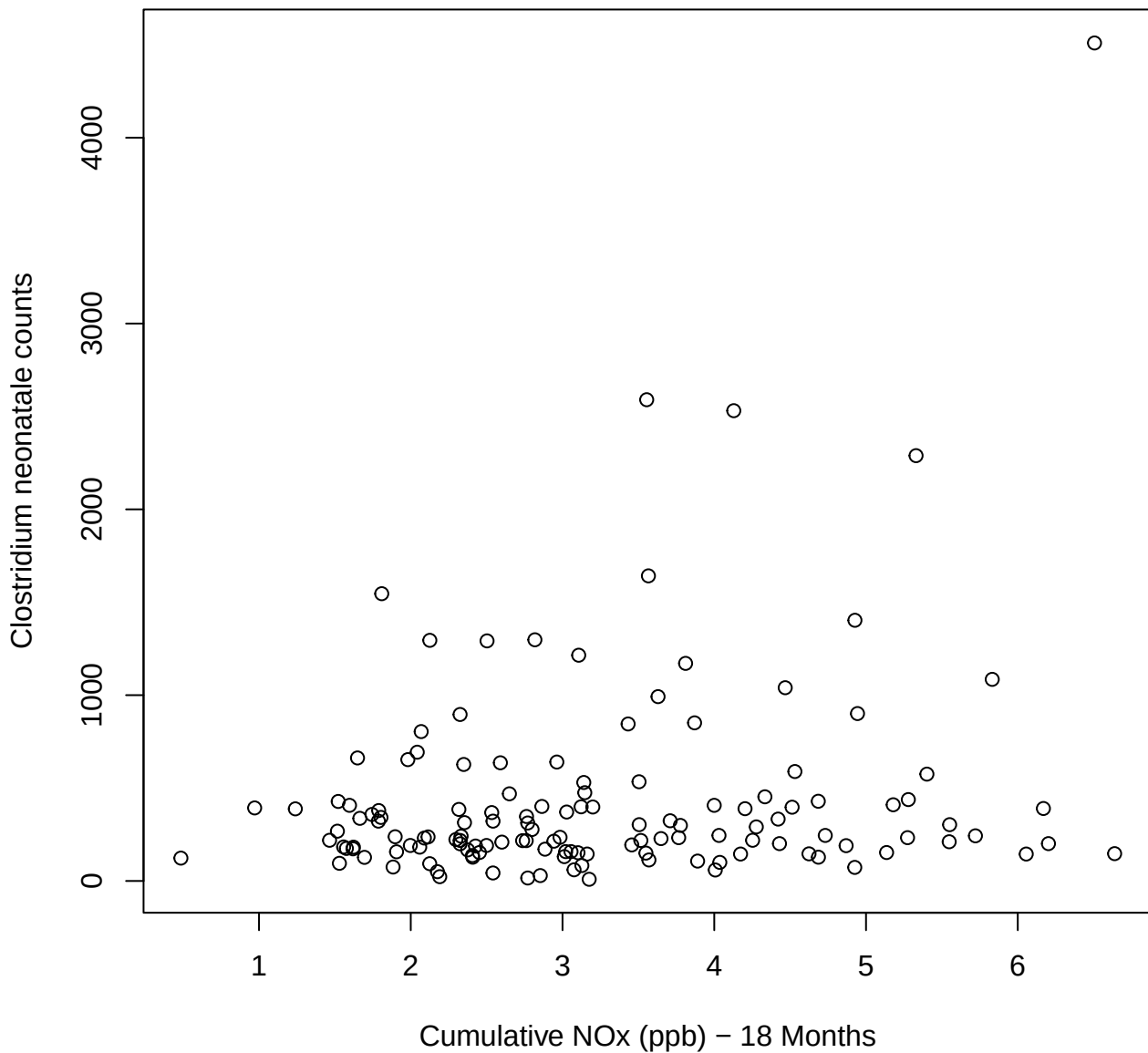
1. *Megasphaera massiliensis*



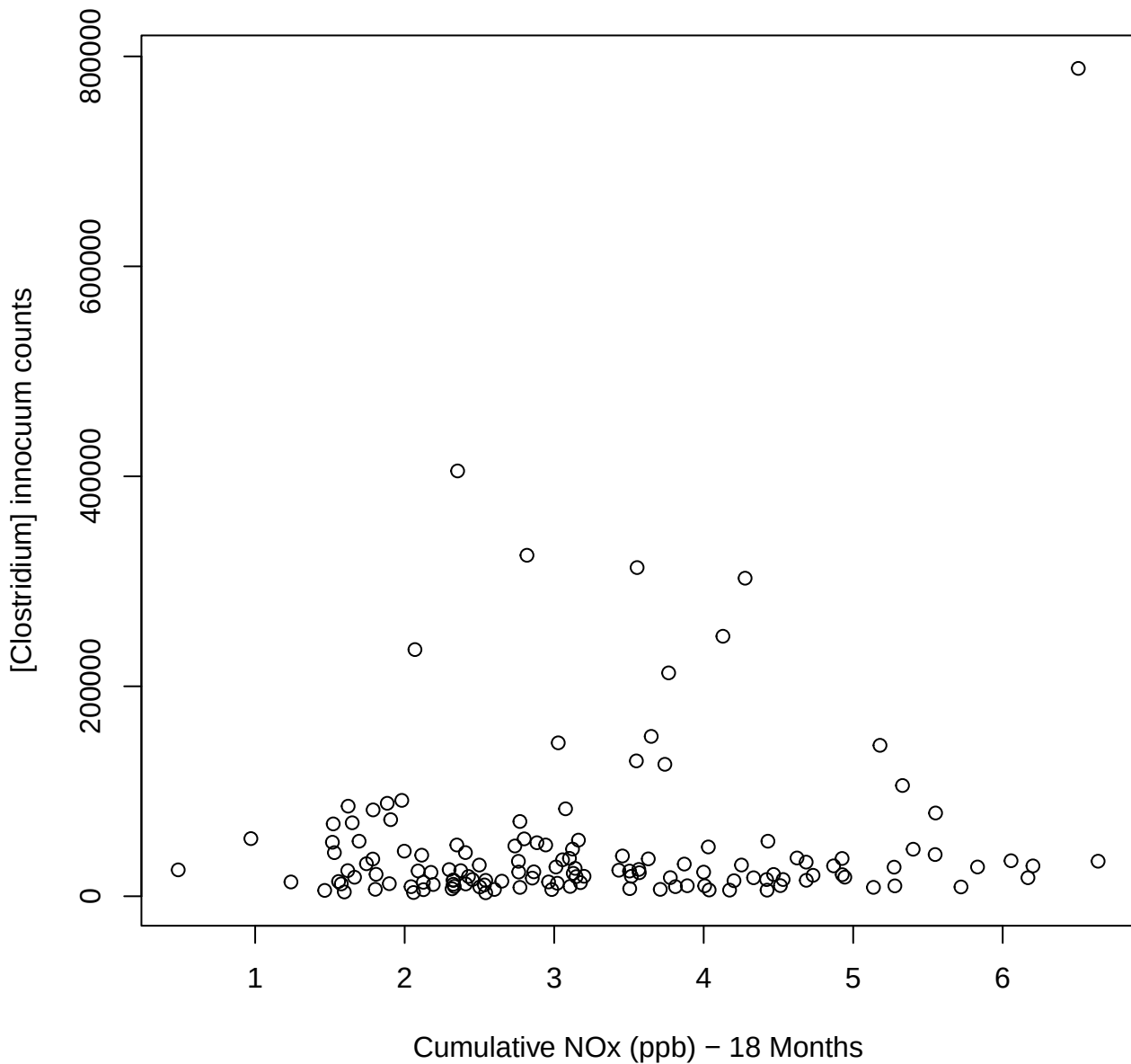
2. Streptococcus anginosus



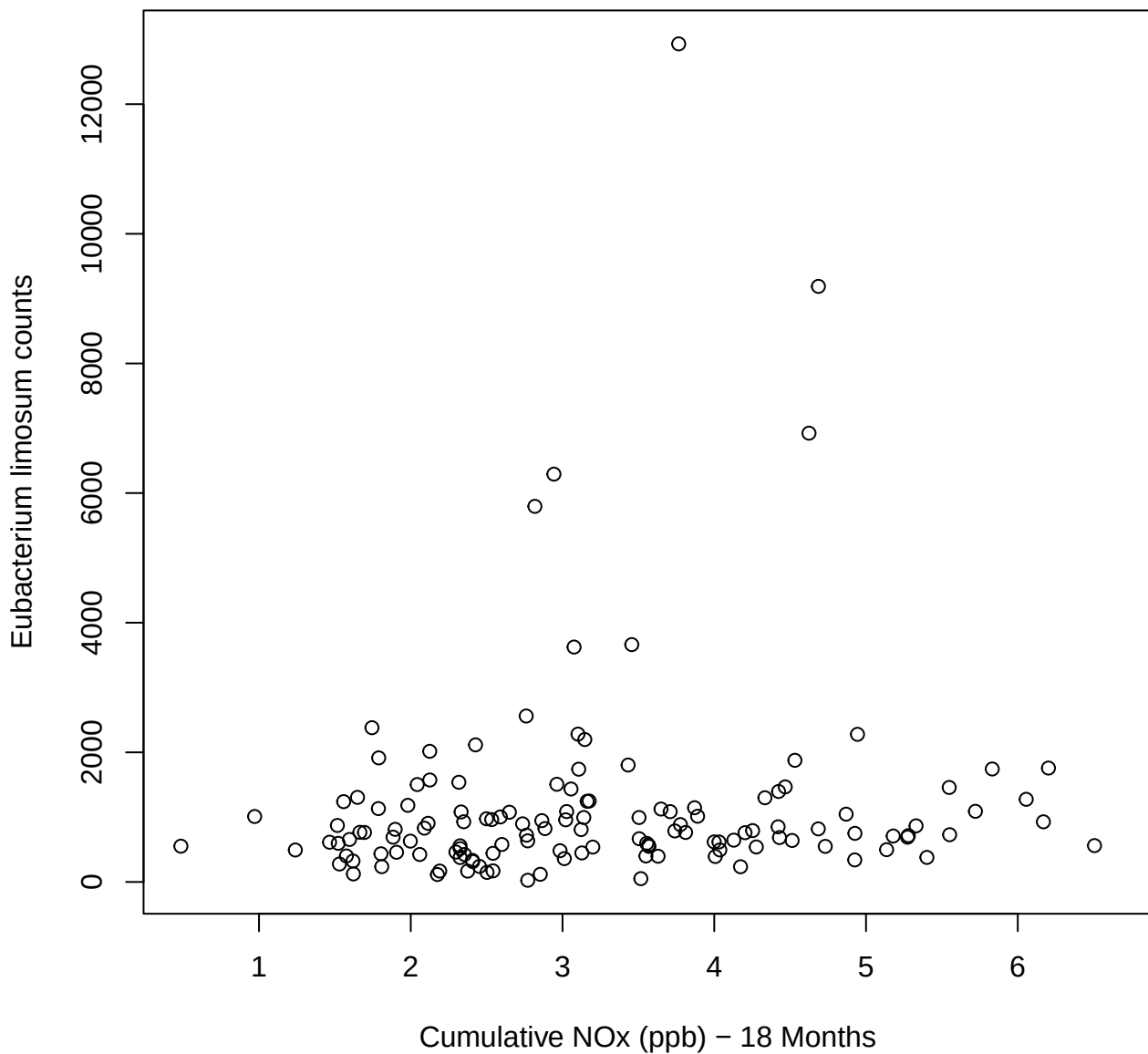
3. Clostridium neonatale



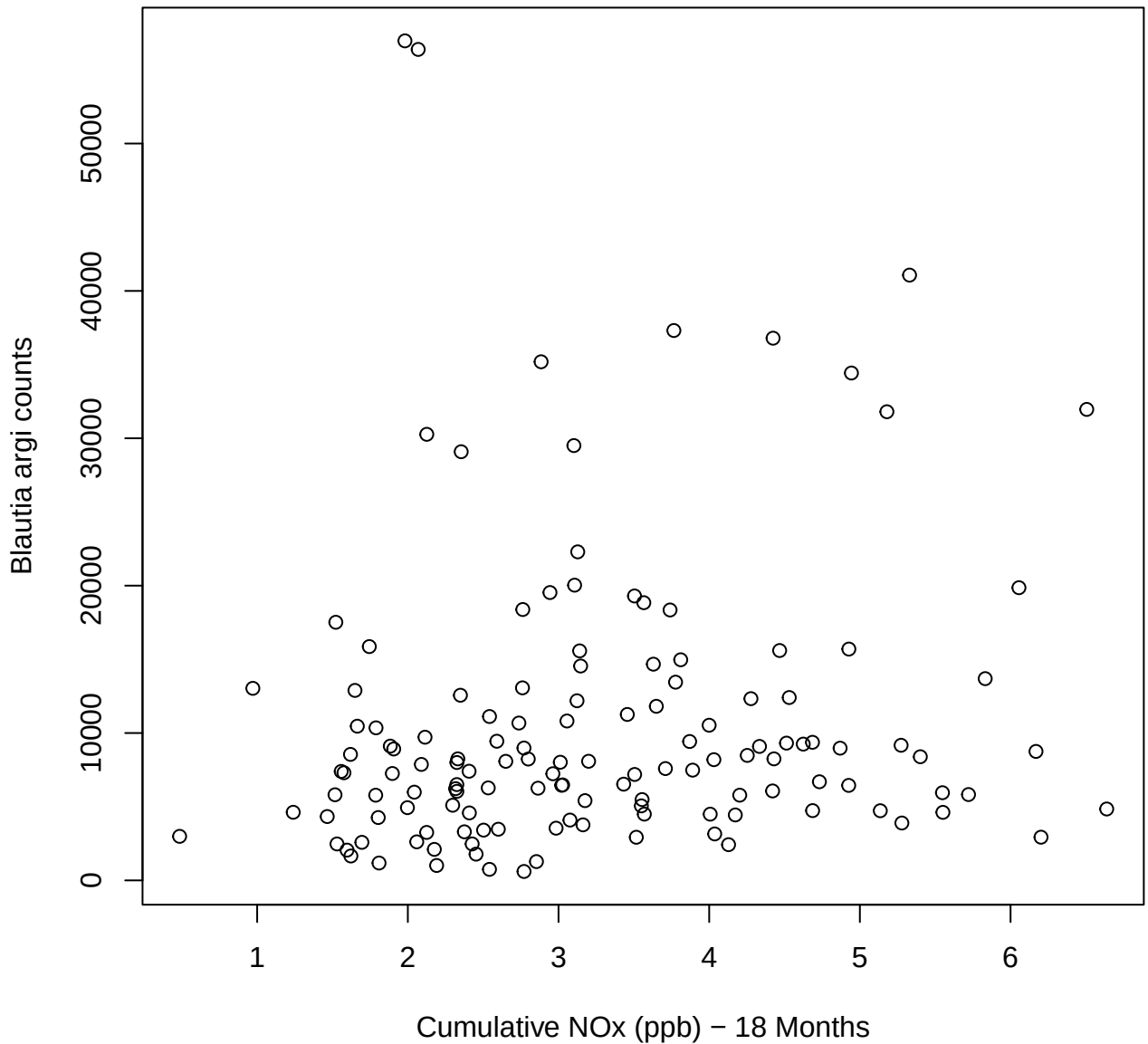
4. [Clostridium] innocuum



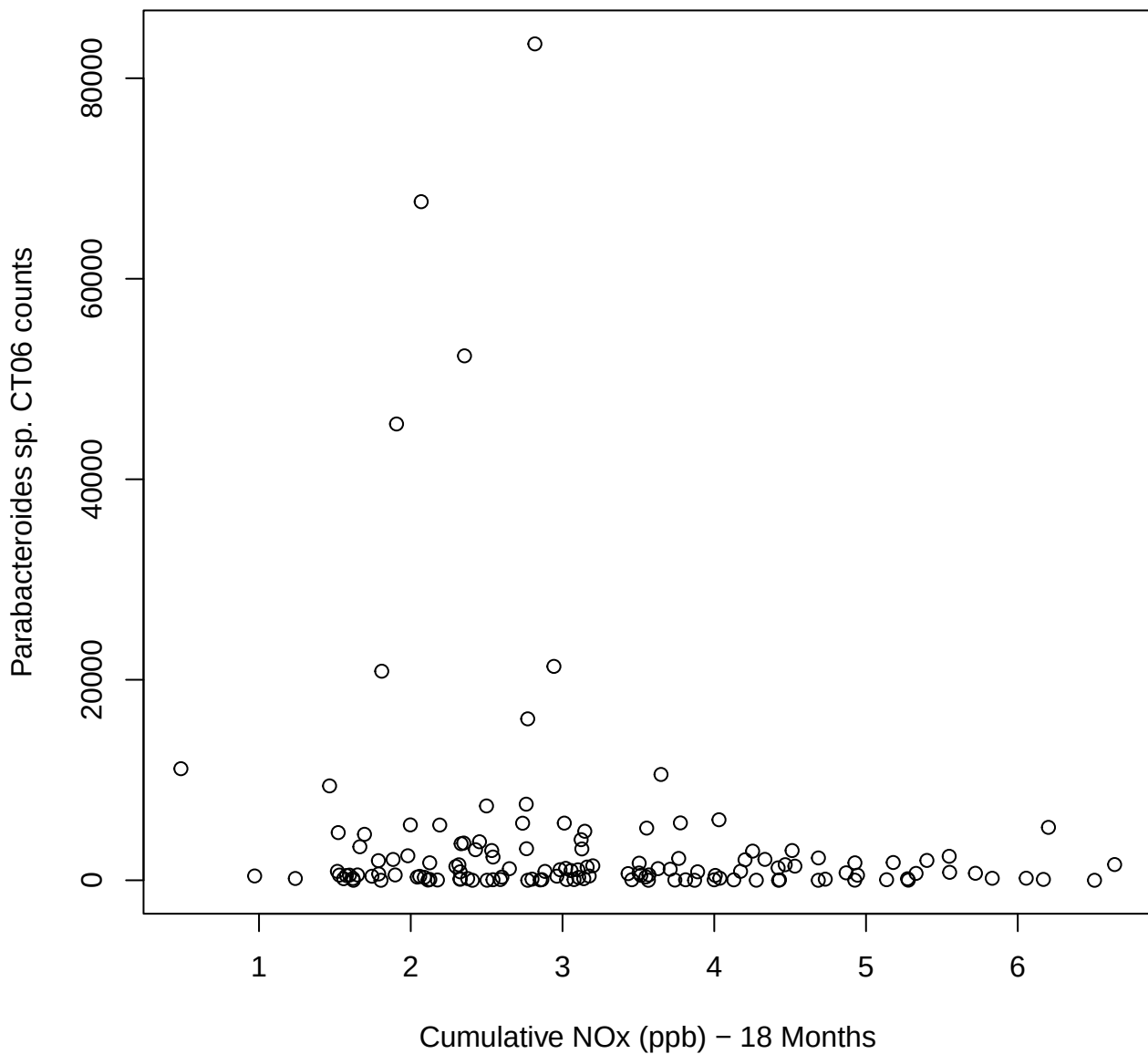
5. *Eubacterium limosum*



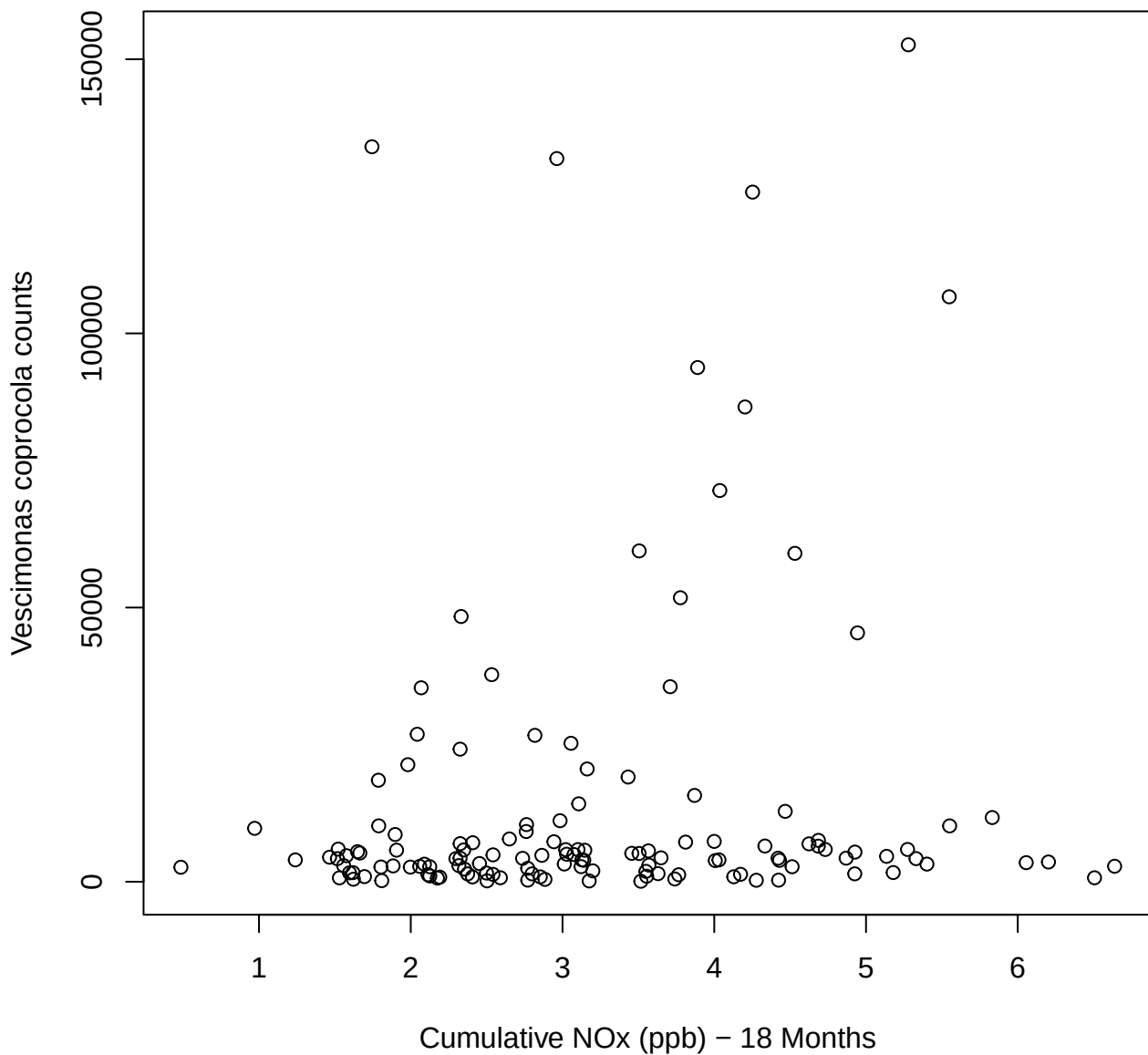
6. *Blautia argi*



7. Parabacteroides sp. CT06



8. *Vescimonas coprocola*

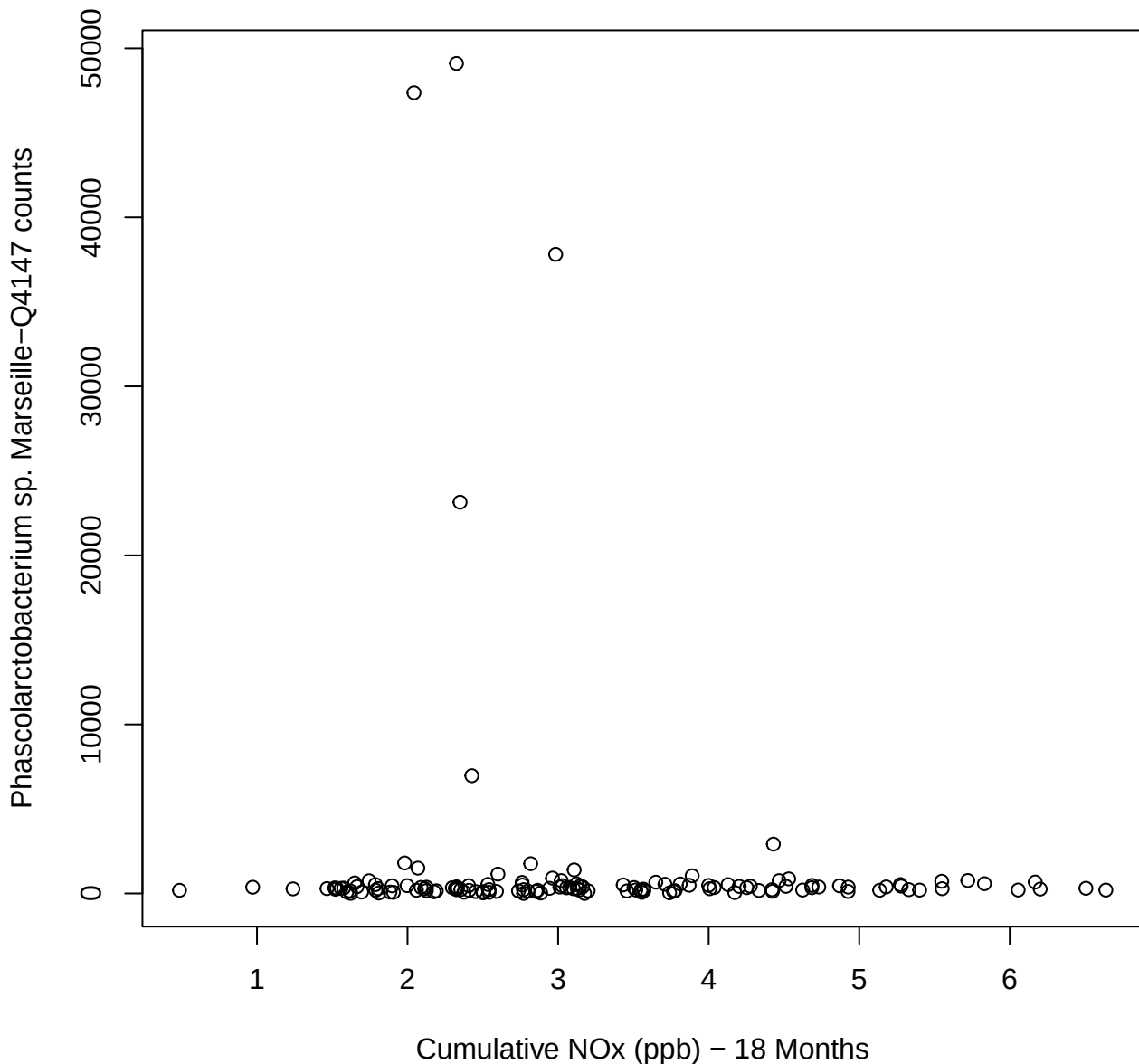


9. Bacteroides ovatus

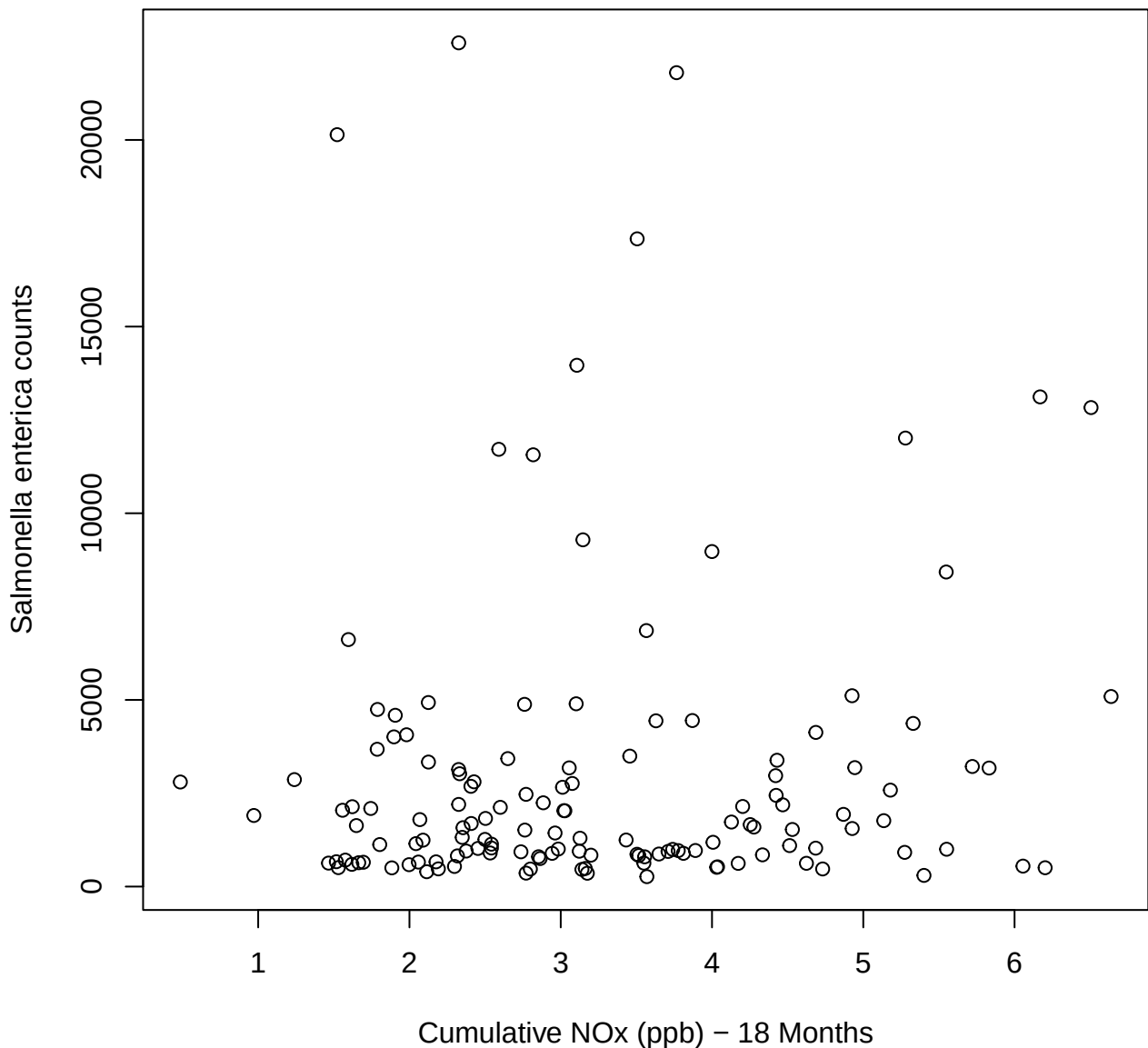


Cumulative NOx (ppb) – 18 Months

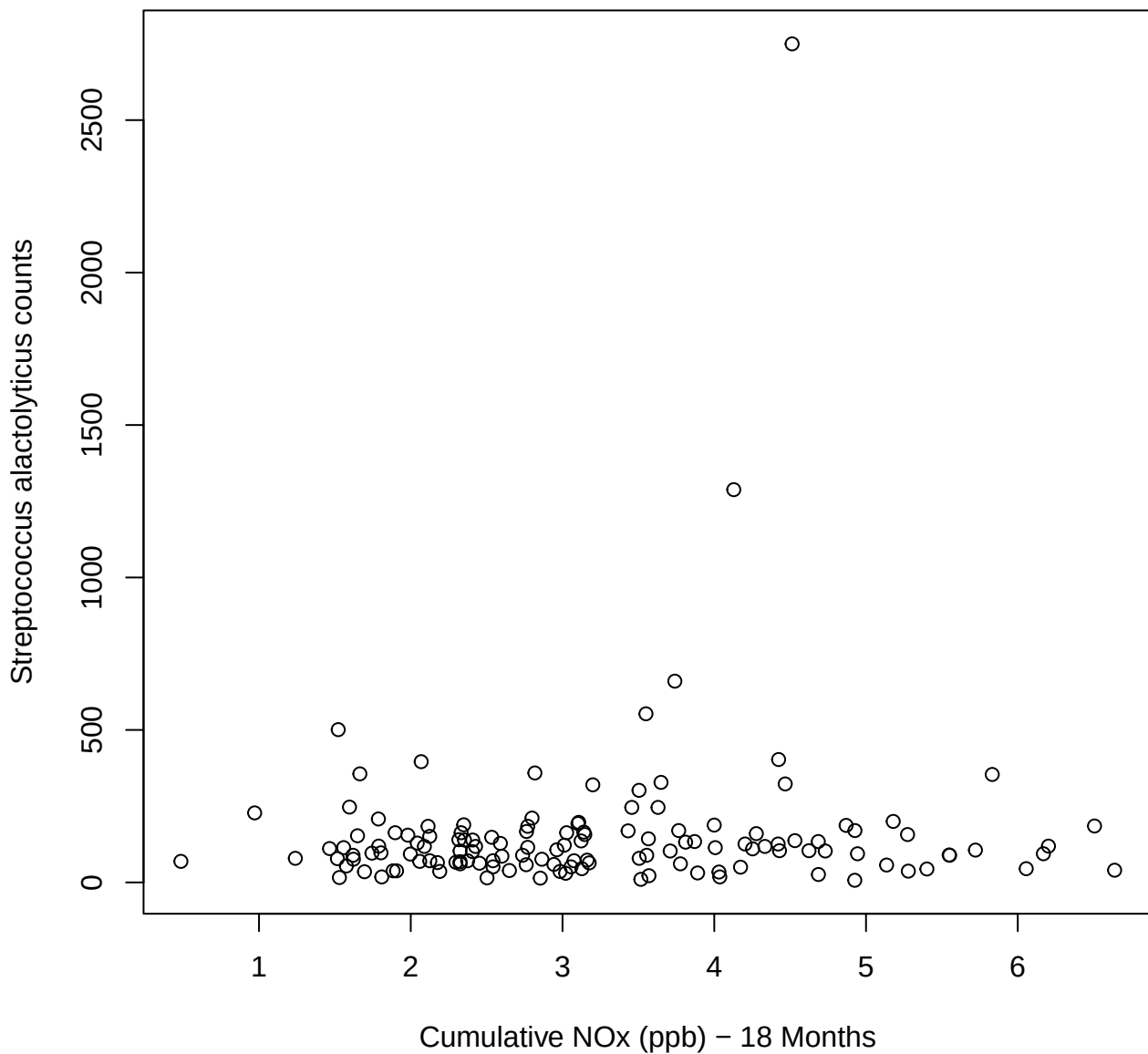
10. *Phascolarctobacterium* sp. Marseille-Q4147



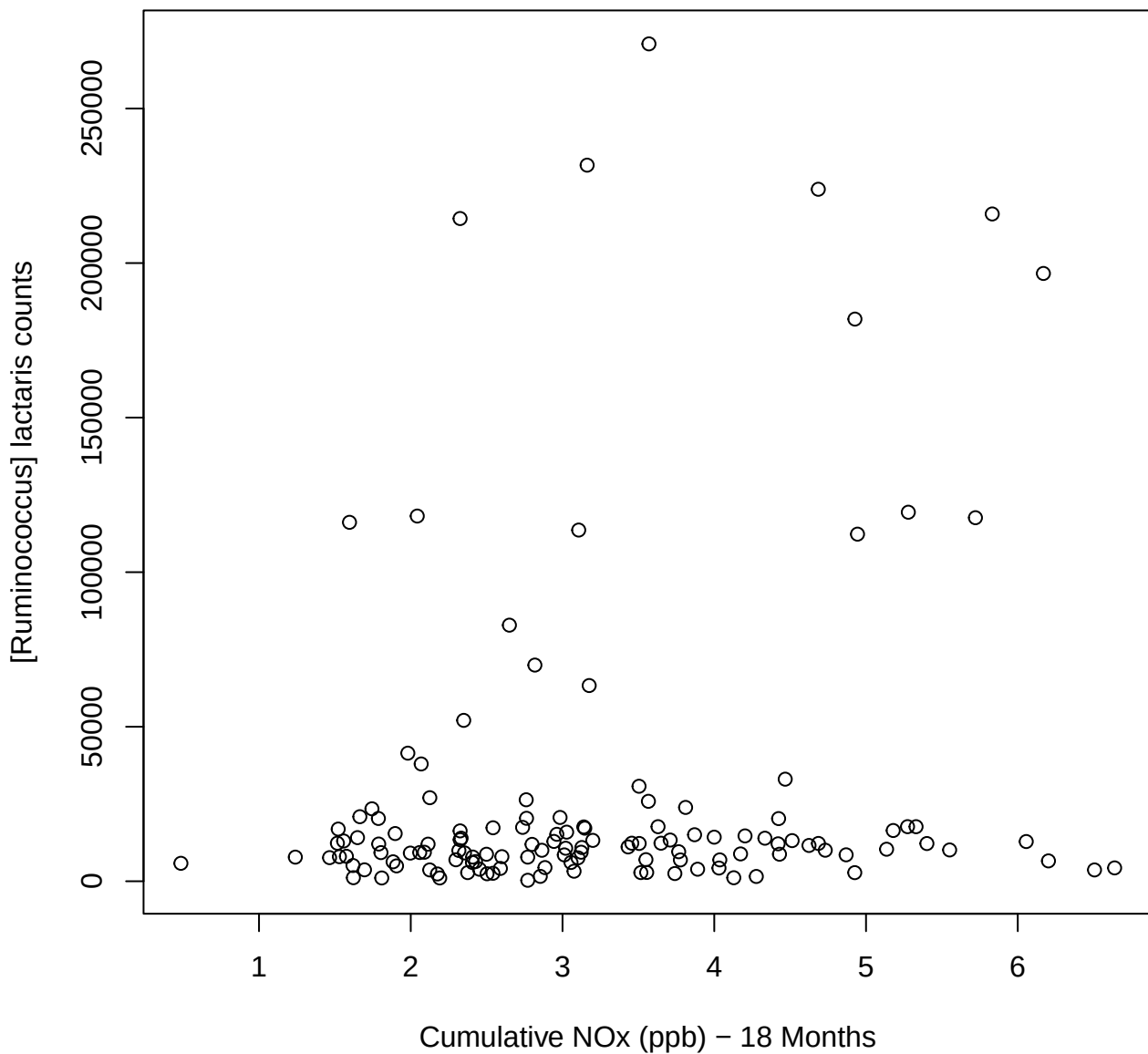
11. *Salmonella enterica*



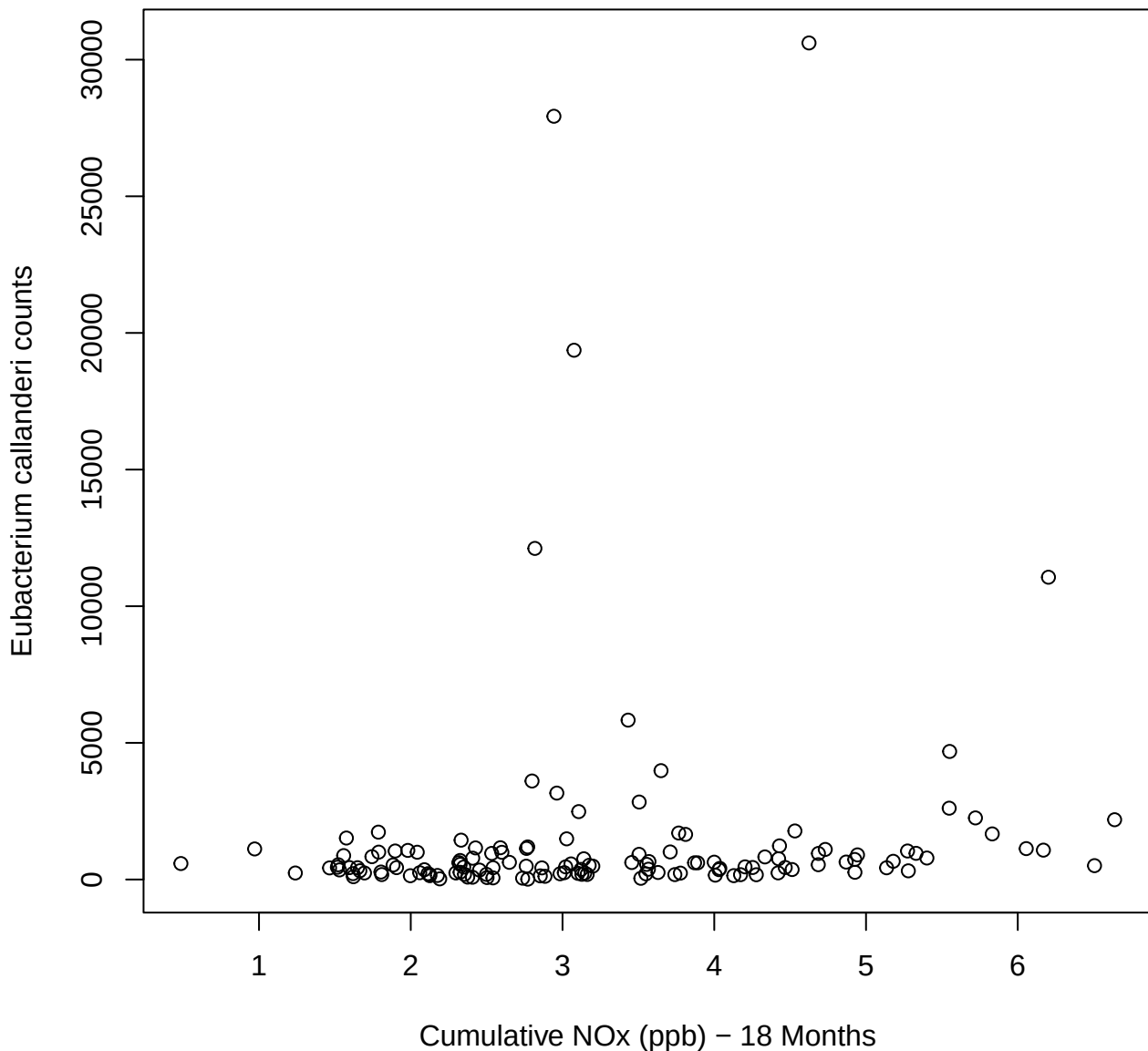
12. Streptococcus alactolyticus



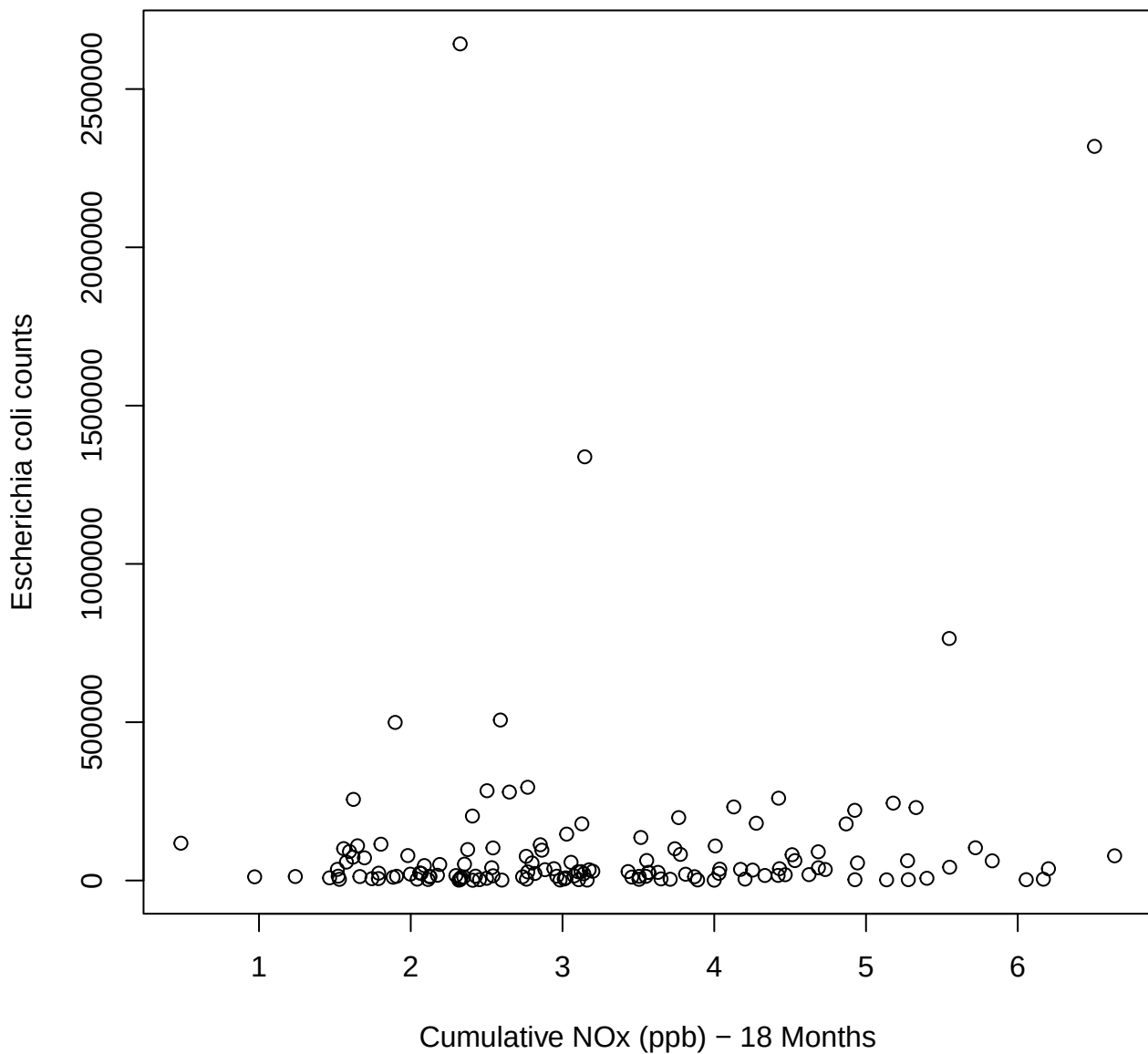
13. [Ruminococcus] lactaris



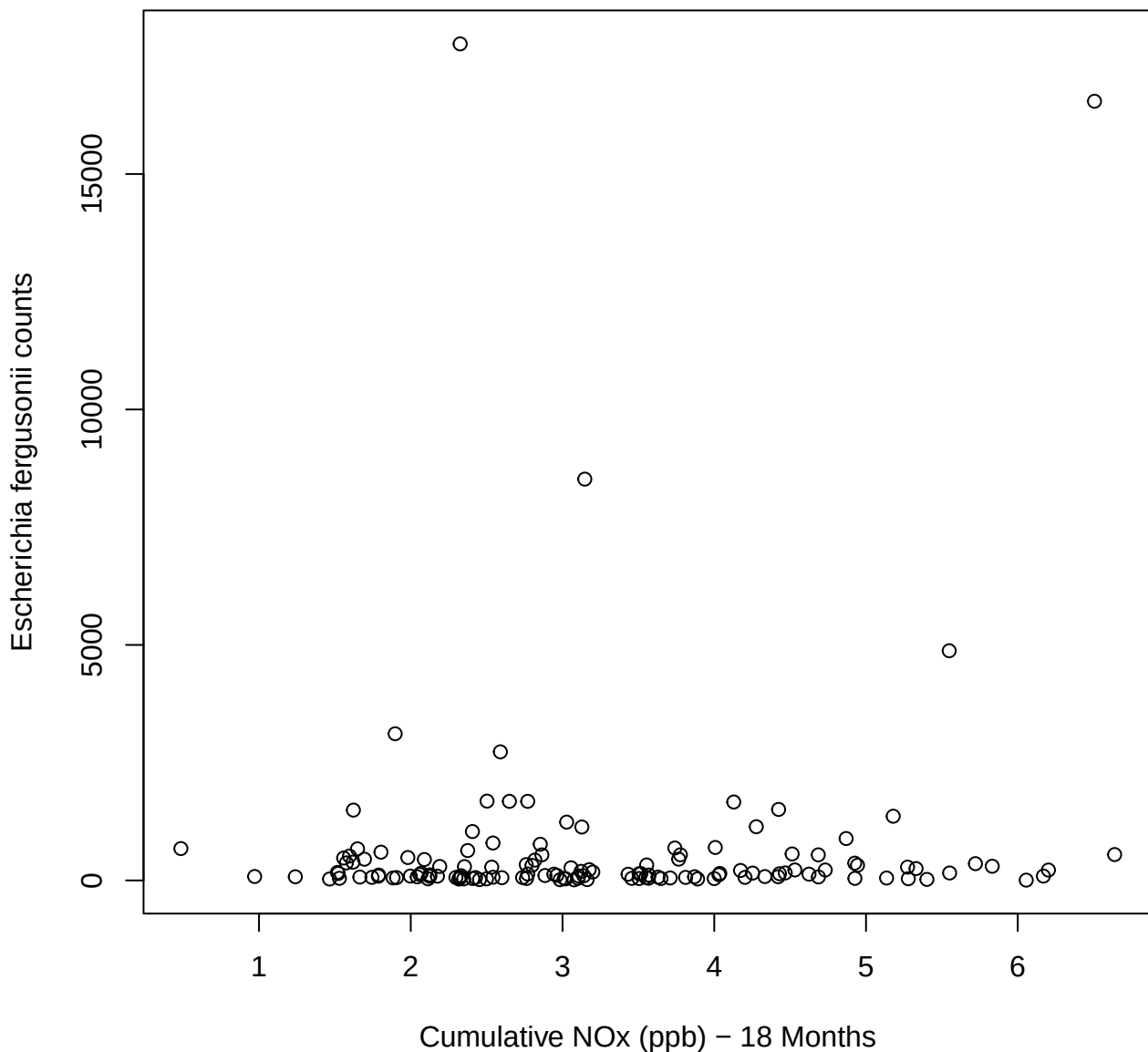
14. *Eubacterium callanderi*



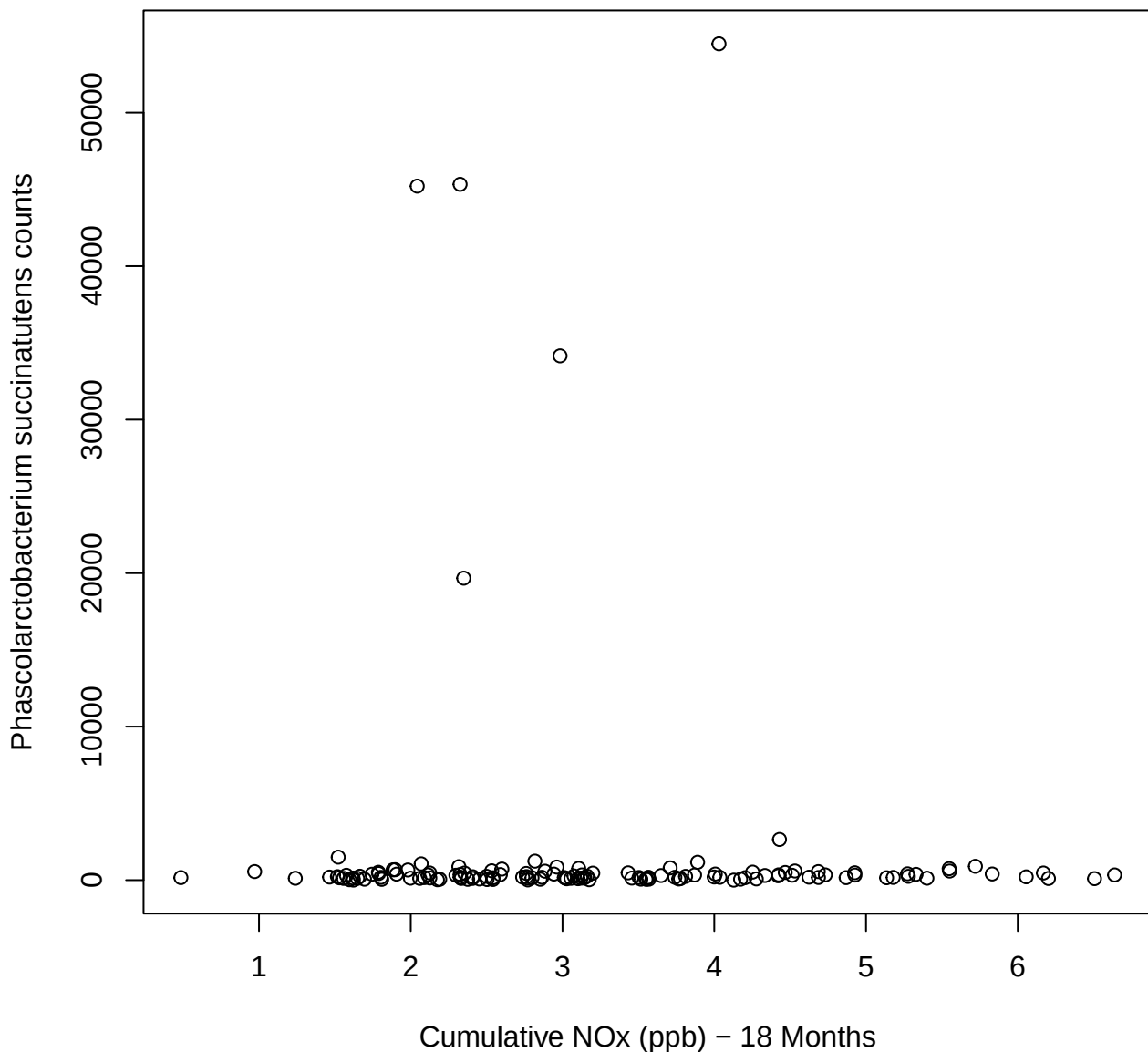
15. Escherichia coli



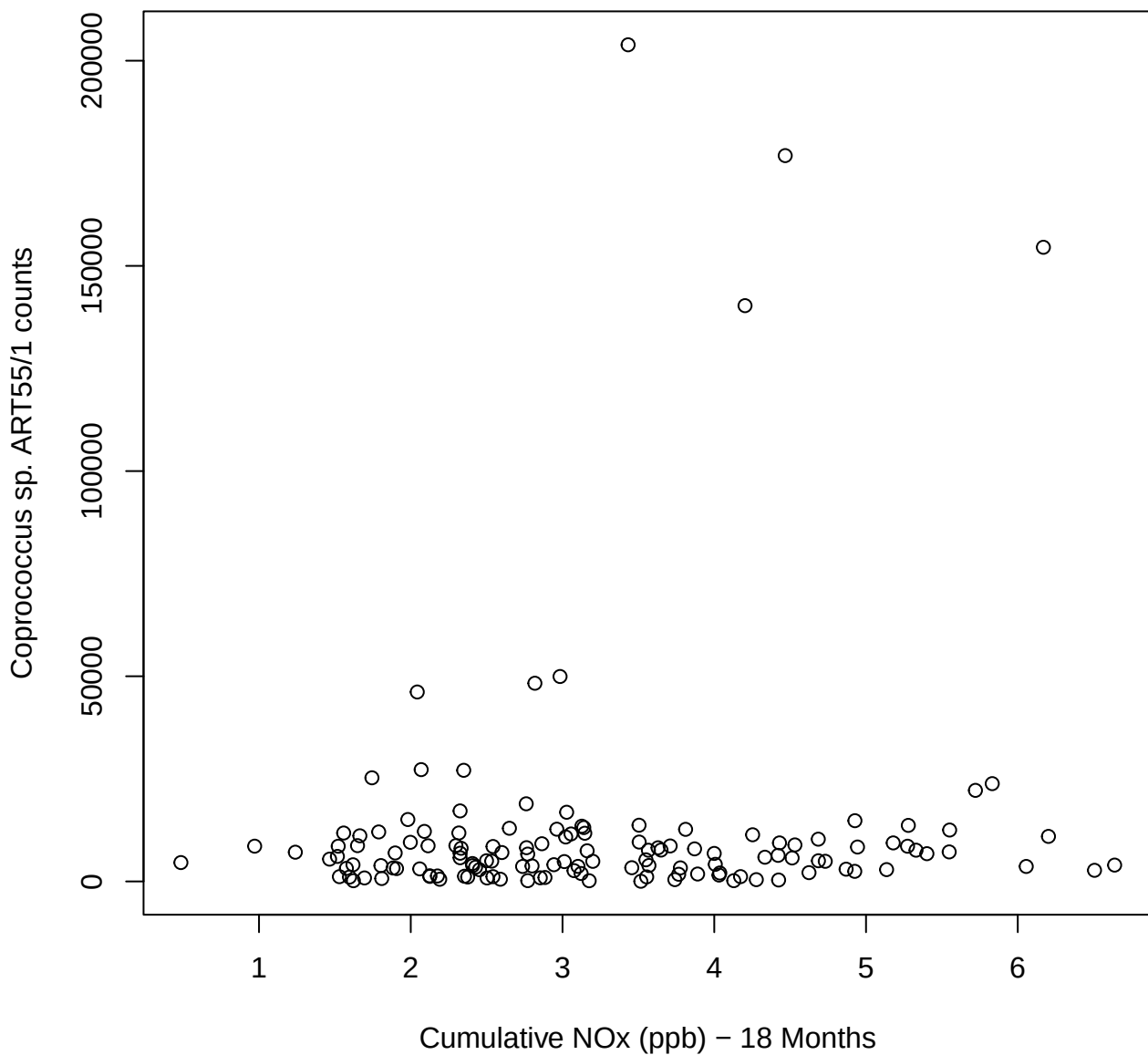
16. *Escherichia fergusonii*



17. *Phascolarctobacterium succinatutens*



18. Coprococcus sp. ART55/1



19. *Megasphaera elsdenii*

