

Module 3: Project 2 by Team 5

Charting the distributed nitrogen cycle

Karl Abuan

May Ho

Jonah Lin

Leilynaz Malekafzali

Tiffany Yang

Abdur Rahman M. A. Basher

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Abstract

This is the abstract. It consists of two paragraphs.

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1 Introduction

2 Materials and Experimental Configuration

2.1 Experimental Protocols

To understand the correlation of microbial diversity and oxygen concentration across samples, we report four experimentally designed test protocols:

P1. Analysis of the DNA abundance of norB with depth.

P2. Analysis of similarities between the RNA abundance and the DNA abundance of norB with depth.

P3. Reconstructing the taxa associated with norB and Analyze variance based on depth, DNA, and RNA.

P4. Analysis of the abundance of norB in relation to nitrogen species in Saanich.

2.2 Dataset

2.3 Data Preporocessing

Manipulate the data into a single data frame

3 Results

3.1 Analysis of the DNA abundance of norB with depth

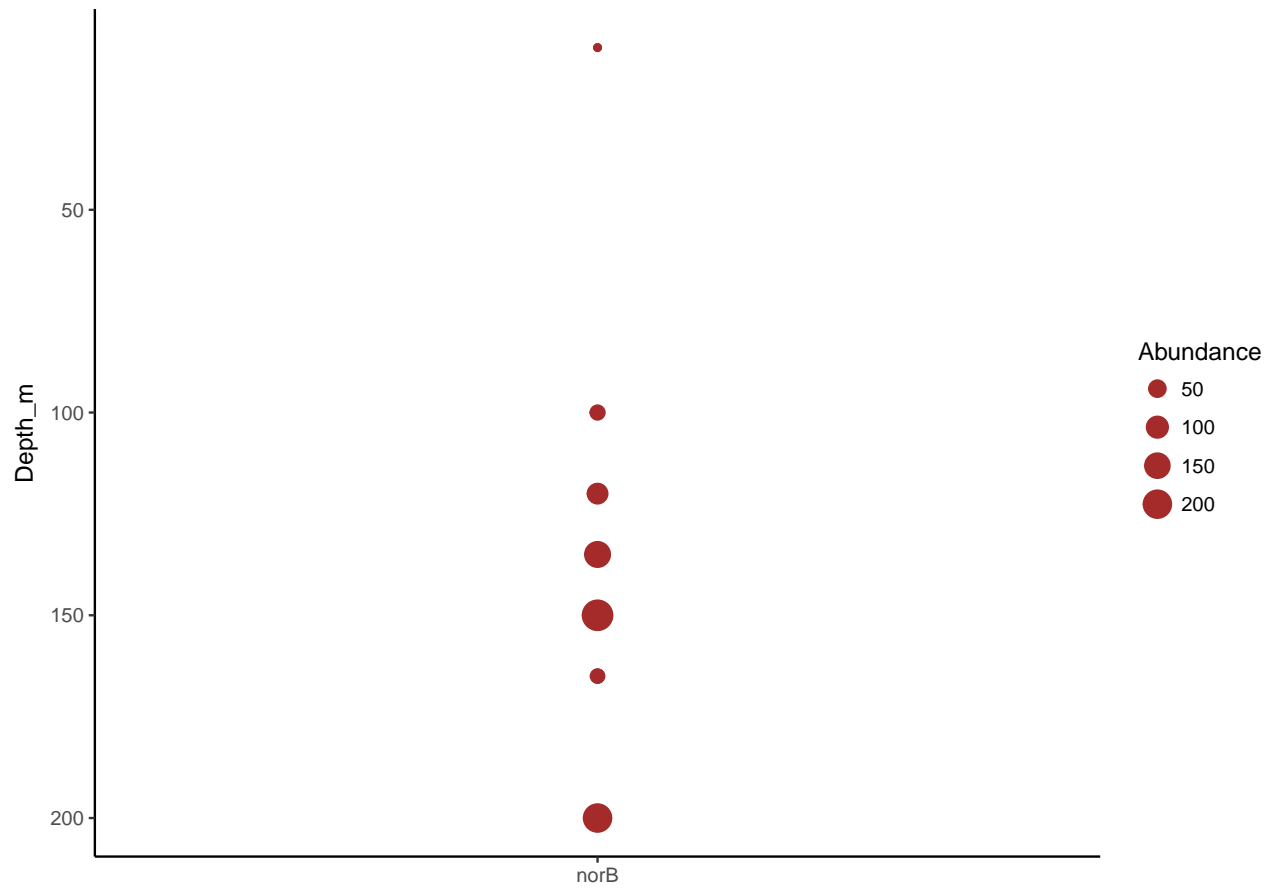


Figure 1: Abundance of the norB gene (DNA) at different depths

3.2 Analysis of similarities between the RNA abundance and the DNA abundance of norB with depth

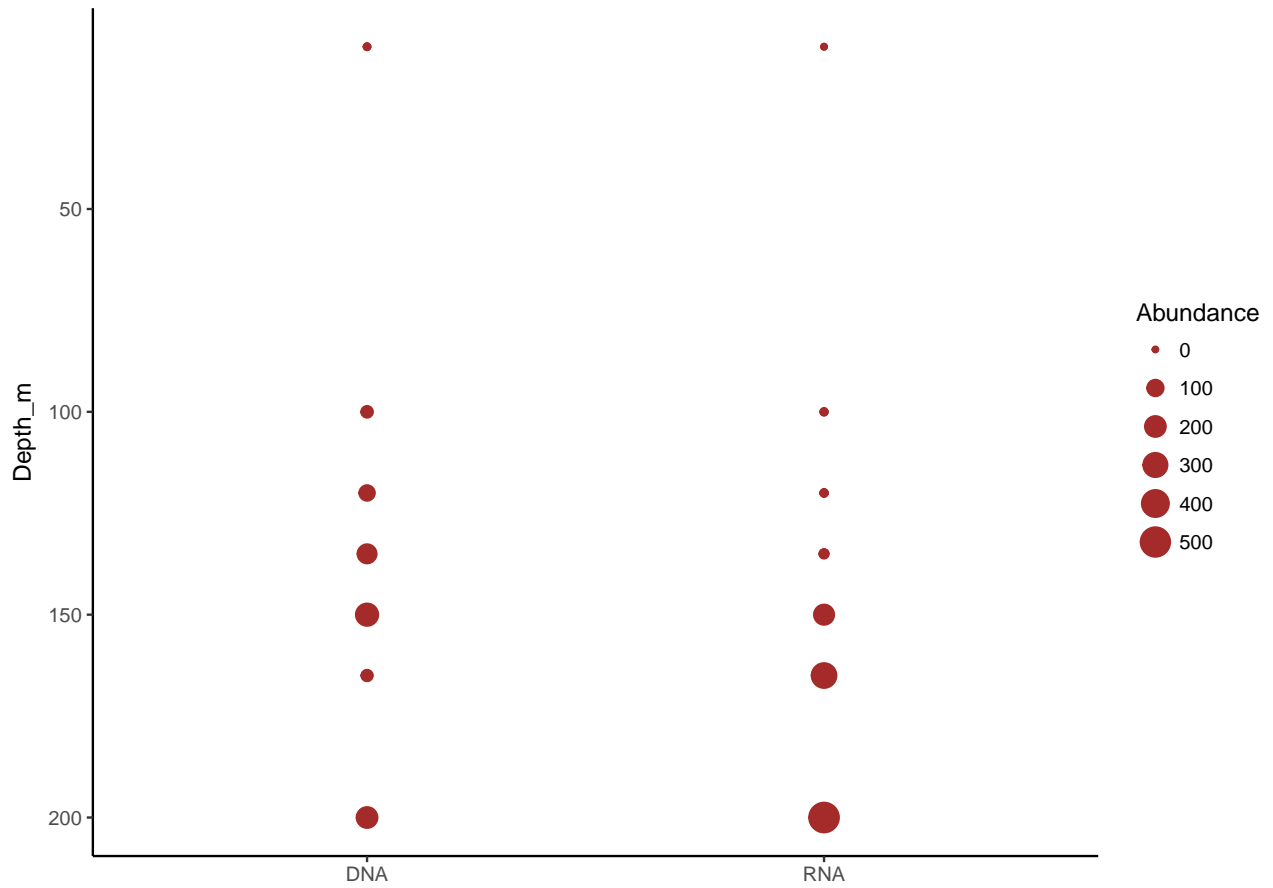


Figure 2: Abundance of the norB gene (DNA vs. RNA) at different depths

3.3 Reconstructing the taxa associated with norB and Analyze variance based on depth, DNA, and RNA

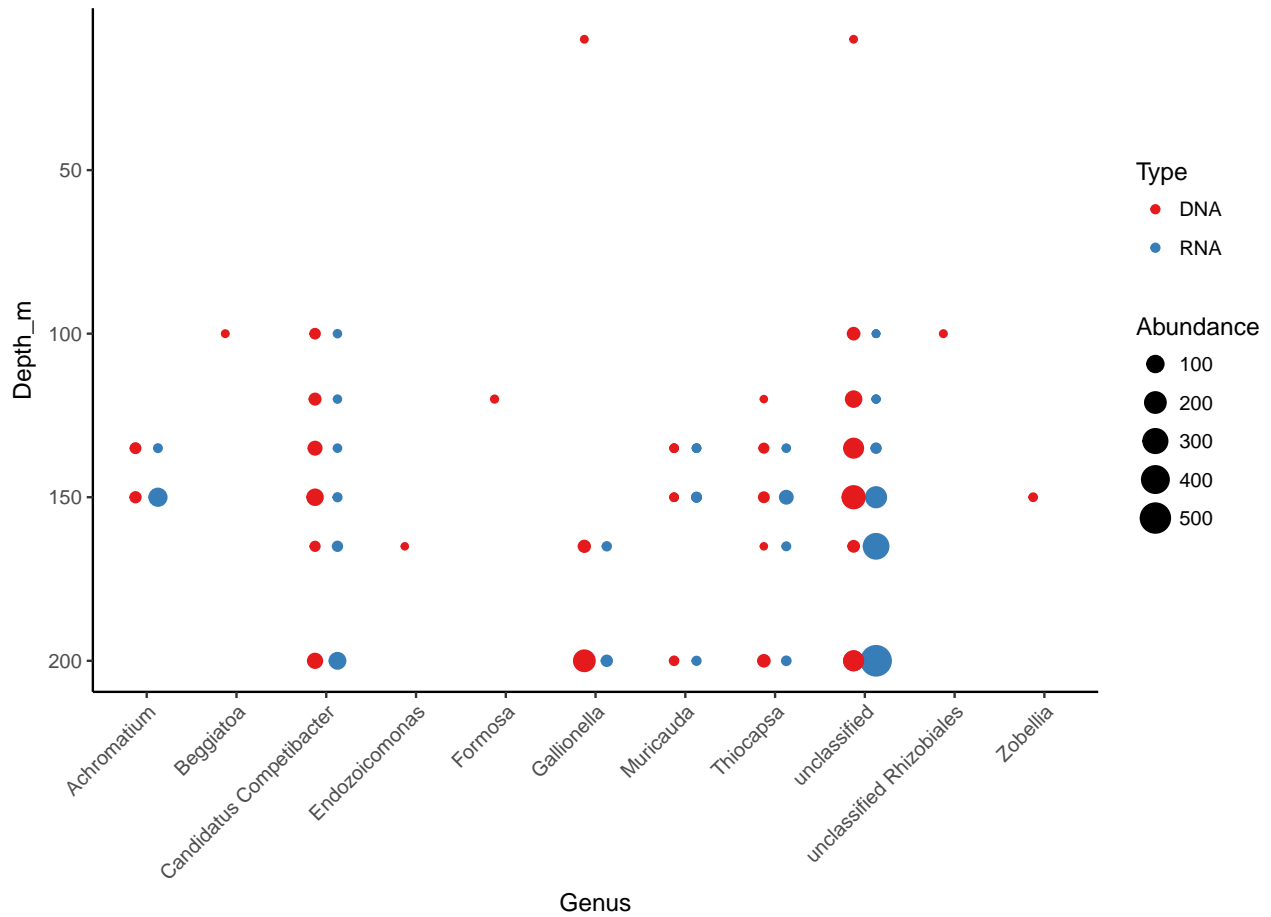


Figure 3: Abundance of genera with norB (DNA vs. RNA) at different depths

Another way to represent the same figure above:

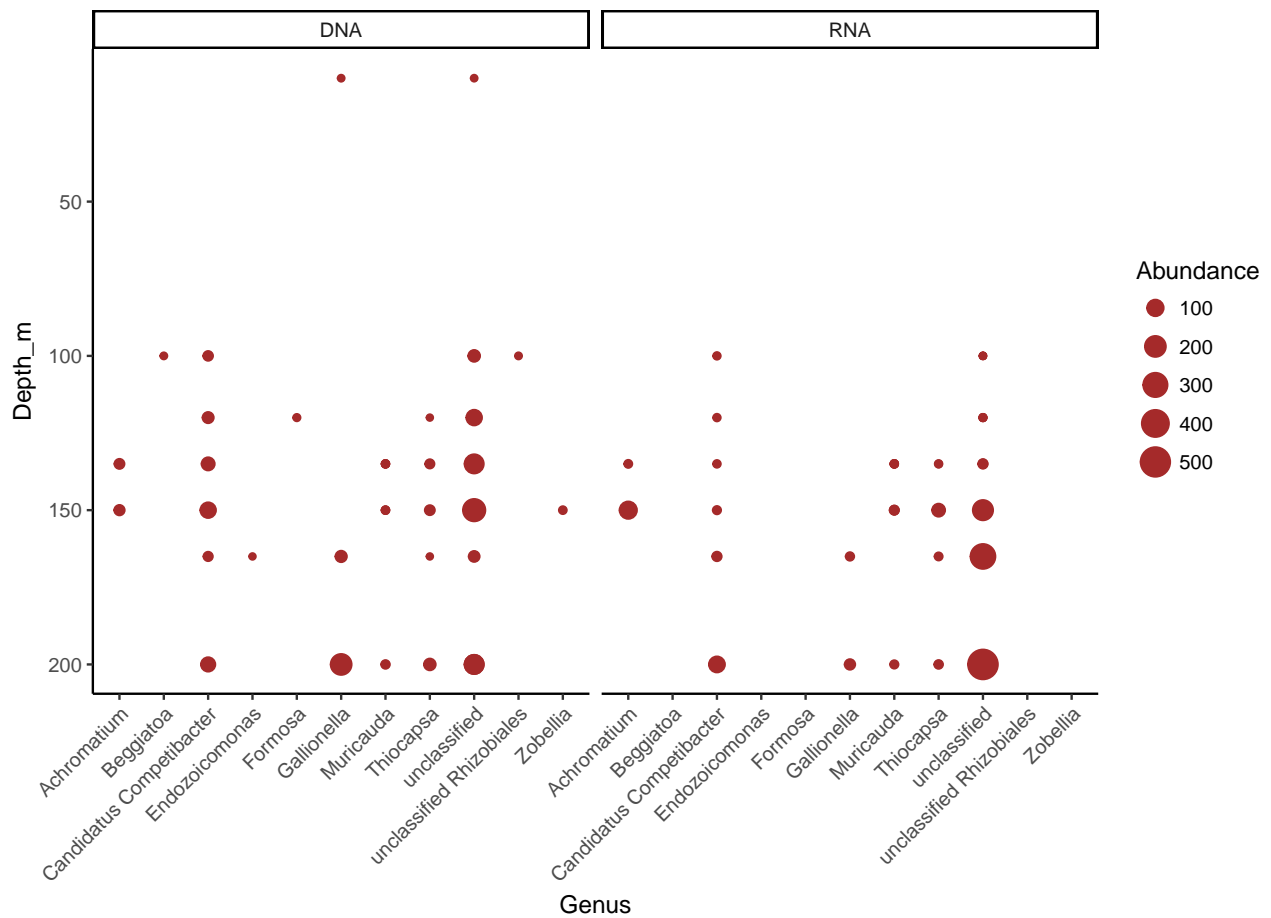


Figure 3a: Abundance of genera with *norB* (DNA vs. RNA) at different depths

3.4 Analysis of the abundance of *norB* in relation to nitrogen species in Saanich

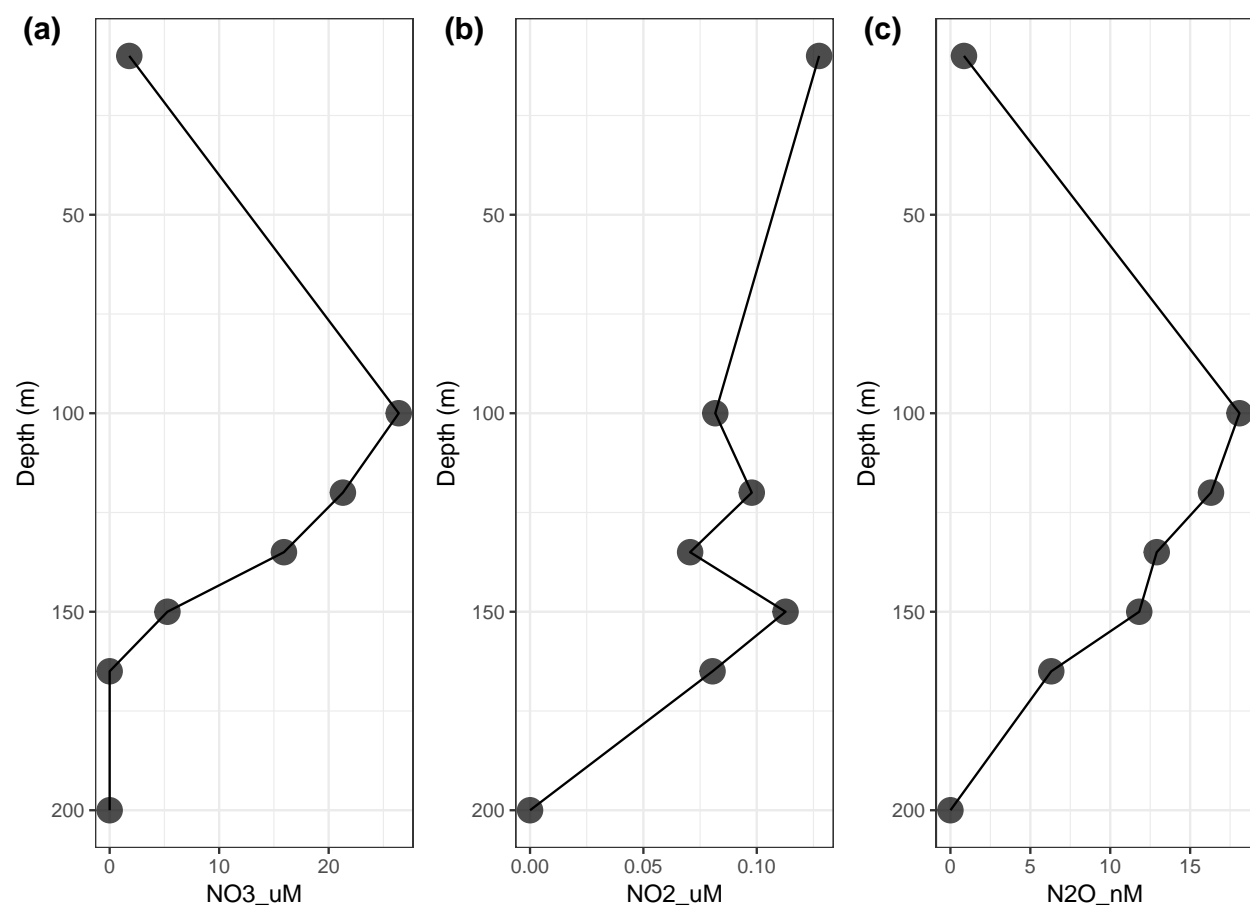


Figure 4: Abundance of norB in relation to nitrogen species across depths

4 Discussion

5 References