UNIVERSITY OF NEBRASKA-LINCOLN

Introduction

- The first step in nitrification is often rate limiting (conversion of ammonia to nitrite) and until recently was thought to solely be facilitated by ammonia oxidizing bacteria (AOB). It is now known that ammonia oxidizing archaea (AOA) also possess the ammonia monooxygenase gene (amoA), the functional gene necessary for the oxidation of ammonia to nitrite (Venter et al. 2004; Treusch et al. 2005), but the extent of its role in soils is unknown.
- Much of the literature from cultivated soils has found that AOA abundance greatly outnumbers AOB abundance and that AOB abundance is less resilient to changes in soil conditions.
- Xu et al (2012) found that in acidic soils the abundance of AOB decreased while AOA abundance was unaffected. Another study on the effect of various fertilizer types found that AOB abundance was also affected while AOA abundance was not (Shen et al. 2008).
- Currently more studies are looking into the specialized roles of AOB and AOA in nitrification.
- Our study is a long-term (25 yr+) fertilized, monoculture maize field with no-tilled and disk-tilled plots. The average field pH is around 5.6 and acidity is postulated to be one of the drivers of AOA abundance over AOB.
- This long-term study allows for unique look into the habitat drivers of AOB and AOA abundance. Herein we report on 1) the long term abundance versus the change in abundance after planting of maize 2) the effect of habitat modification (no tillage versus disk tillage) and 3) the impact of long-term N fertilization rate on AOB and AOA abundance.

Materials and Methods

Field Site:

Two sampling dates: Pre-Plant and 4 days after planting (post-plant) Disk Tillage versus No Tillage

Continuous Corn

0, 40, 80, 120, 160 kg N ha⁻¹ yr⁻¹

DNA Extractions:

500mg of Freeze-Dried Soil

MoBio Ultra Clean® soil DNA isolation kits

Tubes were held in the MoBio Vortex Adapter tube holder. Modifications included: Tubes were incubated at 70°C for 10 mins in a water bath after solution S1 was added.

Followed by the 'Alternative Protocol for maximum yields' according to manufacturer's specifications.

Standard Curve:

Cells derived from Dr. Kim Cook's lab (USDA-ARS, Bowling Green, KY) were grown in LB broth and purified using Wizard Plus Minipreps DNA purification system.

Concentrations were calculated using NanoDrop

10-fold dilutions between 10⁸-10² were used

AOB efficiencies between 94% and 115% % R² values of 0.981-0.999 AOA efficiencies between 91% and 99% with R² values of 0.990-0.998

Quantitative Real-Time PCR:

AOB primers: AmoA 1F/2R

AOA primers: AmoA 19F/643R

Samples were carried out in triplicate 20µL reactions using Sybr Green I dye Single PCR products were validated using Melt Curve Analysis and Gel Electrophoresis

References / Acknowledgements

Venter, J.C., et al. (2004). Environmental genome shotgun sequencing of the Sargasso Sea. Science 304(66). Treusch, A.H., et al. (2005). Novel genes for nitrite reductase and Amo-related proteins indicate a role of uncultivated mesophilic crenarchaeota in nitrogen cycle. Environmental Microbiology 7(12). Xu, Y., Yu, W., Qiang, M., Zhou, H. (2012). Responses of bacterial and archaeal ammonia oxidisers of an acidic luvisols

soil to different nitrogen fertilization rates after 9 years. Biol Fertil Soils 48(7). Shen, J., Zhang, L., Zhu, Y., Zhang, J., He, J. (2008). Abundance and composition of ammonia-oxidizing bacteria and

ammonia-oxidizing archaea communities of an alkaline sandy loam. Environmental Microbiology 10(6). We would like to give special acknowledgement to Dr. Kim Cook (USDA-ARS) for providing her services in making our standard curve and Ryan McGhee (USDA-ARS) for technical support on all aspects of the project.

Abundance of Ammonia-Oxidizing Bacteria and Archaea Under Long-Term Maize Cropping Systems

Lauren Segal*; Rhae Drijber*; Daniel Miller**; Terrance Loecke***

*Dept. of Agronomy and Horticulture **USDA-ARS ***Dept. of Natural Resources







