

Nitrifying Community and N-cycle Activity are Altered by Increasing Timber Harvest Intensity in Surface and Subsurface Soils of the Western Gulf Coastal Plain

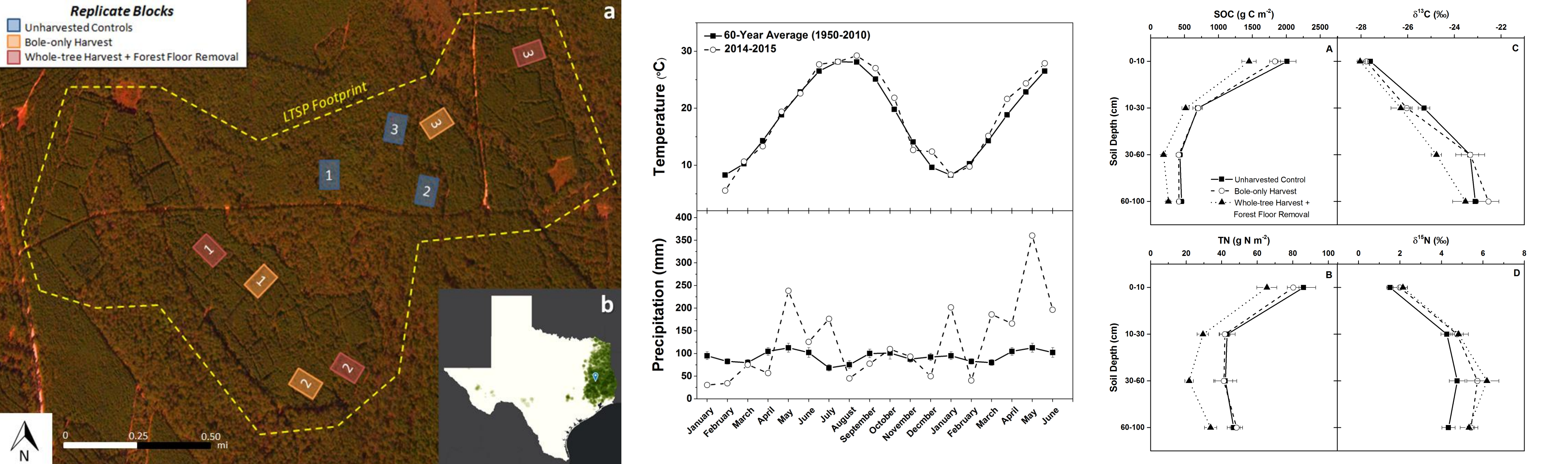
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Introduction

Timber harvest has been shown to alter N-biogeochemistry through alterations in soil N stocks; however, soil pools of plant available N have not been quantified, especially in deeper horizons. Furthermore, microbes that mediate N-transformations are likely to be affected by increased harvest practices. The purpose of this study was to determine the effect of differing forest harvest intensities (i.e., no harvest, bole-only harvest, and whole-tree harvest + forest floor removal), 18-years post harvest, on microbial ecology associated with N-biogeochemistry. The Groveton, TX Long-term Soil Productivity site was utilized for this study. We hypothesize that microbial biomass and gene copy number of *amoA* will be significantly reduced by increased harvest intensity resulting in smaller pools NO_3^- .

Groveton Long-term Soil Productivity Site

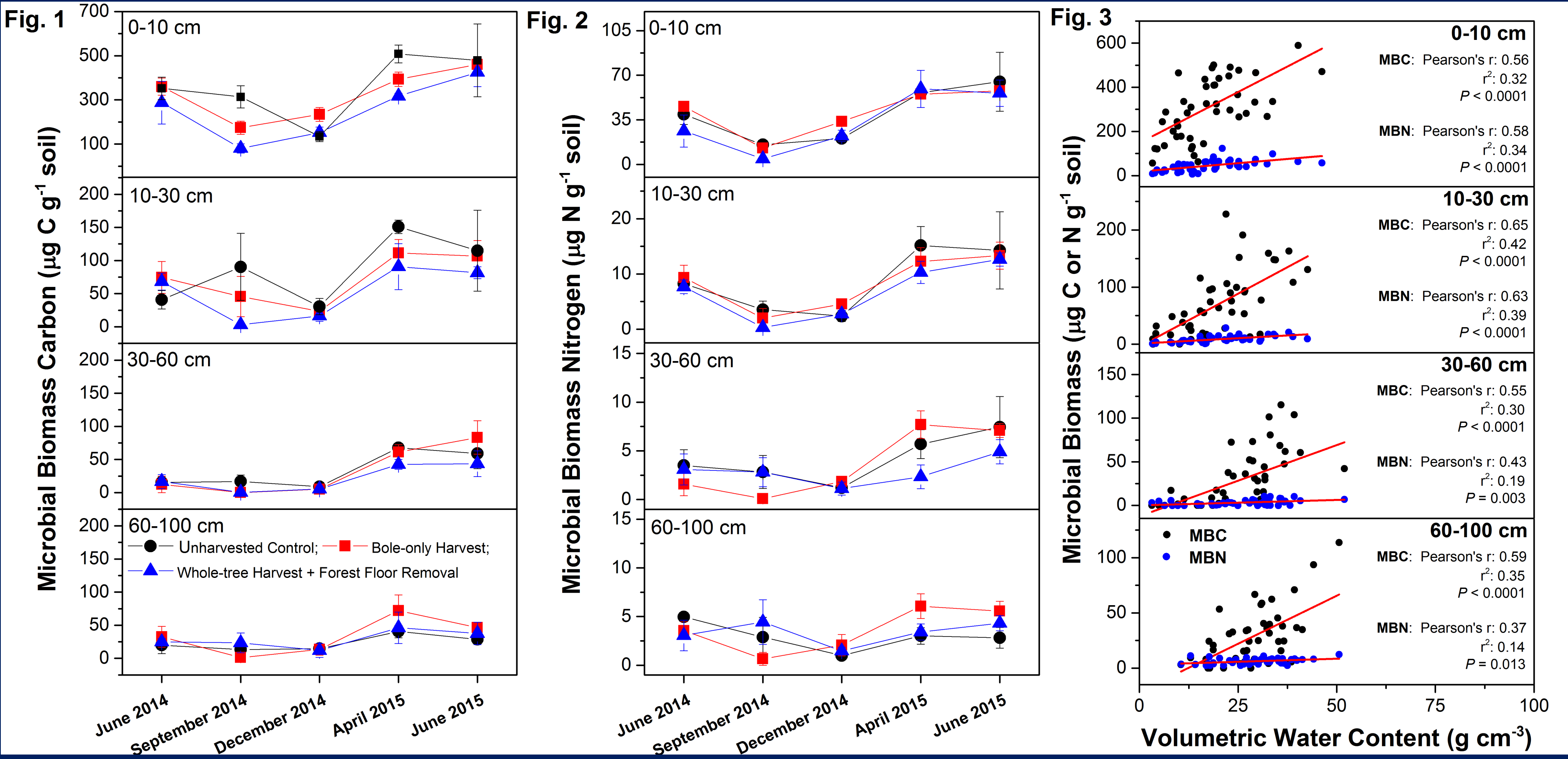


Harvest Method	Description	Soil Physicochemical Characteristics
Unharvested Control	No harvest methods employed. Tree age: 55+ years	Bulk Density (g cm ⁻³) Soil pH Soil Texture
Bole-only Harvest	Tree boles were removed 18 years ago. Crowns, felled woody and herbaceous understorey and forest floor retained	Soil Depth (cm) Averaged across all treatments & sampling points (n=45)
Whole-tree Harvest + Forest Floor Removal	All aboveground biomass was removed 18 years ago. Bare soil exposed	0-10 10-30 30-60 60-100

Methods

Soil was sampled to a depth of 1-meter, seasonally for five sequential time points. All analyses were carried out either in the Stable Isotopes for Biosphere Sciences Laboratory or the Soil and Aquatic Microbiology Laboratory, both located on the campus of Texas A&M University. Microbial Biomass C and N was extracted using the chloroform fumigation method and quantified on a Shimadzu TOC/TN Analyzer. $\text{NH}_3/\text{NH}_4^+$ and NO_3^- were extracted from soil using 2M KCl and analyzed on a Seal Analytical discrete chemistry analyzer. Copy number of archaeal and bacterial *amoA* were quantified on an Eppendorf Masterplex using SYBR florescent chemistry. Primers used: Arch-*amoA*F & Arch-*amoA*R (AOA); *amoA*-1F & *amoA*-2R (AOB).

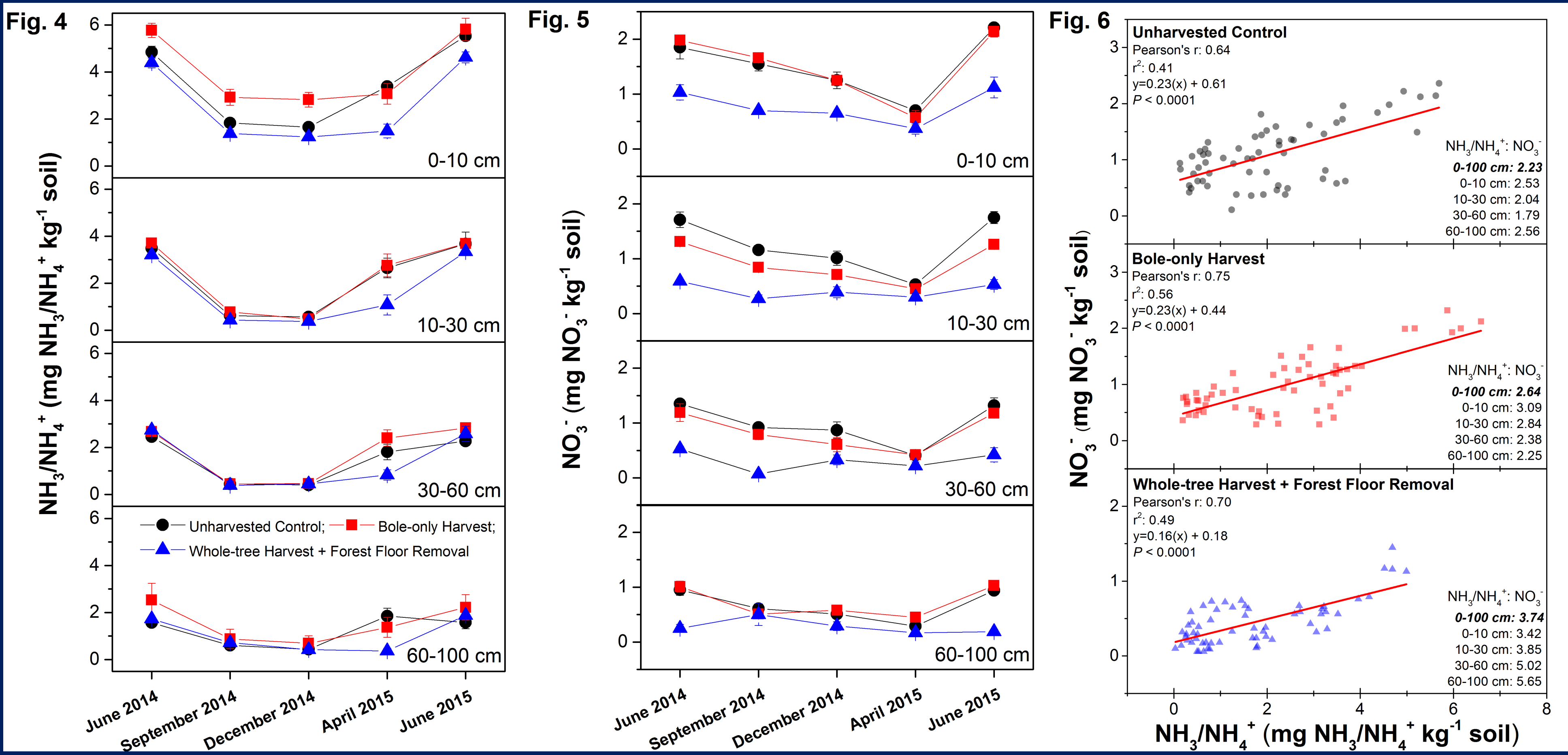
Microbial Biomass



Microbial Biomass Carbon (μg C g ⁻¹ soil)	Harvest Method	Season	Harvest Method * Season	Microbial Biomass Nitrogen (μg N g ⁻¹ soil)	Harvest Method	Season	Harvest Method * Season
Soil Depth (cm) ^a		p-values ^b		Soil Depth (cm) ^a		p-values ^b	
0-10a	0.0291*	<0.0001***	0.4889	0-10a	0.4113	<0.0001***	0.8987
10-30b	0.1565	<0.0001***	0.6659	10-30b	0.4219	<0.0001***	0.9798
30-60b	0.2537	<0.0001***	0.6300	30-60b	0.3634	0.0003***	0.2577
60-100b	0.5037	<0.0001***	0.7679	60-100b	0.6771	0.0318*	0.1497

^a Different letters indicate statistical difference (p<0.05) between soil depth (Tukey Kramer). ^b * p<0.05; ** p<0.01; *** p<0.001.

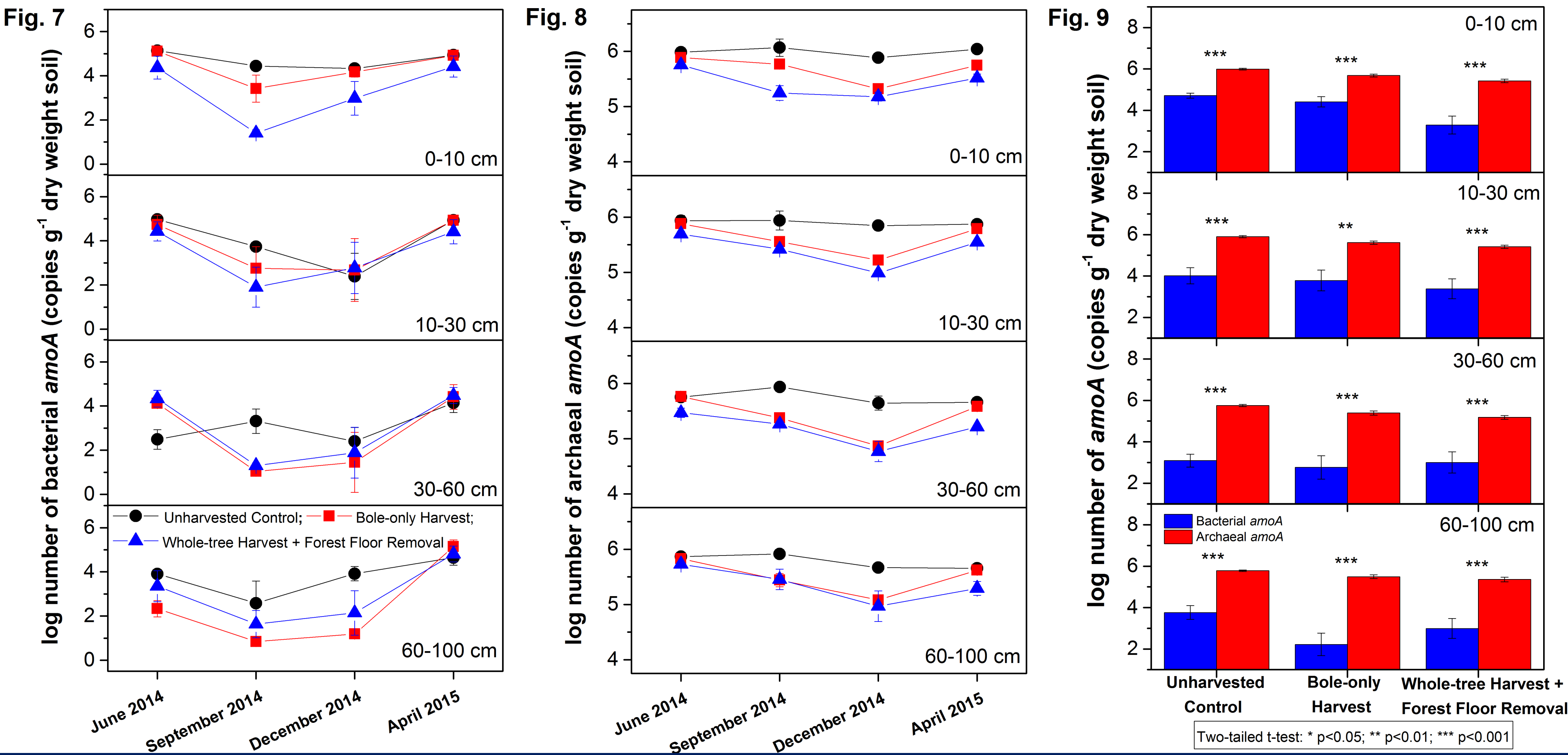
Soil $\text{NH}_3/\text{NH}_4^+$ & NO_3^- Pools



Soil $\text{NH}_3/\text{NH}_4^+$ Pool (mg $\text{NH}_3/\text{NH}_4^+$ kg ⁻¹ soil)	Harvest Method	Season	Harvest Method * Season	Soil NO_3^- Pool (mg NO_3^- kg ⁻¹ soil)	Harvest Method	Season	Harvest Method * Season
Soil Depth (cm) ^a		p-values ^b		Soil Depth (cm) ^a		p-values ^b	
0-10a	<0.0001***	<0.0001***	0.1019	0-10a	<0.0001***	<0.0001***	0.0230*
10-30b	0.0018**	<0.0001***	0.1013	10-30b	<0.0001***	<0.0001***	<0.0001***
30-60bc	0.0236*	<0.0001***	0.0102*	30-60bc	<0.0001***	<0.0001***	0.0141*
60-100c	0.0500	<0.0001***	0.1739	60-100c	<0.0001***	<0.0001***	0.0007***

^a Different letters indicate statistical difference (p<0.05) between soil depth (Tukey Kramer). ^b * p<0.05; ** p<0.01; *** p<0.001.

Bacterial & Archaeal *amoA* Quantification



Bacterial <i>amoA</i> (log (copies g ⁻¹ dry soil))	Harvest Method	Season	Harvest Method * Season	Archaeal <i>amoA</i> (log (copies g ⁻¹ dry soil))	Harvest Method	Season	Harvest Method * Season
Soil Depth (cm) ^a		p-values ^b		Soil Depth (cm) ^a		p-values ^b	
0-10a	<0.0001***	<0.0001***	0.0597	0-10a	<0.0001***	<0.0001***	0.0097**
10-30ab	0.5064	0.0014**	0.8912	10-30ab	<0.0001***	<0.0001***	0.0030**
30-60b	0.7559	<0.0001***	0.0812	30-60b	<0.0001***	<0.0001***	0.0005***
60-100b	0.0116*	<0.0001***	0.3060	60-100ab	0.0001***	<0.0001***	0.1106

^a Different letters indicate statistical difference (p<0.05) between soil depth (Tukey Kramer). ^b * p<0.05; ** p<0.01; *** p<0.001.

Conclusions and Future Directions

18-years post harvest, microbial biomass carbon and nitrogen were generally unaffected by increasing timber harvest intensity; however, $\text{NH}_3/\text{NH}_4^+$ pools, NO_3^- pools, and *amoA* copy number were significantly affected. Even though the relative biomass of the microbial community has generally stabilized, the quantity of archaeal and bacterial nitrifiers has been reduced by more intensive timber harvest methods, thus reducing the soil NO_3^- pool and possibly leading to the increased ratio of $\text{NH}_3/\text{NH}_4^+$ to NO_3^- . This not only has implications for current plant growth, but could affect the viability of future rotations. In regards to a broader microbial ecology perspective, this is one of the first studies to quantify ammonia oxidizers, in surface and subsurface soils, in the southeastern United States. Archaea have been implicated as having a greater impact on ammonia oxidation than bacteria, and our results indicate that this could be true, especially in acidic forest soils. Current efforts are underway to sequence bacterial and archaeal ammonia oxidizer communities from the Groveton-LTSP site as well as quantify their relative activity.