

The Effect of Organic and Conventional Agricultural Production on Fungal Community Assemblage in Ethiopian Green Coffee Beans

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Abstract We are developing coffee production as a model to understand the transport of fungi through the global food trade and to evaluate the ways in which organic and conventional agricultural production influences fungal community assemblage on harvested crops. We used culture-based and culture-independent methods along with molecular techniques to identify fungal communities on Ethiopian green coffee beans. The coffee was either produced by conventional or organic methods. Our findings suggest that farming practices have significant effects on fungal community assemblage in harvested coffee. USDA Certified Organic coffee hosts a lower diversity and abundance of fungal species than conventional coffee. Methods of sampling strongly affect the recovery of species. Culture-based and culture-independent methods resulted in distinct views of fungal community assemblage in green coffee beans.

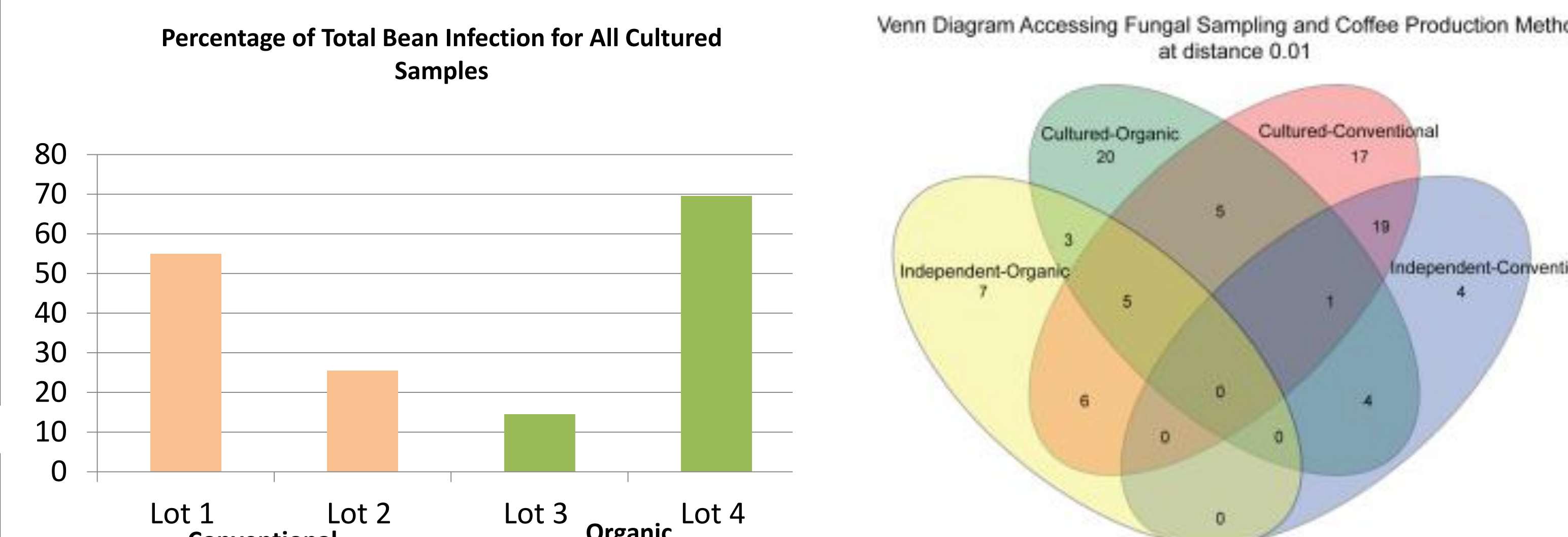
Introduction

As the fourth largest agricultural commodity, the importance of the coffee trade cannot be overstated for the economies of many tropical countries. Development of shade-grown, organic, and fair trade coffee practices may have numerous economic and environmental benefits, but little is known about how these different methods of production affect fungal communities on coffee. Adequately accessing fungal communities in coffee may have human health implications by reducing the risk of mycotoxins. Mycotoxins such as ochratoxin and aflatoxin are mutagenic and carcinogenic secondary metabolites produced by a variety of *Aspergillus*, *Fusarium*, and *Penicillium* species (Edwards, et al. 2002). These contaminants are found in coffee at various stages of production, including cherry ripening, drying and storage (Taniwaki, et al. 2003). USDA Organic certification prevents the use of specific fungicides, pesticides, and chemical fertilizers during the growth and processing of agricultural productions. In this study we use both culture-based and culture-independent methods to determine how organic and conventional production methods affect fungal communities on green coffee beans.

Methodology

Four lots of Ethiopian coffee were purchased. All samples were shade grown and wet-processed. Lots 1 and 2 were produced by conventional methods, and Lots 3 and 4 were certified organic. DNA-barcoding techniques using the ITS rRNA region allowed us to identify fungal species present in the coffee beans. Culture-based methods involved surface-sterilization using a 2% bleach solution, plating 200 beans per lot on MEA++ and DG18 media, isolating pure cultures of fungi, and extracting fungal DNA for PCR and Sanger sequencing. Culture-independent methods involved drying 20 beans per lot in a 60°C oven for 7 days before freezing with liquid nitrogen to grind the beans into fine powder. DNA extraction was then carried out on the coffee powder, and PCR was performed with fungal-specific ITS primers, cloned into a plasmid, and 24 subclones were sequenced per lot. All DNA sequences were analyzed using Sequencher 4.1. Sequences were compared to GenBank using BLAST to determine the closest matched OTUs. Community analyses based on OTU data were conducted using MOTHUR. Significance differences in communities were estimated using Libshuff tests and diversities calculated using Chao and Shannon estimators.

Group	No. Sequences	O.T.U.s	Chao	Shannon
Lot 1 Cultured	94	20	32 (22.66-74.20)	2.05 (1.75-2.35)
Lot 2 Cultured	44	12	22.5 (14.03-66.19)	1.79 (1.44-2.15)
Lot 3 Cultured	27	10	13.3 (10.50-32.07)	1.95 (1.61-2.29)
Lot 4 Cultured	154	24	62.25 (35.59-150.21)	1.15 (0.86-1.43)
Lot 1 Independent	18	8	9.5 (8.18-20.47)	1.73 (1.29-2.18)
Lot 2 Independent	9	8	18.5 (10.03-62.19)	2.04 (1.59-2.49)
Lot 3 Independent	8	4	4.5 (4.030-12.261)	1.21 (0.71-1.71)
Lot 4 Independent	14	9	19.5 (11.03-63.19)	1.96 (1.49-2.44)
Cultured-Conventional	139	53	143 (88.78-279.36)	3.02 (2.74-3.30)
Cultured-Organic	183	38	125.75 (67.88-295.73)	1.88 (1.60-2.16)
Independent-Conventional	52	28	53.5 (36.05-108.79)	3.08 (2.84-3.32)
Independent-Organic	44	21	51.33 (29.18-133.43)	2.68 (2.39-2.98)



Top Blast Match O.T.U.	Occurrence in Conventional n = 191	Top Blast Match O.T.U.	Occurrence in Organic n = 227
<i>Aspergillus niger</i>	55	<i>Aspergillus niger</i>	107
<i>Wickerhamomyces anomalus</i>	30	<i>Aspergillus tubingensis</i>	46
<i>Aspergillus tubingensis</i>	17	<i>Eurotium rubrum</i>	11
<i>Cladosporium sp.</i>	15	<i>Eurotium amstelodami</i>	10
<i>Eurotium rubrum</i>	11	<i>Dipodascaceae sp.</i>	8

Top Blast Match O.T.U.	Occurrence in All Cultured Samples n = 319	Top Blast Match O.T.U.	Occurrence in All Culture-Independent Samples n = 96
<i>Aspergillus niger</i>	163	<i>Wickerhamomyces anomalus</i>	18
<i>Aspergillus tubingensis</i>	65	<i>Cladosporium sp.</i>	16
<i>Eurotium rubrum</i>	23	<i>Pichia pijperi</i>	11
<i>Wickerhamomyces anomalus</i>	12	<i>Eurotium amstelodami</i>	10
<i>Aspergillus fumigatus</i>	9	<i>Dipodascaceae sp.</i>	9
<i>Chaetomium reflexum</i>	7	<i>Debaryomyces hansenii</i>	6
<i>Candida orthopsilosis</i>	5	<i>Epicoccum sp.</i>	4
<i>Cystofilobasidium feraegula</i>	4	<i>Uncultured Basidiomycete</i>	3
<i>Cladosporium sp.</i>	4	<i>Alternaria alternata</i>	2
<i>Fusarium sp.</i>	4	<i>Hanseniaspora uvarum</i>	2

★ At least five species: *Aspergillus fumigatus*, *A. niger*, *A. phoenicis*, *Penicillium brevicompactum*, and *P. citrinum*, overlapped with endophytic species found by Vega, et al. (2010) in coffee leaves, seeds, and fruits.

Discussion

Both production and sampling methods seem to affect which fungal species are recovered from dried green coffee beans. Overall, culture-dependent methods recovered communities dominated by *Aspergillus*, particularly *A. niger*, as has been recorded in other studies (Perrone, et al. 2007). In contrast, culture-independent investigations found a much lower frequency of *Aspergillus*, the fungus largely suspected of causing mycotoxins in coffee. Conventional and organic coffee production results in fungal community assemblages that are significantly different (Libshuff, results not shown), with organic samples hosting lower diversity and abundance of OTUs. The lot to lot (farm to farm) variation in bean infection percentage suggests that farming practices, such as the use of fungicides, fertilizers, or pesticides, may create distinct chemical environments for microorganisms in agroecosystems, resulting in differentiation in habitat suitability. Organic shade coffee farms are often cited to be reservoirs for biodiversity, but this may not extend to the fungi found as contaminants on the harvest, as observed in this study. Using various sampling methods strongly affected the species recovered. Culture-based methods result in a greater number of species being identified, though species recovered through culture-independent methods are non-overlapping and distinct. This might be because culture-independent methods are unbiased to preferential spore germination on media plates, alleviating competition between fungi.

Future Research

We plan to study the ways in which various methods of processing coffee (dry or wet) affects fungal communities on green coffee beans, as well as the effect of sun and shade farming, which enlist different methods of agroforestry. Future projects also involve population genetic analysis of mycotoxigenic *Aspergillus* species on green coffee beans from various countries and coffee-producing geographical regions. Preliminary data from Mexican and Kenyan lots show very different fungal communities than the ones found in Ethiopian beans.

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