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ABSTRACT

We isolated DNA from 12 "true morels" (Morchella) and one "false morel" (Gyromitra) collected in interior Alaska. We PCR amplified the Internally Transcribed Spacer region of the ribosomal DNA genes. After cycle sequencing our amplified gene, we used the Sequencher program to assemble the raw sequences, ClustalW to align sequences across samples, and the GARLI program to create a phylogenetic tree of evolutionary history that compares our samples with all publicly available Morchella sequences, obtained from GenBank. We also included sequences obtained directly from interior AK forest soils and found close matches to aboveground morels. In general we found that our Alaskan samples span much of the diversity of various groups of "black" morels, but we found no "yellow" morels. Interestingly, five of our samples from 3 interior AK burn sites composed a group that appears distinctly Alaskan.

INTRODUCTION

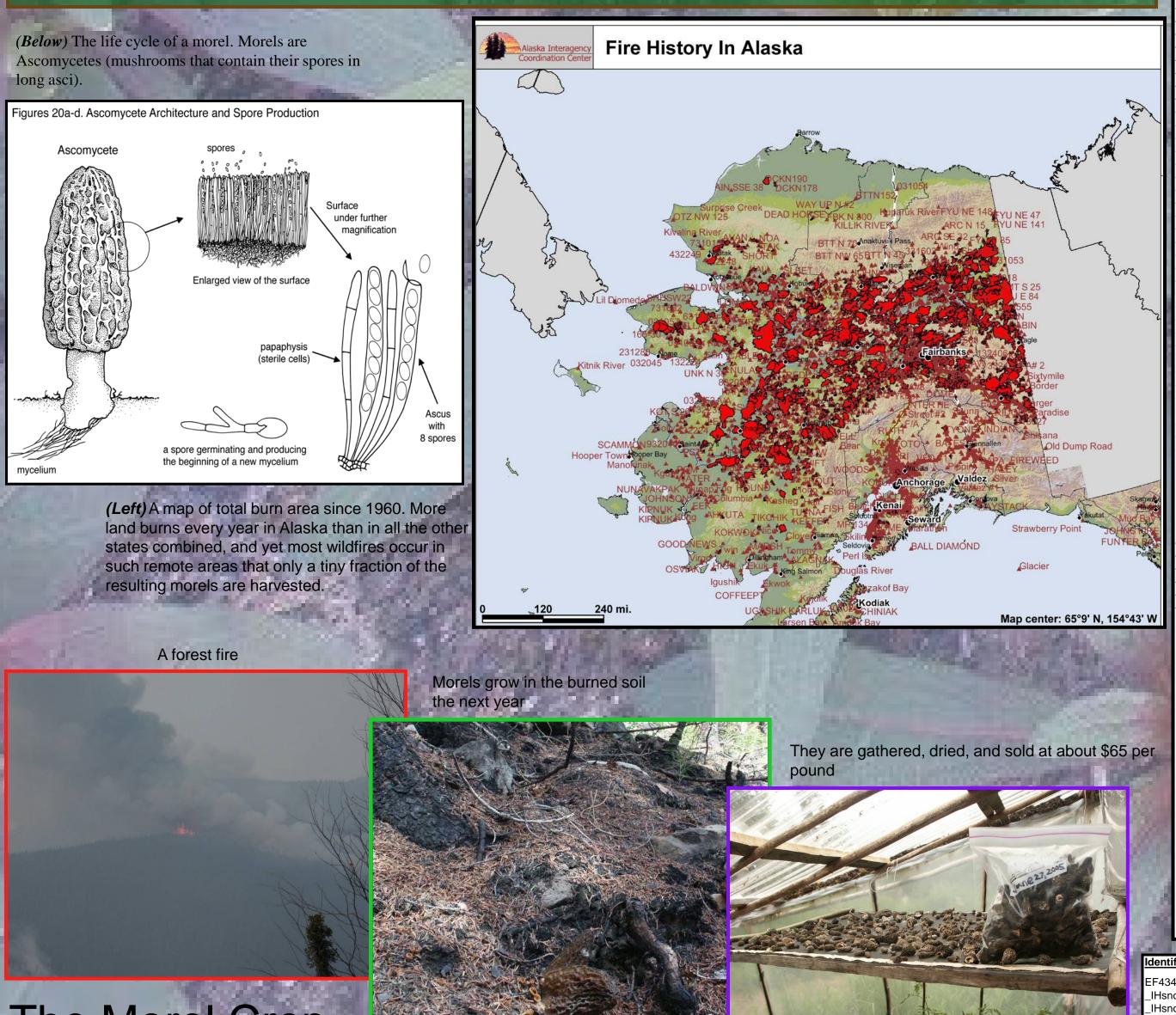
DESCRIPTION AND DIVERSITY

Morels are cone-shaped Ascomycete mushrooms often found growing prolifically in burned soil the summer after a wildfire and occasionally in other disturbed habitats. All morels are choice edibles and easily identifiable as a genus. However, morels exhibit a wide variety of forms and colors, and individual species are difficult to identify. Current estimates put North America with 4 "yellow" and 9 "black" morels (Pilz et. al.), although we provided evidence that an entirely separate group exists (see figure 1).

BIOLOGY

Morels are the sporocarps (fruiting bodies) of underground mycelia, the microscopic fungal threads that spread through the soil and collect water and nutrients. It is poorly understood whether morels are symbionts (organisms that live in a mutually beneficial relationship with other organisms), saprophytes (organisms that decompose decaying matter), or parasites (organisms that feed directly off of other living organisms). It seems likely that across the various species or across the lifespan within a species all three lifestyles may occur. The morel sporocarp itself grows only for about 1-6 days before releasing spores and disintegrating. The spores float in the wind, some eventually landing on an area suitable for germination. Some theories suggest that the spores germinate soon after hitting the soil and remain as mycelia until a fire or other disturbance. Others imply that the spores remain underground for long periods, then germinate after a fire and produce sporocarps the following summer (Pilz et. al.) The flush of organic matter from the roots of dead plants may provide the resources necessary for sporocarp production. However, it has also been suggested that the mycelia produce sporocarps for entirely different reasons—that the burned environment is too harsh for the mycelia, which put their last resources into reproduction before they die. The importance of morels in the ecosystem, especially in

replenishing nutrients to burned soil, is unknown (Pilz et. al.).

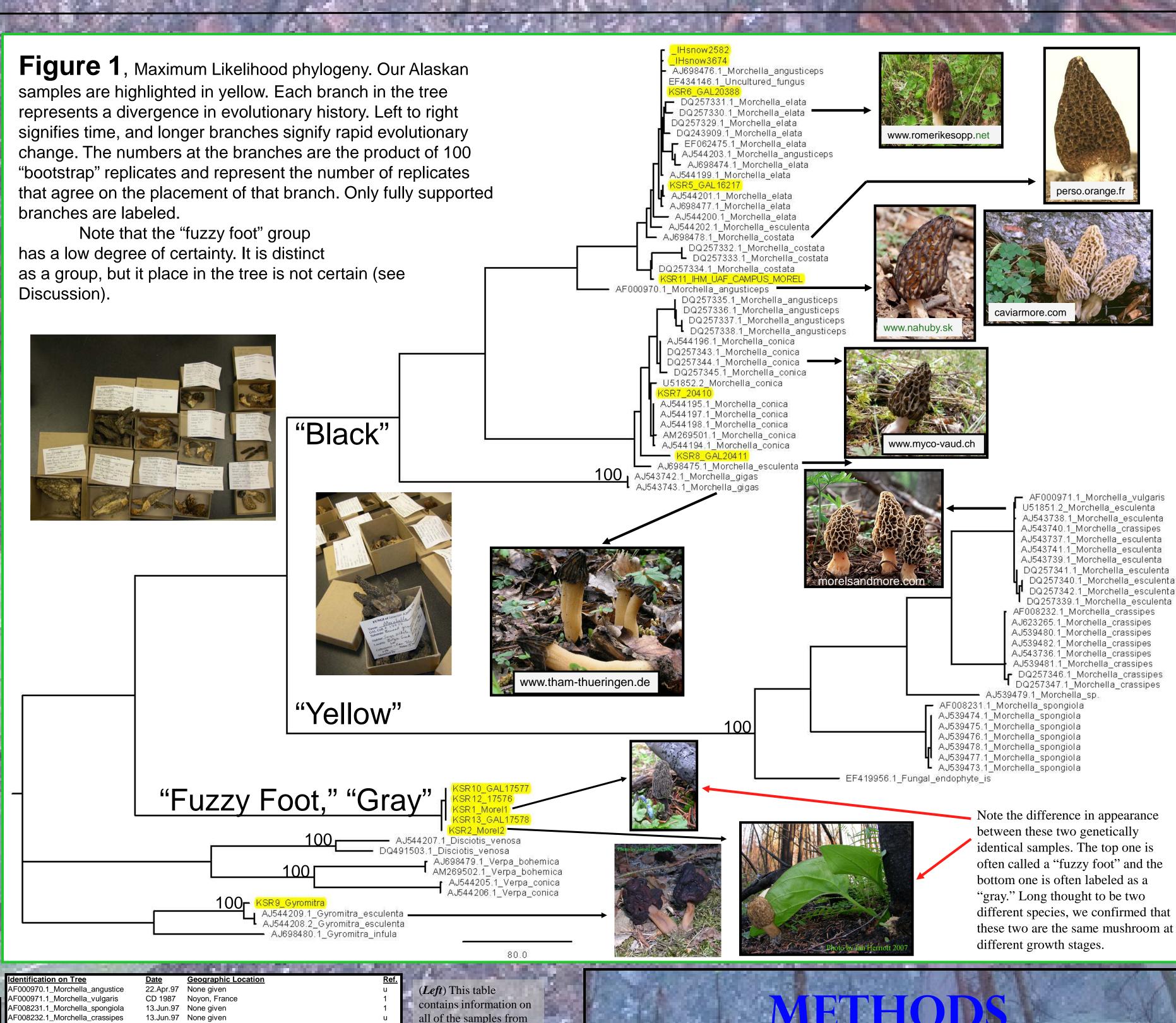


The Morel Crop

Morels are gathered in large quantities where they occur, and result in a sale of \$5 million to \$10 million annually in Western North America (Pilz et.

GENETIC DIVERSITY WITHIN ALASKAN MORELS

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KSR7_{GAL}20410 KSR8_GAL20411 KSR9 Gyromitra

KSR13_GAL17578

SR5_GAL16217

SR6_GAL20388

KSR11 IHM UAF_{GAL20701

SR1 Morel1 {GAL20707}

R2 Morel2 (GAL20705)

13.Jun.97 None given 08232.1 Morchella crassipes 39473.1 Morchella spongiol Thuringia:Jena:Kunitz:Kunitzburg, German 31.Jan.03 Thuringia:Jena:Goeschwitz, Germany ngia:Jena:Goeschwitz, Germany 39475.1 Morchella spongiol 31.Jan.03 ngia:Jena:Goeschwitz, Germany 39476.1 Morchella spongiol gia:Eschenbergen:Fahnersche:Hoehe, Germany as de Calais:Merlimont, France 39478.1 Morchella spongi 31.Jan.03 Solan Himachal Pradesh, India 39479.1 Morchella si 31 Jan 03 huringia:Jena:Drackendorf, Germany 39480.1 Morchella crassin 31.Jan.03 ace:Illkirch-Graffenstaden, France 39481.1 Morchella crassir Saxony:Leipzig:Mark-Kleeberg, Germany 3736.1 Morchella crassipe 10.Feb.03 Thuringia, Jena, Drackendorf, Germany Morchella esculent 0.Feb.03 Alsace, Offendorf, France 3738.1 Morchella esculent 0.Feb.03 Jena, Grance, Germany 43739.1 Morchella esculent 0.Feb.03 543740.1 Morchella crassipe 10.Feb.03 USA 10.Feb.03 North Rhine-Westphalia, Muenster, Germar 543741.1 Morchella esculent 0.Feb.03 Thuringia, Jena, Kunitz, Kunitzburg, Germany 43742.1 Morchella gigas 43743.1 Morchella giga huringia, Jena, Botanischer Garten, Germany 2.Feb.03 Thuringia, Jena, Kritzegraben, Germany 44194.1 Morchella conica 195.1 Morchella conica 12.Feb.03 Thuringia, Jena, Kunitz, Germany 2.Feb.03 Thuringia, Jena, Damenviertel, Germa 2.Feb.03 Thuringia, Jena, Damenvierte, Germanv 44197.1 Morchella conica 544198.1 Morchella conica 12.Feb.03 North Israel J544199.1_Morchella_elata 12.Feb.03 Solan Himachal Pradesh, India 12.Feb.03 Solan Himachal Pradesh, India 544200.1_Morchella_elata 2.Feb.03 Solan Himachal Pradesh. India 44201.1_Morchella_elata 44202.1_Morchella_esculenta 12.Feb.03 USA 12.Feb.03 Canada, BC 44203.1 Morchella angustic 2.Feb.03 Thuringia, Jena, Kunitz, Kunitzburg, Germany 44205.1_Verpa_conica 2.Feb.03 Thuringia, Jena, Kunitz, Kunitzburg, Germany 44206.1 Verpa conica 44207.1_Disciotis_venosa 2.Feb.03 Saxony-Anhalt, Beendorf, Germany 544208.2 Gyromitra esculenta 30.Aug.06 Alsace, Rhinau, France 12.Feb.03 Liverpool, Neston, United Kingdom 544209.1 Gyromitra esculent 29.Jan.04 Lovcen National Park, Serbia, Montenegro J623265.1 Morchella crassipes 26.Apr.04 Skomas Peninsula, Cyprus J698474.1 Morchella elata J698475.1_Morchella_esculenta 26.Apr.04 Huatusco, Veracruz, Mexico J698476.1 Morchella angusticeps 19.Oct.05 Mexico 1698477 1 Morchella elat 26.Apr.04 Huatusco, Veracru, Mexico 698478.1 Morchella_costata 26.Apr.04 El Paraiso, Tlaxco, Tlaxcala, Mexico 698479.1 Verpa bohemica 26.Apr.04 Bienitz, Rueckmarsdorf, Leipzig, German 26.Apr.04 California, Blodgett Forest, USA 98480.1 Gvromitra infula 269501 1 Morchella conica CD 2004 Zakopane, Poland 269502.1_Verpa_bohemica CD 2004 Krakow, Poland Q243909.1 Morchella elata 15.Oct.05 China 257329.1_Morchella_elata 19.Oct.05 China 19.Oct.05 China Q257330.1 Morchella elata Q257331.1_Morchella_elata 19.Oct.05 China 2257332.1_Morchella_costata 19.Oct.05 China 19.Oct.05 China 2257333.1 Morchella costata 19.Oct.05 China 257334.1 Morchella costata 20.Oct.05 China 257335.1 Morchella angustice 20.Oct.05 China 257336.1 Morchella angustice 57337.1 Morchella angustice 20.Oct.05 China 20.Oct.05 China 7338.1 Morchella angustice 20.Oct.05 China 7339.1 Morchella esculenta 20.Oct.05 China 57340.1 Morchella esculenta 257341.1 Morchella esculenta 20.Oct.05 China 257342 1 Morchella esculenta 20.Oct.05 China 20.Oct.05 China 257343.1 Morchella conica 57344.1 Morchella conica 20.Oct.05 China 257345.1 Morchella conica 20.Oct.05 China 2257346.1 Morchella crassipes 20.Oct.05 China 2257347.1 Morchella crassipes 20.Oct.05 China Q491503.1 Disciotis venosa 1.Apr.06 Not given -062475.1 Morchella elata 15.Oct.06 Not given F419956.1_Fungal_endophyte_is 01.Feb.07 USA. Arizona 51851.2_Morchella_esculenta CD 1987 Northern-Alsace, France 51852.2 Morchella conica ntification on Tree ollection Date Geographic Locati 34146.1_Uncultured_fungus 12.Feb.2007 Interior AK, Bonanza Creek LTER, site FP5C May.2005 Interior AK, Estle Connector Trail Long Term site, UAF now3674 May.2005 Interior AK, Estle Connector Trail Long Term site, UAF R12 {GAL}1757 28.Jun.2005 Interior AK. 2004 Boundary fire SR10_GAL17577 28.Jun.2005 Interior AK, 2004 Bolgen Creek fire

28.Jun.2005 Interior AK, 2004 Bolgen Creek fire May.2004 Interior AK, Placer, off Gilmore Trail 31.May.2007 Interior AK, University of Alaska Fairbanks Campus 16.Jun.2007 Interior AK, 2006 Nenana fire 16.Jun.2007 Inerior AK, 2006 Nenana fire None given

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None aiven

None aiven

27.May.2007 Interior AK, 1524 Whistling Swan Drive, Fairbanks

GenBank. Most are from

outside North America.

"CD" before the date of

actually collected on that

represent the GenBank

accession date when no

collection date could be

collection means

date; other dates

(Below) This table contains information on all of our samples. Note that even though we could not determine a definite geographic location for three of the samples, they were probably still from Interior Alaska. The first three samples Ian Herriott collected from the soil in a previous study.

Soil of spruce forest, not burned (3)

Spruce forest, burned

Wild iris beds, not burned

Edge of sidewalk, landscaping

Soil of spruce forest, not burned unpublishe

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2005.101-107 (5) Pilz, David, et. al. "Ecology and Management of Morels Harvested from Western North America." United Reference States Department of Agriculture. March 2007. lor, D.L., Herriott, I.C., Long, J. and O'Neill, K. "TOPO TA is A-OK: a test of phylogenetic bias in fungal Soil of spruce forest, not burned unpublished This study environmental clone library construction." Environmental Microbiology 9 (5). 2007. 1329-1334 This study This study of, D., Munch, J.C., Botton, B. and Buscot, F. "DNA polymorphism in morels: complete sequences of the This study internal transcribed spacer of genes coding for rRNA in Morchella esculenta (yellow morel) and This study This study Morchella conica (black morel)."Appl. Environtal Microbiology 62 (9). 1996. 541-3543. This study This study

We extracted DNA from 13 dried morel samples from Dr. Gary Laursen's herbarium using the ENZA fungal kit, and copied a fragment of DNA using Polymerase Chain Reaction (PCR). We then conducted gel electrophoresis, a method used to show the amount and quality of DNA in a PCR sample. We purified the PCR products using the Zymo DNA Clean and Concentrator[™]-5 kit a cycle sequencing reaction, in which the order nitrogenous bases (the genetic code) is determined. We ran les and then purified the results, and submitted the results to the AF DNA core lab facility, where the samples were run on a OXL genetic analyzer. After receiving the data, we imported it to a computer and assembled it using Sequencher, then aligned it sing Clustal. We also gathered additional morel samples from GenBank, an online database containing tens of millions of DNA sequences. Our final step involved building a phylogenetic tree of our samples along with those from GenBank using various algorithms (e.g. Parsimony, Neighbor Joining and Maximum Liklihood). (Background photo by

) Wipf, D. Polymorphisms proteique et genomique qu sein des Morchellaceae-Mise au point d'un outil moleculaire adapte a l'etude de l'ecologie du genre Morchella en milieu forestier. Thesis (1997) University H Poincare N I, France.

d Kellner, Carsten Renker, Francois Buscot. "Species diversity within the Morchella esculenta group (Ascomycota: Morchellaceae) in Germany and France." Organisms, Diversity & Evolution 5. Our phylogenetic tree is displayed in Figure 1. There is strong support for a monophyletic (all descendents from a single common ancestor) Morchella genus with Gyromytra as our outgroup. We found that Verpa and Discoti, sister genera to Morchella, had only two and one named species, respectively, in GenBank. Within Morchella, major subdivisions exist. We have samples closely related to the major group containing what are called *M. conica*, M. elata, and M. costata, often called "Blacks." Interestingly, none of our samples are contained within the major goup known as "Yellows," and we did not have any members of *M. gigas*. Within the "black" goup, we found close relationships between DNA isolated directly from soil (IHsnow2582, IHsnow3674, EF434146.1) with DNA from sporocarps (KSR5_GAL16217, KSR6_GAL20388). KSR7_GAL20410 and KSR8_GAL2411 group with Genbank samples identified as M. conica. Samples 1, 10, 12 and 13 form a tight, distinct group. One member of the this group looks much different from the others and was identified as what has recently been called a "fuzzy foot morel," while the rest appeared to be "gray morels." However, our tree shows there is no genetic distinction, and the "fuzzy foot" is simply an immature gray morel, which supports a recent field observation (Pilz et al). We also discovered several samples from GenBank whose species attributions appear incorrect.

The absence of yellow morels in our samples hints that there are few, if any, in Alaska, though the geographic extent and total number of samples in this study is small. During 2005 in a burn just north of the Yukon near Canada, I picked a thousand or more morels and have never seen anything resembling *M. esculenta*, which is one of the most distinctive species groups, nor have I heard of any being obtained by any fellow morel gatherers. However, absence of evidence is not evidence of absence, and extensive further sampling is needed throughout Alaska and Canada before a "complete" picture of morel geographic diversity can be painted. This study represents a first step in that direction.

To our knowledge this is the first report of DNA sequences from a morel species to be found both aboveground and belowground. Detecting and identifying morel DNA from soil is important to understanding their ecology during the majority of their lifecycle. All soil sequences reported here were closely related and, interestingly, grouped with the only two aboveground morels that came from non-burned habitats (KSR5_GAL16217 – from wild iris beds near an old log pile, and KSR11_(GAL20701) from bare soil near a landscaped area on the UAF campus). However one of the 6 total Alaskan samples from this group did come from a burn (KSR5_GAL20388). Much further Alaskan sampling above and belowground, in burns and non-burns, coupled with habitat information is needed to establish confidence in any possible ecological patterns.

ORIGIN OF THE FUZZY FOOT/GRAY MORELS

The phylogenetic algorithms used to infer our phylogenies all resulted in similar trees. The Maximum Liklihood tree we show in Figure 1 is representative. However, different trees often placed the fuzzy foot group on different branches. Some placed the group near the "trunk" (as in Figure 1). Since branches closer to the base mean older in evolutionary history, this suggests that the group diverged early. However, other trees placed the group in with the "blacks" but on an unusually long branch (meaning fast evolution occured). Because small, isolated populations tend to evolve faster than other populations, one hypothesis could be that the fuzzy foot group was trapped in refugia and underwent a genetic "bottleneck", possibly during the last Ice Age. However, fuzzy foots are found in the contiguous U.S., and a comparison between these two groups is needed to test this and other hypotheses.

CONCLUSION

Some samples from GenBank have inconsistenly applied "species names," shown by the fact that hey are more similar genetically to members of different evolutionary groups than to the species" to which they are attributed. This shows how little we know about morel taxonomy and the difficulty of applying morphological identifications to morels that vary widely in appearance across their lifespan. While much needs to be learned about morel biology and ecology, we first must understand what species we are discussing, or otherwise communicating study results is useless. Future studies on morel genetic diversity should involve sequencing a different gene as well as a larger and more diverse array of samples, allowing us to obtain a clearer and more concise picture of emerging morel evolution and ecology.

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INBR

DISCUSSION

THE REPORT OF A DESCRIPTION OF A

WHERE ARE THE ESCULENTAS AND OTHER YELLOW MORELS?

(BURNED vs. DISTURBED HABITATS, ABOVE vs. BELOWGROUND)