























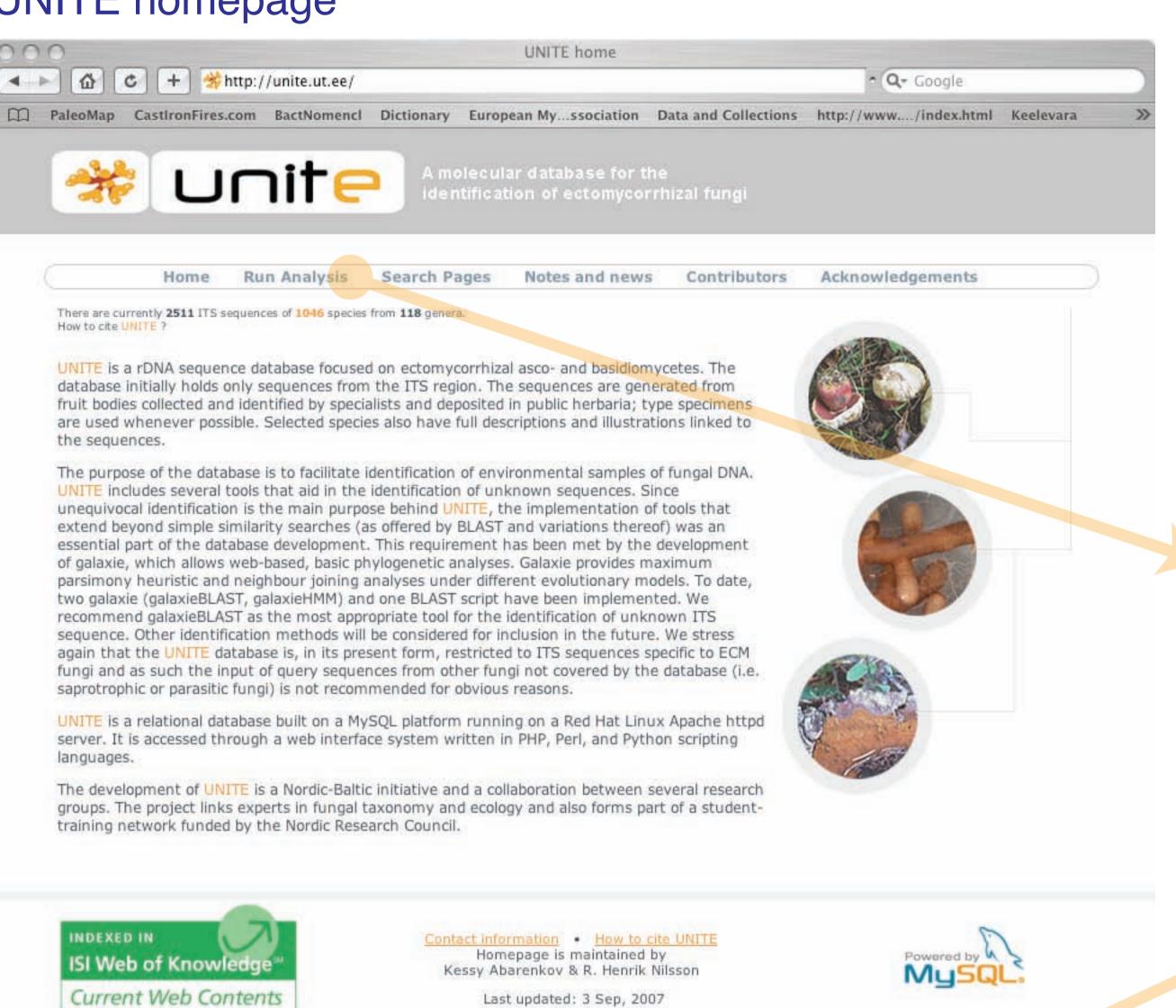
## A database providing web based tools for the molecular identification of ectomycorrhizal fungi - DNA barcoding in action

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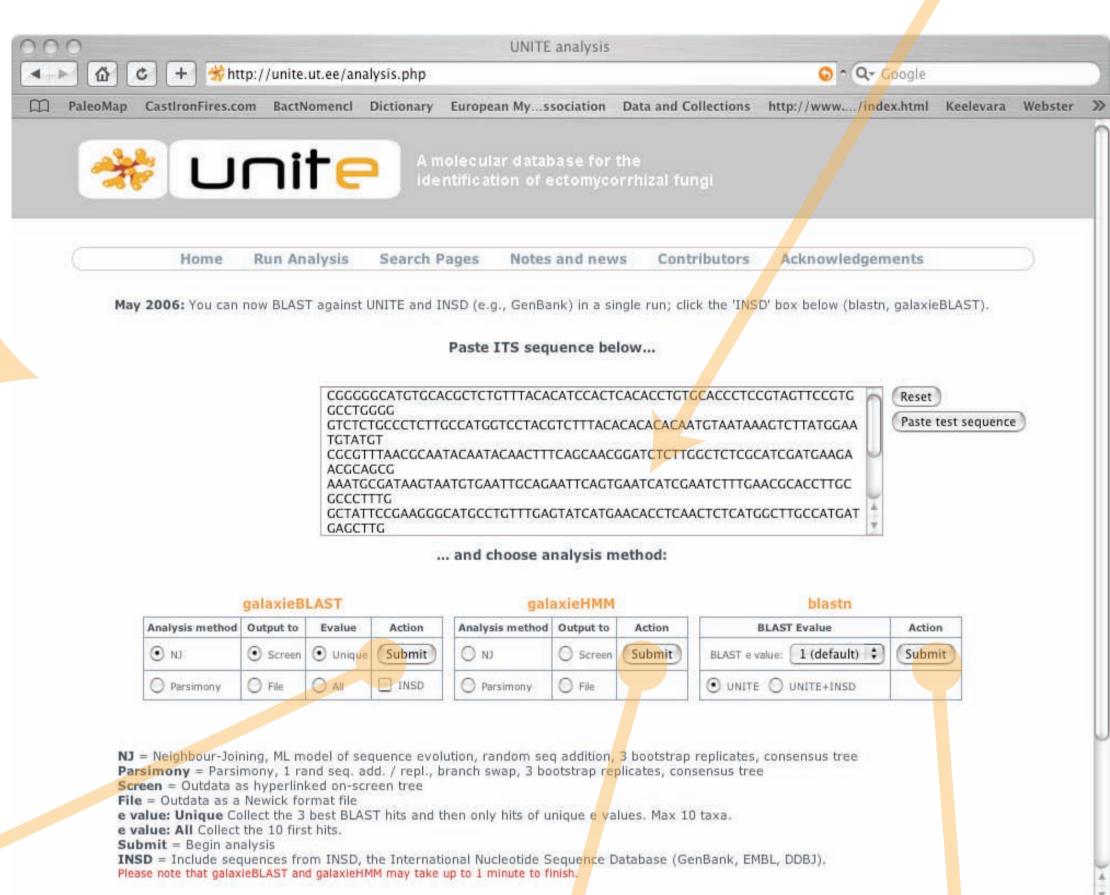
1 University of Tartu, Estonia; 2 Göteborg University, Sweden; 3 University of Aberdeen, UK; 4 Macaulay Institute, UK; 5 Centraalbureau voor Schimmelcultures, Netherland; 6 Lund University, Sweden; 7 University of Copenhagen, Denmark; 8 Finnish Forest Research Institute, Finland; 9 Manchester Metropolitan University, UK; 10 Swedish Univ. of Agricultural Sciences, Sweden; 11 University of Helsinki, Finland; 12 University of Oslo, Norway; 13 Karolinska Institutet, Sweden

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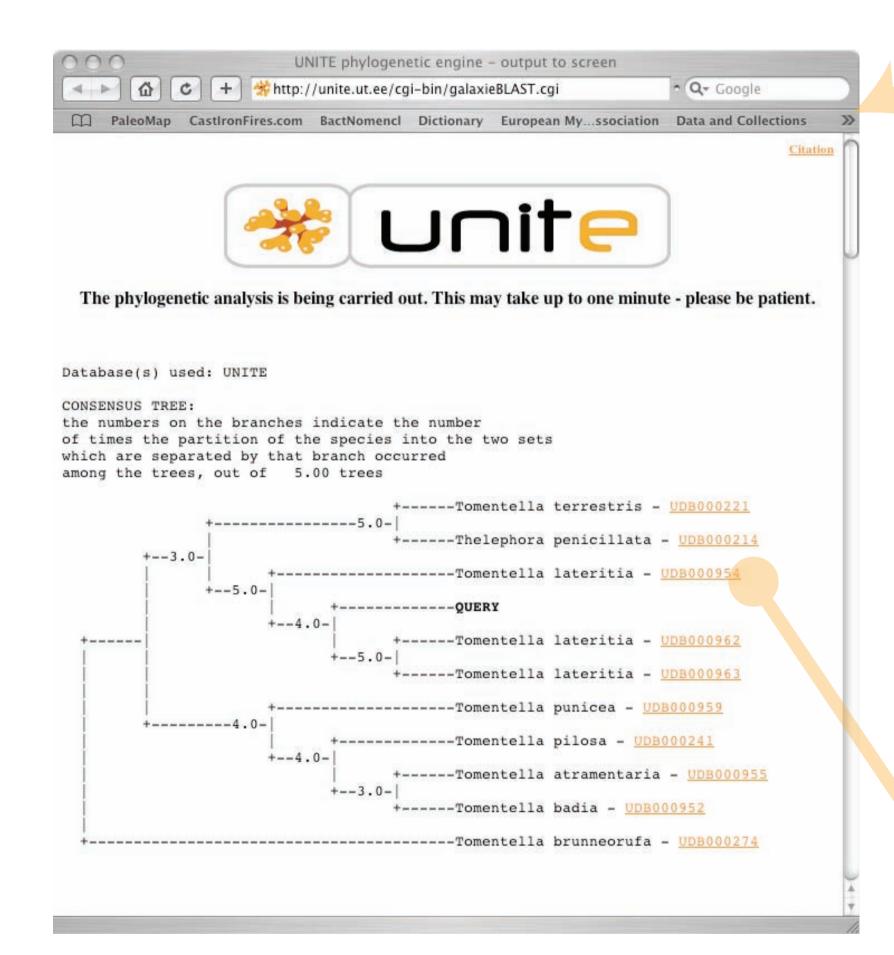
## **UNITE** homepage











The phylogenetic analysis is being carried out. This may take up to one minute - please be patient.

This ITS matrix contains species from the genera Tomentella and Thelephora.

Consensus year:
the numbers on the branches indicate the number of times the partition of the species into the two sets which are separated by that branch occurred along the treas, out of 5.00 trees

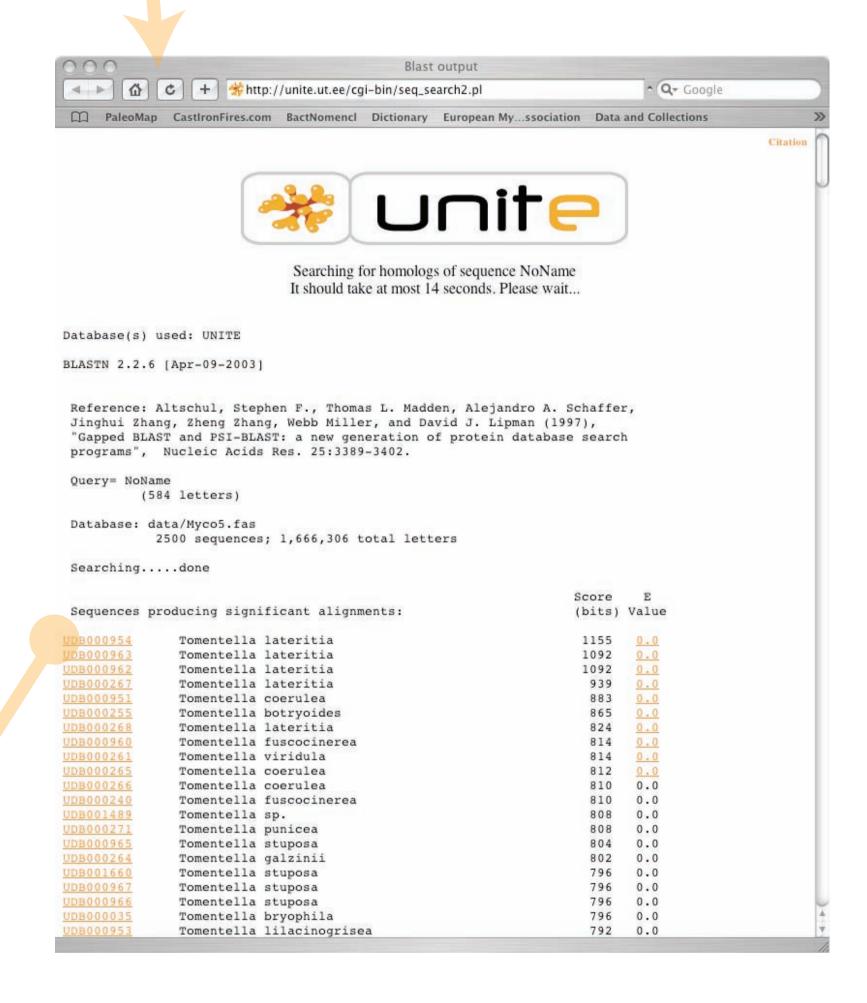
Another trees, out of 5.00 trees

Thelephora numbers of the property of the sets of the sets which are separated by that branch occurred along the trees, out of 5.00 trees

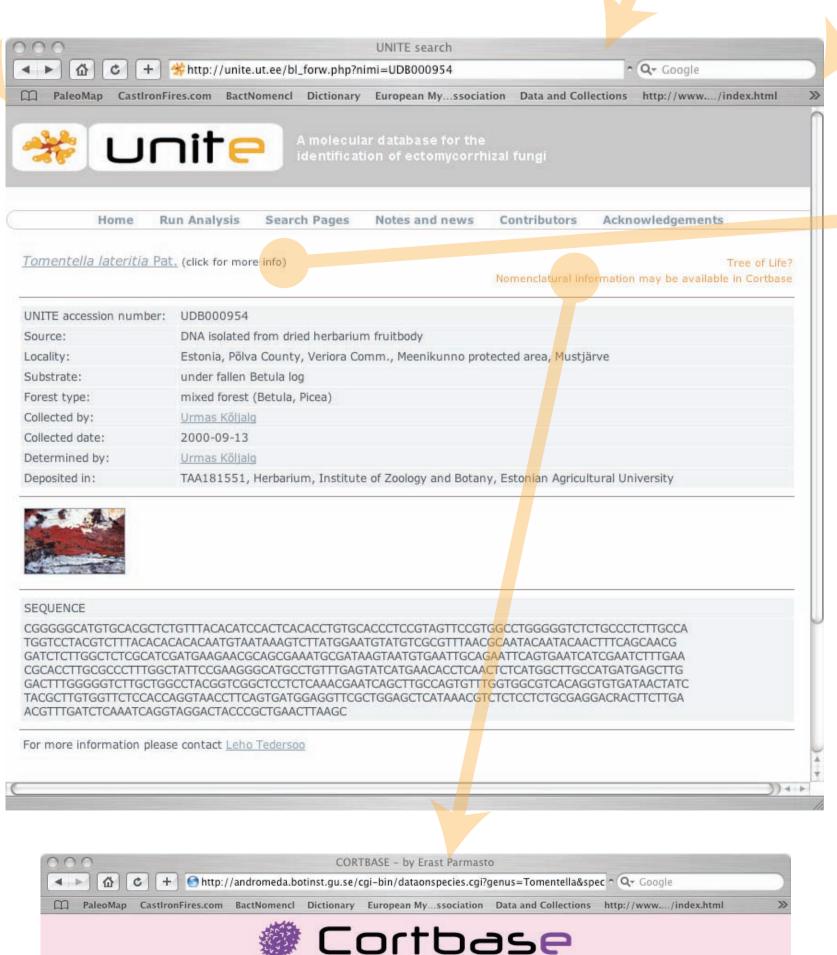
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UNITE phylogenetic engine - output to screen

▲ ▶ G C + 参http://unite.ut.ee/cgi-bin/galaxieHMM.cgi



Composite of "screenshots" displaying some features of UNITE. In the home page of UNITE you can run phylogenetic analyses by using galaxie-BLAST or galaxieHMM as well as BLAST your ITS sequences against the UNITE database. Support for INSD (GenBank, EMBL, DDBJ) ITS sequences is enabled. This means that you, in a single BLAST or galaxieBLAST run, can compare your sequences with those in UNITE and INSD. All you have to do is to click the 'INSD' box next to the 'Submit' button. We use emerencia technology for the INSD support. In galaxie and BLAST search results pages the terminal taxa and best matches respectively have interactive links to the specimen data. The species name in the last page has an interactive link to the species description if available. The species description includes text as well as illustrative material and a link to the nomenclatural database Cortbase. There are also search pages where you can browse UNITE sequences by taxon name.



Parmasto E, Nilsson H, Larsson K-H. 2004. Cortbase version 2. Extensive updates of a nomenclatural database for corticioid fungi (Hymenomycetes).

No matches retrieved. Exiting.
You may have forgotten to specify a name

END OF CORTBASE OUTDATA



▲ ► 🏠 🌣 + 👋 http://unite.ut.ee/sp\_desc.php?sp\_id=AA054

Tomentella lateritia Pat

PaleoMap CastlronFires.com BactNomencl Dictionary European My...ssociation Data and Collections http://www.../index.html

Home Run Analysis Search Pages Notes and news Contributors Acknowledgements

nophore near Brick or Umber (near 2.5YR 4-5/6 or 5.0YR 5/6) when dry and near Rust (near 10.0R 5/8)

Pomentella lateritia Pat., J. Bot. 8: 221. 1894 (and as a \"new species\" in Catal. Raissonne

. Basidiocarps resupinate, adherent to the substratum, arachnoid or mucedinoid, continuous

Hyphal cords absent. Subicular hyphae clamped, 2.5-4 μm diam., thin-walled, without

when fresh, granulose, turning darker than subiculum (often subiculum almost absent or very thin

encrustation, hyaline in 3% KOH.
Subhymenial hyphae clamped, 2.5-4.5  $\mu m$  diam., thin-walled, hyaline in 3% KOH. Cystidia absent.

 Basidia 35-55 μm long and 6-7.5 μm diam. at apex, clamped at base, clavate or utriform, not stalked, sinuous, rarely with transverse septa, hyaline or ochraceous in 3% KOH, hyaline or red in

part) or rarely lobed frontal and ellipsoid lateral face, echinulate, pale brown in 3% KOH and in distilled water. Chlamydospores absent.

Basidiospores 7-8.5 μm long in lateral and frontal face, triangular (with widened proxima.

Synonym: Hypochnus subvinosus Burt, Ann. Missouri Bot. Gard. 3: 231. 1916. (FH)
Tomentella subvinosa (Burt) Bourdot & Galzin, Bull. Soc. Mycol. France 40: 146. 1924

