

Supporting Information

An integrated chromatographic strategy for the large-scale extraction of ergosterol from *Tulasnellaceae* sp

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File S1. Species identification of the Endophytic Fungus *Tulasnellaceae* sp. from

Gymnadenia orchidis

Tulasnellaceae sp. CBS 508.93 18S ribosomal RNA gene, partial sequence; internal transcribed spacer 1, complete sequence; and 5.8S ribosomal RNA gene, partial sequence

LOCUS 272 bp DNA linear PLN 30-SEP-2013

DEFINITION Tulasnellaceae sp. CBS 508.93 18S ribosomal RNA gene, partial sequence; internal transcribed spacer 1, complete sequence; and 5.8S ribosomal RNA gene, partial sequence.

ACCESSION KF267025

VERSION KF267025.1

KEYWORDS .

SOURCE Tulasnellaceae sp. CBS 508.93

ORGANISM [Tulasnellaceae sp. CBS 508.93](#)

Eukaryota; Fungi; Dikarya; Basidiomycota; Agaricomycotina;
Agaricomycetes; Cantharellales; Tulasnellaceae.

REFERENCE 1 (bases 1 to 272)

AUTHORS Waud,M.J., Busschaert,P., Ruyters,S., Jacquemyn,H. and Lievens,B.

TITLE Impact of primer choice on characterization of orchid mycorrhizal communities using 454 pyrosequencing

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REFERENCE 2 (bases 1 to 272)

AUTHORS Waud,M.J., Busschaert,P., Ruyters,S., Jacquemyn,H. and Lievens,B.

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Biology Department, KU Leuven, Kasteelpark Arenberg 31, Heverlee,

Flemish Brabant B-3001, Belgium

COMMENT ##Assembly-Data-START##

Sequencing Technology :: Sanger dideoxy sequencing

##Assembly-Data-END##

FEATURES Location/Qualifiers

source

1..272

/organism="Tulasnellaceae sp. CBS 508.93"

/mol_type="genomic DNA"

/strain="CBS 508.93"

/culture_collection="CBS:[508.93](#)"

/db_xref="taxon:[1403782](#)"

/PCR_primers="fwd_name: NS7, fwd_seq:

gaggcaataacaggctctgatgc, rev_name: ITS2, rev_seq:

gctgcgttctcatcgatgc"

[misc_RNA](#)

<1..>272

/note="contains 18S ribosomal RNA, internal transcribed spacer 1, and 5.8S ribosomal RNA"

ORIGIN

1 aactcggcca ttagaggaa gtaaaagtcg taacaagggt tccgtaggtg aacctgcgga

61 aggatcatta atgagtttgc gcggtcgtg ctggttcta caagtcaca cctgtcttt

121 cttatccacc ccaactgtgca tcccgtacg ctttcgagcg tagtccttt acacacaacc

181 attgtaattg aattagaacg tgcgcgaatg ataataataa tacaactatc aacaacggat

241 ctcttgcat ccaactgat gaagaacgca gc

Figure S1. ESI mass spectrum of ergosterol.

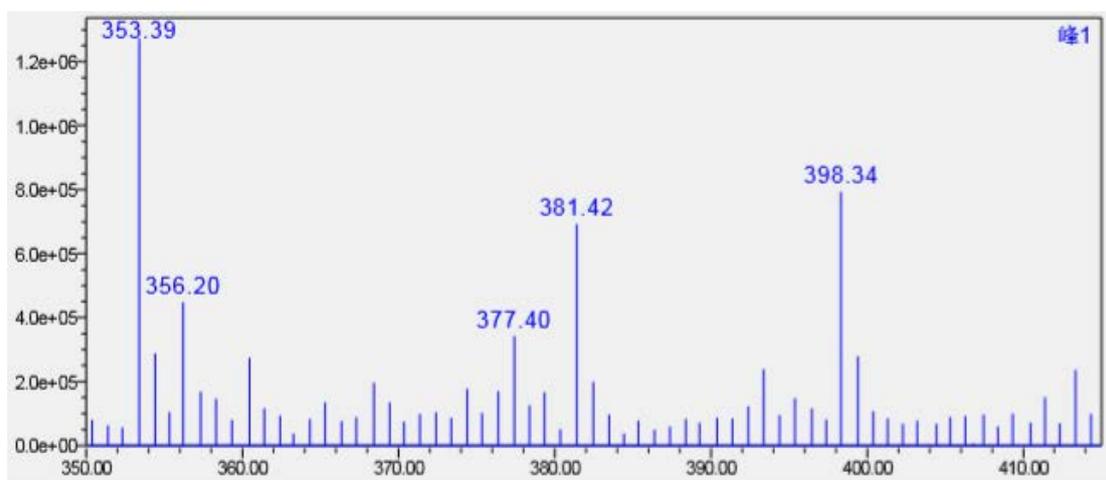


Figure S2. ^1H NMR Spectrum (600 MHz) of ergosterol on the ReproSil C18 analytical column in water/ethanol (in CDCl_3).

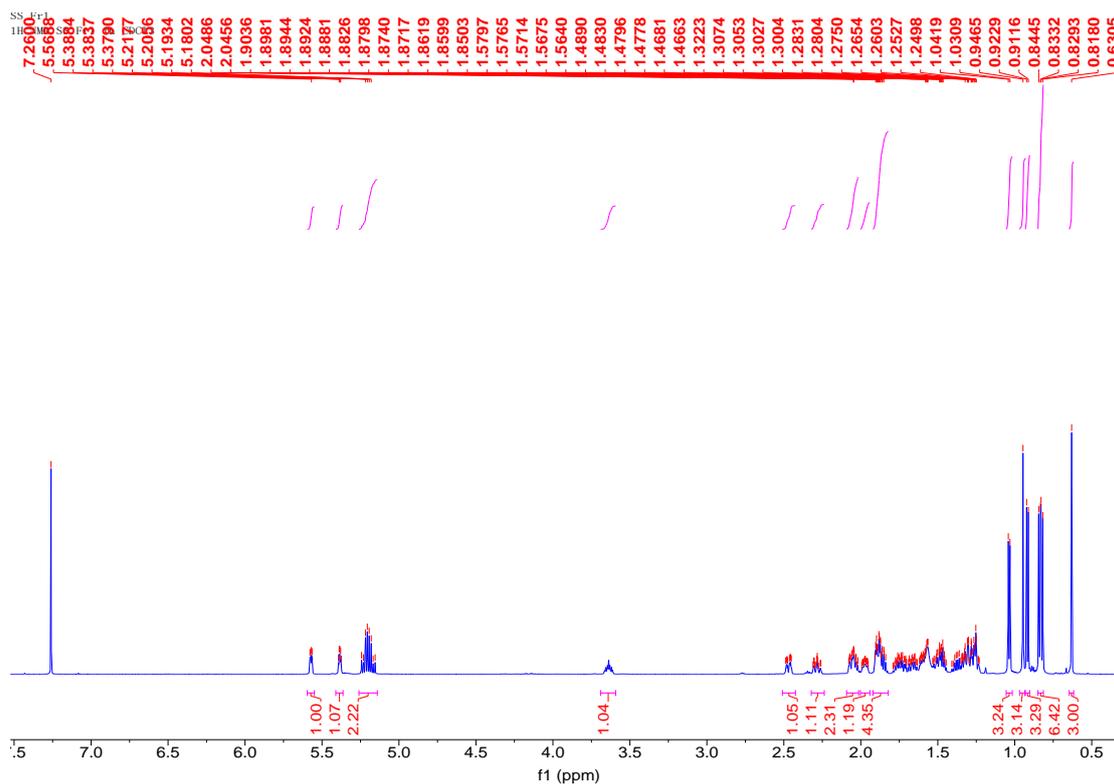


Figure S3. ^{13}C NMR Spectrum (151 MHz) of ergosterol on the ReproSil C18 analytical column in water/ethanol (in CDCl_3).

SS Fr1
13C NMR SS Fr1 in CDCl_3

