Moniliophthora perniciosa strain 1188 18S ribosomal RNA, partial sequence; internal transcribed spacer 1, 5.8S ribosomal RNA, and internal transcribed spacer 2, complete sequence; and 28S ribosomal RNA, partial sequence

GenBank: EU047936.1

FASTA Graphics

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LOCUS EU047936 770 bp DNA linear PLN 18-AUG-2007
DEFINITION Moniliophthora perniciosa strain 1188 18S ribosomal RNA, partial sequence; internal transcribed spacer 1, 5.8S ribosomal RNA, and internal transcribed spacer 2, complete sequence; and 28S ribosomal RNA, partial sequence.
ACCESSION EU047936
VERSION EU047936.1
KEYWORDS .
SOURCE Moniliophthora perniciosa (Crinipellis perniciosa)
ORGANISM Moniliophthora perniciosa
Eukaryota; Fungi; Dikarya; Basidiomycota; Agaricomycotina; Agaricomycetes; Agaricomycetidae; Agaricales; Marasmiineae; Marasmiaceae; Moniliophthora.
REFERENCE 1 (bases 1 to 770)
AUTHORS Marelli,J.-P., Gramacho,K.P., Kang,S., Maximova,S.N. and Guiltinan,M.J.
TITLE Tomato as a model system to study pathogenicity mechanisms of Moniliophthora perniciosa, the causal agent of witches' broom disease of cacao
JOURNAL Unpublished
REFERENCE 2 (bases 1 to 770)
AUTHORS Marelli,J.-P.
TITLE Direct Submission
JOURNAL Submitted (22-JUL-2007) Plant Pathology, The Pennsylvania State University, 312 Buckhout Lab, University Park, PA 16802, USA
FEATURES Location/Qualifiers
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misc_RNA 59..272
/product="internal transcribed spacer 1"
rRNA 273..433

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