

Company: AccuCompany

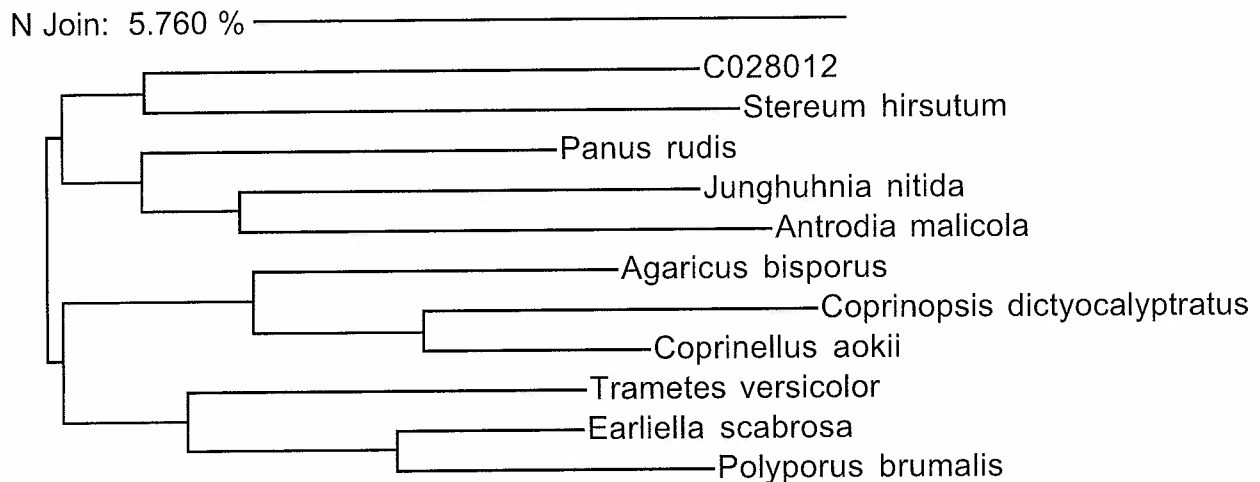
Sample: C028012 - EM sample 19702 suite 101

Date: Wednesday, April 9, 2008 5:35 PM

Sequence Alignment

Alignment: 354 C028012
10.32 % 349 Earliella scabrosa
11.14 % 350 Stereum hirsutum
11.43 % 350 Panus rudis
11.68 % 351 Polyporus brumalis
11.89 % 349 Junghuhnia nitida
12.32 % 349 Trametes versicolor
12.32 % 349 Agaricus bisporus
12.57 % 350 Antrodia malicola
13.47 % 349 Coprinellus aokii
13.47 % 349 Coprinopsis dictyocalyptratus

Neighbor Joining Tree



Accugenix Database Search Result

Generated by: _____

Identification: No Match

Reviewed by: _____

Confidence Level: No Match

QA Review: _____

Batch #: 80410015

ACC1 - Due 4/10/08

NOT FOR USE IN INVITRO DIAGNOSTICS

SOP-GEN-049
Rev. 01Apr08 JM