

Appendix 4.0 Bootstrap maximum parsimony (MP) tree of the ITS region

Tree number 2 (rooted using default outgroup)

Tree length = 178
Consistency index (CI) = 1.0000
Homoplasy index (HI) = 0.0000
CI excluding uninformative characters = 1.0000
HI excluding uninformative characters = 0.0000
Retention index (RI) = 1.0000
Rescaled consistency index (RC) = 1.0000

```
/ 2954
+ 2953
+ 2952
+ 0362
+ 0361
+ 5059
/--+ 1350
| + 3533
| + 3532
| + 1349
/-----+
+ 0301
|
| + 0201
|
| \-- GQ203796
|
+ 0658
|
\ 0657
\-----
--- Phoma sp. HQ608114

/----- 2954
+----- 2953
+----- 2952
+----- 0362
+----- 0361
```

+----- 5059
/-----
-----+-----
--- 1350
|
+----- 3533
|
+----- 3532
|
+----- 1349
/-----+
+----- 0301
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+----- 0201
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\\----- GQ203796
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--- Phoma sp. HQ608114