

# Infer\_ST\_Bootstrap Command

## Description

Infers the species tree using bootstrap with existing methods.

## Usage

```
Infer_ST_BOOTSTRAP num [-s threshold] other_inferST_command
```

|                              |  |           |
|------------------------------|--|-----------|
| <i>num</i>                   | The number of repetitions.   | mandatory |
| <i>-s threshold</i>          | Threshold value. If unspecified, defaults to 0.5.                              | optional  |
| <i>other_inferST_command</i> | The command name (and parameters) for any other Phylonet InferST_XXXX command. | mandatory |

## Examples

```
#NEXUS

BEGIN NETWORKS;

Network g1 = (((a:5,b:5):4,c:9):3,d:12):3,e:15);
Network g2 = ((a:6,b:6):11,((c:12,e:12):2,d:14):3);
Network g3 = ((a:8,c:8):7,((b:14,e:14):1,d:15));

END;

BEGIN PHYLONET;

Infer_ST_BOOTSTRAP 3 -s .6 Infer_ST_MDC_Time (g1, g2, g3);

END;
```

## See Also

- [List of PhyloNet Commands](#)