

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 <i>InferST_XXXX</i> 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](#)