

Table 1: Responders to both ¹³C-xylose ¹³C-cellulose BLAST
against Living Tree Project

| OTU ID | Amendment | Fold change _a | Day ^b | All days _c | Top BLAST hits | BLAST %ID | Phylum;Class;Order |
|---------|-----------|--------------------------------|------------------|--------------------------|--|-----------|---|
| OTU.11 | 13CXPS | 5.25 | 7 | 7 | <i>Stenotrophomonas pavanii</i> , <i>Stenotrophomonas maltophilia</i> , <i>Pseudomonas geniculata</i> | 99.54 | <i>Proteobacteria</i> <i>Gammaproteobacteria</i> <i>Xanthomonadales</i> |
| OTU.11 | 13CCPS | 3.41 | 14 | 14 | <i>Stenotrophomonas pavanii</i> , <i>Stenotrophomonas maltophilia</i> , <i>Pseudomonas geniculata</i> | 99.54 | <i>Proteobacteria</i> <i>Gammaproteobacteria</i> <i>Xanthomonadales</i> |
| OTU.150 | 13CCPS | 4.06 | 14 | 14 | No hits of at least 90% identity | 86.76 | <i>Planctomycetes</i> <i>Planctomycetacia</i> <i>Planctomycetales</i> |
| OTU.150 | 13CXPS | 3.08 | 14 | 14 | No hits of at least 90% identity | 86.76 | <i>Planctomycetes</i> <i>Planctomycetacia</i> <i>Planctomycetales</i> |
| OTU.165 | 13CCPS | 3.1 | 14 | 14 | <i>Rhizobium skierniewicense</i> , <i>Rhizobium vignae</i> , <i>Rhizobium larrymoorei</i> , <i>Rhizobium alkalisoli</i> , <i>Rhizobium galegae</i> , <i>Rhizobium huautlense</i> | 100.0 | <i>Proteobacteria</i> <i>Alphaproteobacteria</i> <i>Rhizobiales</i> |
| OTU.165 | 13CXPS | 2.38 | 3 | 3 | <i>Rhizobium skierniewicense</i> , <i>Rhizobium vignae</i> , <i>Rhizobium larrymoorei</i> , <i>Rhizobium alkalisoli</i> , <i>Rhizobium galegae</i> , <i>Rhizobium huautlense</i> | 100.0 | <i>Proteobacteria</i> <i>Alphaproteobacteria</i> <i>Rhizobiales</i> |
| OTU.19 | 13CCPS | 2.44 | 14 | 14 | <i>Rhizobium alamii</i> , <i>Rhizobium mesosinicum</i> , <i>Rhizobium mongolense</i> , <i>Arthrobacter viscosus</i> , <i>Rhizobium sullae</i> , <i>Rhizobium yanglingense</i> , <i>Rhizobium loessense</i> | 99.54 | <i>Proteobacteria</i> <i>Alphaproteobacteria</i> <i>Rhizobiales</i> |
| OTU.19 | 13CXPS | 2.14 | 7 | 7 | <i>Rhizobium alamii</i> , <i>Rhizobium mesosinicum</i> , <i>Rhizobium mongolense</i> , <i>Arthrobacter viscosus</i> , <i>Rhizobium sullae</i> , <i>Rhizobium yanglingense</i> , <i>Rhizobium loessense</i> | 99.54 | <i>Proteobacteria</i> <i>Alphaproteobacteria</i> <i>Rhizobiales</i> |
| OTU.241 | 13CXPS | 3.38 | 3 | 3, 14 | No hits of at least 90% identity | 87.73 | <i>Verrucomicrobia</i> <i>Spartobacteria</i> <i>Chthoniobacterales</i> |
| OTU.241 | 13CCPS | 2.66 | 14 | 14 | No hits of at least 90% identity | 87.73 | <i>Verrucomicrobia</i> <i>Spartobacteria</i> <i>Chthoniobacterales</i> |
| OTU.32 | 13CXPS | 3.0 | 3 | 3, 7, 14 | <i>Sandaracinus amylolyticus</i> | 94.98 | <i>Proteobacteria</i> <i>Deltaproteobacteria</i> <i>Myxococcales</i> |

Table 1 – continued from previous page

| OTU ID | Amendment | Fold change | Day | All days | Top BLAST hits | BLAST %ID | Phylum;Class;Order |
|--------|-----------|-------------|-----|----------|--|-----------|---|
| OTU.32 | 13CCPS | 2.34 | 3 | 3 | <i>Sandaracinus amylolyticus</i> | 94.98 | <i>Proteobacteria</i> <i>Deltaproteobacteria</i> <i>Myxococcales</i> |
| OTU.5 | 13CXPS | 3.69 | 7 | 7 | <i>Delftia tsuruhatensis</i> , <i>Delftia lacustris</i> | 100.0 | <i>Proteobacteria</i> <i>Betaproteobacteria</i> <i>Burkholderiales</i> |
| OTU.5 | 13CCPS | 2.69 | 14 | 14 | <i>Delftia tsuruhatensis</i> , <i>Delftia lacustris</i> | 100.0 | <i>Proteobacteria</i> <i>Betaproteobacteria</i> <i>Burkholderiales</i> |
| OTU.6 | 13CCPS | 3.62 | 7 | 3, 7, 14 | <i>Cellvibrio fulvus</i> | 100.0 | <i>Proteobacteria</i> <i>Gammaproteobacteria</i> <i>Pseudomonadales</i> |
| OTU.6 | 13CXPS | 3.24 | 3 | 3 | <i>Cellvibrio fulvus</i> | 100.0 | <i>Proteobacteria</i> <i>Gammaproteobacteria</i> <i>Pseudomonadales</i> |

^a Maximum observed \log_2 of fold change.^b Day of maximum fold change.^c All response days.