

**Supplemental Material S1- Prominent bacterial families (%) present in endophyte infected and endophyte free tall fescue soil**

Phylum	Order (EI, EF)	Family	Endophyte Infected (EI)	Endophyte Free (EF)
Proteobacteria	Rhizobiales (38, 37)	<i>Bradyrhizobiaceae</i> ( <i>N-fix</i> ) ( <i>Myc-Symb</i> )	26	26
		<i>Hyphomicrobiaceae</i> ( <i>Pht</i> )	13	18
		<i>Rhizobiales_Incertae_Sedis</i> ( <i>N-Fix</i> )	10	10
		<i>Xanthobacteraceae</i> ( <i>N-Fix</i> )	35	30
		<i>Rhodobiaceae</i> ( <i>N-Fix</i> )	4	6
		<i>Phyllobacteriaceae</i>	4	Abs
		<i>Methylbacteriaceae</i> ( <i>PGP</i> )	3	Abs
	Rickettsiales (13, 12)	<i>Mitochondria</i>	86	77
	Xanthomonadales (6, 14)	<i>Xanthomonadales_Incertae_Sedis</i>	80	75
		<i>Xanthomonadaceae</i> ( <i>PP</i> )	5	9
	Rhodospirillales (7, 6)	<i>Rhodospirillales_Incertae_Sedis</i>	20	33
		<i>Rhodospirillaceae</i> ( <i>N-fix</i> )	14	22
		<i>Acetobacteraceae</i>	26	11
	Nitrosomonadales (8, 9)	<i>Nitrosomonadaceae</i> ( <i>AOB</i> )	100	100
	Burkholderiales (7, 3)	<i>Comamonadaceae</i> ( <i>Dntf</i> ) ( <i>Myc-Symb</i> )	38	54
		<i>Burkholderiaceae</i> ( <i>PSB</i> ) ( <i>ZnSb</i> ) ( <i>DrtTol</i> )	38	23
		<i>Oxalobacteraceae</i> ( <i>Myc-Symb</i> )	24	15
	Sphingomonadal es (3, 25)	<i>Sphingomonadaceae</i> ( <i>Biocon</i> ) ( <i>PhtRem</i> )	96	100
	Desulfuromonadales (1, 5)	<i>Geobacteraceae</i> ( <i>FeRd</i> ) ( <i>SlfRB</i> )	100	100
	Desulfurellales (5, 4)	<i>Desulfurellaceae</i> ( <i>SlfRB</i> )	100	100
Planctomycetes	Planctomycetales (88, 100)	<i>Planctomycetaceae</i> ( <i>NH3-Ox</i> )	100	95
		<i>Phycisphaeraceae</i>	18	5
Verrucomicrobia	Chthoniobacterial es (99,98)	<i>DA101 soil group</i>	86	86
		<i>Xiphinematobacteraceae</i>	3	7
		<i>Chthoniobacteraceae</i>	7	7
	Verrucomicrobia les (6, 2)	<i>Verrucomicrobiaceae</i>	100	100

Bacteroidetes	Sphingobacterial es (68, 87)	<i>Sphingobacteriaceae (Biocon)</i>	10	10
		<i>Chitinophagaceae</i>	84	89
	Cytophagales (7, 8)	<i>Cytophagaceae</i>	100	100
		<i>Flavobacteriaceae (PGPR)</i>	98	80
Nitrospirae	Flavobacteriales (25, 4)	<i>NS9 marine group</i>	2	20
		<i>Nitrospiraceae (NB)</i>	69	70
	Nitrospirales (100, 100)	<i>0319-6A21 (NB)</i>	31	30
		<i>Blastocatellales (67, 79)</i>	<i>Blastocatellaceae (Subgroup_4)</i>	100
Acidobacteria	Acidobacteriales (25, 0)	<i>Acidobacteriaceae (Subgroup_1) (NitR)</i>	100	Abs
		<i>Solibacteriales (1, 0)</i>	<i>Solibacteraceae (Subgroup_3)</i>	100
Gemmatimonad etes	Gemmatimonada les (100, 100)	<i>Gemmatimonadaceae</i>	100	100
Actinobacteria	Solirubrobacterial es (27, 43)	<i>Elev-16S-1332</i>	81	61
		<i>0319-6M6</i>	11	11
		<i>Solirubrobacteraceae</i>	Abs	11
	Micrococcales (17, 21)	<i>Micrococcaceae (SlntRlf)</i>	91	100
	Gaiellales (20, 10)	<i>Gaiellaceae</i>	20	25
	Acidobacteriales (0, 7)	<i>Acidobacteriaceae (Subgroup_1)</i>	Abs	100
	Corynebacteriale s (10, 0)	<i>Mycobacteriaceae</i>	100	Abs
	Acidimicrobiales (14, 0)	<i>Acidimicrobiaceae</i>	5	Abs
Chloroflexi	Anaerolineales (61, 53)	<i>Anaerolineaceae</i>	100	100
	Ktedonobacterial es (23, 23)	<i>HSB_OF53-F07</i>	22	39
		<i>Ktedonobacteraceae</i>	48	16
		<i>Thermosporotrichaceae</i>	5	Abs
Firmicutes	Chloroflexales (5, 5)	<i>Roseiflexaceae</i>	100	100
	Bacillales (84, 65)	<i>Bacillaceae (PSB)</i>	80	84
		<i>Paenibacillaceae (PSB)</i>	8	10
		<i>Alicyclobacillaceae</i>	4	6
	Clostridiales (6, 13)	<i>Clostridiaceae I (PP)</i>	36	40
		<i>Gracilibacteraceae</i>		10
		<i>Peptostreptococcaceae</i>	Abs	30
		<i>Ruminococcaceae</i>	Abs	10
		<i>Lachnospiraceae</i>	64	10
	Erysipelotrichale s (10, 22)	<i>Erysipelotrichaceae</i>	100	100

Abbreviations for bacterial functions within EI and EF rhizosphere soil: AnPhtB, anaerobic photosynthetic bacteria; AOB, Ammonia oxidizing bacteria; BC, bio-controller; DnB, denitrifying bacteria; FBC, fungal bio-controller; FeRB, iron reducing bacteria; N-fix, nitrogen fixing bacteria; NOA, nitrite oxidizing Archaea; NOB, nitrite oxidizing bacteria; NitR, nitrate reducers; ; Pdtx, Phosphate detoxifier; PGP, plant growth promoter; Pht, phototrophic Bacteria; PP, Plant Pathogen; PrSlfB, purple sulfur reducing bacteria; PSB, phosphorus solubilizing bacteria; SlfRB, sulfur reducing bacteria and UD, undetermined. \*Abs = Absent. Percent abundance of order is given in parenthesis for respective families.