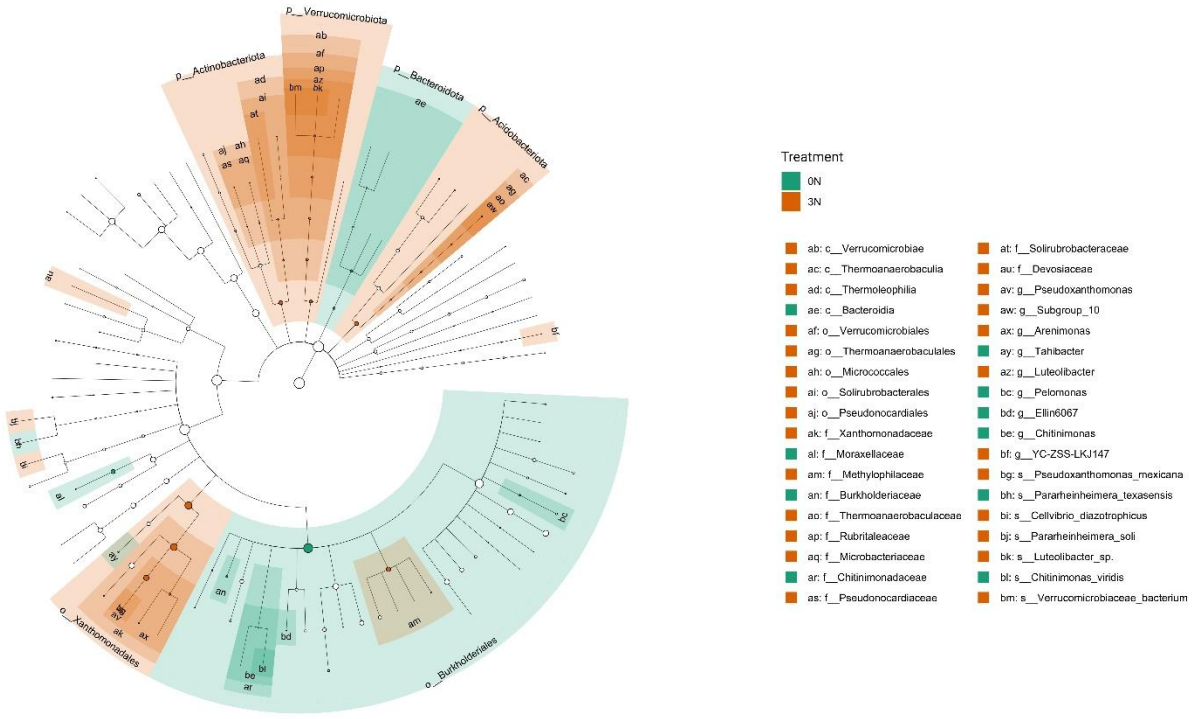
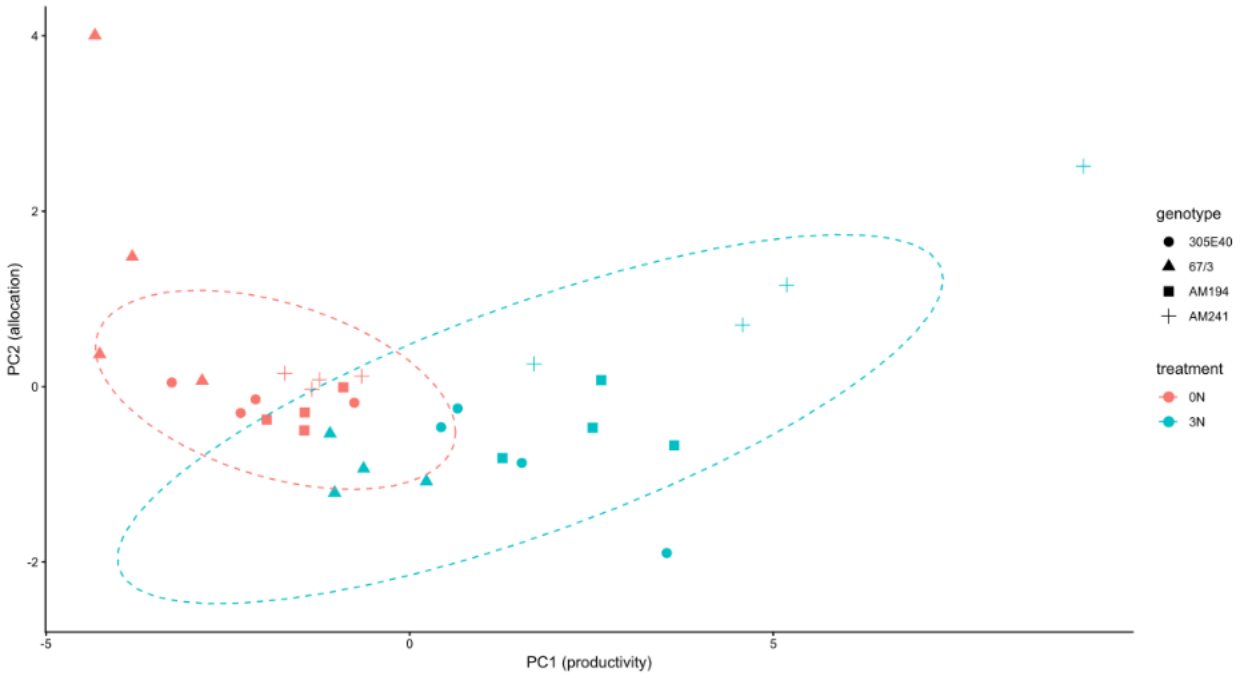
Supplementary Figures



Supplementary Figure 1 - LEfSe cladogram showing taxonomic lineages differentially enriched between N treatments (0N; Green and 3N; Orange). Nodes represent taxa significantly associated with each genotype.



Supplementary Figure 2 - PCA was performed on scaled production traits. PC1 (75.0%) represents an overall productivity gradient, while PC2 (9.9%) reflects differences in biomass allocation.

Supplementary Tables

Supplementary Table 1 - Raw per-plant growth and yield trait measurements for four eggplant genotypes (67/3, AM241, AM194, 305E40) grown under two nitrogen regimes (0N, 3N), four replicates each, underlying the principal component analysis reported in Supplementary Table 3 and Supplementary Figure 2. Variables include fruit number (count), total and average fruit weight (g), plant height (cm), shoot diameter (mm), fresh and dry biomass of shoot tissue (g) and shoot dry matter content (dry:fresh ratio), together with the corresponding leaf-tissue measurements.

Sample-id	Genotype	Treatment	Replicate	fruit number	total weight (g)	average weight (g)	plant height (cm)	shoot diameter (mm)	fresh weight (g)	fresh weight leaf (g)	dry weight leaf	leaf dry matter content	fresh weight shoot (g)	dry weight shoot (g)	shoot dry matter
6730N1	67/3	0N	1	2,00	114,50	60,0	74,0	8,29	38,75	13,66	-35,25	-4,22	11,02	2,68	0,24
6730N2	67/3	0N	2	2,50	141,25	54,5	56,5	7,58	24,75	0,43	-31,75	-4,32	9,69	2,27	0,23
6730N3	67/3	0N	3	2,50	258,25	109,8	82,0	9,08	44,75	0,46	-37,25	-4,13	13,93	3,26	0,23
6730N4	67/3	0N	4	2,33	171,33	74,8	70,8	8,31	36,08	4,85	-34,75	-4,22	11,55	2,73	0,24
AM2410N1	AM241	0N	1	3,50	345,75	93,4	112,5	11,17	118,00	0,91	5,50	0,50	15,80	4,74	0,30
AM2410N2	AM241	0N	2	3,50	265,50	77,1	128,0	11,24	140,00	0,97	12,00	1,04	22,23	6,97	0,31
AM2410N3	AM241	0N	3	4,00	280,75	70,2	114,0	10,29	98,75	0,79	-15,25	-1,45	18,69	5,17	0,27
AM2410N4	AM241	0N	4	3,67	297,33	80,2	118,2	10,90	118,92	0,89	0,75	0,03	18,90	5,62	0,30
AM1940N1	AM194	0N	1	4,00	334,25	83,6	112,0	9,96	103,00	0,69	-9,00	-0,93	15,37	5,15	0,33
AM1940N2	AM194	0N	2	5,00	332,50	66,5	99,5	8,78	93,50	0,76	-6,00	-0,91	14,36	4,67	0,32
AM1940N3	AM194	0N	3	6,00	522,75	90,5	108,5	10,44	137,75	0,52	29,25	3,01	18,02	5,09	0,28
AM1940N4	AM194	0N	4	5,00	396,50	80,2	106,7	9,73	111,42	0,65	4,75	0,39	15,92	4,97	0,31
305E400N1	305E40	0N	1	6,00	309,25	49,7	98,0	8,83	80,75	0,66	-17,25	-2,04	13,62	4,17	0,30
305E400N2	305E40	0N	2	5,50	409,75	72,9	115,0	10,31	129,00	0,55	14,00	1,18	20,93	6,67	0,32
305E400N3	305E40	0N	3	5,50	201,25	36,9	89,0	8,31	56,50	0,53	-32,50	-3,88	12,05	3,12	0,26
305E400N4	305E40	0N	4	5,67	306,75	53,2	100,7	9,15	88,75	0,58	-11,92	-1,58	15,53	4,65	0,29
6733N1	67/3	3N	1	6,00	633,25	111,7	98,0	10,33	62,25	0,49	-35,75	-4,23	15,91	5,39	0,36

6733N2	67/3	3N	2	6,50	669,00	103,4	105,0	10,55	81,00	0,55	-24,00	-2,29	17,28	5,09	0,29
6733N3	67/3	3N	3	9,50	991,50	107,3	97,5	12,51	93,00	0,31	-4,50	-0,75	19,46	6,85	0,36
6733N4	67/3	3N	4	7,33	764,58	107,5	100,2	11,13	78,75	0,48	-21,42	-2,42	17,55	5,77	0,34
AM2413N1	AM241	3N	1	8,00	799,25	111,8	128,0	12,74	273,75	0,92	145,75	10,64	24,30	8,14	0,33
AM2413N2	AM241	3N	2	12,00	1306,75	107,6	150,5	15,38	474,50	0,77	324,00	20,40	28,54	10,55	0,36
AM2413N3	AM241	3N	3	14,00	2005,75	141,8	153,0	22,59	817,00	0,93	664,00	29,27	43,97	15,95	0,36
AM2413N4	AM241	3N	4	11,33	1370,58	120,4	143,8	16,90	521,75	0,87	377,92	20,11	32,27	11,54	0,35
AM1943N1	AM194	3N	1	9,50	1233,00	131,4	132,5	12,96	310,75	0,96	178,25	14,16	23,90	8,03	0,34
AM1943N2	AM194	3N	2	11,00	1375,00	125,9	116,0	12,87	127,00	0,58	11,00	0,74	24,21	7,62	0,32
AM1943N3	AM194	3N	3	11,50	1641,50	145,1	151,5	15,19	267,00	0,96	115,50	7,58	27,07	9,82	0,36
AM1943N4	AM194	3N	4	10,67	1416,50	134,1	133,3	13,67	234,92	0,83	101,58	7,50	25,06	8,49	0,34
305E403N1	305E40	3N	1	10,00	909,75	90,7	130,5	10,76	157,75	0,75	27,25	2,50	21,83	6,73	0,31
305E403N2	305E40	3N	2	9,00	847,25	95,8	139,0	11,33	226,50	0,73	87,50	7,66	17,53	5,50	0,31
305E403N3	305E40	3N	3	13,50	1662,75	117,9	141,5	12,52	283,25	0,52	141,75	10,92	19,77	8,85	0,49
305E403N4	305E40	3N	4	10,83	1139,92	101,5	137,0	11,53	222,50	0,66	85,50	7,03	19,71	7,03	0,37

Supplementary Table 2 - Loadings of growth and production traits on the first two principal components (PC1 and PC2) from a principal component analysis (PCA) summarizing plant production traits (see Supplementary Figure 2). Twelve scaled variables were included: fruit number, total and average fruit weight, plant height, shoot diameter, fresh and dry weight of leaves and shoots, leaf and shoot dry matter content. PC1 (75.0% of total variance) loaded consistently and positively across nearly all traits (loadings ≈ 0.24–0.33), reflecting overall plant productivity and vigor, with the notable exception of leaf fresh weight, which showed a near-zero loading (−0.097) — this trait instead loads almost entirely onto PC2. PC2 (9.95% of total variance) was driven primarily by leaf fresh weight (0.679, the strongest loading on this axis) and, to a lesser extent, leaf dry weight (0.319), with shoot dry matter content contributing the strongest negative loading (−0.423), reflecting variation in biomass allocation and tissue water content rather than overall yield.

Variable	**PC1**	**PC2**
fruit number	0,29879466	-0,20634106
total weight	0,31060589	-0,12578968
average weight	0,26328158	-0,15470329
plant height	0,29185338	-0,14969437
shoot diameter	0,32031269	0,1475904
fresh weight	0,31094146	0,26275289
fresh weight leaf	-0,09657357	0,67859776
dry weight leaf	0,30038935	0,31880005
leaf dry matter content	0,31242057	0,20036257
fresh weight shoot	0,31530958	0,14141074
dry weight shoot	0,32732772	0,05935163
shoot dry matter	0,23819096	-0,42266256

Supplementary Table 3 – Taxonomic identity and metadata for the 116 bacterial strains isolated from eggplant roots in this study, spanning four genotypes (67/3, n=30; AM241, n=35; AM194, n=35; 305E40, n=15) — annotated by their previously characterized nitrogen-use efficiency (high: 67/3, AM241; low: AM194, 305E40) — and two nitrogen regimes (0N, n=41; 3N, n=75). Isolates were recovered using two complementary strategies: non-selective isolation on R2A agar (n=80) and an Actinobacteria-selective enrichment approach (n=36; see Methods). Strains span 26 genera, identified by 16S rRNA gene sequencing to the genus or species level where resolvable (isolates identifiable only to a higher taxonomic rank are noted accordingly). All isolates were recovered from root tissue collected in 2022 from the same field site (Italy, Lodi, Montanaso Lombardo) and are preserved in the CREA – Research Centre for Viticulture and Enology microbial culture collection.

Strain identifier	Host genotype	Nitrogen treatment	Genus	Specific epithet	Geographical origin	Tissue	Collection year
AM241-0N-B1	AM241	0	*Metabacillus*	*idriensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-B2	AM241	0	*Shinella*	*zoogloeoides*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-B3	AM241	0	*Neobacillus*	*niacini*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-B4	AM241	0	*Bacillus*	*barbaricus/enclensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-B5	AM241	0	*Sphingobium*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-B6	AM241	0	*Agrobacterium*	*radiobacter*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-B7	AM241	0	*Bacillus*	*velezensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-B9	AM241	0	*Mesobacillus*	*subterraneus*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-B10	AM241	0	*Fictibacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-B11	AM241	0	*Mesobacillus*	*subterraneus*	Italy, Lodi, Montanaso Lombardo	root	2022

AM241-0N-B12	AM241	0	*Bacillus*	*velezensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-B13	AM241	0	*Bacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-B14	AM241	0	*Lentzea*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-B1	AM241	3	*Sphingopyxis*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-B2	AM241	3	*Fictibacillus*	*enclensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-B3	AM241	3	*Mesobacillus*	*subterraneus*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-B4	AM241	3	*Rhizobium*	*tropici*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-B5	AM241	3	*Sphingopyxis*	*macrogoltabida*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-B6	AM241	3	*Fictibacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-B7	AM241	3	*Fictibacillus*	*barbaricus*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-B8	AM241	3	*Pantoea*	*agglomerans*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-B9	AM241	3	*Fictibacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-B10	AM241	3	*Fictibacillus*	*nanhaiensis*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-0N-B1	305E40	0	*Agrobacterium*	*tumefaciens*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-0N-B2	305E40	0	*Bacillus*	*velezensis*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-0N-B3	305E40	0	*Fictibacillus*	*enclensis*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-0N-B4	305E40	0	*Pseudomonas*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022

305E40-3N-B1	305E40	3	*Fictibacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-3N-B2	305E40	3	*Cytobacillus*	*firmus*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-3N-B3	305E40	3	*Fictibacillus*	*barbaricus*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-3N-B4	305E40	3	*Fictibacillus*	*barbaricus*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-3N-B5	305E40	3	*Fictibacillus*	*barbaricus*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-3N-B7	305E40	3	*Fictibacillus*	*enclensis*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-3N-B8	305E40	3	*Fictibacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-3N-B9	305E40	3	*Fictibacillus*	*barbaricus*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-3N-B10	305E40	3	*Bacillus*	*niacini*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-3N-B12	305E40	3	*Neobacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
305E40-3N-B13	305E40	3	*Cytobacillus*	*firmus*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-0N-B1	67/3	0	*Sphingobium*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-0N-B2	67/3	0	*Micrococcus*	*yunnanensis/luteus*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-0N-B3	67/3	0	*Bacillus*	*paramycoides*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-0N-B5	67/3	0	*Bosea*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-0N-B6	67/3	0	*Fictibacillus*	*barbaricus*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-0N-B7	67/3	0	*Fictibacillus*	*barbaricus*	Italy, Lodi, Montanaso Lombardo	root	2022

67/3-0N-B8	67/3	0	*Fictibacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-0N-B9	67/3	0	*Agrobacterium*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-0N-B11	67/3	0	*Fictibacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-B1	67/3	3	*Actinomycetes*	*bacterium*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-B2	67/3	3	*Fictibacillus*	*barbaricus*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-B3	67/3	3	*Sphingobium*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-B4	67/3	3	*Fictibacillus*	*barbaricus*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-B6	67/3	3	*Bacillus*	*subtilis*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-B7	67/3	3	*Bacillus*	*subtilis*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-B8	67/3	3	*Agrobacterium*	*tumefaciens*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-B9	67/3	3	*Fictibacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-B10	67/3	3	*Metabacillus*	*indicus*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-B11	67/3	3	*Peribacillus*	*muralis*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-B13	67/3	3	*Mesobacillus*	*subterraneus*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-0N-B1	AM194	0	*Paracoccus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-0N-B6	AM194	0	*Bacillus*	*altitudinis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-0N-B7	AM194	0	*Fictibacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022

AM194-0N-B8	AM194	0	*Fictibacillus*	*barbaricus*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-0N-B9	AM194	0	*Rhizobium*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-0N-B10	AM194	0	*Fictibacillus*	*enclensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-0N-B11	AM194	0	*Sphingobium*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-0N-B12	AM194	0	*Bacillus*	*velezensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B1	AM194	3	*Bacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B2	AM194	3	*Microbacterium*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B3	AM194	3	*Mesobacillus*	*subterraneus*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B4	AM194	3	*Bacillus*	*velezensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B5	AM194	3	*Rhizobium*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B6	AM194	3	*Fictibacillus*	*enclensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B7	AM194	3	*Fictibacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B8	AM194	3	*Bacillus*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B9	AM194	3	*Cytobacillus*	*firmus*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B10	AM194	3	*Lysobacter*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B11	AM194	3	*Microbacterium*	*yannicii*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B12	AM194	3	*Bacillus*	*flexus*	Italy, Lodi, Montanaso Lombardo	root	2022

AM194-3N-B13	AM194	3	*Bacillus*	*thuringiensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-B14	AM194	3	*Bacillus*	*amyloliquefaciens*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-ACT1	AM241	0	*Staphylococcus*	*cohnii*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-0N-ACT2	AM241	0	*Staphylococcus*	*warneri*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-ACT1	AM241	3	*Rhizobium*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-ACT2	AM241	3	*Agrobacterium*	*tumefaciens*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-ACT3	AM241	3	*Streptomyces*	*rishiriensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-ACT5	AM241	3	*Bacillus*	*siamensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-ACT6	AM241	3	*Streptomyces*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-ACT8	AM241	3	*Streptomyces*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-ACT9	AM241	3	*Nocardia*	*rhizosphaerihabitans*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-ACT10	AM241	3	*Streptomyces*	*gancidicus*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-ACT13	AM241	3	*Serratia*	*ficaria*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-ACT14	AM241	3	*Rhizobium*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM241-3N-ACT15	AM241	3	*Bacillus*	*velezensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-0N-ACT1	AM194	0	*Agrobacterium*	*tumefaciens*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-0N-ACT2	AM194	0	*Staphylococcus*	*warneri*	Italy, Lodi, Montanaso Lombardo	root	2022

AM194-0N-ACT3	AM194	0	*Streptomyces*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-0N-ACT4	AM194	0	*Streptomyces*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-0N-ACT6	AM194	0	*Nocardia*	*rhamnosiphila*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-ACT1	AM194	3	*Pseudomonas*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-ACT2	AM194	3	*Streptomyces*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-ACT3	AM194	3	*Nocardioides*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-ACT5	AM194	3	*Bacillus*	*thuringiensis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-ACT8	AM194	3	*Pseudomonas*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-ACT9	AM194	3	*Bacillus*	*altitudinis*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-ACT11	AM194	3	*Pseudomonas*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
AM194-3N-ACT12	AM194	3	*Streptomyces*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-ACT1	67/3	3	*Pseudomonas*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-ACT2	67/3	3	*Bacillus*	*subtilis*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-ACT3	67/3	3	*Pseudomonas*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-ACT4	67/3	3	*Streptomyces*	*rishiriensis*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-ACT6	67/3	3	*Agrobacterium*	*tumefaciens*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-ACT7	67/3	3	*Bacillus*	*subtilis*	Italy, Lodi, Montanaso Lombardo	root	2022

67/3-3N-ACT8	67/3	3	*Staphylococcus*	*warneri*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-ACT9	67/3	3	*Streptomyces*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-ACT11	67/3	3	*Streptomyces*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022
67/3-3N-ACT13	67/3	3	*Nocardia*	*sp.*	Italy, Lodi, Montanaso Lombardo	root	2022