

Exploring the rhizosphere of perennial wheat: potential for plant growth promotion and biocontrol applications

Gianluigi Giannelli¹, Lorenzo Del Vecchio¹, Martina Cirlini¹, Marco Gozzi¹, Laura Gazza², Gianni Galaverna¹, Silvia Potestio³ and Giovanna Visioli^{3*}

¹Department of Food and Drug, University of Parma, Parma, Italy; gianluigi.giannelli@unipr.it (G.Gi.); lorenzo.delvecchio@unipr.it (L.D.V.); martina.cirlini@unipr.it (M.C.); marco.gozzi@unipr.it (M.G.); gianni.galaverna@unipr.it (G.Ga.);

²CREA, Research Centre for Engineering and Agro-Food Processing, Rome, Italy; laura.gazza@crea.gov.it (L.G.)

³Department of Chemistry, Life Sciences and Environmental Sustainability, University of Parma, Parma, Italy; giovanna.visioli@unipr.it (G.V.); silvia.potestio@unipr.it (S.P.)

*Corresponding author: giovanna.visioli@unipr.it (G.V.)

Table S1. Identification and plant growth promotion traits of the forty-three bacterial strains isolated from the rhizosphere of the OK72 perennial wheat line

Strain	Homology (%)	GenBank accession no.	IAA ($\mu\text{g/mL}$)	PSU (%)	Biofilm formation (Abs units)	ACC-deaminase activity ^(a)	Protease activity (SI) ^(b)	Phosphate solubilization index (SI) ^(c)
OK_1	<i>Variovorax boronicumulans</i> (99.76)	PQ015127	0	66.98 $\pm$ 3.11	0.092 $\pm$ 0.033	+	-	-
OK_2	<i>Microbacterium yannicii</i> (100)	PQ015128	6.07 $\pm$ 2.89	58.07 $\pm$ 4.73	0.122 $\pm$ 0.043	-	-	-
OK_4	<i>Microbacterium proteolyticum</i> (100)	PQ015129	36.68 $\pm$ 7.95	44.93 $\pm$ 6.22	0.080 $\pm$ 0.011	-	2	-
OK_5	<i>Microbacterium hibisci</i> (98.69)	PQ015130	3.27 $\pm$ 1.86	56.64 $\pm$ 0.80	0.144 $\pm$ 0.068	-	-	-
OK_6	<i>Pedobacter petrophilus</i> (98.93)	PQ015131	10.96 $\pm$ 1.73	50.62 $\pm$ 9.45	0.084 $\pm$ 0.005	-	3.1 $\pm$ 0.1	-
OK_7	<i>Pedobacter borealis</i> (99.55)	PQ015132	15.38 $\pm$ 2.13	72.62 $\pm$ 12.36	0.258 $\pm$ 0.065	-	2.9 $\pm$ 0.1	-
OK_8	<i>Dyadobacter luteus</i> (97.21)	PQ015133	3.32 $\pm$ 0.34	50.94 $\pm$ 1.10	0.278 $\pm$ 0.165	-	-	-
OK_9	<i>Plantibacter auratus</i> (99.78)	PQ015134	99.39 $\pm$ 7.77	74.71 $\pm$ 12.38	0.504 $\pm$ 0.020	-	2.6 $\pm$ 0.2	-
OK_11	<i>Caulobacter soli</i> (99.78)	PQ015135	8.14 $\pm$ 3.55	38.49 $\pm$ 3.99	0.068 $\pm$ 0.001	-	-	-
OK_12	n.d.	n.d.	1.65 $\pm$ 0.24	14.66 $\pm$ 0.14	0.218 $\pm$ 0.025	-	2.9 $\pm$ 0.3	-
OK_13	n.d.	n.d.	0	18.41 $\pm$ 2.23	0.585 $\pm$ 0.086	-	2.8 $\pm$ 0.3	-
OK_14	<i>Dyadobacter ginsengisoli</i> (98.24)	PQ015136	0	38.31 $\pm$ 4.15	0.177 $\pm$ 0.049	-	-	-
OK_15	<i>Flavobacterium bizetiae</i> (99.17)	PQ015137	0	88.94 $\pm$ 2.65	0.112 $\pm$ 0.023	-	-	-
OK_16	<i>Variovoax paradoxus</i> (99.70)	PQ015138	0	90.43 $\pm$ 2.48	0.373 $\pm$ 0.015	-	-	-
OK_17	<i>Pseudomonas azotoformans</i> (99.89)	PQ015139	0.92 $\pm$ 0.35	87.54 $\pm$ 4.25	0.838 $\pm$ 0.078	-	3.5 $\pm$ 0.9	2.7 $\pm$ 0.1
OK_18	<i>Microbacterium oleivorans</i> (100)	PQ015140	61.026 $\pm$ 10.49	58.64 $\pm$ 5.85	0.107 $\pm$ 0.029	-	2	-
OK_20	<i>Paenibacillus silagei</i> (99.67)	PQ015141	6.46 $\pm$ 2.57	48.55 $\pm$ 0.64	0.338 $\pm$ 0.038	-	-	-
OK_21	<i>Epilithonimonas lactis</i> (97.43)	PQ015142	7.45 $\pm$ 2.6	14.90 $\pm$ 1.72	0.064 $\pm$ 0.004	-	2.5 $\pm$ 0.4	-
OK_22	<i>Frigoribacterium endophyticum</i> (99.30)	PQ015143	48.22 $\pm$ 10.59	50.05 $\pm$ 5.30	0.255 $\pm$ 0.071	-	-	-
OK_23	<i>Pseudomonas petroselini</i> (100)	PQ015144	0	86.94 $\pm$ 5.72	0.083 $\pm$ 0.002	-	-	3.8 $\pm$ 0.1
OK_26	<i>Aeromicrobium ginsengisoli</i> (97.99)	PQ015145	0	72.21 $\pm$ 10.52	0.831 $\pm$ 0.064	-	5.6 $\pm$ 0.6	-
OK_27	<i>Microbacterium aurantiacum</i> (99.86)	PQ015146	19.05 $\pm$ 5.70	38.18 $\pm$ 9.55	0.098 $\pm$ 0.008	-	2.3 $\pm$ 0.8	-
OK_28	<i>Curtobacterium herbarum</i> (99.75)	PQ015147	0	42.89 $\pm$ 8.56	0.085 $\pm$ 0.021	-	3.1 $\pm$ 0.5	-
OK_30	<i>Pseudomonas brassicacearum</i> (98.84)	PQ015148	2.78 $\pm$ 1.49	95.80 $\pm$ 6.09	0.086 $\pm$ 0.016	-	-	2.8 $\pm$ 0.1
OK_31	<i>Pseudomonas nabeulensis</i> (100)	PQ015149	4.03 $\pm$ 1.62	92.93 $\pm$ 9.77	0.097 $\pm$ 0.006	-	2.6 $\pm$ 0.3	2.4 $\pm$ 0.1
OK_32	<i>Flavobacterium olei</i> (99.53)	PQ015150	0	47.05 $\pm$ 5.10	0.067 $\pm$ 0.016	-	2.7 $\pm$ 0.1	-
OK_33	<i>Rhodococcus qingshengii</i> (100)	PQ015151	44.18 $\pm$ 6.88	42.91 $\pm$ 12.99	0.086 $\pm$ 0.013	+	-	-
OK_35	<i>Okibacterium fritillariae</i> (99.82)	PQ015152	57.52 $\pm$ 6.12	65.71 $\pm$ 7.06	0.045 $\pm$ 0.003	-	-	-
OK_39	<i>Pseudomonas lactis</i> (100)	PQ015153	7.9 $\pm$ 0.74	85.65 $\pm$ 4.27	0.554 $\pm$ 0.033	-	+	2.5 $\pm$ 0.1

OK_41	<i>Acinetobacter guillouiae</i> (100)	PQ015154	0	7.76 ± 4.98	0.800 ± 0.012	-	-	-
OK_42	<i>Pseudomonas canavaninivorans</i> (100)	PQ015155	12.51 ± 3.68	89.91 ± 0.30	0.096 ± 0.014	+	-	4.1 ± 0.1
OK_44	<i>Agrobacterium arsenijevicei</i> (99.88)	PQ015156	4.45 ± 0.66	14.33 ± 1.47	0.093 ± 0.032	-	+	-
OK_45	<i>Pseudomonas lactis</i> (99.58)	PQ015157	17.02 ± 2.46	73.02 ± 0.25	n.d.	-	2.3 ± 0.4	-
OK_46	n.d.	n.d.	33.92 ± 3.18	48.61 ± 0.55	0.061 ± 0.010	-	+	-
OK_47	<i>Caulobacter rhizosphaerae</i> (99.88)	PQ015158	0	23.44 ± 0.98	0.050 ± 0.010	-	-	-
OK_50	n.d.	n.d.	3.82 ± 0.46	5.14	n.d.	-	-	-
OK_52	<i>Pseudoxanthomonas wuyuanensis</i> (99.10)	PQ015159	1.95 ± 0.04	21.41 ± 1.29	n.d.	-	3.6 ± 0.5	-
OK_54	<i>Pseudomonas brassicacearum</i> (98.17)	PQ015160	88.58 ± 1.05	88.43 ± 1.44	0.078 ± 0.004	-	-	3.3 ± 0.1
OK_58	<i>Polaromonas ginsengisoli</i> (99.90)	PQ015161	0	n.d.	0.143 ± 0.023	-	-	-
OK_59	<i>Cellulosimicrobium funkei</i> (99.44)	PQ015162	n.d.	n.d.	0.056 ± 0.023	+	n.d.	n.d.
OK_64	<i>Pseudomonas brassicacearum</i> (99.66)	PQ015163	9.5 ± 1.1	79.68 ± 1.05	0.058 ± 0.002	-	+	3.3 ± 0.3
OK_71	n.d.	n.d.	15.94 ± 0.54	2.59 ± 1.71	n.d.	-	-	-
OK_90	<i>Plantibacter flavus</i> (99.55)	PQ015164	44.43 ± 3.96	40.46 ± 6.17	0.055 ± 0.009	-	-	-

^(a) ACC-deaminase activity: (-) no bacterial growth on medium containing 1-aminocyclopropane-1-carboxylate as the only N source; (+) bacterial growth on medium containing 1-aminocyclopropane-1-carboxylate as the only source of N.

^(b) (-) no protease activity detected while (+) indicates positive protease activity but the morphology of the colony did not allow the measurement to be made.

^(c) (-) no phosphate solubilization detected.

n.d. Not determined.

Data are average of three independent experiments ± standard deviation (S.D.).

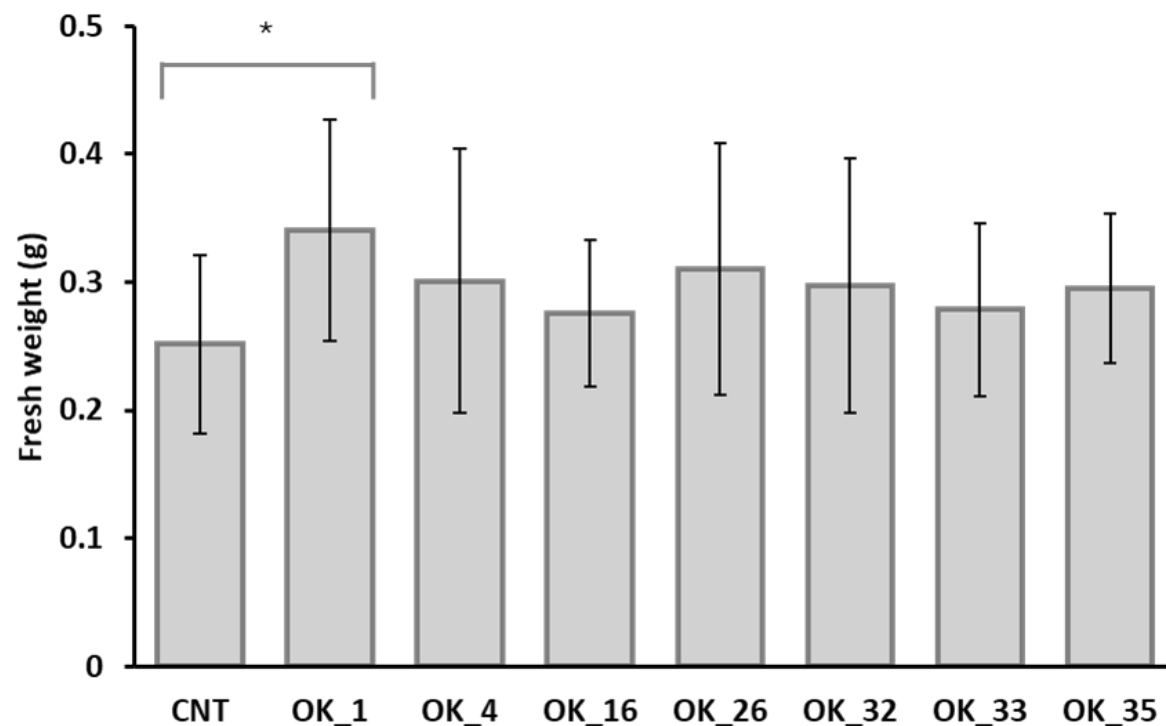


Figure S1. Growth-promoting activity exerted by bacteria on the fresh weight of the commercial wheat line. (A) Effect of the inoculation of the commercial wheat (cv Bologna) seeds with bacterial strains on fresh weight after ten days of growth. Data are presented as mean of thirty plantlets $\pm$ S.D. All the data showed significant differences compared to control conditions according to One-way ANOVA and Tukey's test ($p < 0.05$). Asterisk indicates significant differences between plants inoculated with respect to control condition (uninoculated) (One-way ANOVA and Tukey's test, $*p < 0.05$)

