

Planococcus maritimus ML1206 Isolated from Wild Oysters Enhances the Survival of Caenorhabditis elegans against Vibrio anguillarum

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Method

1. C. elegans Length Determination

Ten synchronized *C. elegans* N2 in L1 stage were transferred to NGM medium coated with OP50 and ML1206, respectively. *C. elegans* were cultured at 20 °C to the 2-days adult stage. *C. elegans* were picked to slides covered with agarose, anesthetized with levamisole hydrochloride, and placed under a stereomicroscope to measure body length with a micrometer. The head (oral position) of the *C. elegans* was taken as the starting point, and following the middle part of the body to the tail part of the *C. elegans* was taken as the measurement endpoint. NIS elements software was used to calculate the body of the *C. elegans* body ^[1]. The test was performed three times.

2. C. elegans Motility Determination

Referring to the methods of Tsalik and Hobert ^[2], the effect of ML1206 on *C. elegans* motility was tested. Ten synchronized *C. elegans* N2 in L1 stage were transferred to NGM medium coated with OP50 and ML1206, respectively. *C. elegans* were cultured at 20 °C to the 2-days adult stage. Then the *C. elegans* were transferred to new NGM plates. After the free movement of *C. elegans* in the first minute, the bending times of the *C. elegans* within 30 s were observed and recorded under a stereoscopic microscope as the indicator of bending frequency. One bending of the body is defined as a change in the movement of the body along the corresponding X-axis during *C. elegans* crawling, assuming that the direction along the pharyngeal pump is the Y-axis ^[3]. The test was performed three times.

3. C. elegans Pharyngeal Pumping Experiment

Ten synchronized *C. elegans* N2 in L1 stage were transferred to NGM medium coated with OP50 and ML1206, respectively. *C. elegans* were cultured at 20 °C to the 2-days adult stage. Then the *C. elegans* were transferred to new NGM plates, the pump activity of *C. elegans* pharyngeal terminal bulb was observed through stereomicroscope (×400), and the number of pharyngeal pump activities within 1 min was recorded, the swallowing rate of *C. elegans* ^[1]. The test was performed three times.

4. C. elegans Reproductive Capacity Determination

Ten synchronized L1 N2 were selected to NGM medium coated with OP50 and ML1206, respectively. One plate was for one worm and kept *C. elegans* lay eggs at 20 °C. The parent *C. elegans* were transferred to a new NGM medium coated with OP50 and ML1206 every 12 h, and the number of eggs in each old culture dish was recorded until the *C. elegans* stopped laying eggs. The sum of eggs in plates of each group was the total number of eggs laid by 10 *C. elegans*, and the averages of offspring produced by each *C. elegans* were calculated to characterize the reproductive capacity of the parent *C. elegans* ^[1]. The test was performed three times.

Results

Table S1. Stains from the intestine of oyster and perch

number	Stain number	Putative identity
1	M0101	<i>Pseudoalteromonas haloplanktis</i> 98.2%

2	M0102	<i>Marinobacter hydrocarbonoclasticus</i> 98.8%
3	M0103	<i>Psychrobacter celer</i> 98.8%
4	M0104	<i>Pseudoalteromonas nigrifaciens</i> 98.1%
5	M0106	<i>Lacinutrix undariae</i> 98.0%
6	M0107	<i>Pseudoalteromonas translucida</i> 98.2%
7	M1201	<i>Bacillus siamensis</i> 98.0%
8	ML1202	<i>Bacillus tequilensis</i> 98.9%
9	ML1206	<i>Planococcus maritimus</i> 99.2%
10	ML1209	<i>Pseudoalteromonas fuliginea</i> 99.2%
11	ML1210	<i>Sulfitobacter pontiacus</i> 98.3%
12	ML1211	<i>Olleya algicola</i> 99.9%
13	ML1212	<i>Bacillus flexus</i> 98.8%
14	ML1216	<i>Bacillus firmus</i> 97.5%
15	ML1224	<i>Pseudoalteromonas neustonica</i> 97.3%
16	ML1226	<i>Pseudovibrio japonicus</i> 99.0%
17	ML1227	<i>Bacillus altitudinis</i> 98.4%
18	ML1229	<i>Ruegeria atlantica</i> 99.3%
19	YLY02	<i>Vibrio alfacensis</i> 99.5%
20	YLY03	<i>Sunxiuqinia elliptica</i> 99.1%
21	YLY04	<i>Maribius pontilimi</i> 95.33%
22	YLY05	<i>Idiomarina sediminum</i> 98.9%
23	YLY06	<i>Vibrio orientalis</i> 100%
24	YLY07	<i>Brumimicrobium mesophilum</i> 95.2%
25	YLY08	<i>Oceaniglobus indicus</i> 93.4%
26	YLY09	<i>Nitratireductor aquimarinus</i> 98.1%
27	YLY10	<i>Microbacterium esteraromaticum</i> 98.1%
28	YLY13	<i>Bacillus halosaccharovorans</i> 98.8%
29	YLY14	<i>Roseovarius pacificus</i> 99.6%
30	YLY17	<i>Shewanella indica</i> 98.6%
31	YLY18	<i>Vibrio atypicus</i> 99.9%
32	YLY20	<i>Ornithinimicrobium kibberense</i> 95.7%
33	YLY21	<i>Litoreibacter arenae</i> 99.8%
34	YLY22	<i>Halobacillus alkaliphilus</i> 97.8%
35	YLY23	<i>Jeotgalibacillus campisalis</i> 99.0%
36	YLY25	<i>Marinobacter pelagius</i> 98.9%
37	YLY26	<i>Stappia stellulata</i> 99.1%
38	YLY27	<i>Idiomarina aestuarii</i> 99.4%
39	YLY28	<i>Roseovarius halotolerans</i> 98.9%
40	YLY29	<i>Shimia biformata</i> 97.8%
41	YLY32	<i>Halomonas aestuarii</i> 96.8%

Table S2. Hemolysis of strains

Strain number	Hemolysis	Strain number	Hemolysis
M0101	+	YLY06	-
M0102	-	YLY07	-
M0103	+	YLY08	-
M0104	+	YLY09	-
M0106	+	YLY10	-
M0107	+	YLY13	+
ML1201	+	YLY14	+
ML1202	+	YLY17	+
ML1206	-	YLY18	+
ML1209	+	YLY20	-
ML1210	-	YLY21	-
ML1211	+	YLY22	+
ML1212	+	YLY23	+
ML1216	+	YLY25	-
ML1224	+	YLY26	-
ML1226	+	YLY27	-
ML1227	+	YLY28	-
ML1229	-	YLY29	+
YLY02	-	YLY32	-
YLY03	-	SDUM002245	+
YLY04	-	LGG	-
YLY05	-		

Note: "+" means positive; "-" stands means negative

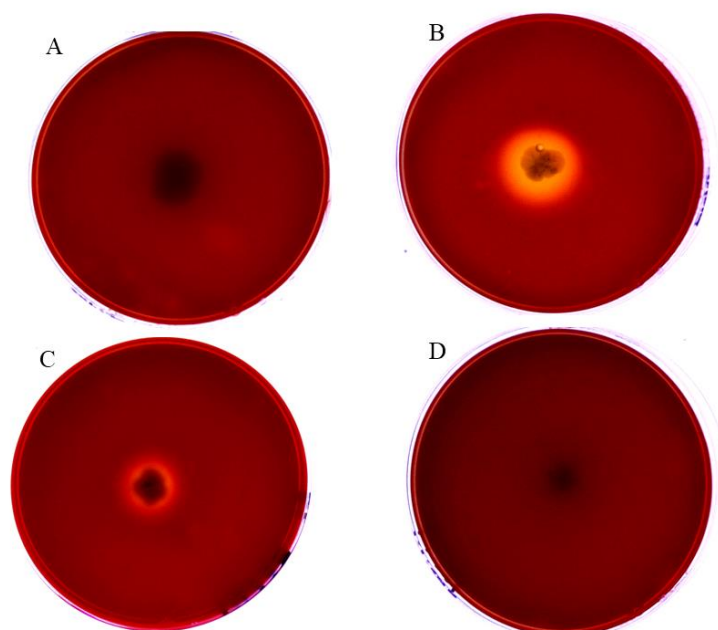


Figure S1. Hemolysis of some strains on 5% sheep blood agar (partial)

Note: A, strain ML1206; B, strain YLY14; C, control (*V. anguillarum*); D, negative control (strain LGG)

Table S3 Low pH tolerance ability of twenty intestinal bacteria (Mean±SD)

Strain number	OD ₆₀₀ value		Acid-resistant survival rate (%)
	pH 7.4 (pH5.6 for LGG)	pH3.0	

M0102	0.49±0.03 ^{cd,A}	0.05±0.05 ^{gh,B}	10.30±0.02 ^{gf}
ML1206	0.47±0.03 ^{cde,A}	0.37±0.02 ^{c,A}	79.51±0.03 ^b
ML1210	0.41±0.02 ^{defg,A}	0.07±0.05 ^{gh,A}	16.33±0.02 ^{gf}
ML1229	0.68±0.02 ^{b,A}	0.08±0.01 ^{gh,A}	11.80±0.02 ^{gf}
YLY02	0.44±0.04 ^{deg,A}	0.06±0.00 ^{gh,A}	13.73±0.01 ^{gf}
YLY03	0.81±0.02 ^{a,A}	0.09±0.02 ^{g,A}	11.12±0.02 ^{gf}
YLY04	0.38±0.03 ^{fg,A}	0.07±0.01 ^{gh,A}	18.45±0.04 ^{gf}
YLY05	0.45±0.02 ^{de,A}	0.28±0.03 ^{e,A}	61.14±0.05 ^e
YLY06	0.68±0.01 ^{b,A}	0.13±0.01 ^{f,A}	18.69±0.02 ^{gf}
YLY07	0.39±0.06 ^{efg,A}	0.06±0.01 ^{gh,B}	16.67±0.03 ^{gf}
YLY08	0.20±0.01 ^{j,A}	0.05±0.01 ^{i,A}	23.89±0.04 ^f
YLY09	0.38±0.03 ^{gh,A}	0.21±0.03 ^{e,A}	54.62±0.04 ^d
YLY10	0.32±0.02 ^{hi}	0.17±0.01 ^e	53.77±0.05 ^d
YLY20	0.43±0.02 ^{defg}	0.07±0.01 ^{gh}	16.91±0.01 ^{gf}
YLY21	0.45±0.01 ^{def}	0.05±0.01 ^{gh}	11.77±0.01 ^{gf}
YLY25	0.31±0.03 ^{i,A}	0.26±0.02 ^{d,A}	85.24±0.04 ^b
YLY26	0.29±0.01 ^{i,A}	0.07±0.01 ^{gh,A}	25.02±0.02 ^f
YLY27	0.30±0.02 ^{i,A}	0.07±0.00 ^{gh,B}	23.74±0.02 ^f
YLY28	0.44±0.02 ^{de,A}	0.06±0.00 ^{gh,B}	13.57±0.01 ^{gf}
YLY32	0.76±0.06 ^{a,A}	0.49±0.04 ^{b,A}	65.56±0.10 ^c
LGG	0.52±0.03 ^{c,A}	0.62±0.02 ^{a,A}	118.71±0.02 ^a

Note: different lowercase letters on superscript represent significant differences between different strains ($p < 0.05$); Different capital letters represent significant differences under different pH ($p < 0.05$).

Table S4. Bile salt tolerance ability of twenty intestinal bacteria (Mean±SD)

Strain number	D ₆₀₀ value		Bile salt tolerance survival rate (%)
	Bile salt 0.0%	Bile salt 0.3%	
M0102	0.41±0.08 ^{ghij,A}	0.07±0.01 ^{ghi,B}	17.75±0.03 ^{efg}
ML1206	0.70±0.02 ^{c,A}	0.57±0.03 ^{a,A}	81.84±0.04 ^{ab}
ML1210	0.46±0.04 ^{fgh,A}	0.07±0.01 ^{ghi,B}	15.96±0.01 ^{efgh}
ML1229	0.46±0.02 ^{fgh,A}	0.06±0.01 ^{hi,B}	12.46±0.01 ^{fgh}
YLY02	0.34±0.02 ^{ijkl,A}	0.20±0.02 ^{e,A}	57.06±0.06 ^c
YLY03	0.42±0.02 ^{ghi,A}	0.10±0.01 ^{g,A}	24.91±0.03 ^e
YLY04	0.57±0.02 ^{de,B}	0.05±0.01 ^{hi,B}	9.29±0.01 ^h
YLY05	0.51±0.01 ^{ef,A}	0.38±0.03 ^{b,A}	74.45±0.04 ^b
YLY06	0.70±0.02 ^{c,A}	0.56±0.03 ^{a,A}	79.60±0.03 ^{ab}
YLY07	0.32±0.02 ^{l,B}	0.04±0.01 ^{hi,B}	13.46±0.02 ^{fgh}
YLY08	0.32±0.02 ^{l,B}	0.04±0.00 ^{hi,B}	12.56±0.01 ^{fgh}
YLY09	0.43±0.02 ^{gh,A}	0.27±0.03 ^{c,A}	63.90±0.07 ^c
YLY10	0.33±0.04 ^{kl,A}	0.04±0.01 ^{i,B}	11.32±0.03 ^{fgh}
YLY20	0.39±0.03 ^{hijkl,A}	0.05±0.01 ^{hi,B}	13.01±0.03 ^{fgh}
YLY21	0.40±0.02 ^{ghijk,A}	0.08±0.01 ^{gh,B}	20.16±0.03 ^{ef}
YLY25	0.45±0.05 ^{fgh,A}	0.15±0.02 ^{f,A}	35.12±0.07 ^d
YLY26	0.80±0.01 ^{b,A}	0.06±0.00 ^{hi,B}	7.50±0.01 ^h
YLY27	0.60±0.01 ^{d,A}	0.04±0.00 ^{hi,B}	6.67±0.00 ^h
YLY28	0.47±0.03 ^{fg,A}	0.05±0.01 ^{hi,B}	11.45±0.01 ^{fgh}
YLY32	0.97±0.03 ^{a,A}	0.23±0.04 ^{d,A}	24.10±0.04 ^e
LGG	0.35±0.02 ^{ijkl,A}	0.30±0.02 ^{c,A}	87.42±0.05 ^a

Note: different lowercase letters on superscript represent significant differences between different strains ($p < 0.05$);

Different capital letters represent significant differences under different bile salt concentration values ($p < 0.05$).

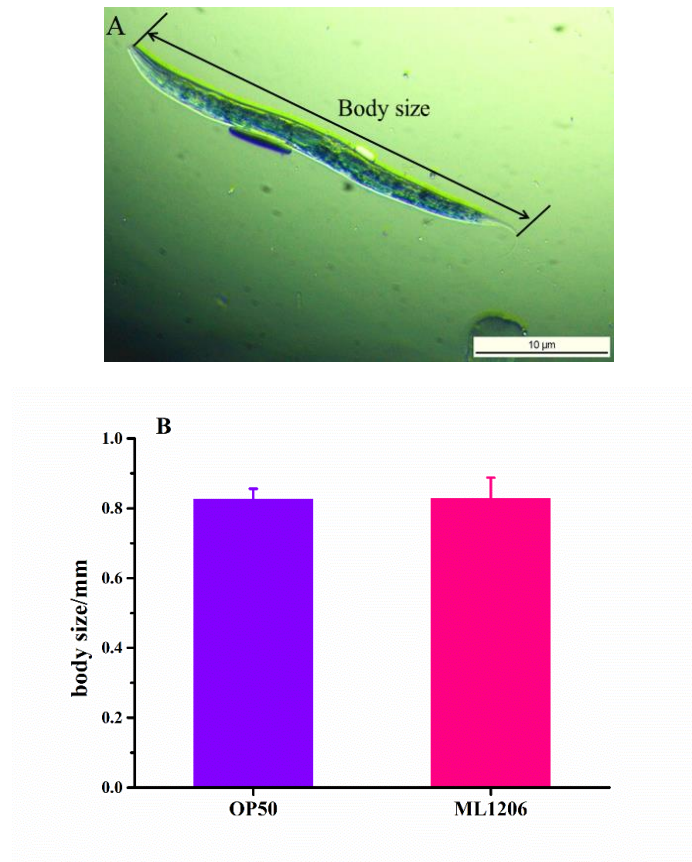


Figure S2. Effect of ML1206 strain on *C. elegans* growth: (a) Schematic diagram of *C. elegans* body length calculated with NIS elements software (b) Worms were grown in the presence of OP50 or ML1206 to the 2-day adult stage *C. elegans*. After anesthesia with levamisole hydrochloride, their body length was measured under a stereoscopic microscope ($\times 400$) with a micrometer. Data are representative of three independent experiments and presented as means \pm SD.

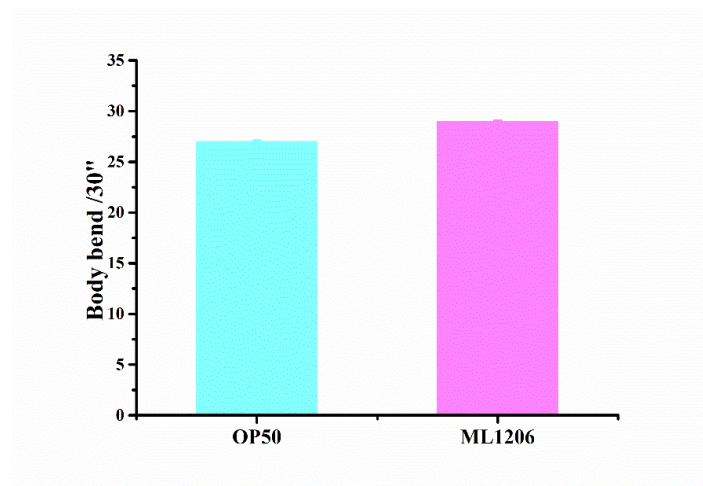


Figure S3. Effect of ML1206 strain on *C. elegans* body bending frequency: Worms were grown in the presence of OP50 or ML1206 to the 2-day adult stage *C. elegans*, and the body bending times of *C. elegans* within 30 S were recorded under a stereoscopic microscope ($\times 400$). Data are representative of three independent experiments and presented as means \pm SD.

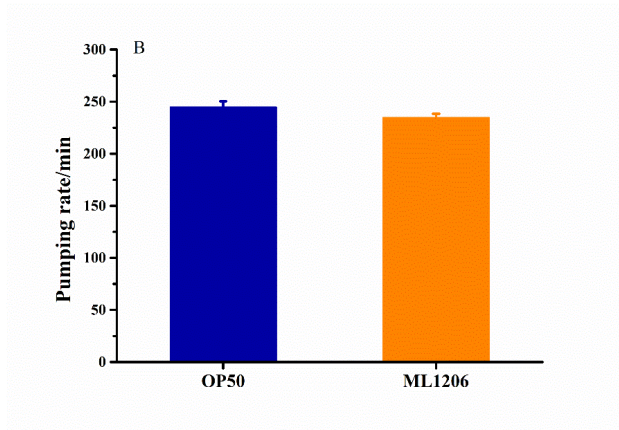


Figure S4. Effect of ML1206 strain on *C. elegans* pharyngeal pumping rate: Worms were grown in the presence of OP50 or ML1206 to the 2-day adult stage *C. elegans*. The worm was recorded within 1 min. Data are representative of three independent experiments and presented as means \pm SD.

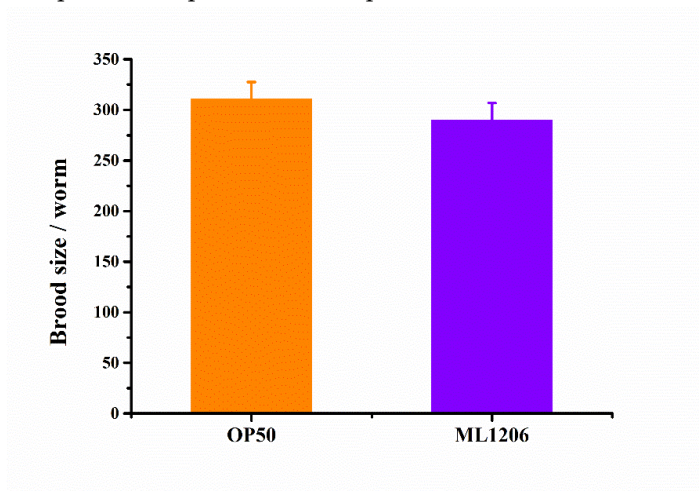


Figure S5. Effect of ML1206 strain on *C. elegans* reproduction ability: Average egg production per worm fed with OP50 or ML1206. Data are representative of three independent experiments and presented as means \pm SD.

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