

Supplementary Material

Isolation, Genomic Analysis, and Functional Characterization of Enterococcus rotai CMTB-CA6, a Potential Probiotic Strain from a Medicinal Plant Centella asiatica

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1. Supplementary Figure

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A API 50CHE

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	30	31	32	33	34	35	*	37	38	39
			F	F	Ĵ	Ĵ	Ĵ	Ĵ	Ĵ	F
40 41 42 43 44 45 46 47 49 49	40	41.	42	43	44	45	46	47	48	49

B

API 20E	CMTB-CA6		
characteristic			
β-galactosidase	5		
Arginine dihydrolase	+		
Lysine decarboxylase	н		
Ornithine decarboxylase	2		
Citrate utilization	-		
H ₂ S production	.		
Urease	×		
Tryptophane desaminase	2		
Indole production	5		
Acetoin production	+		
Gelatinase	2		
Glucose	+		
Mannitol	+		
Inositol	+		
Sorbitol	+		
Rhamnose	+		
Sucrose	+		
Melibiose	+		
Arabinose	+		
Cytochrome oxidase	<u>12</u> 2		

API 50CHE characteristic	CMTB-CA6
Giverni	+
Frythritol	
Arabinose	
Arabinose	-
2-Arabinose Riboos	-
Note:	1
J-Aylose	-
L-Aylose	
Adonitol	-
3-Methyl-Xyloside	-
Jalactose	Ť
J-Glucose	d.
D-Fructose	+
J-Mannose	+
,-Sorbose	2
Chamnose	5
Juicitol	-
nositol	-
Mannitol	1
Sorbitol	-
-Methyl-D-Mannose	52
-Methyl-D-Glucoside	-
I-Acetyl glucosamine	+
mygdaline	+
arbutine	+
sculine	÷
alicine	+
Cellobiose	+
faltose	+
actose	-
felibiose	Ξ.
accharose	+
rehalose	+
nuline	
felezitose	+
0-Raffinose	-
midon	-
Hycogen	2
[ylito]	8
-Gentiobiose	+
-Turanose	-
D-Lvxose	-
-Tagatose	2
)-Fucose	-
-Fucose	-
D-Arabitol	-
-Arabitol	
Juconate	
ceto-gluconate	-
ceto gluconate	8
ceto-gluconate	-

Supplementary Figure S1. Biochemical tests for rapid identification of the CMTB-CA6 strain. Using the analytical profile index (API) 50CHE (**A**, **B**) and 20E (**C**) systems, the enzymatic activities of the CMTB-CA6 strain were assessed. The reactions occurred under anaerobic conditions, and the results were interpreted after 48 h of incubation. A positive reaction is indicated by "+," whereas a negative reaction is indicated by "-."

2. Supplementary Table

Supplementary Table S1. The nucleotide identity scores for comparison of the 16S ribosomal RNA (rRNA) gene sequences between CMTB-CA6 and *Enterococcus* spp. strains.

Rank	Matched bacterial strains	Accession Id	% identity	Total Score	E value
1	Enterococcus rotai strain LMG 26678	CP013655.1	99.8	2745	0
2	Enterococcus rotai strain CCM 4630	NR_108137.1	99.8	2745	0
3	Enterococcus sp. P7644	AJ276462.1	99.8	2745	0
4	Enterococcus silesiacus strain LMG 23085	CP013614.1	99.73	2739	0
5	Enterococcus silesiacus strain R-23712	NR_042405.1	99.73	2739	0
6	Enterococcus silesiacus LMG 23084	AM039967.1	99.67	2736	0
7	Enterococcus crotali ETRF1	NR_156980.1	99.93	2730	0
8	Enterococcus ureilyticus CCM 4629	NR_125485.1	99.6	2728	0
9	Enterococcus caccae 2215-02	NR_043285.1	99.87	2724	0
10	Enterococcus sp. HAMBI3059	FN822762.1	99.86	2721	0