

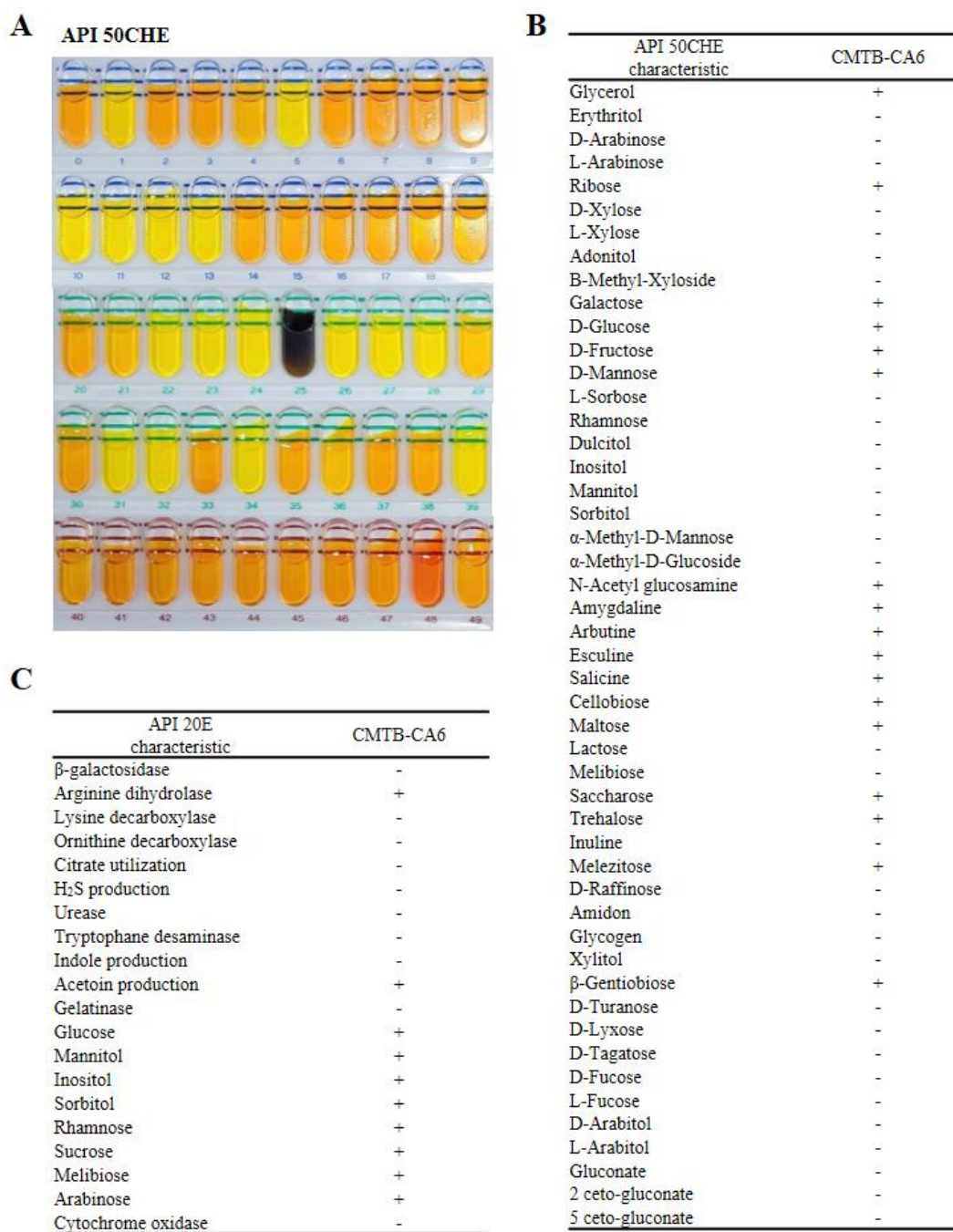
Supplementary Material

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

Yunsik Kim, Jin Hee Lee, Jimyeong Ha, Eun-Gyung Cho*

*** Correspondence:** Eun-Gyung Cho: egcho@chamc.co.kr

1. Supplementary Figure

**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	<i>Enterococcus rotai</i> strain LMG 26678	CP013655.1	99.8	2745	0
2	<i>Enterococcus rotai</i> strain CCM 4630	NR_108137.1	99.8	2745	0
3	<i>Enterococcus</i> sp. P7644	AJ276462.1	99.8	2745	0
4	<i>Enterococcus silesiacus</i> strain LMG 23085	CP013614.1	99.73	2739	0
5	<i>Enterococcus silesiacus</i> strain R-23712	NR_042405.1	99.73	2739	0
6	<i>Enterococcus silesiacus</i> LMG 23084	AM039967.1	99.67	2736	0
7	<i>Enterococcus crotali</i> ETRF1	NR_156980.1	99.93	2730	0
8	<i>Enterococcus ureilyticus</i> CCM 4629	NR_125485.1	99.6	2728	0
9	<i>Enterococcus caccae</i> 2215-02	NR_043285.1	99.87	2724	0
10	<i>Enterococcus</i> sp. HAMBI3059	FN822762.1	99.86	2721	0