

Probiotic potential of a high GABA producing strain, Lactobacillus brevis FEM 1874, isolated from traditional "wild" Alpine cheese.



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Introduction

"Wild" Alpine cheese isolate *Lactobacillus brevis* FEM 1874 was characterized for potential probiotics traits and GABA production. Acid, bile and pancreatic fluid resistance was assessed and the genetic loci for GABA production and export characterized for the subsequent knock-out vectors construction.

Probiotics and their metabolites can act as brain modulators through the gut:brain axis system. In this respect γ -aminobutyric acid (GABA) gut bacteria produced, has been identified as a possible factor improving brain function, inhibiting inflammation and regulating energy metabolism.

1) FEM 1874 survives in gastrointestinal phisicochemical condition

Results

2) FEM 1874 produced high GABA level

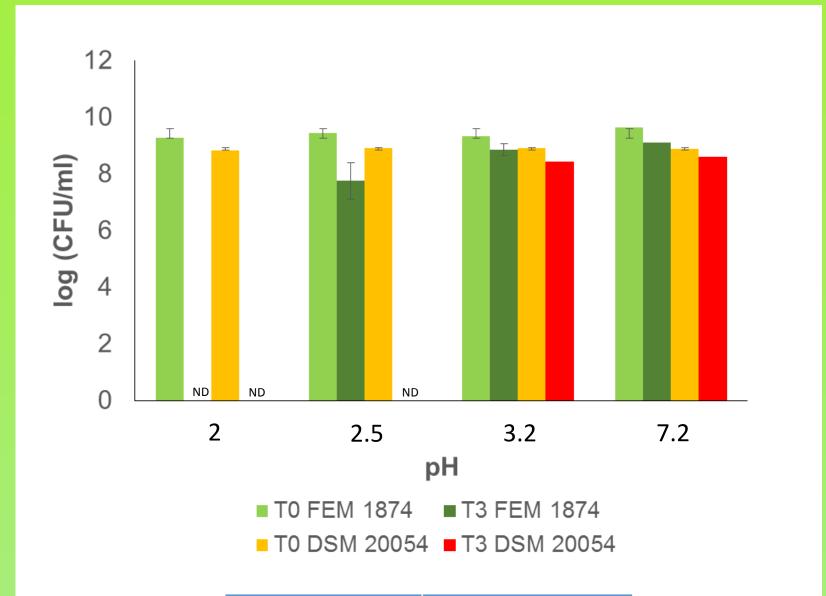
Acid tolerance (pH 2, 2.5, 3.2). Acid treatment performed in PBS at 0 and 3h (T0 and T3 resp). Cell counts compared to type strain grown in the same condition (mean \pm sd, N = 3).

Bile Salt Hydrolysis (BSH) assay. BSH activity perfomed on MRS agar plates supplemented with 0.5% of bile acids. (mean ± sd, N = 3).

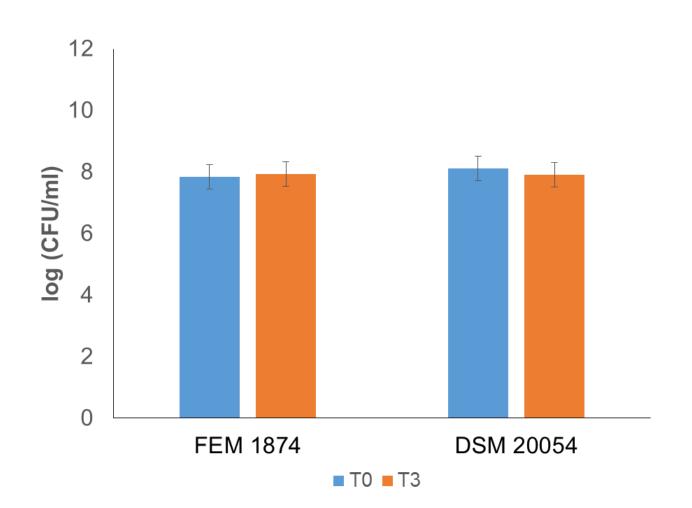
Pancreatic fluid tolerance. Tolerance test perfomed in $NaHCO_3$ with 1.9 mg/ml pancreatin for 3h. Cell counts compared to type strain grown in the same condition (mean \pm sd, N = 3).

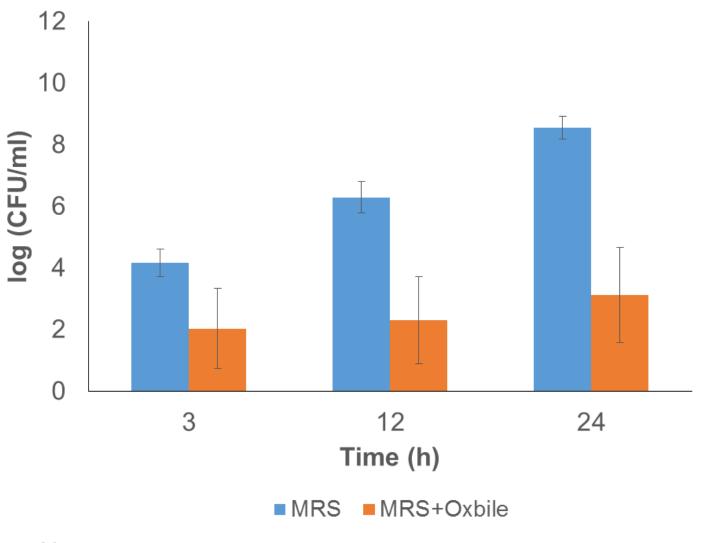
Effect of bile salts after low pH treatment. Bile acid (Oxbile) treatment performed in MRS over time after pH 2.5 exposure (3h) (mean \pm sd, N = 3).

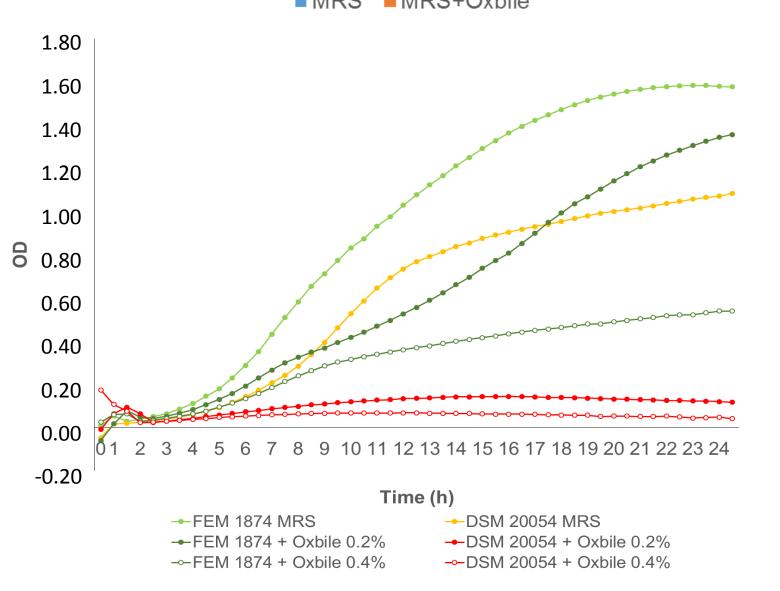
Growth rate with bile salts. Bile acid (Oxbile) treatment performed in MRS over time (mean ± sd, N = 3).

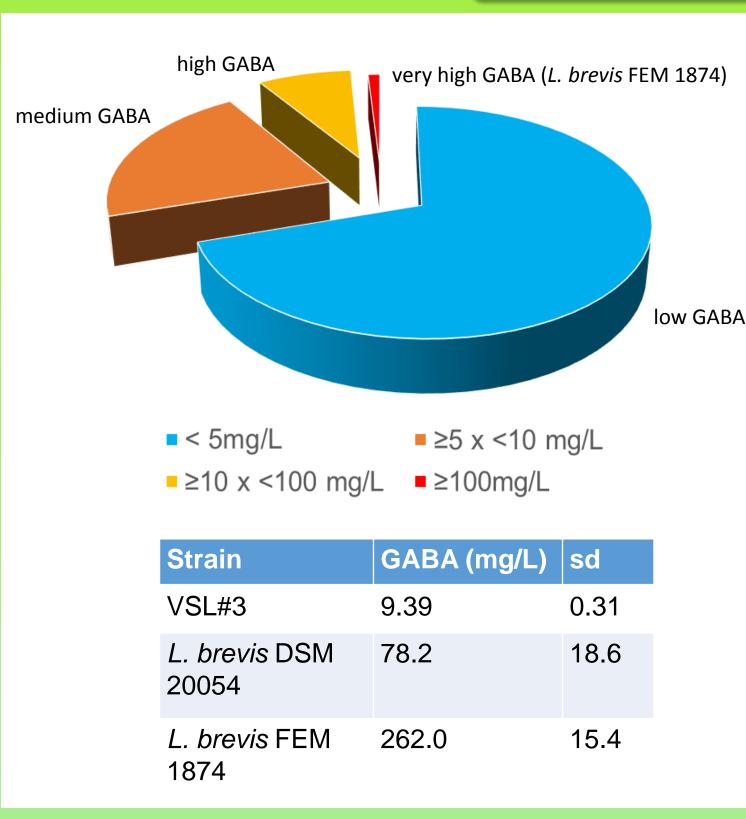


Strain	BSH activity
<i>L. brevis</i> DSM 20054	-
L. brevis FEM 1874	+





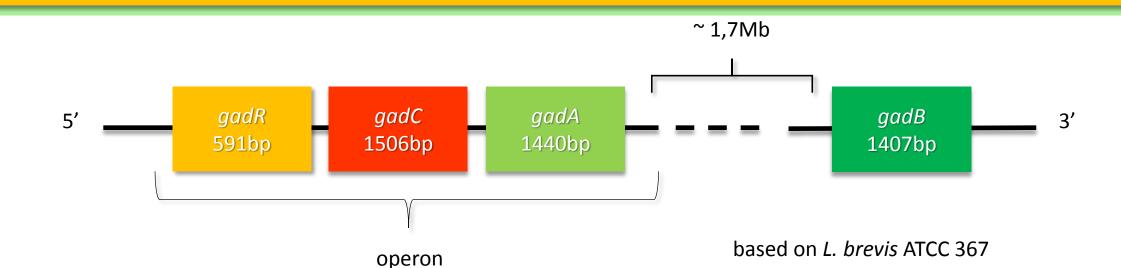




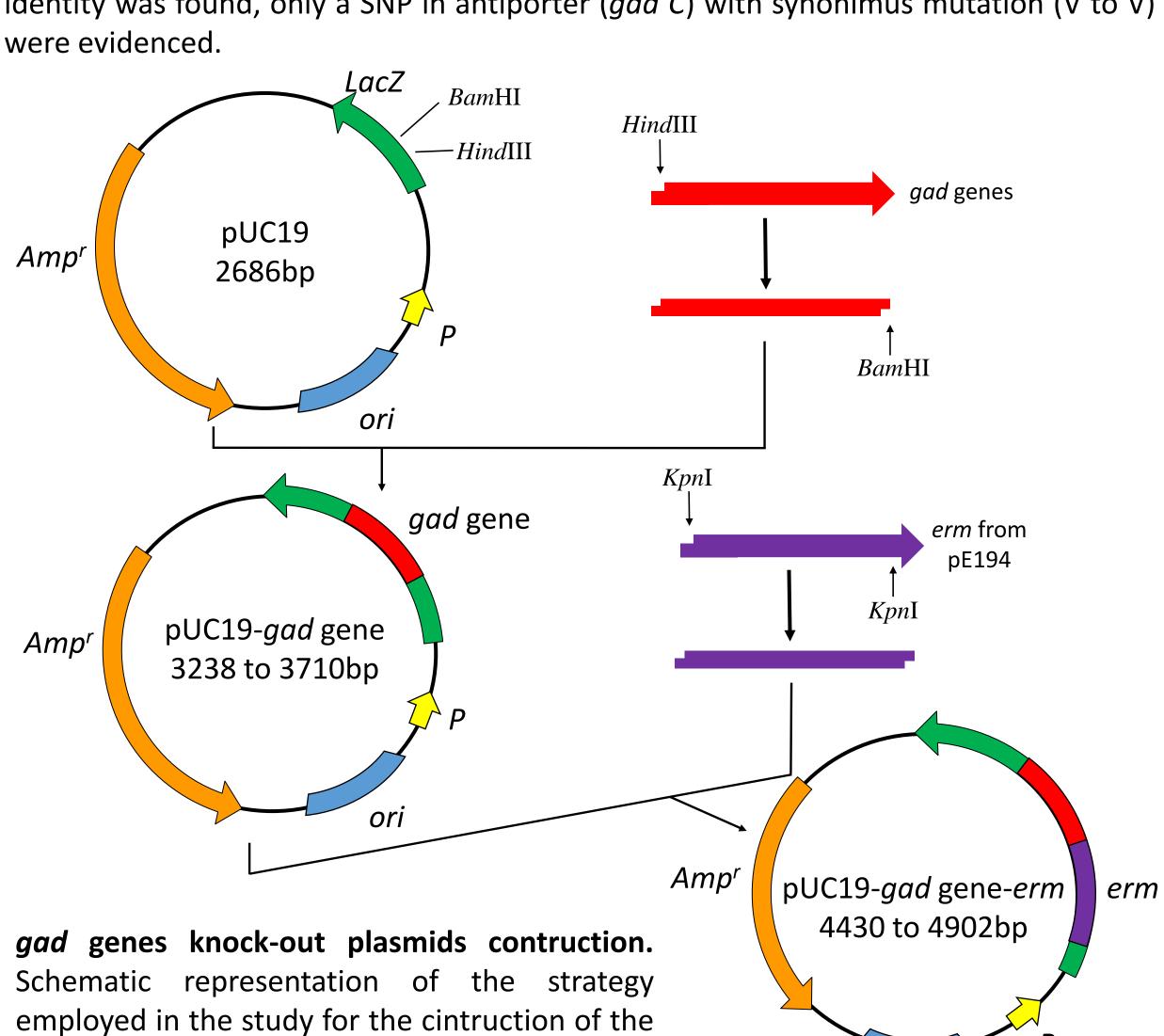
«Wild» Alpine cheese isolates GABA production.
GABA production (mg/L) after 24h in experimental buffer + glutamate [1mg/ml]. Total isolate considered 110. Mean ± sd, N = 3.

GABA production. GABA production after 24h in experimental buffer + glutamate [1mg/ml]. Mean ± sd, N = 3

3) FEM 1874 GABA genes sequences analysis and knock-out generation strategy



FEM 1874 GABA genes sequenced: 100% identity. Based on *L. brevis* ATCC 367 gene sequencing, nine sets of primer were used to sequence all GABA genes. 100% of identity was found, only a SNP in antiporter (*gad C*) with synonimus mutation (V to V) were evidenced.



pUC19-gad gene-erm vectors.