

SUPPLEMENTARY DATA - Bosma *et al.*

Table S1. Growth, end product formation and genetic accessibility of selected isolates.

Strain	Identification ¹	Isol. ²	Sugars supporting growth ³				Products (g/L) ⁴										CFU ⁵
			G	S	X	A	Lac	Ace	Suc	Mal	Eth	For	2,3-BDO	Prop	Total	OD ₆₀₀	
ET 138	<i>B. smithii</i>	3	+	+	+	+/-	6.74 ±0.18	0.81 ±0.02	0.07 ±0.00	nd	nd	nd	nd	nd	7.61 ±0.16	1.425 ±0.058	3
DSM 1	<i>B. coagulans</i>	T	+ ⁷	- ⁷	- ⁷	- ⁷	6.37 ±0.76	0.36 ±0.05	0.13 ±0.05	0.18 ±0.30	0.04 ±0.07	nd	0.10 ±0.02	nd	7.17 ±1.16	1.467 ±0.147	20 ⁶
ET 236	<i>G. thermodenitrificans</i>	5	+/-	+	+	+	5.37 ±0.60	1.15 ±0.07	0.13 ±0.01	0.17 ±0.11	0.04 ±0.02	0.07 ±0.04	0.05 ±0.01	nd	6.97 ±0.85	0.833 ±0.006	0
ET 239	1: <i>A. pallidus</i> 2: <i>B. coagulans</i>	5	+/-	+/-	++	+/-	5.03 ±0.22	0.87 ±0.19	0.15 ±0.04	nd ¹	0.07 ±0.00	nd	0.04 ±0.03	nd	6.68 ±0.27	0.928 ±0.041	1: 0 2: bg
ET 224	<i>B. coagulans</i>	5	+/-	+/-	+	+/-	4.47 ±0.12	0.72 ±0.02	0.20 ±0.00	nd ¹	nd	nd	0.06 ±0.01	nd	6.20 ±0.03	0.852 ±0.011	0
ET 251	<i>G. thermodenitrificans</i>	5	+	+/-	+/-	+/-	4.89 ±0.03	0.85 ±0.02	0.13 ±0.00	0.11 ±0.01	0.13 ±0.02	0.00 ±0.00	0.07 ±0.01	nd	6.18 ±0.03	0.876 ±0.002	4
ET 131	1: <i>G. caldxylosilyticus</i> 2: <i>G. thermodenitrificans</i>	2	++	+/-	++	+/-	4.71 ±0.37	0.91 ±0.00	0.14 ±0.04	0.12 ±0.02	nd	0.02 ±0.01	nd	nd	5.91 ±0.39	0.538 ±0.076	1/2: 0
DSM 2542	<i>G. thermoglucosidans</i>	T	+ ⁸	+ ⁸	+ ⁸	- ⁸	4.39 ±0.19	0.92 ±0.03	0.21 ±0.05	0.03 ±0.02	0.09 ±0.13	0.14 ±0.15	nd	0.02 ±0.01	5.81 ±0.26	0.695 ±0.195	2300
ET 130	<i>G. caldxylosilyticus</i>	2	++	++	+	+/-	4.59 ±0.37	0.76 ±0.04	0.18 ±0.01	0.08 ±0.01	nd	0.04 ±0.00	nd	nd	5.64 ±0.40	0.578 ±0.048	1: bg 2: 0
ET 186	<i>G. thermodenitrificans</i>	4	+/-	+	+/-	+/-	4.12 ±0.39	0.92 ±0.01	0.21 ±0.07	0.07 ±0.03	nd	0.02 ±0.00	0.04 ±0.06	nd	5.37 ±0.49	0.803 ±0.168	0
ET 144	1: <i>B. thermocopriae</i> 2: <i>G. thermodenitrificans</i>	3	+/-	+/-	+	+/-	3.63 ±0.34	1.02 ±0.03	0.15 ±0.02	0.12 ±0.10	0.21 ±0.18	0.11 ±0.02	0.08 ±0.01	nd	5.32 ±0.06	0.817 ±0.101	1: 0 2: 6
ET 241	<i>G. thermodenitrificans</i>	5	+/-	+/-	+	+/-	3.99 ±0.89	0.98 ±0.07	0.16 ±0.05	0.01 ±0.01	nd	0.06 ±0.05	0.03 ±0.05	nd	5.23 ±1.00	0.662 ±0.018	0
ET 050	<i>G. thermodenitrificans</i>	1	+/-	+	++	-	3.72 ±0.49	1.01 ±0.24	0.12 ±0.02	0.09 ±0.04	nd	0.02 ±0.00	0.05 ±0.07	nd	5.00 ±0.85	0.611 ±0.125	bg

ET 208	<i>G. caldxylosilyticus</i>	4	+/-	+/-	-	+/-	3.14 ±0.80	1.06 ±0.14	0.57 ±0.20	0.16 ±0.03	nd	0.04 ±0.01	nd	nd	4.98 ±1.18	0.881 ±0.081	bg
ET 129	1: <i>G. caldxylosilyticus</i> 2: <i>G. thermodenitrificans</i>	2	++	+++	+	-	3.56 ±0.48	0.99 ±0.06	0.16 ±0.04	0.08 ±0.02	nd	0.02 ±0.01	nd	nd	4.81 ±0.36	0.579 ±0.082	1: 0 2: bg
ET 143	<i>B. thermocopriae</i>	3	+	+	+	+/-	2.81 ±0.33	1.07 ±0.17	0.15 ±0.04	nd	0.26 ±0.15	0.20 ±0.12	0.01 ±0.01	0.09 ±0.08	4.60 ±0.90	0.719 ±0.095	res
ET 072	<i>G. thermodenitrificans</i>	2	+	+	++	-	3.48 ±0.41	0.81 ±0.18	0.10 ±0.01	nd	nd	0.01 ±0.00	0.11 ±0.15	nd	4.52 ±0.45	0.356 ±0.020	1/2: bg
ET 157	<i>G. thermodenitrificans</i>	3	+	+/-	+/-	-	3.45 ±0.03	0.94 ±0.06	0.07 ±0.01	nd	nd	0.01 ±0.00	0.03 ±0.02	nd	4.49 ±0.03	0.553 ±0.028	0
ET 136	<i>G. thermodenitrificans</i>	3	+	++	+	+	2.46 ±0.07	1.14 ±0.01	0.16 ±0.03	nd	0.50 ±0.42	0.16 ±0.04	nd	0.03 ±0.02	4.45 ±0.58	0.564 ±0.090	0
ET 244	<i>B. thermocopriae</i>	5	+	+/-	+/-	+/-	3.46 ±0.38	0.36 ±0.02	0.01 ±0.00	nd ¹	nd	nd	0.04 ±0.00	0.02 ±0.00	3.89 ±0.41	0.403 ±0.054	0
ET 226	<i>B. thermocopriae</i>	5	+/-	+/-	+++	+/-	3.41 ±0.18	0.38 ±0.01	0.02 ±0.01	nd	nd	nd	0.01 ±0.01	0.02 ±0.00	3.84 ±0.21	0.663 ±0.020	0
ET 225	<i>A. pallidus</i>	5	+/-	+/-	+	+/-	3.07 ±0.87	0.63 ±0.00	0.07 ±0.03	nd	nd	nd	nd	0.03 ±0.01	3.79 ±0.89	0.558 ±0.033	bg
ET 039	<i>G. thermodenitrificans</i>	1	+	+	+	+	2.57 ±0.08	0.92 ±0.07	0.07 ±0.03	0.04 ±0.01	nd	0.02 ±0.00	nd	nd	3.62 ±0.19	0.481 ±0.010	bg
ET 261	1+2: <i>B. thermocopriae</i> 3: <i>G. thermodenitrificans</i>	5	+	+/-	+	+	1.96 ±0.17	1.14 ±0.10	0.06 ±0.02	nd	0.23 ±0.01	0.20 ±0.09	nd	0.03 ±0.00	3.62 ±0.02	0.613 ±0.014	1: bg 2/3: 0
ET 145	<i>G. thermodenitrificans</i>	3	+	+/-	+	+/-	1.91 ±0.03	1.18 ±0.09	0.11 ±0.06	nd	0.24 ±0.05	0.10 ±0.02	0.01 ±0.01	nd	3.54 ±0.14	0.505 ±0.029	0
ET 200	<i>G. thermodenitrificans</i>	4	+	+/-	+	+/-	2.65 ±0.01	0.64 ±0.25	0.16 ±0.04	0.07 ±0.05	nd	nd	nd	nd	3.53 ±0.34	0.729 ±0.055	0
ET 036	<i>G. thermodenitrificans</i>	1	+	+	+	+	2.50 ±1.41	0.80 ±0.31	0.08 ±0.07	0.12 ±0.11	nd	0.02 ±0.01	nd	nd	3.52 ±1.69	0.456 ±0.052	bg
ET 156	<i>G. thermodenitrificans</i>	3	++	++	+	+/-	2.34 ±1.02	0.90 ±0.20	0.16 ±0.11	0.02 ±0.02	nd	0.01 ±0.01	0.04 ±0.05	nd	3.46 ±1.36	0.448 ±0.083	0
ET 042	<i>G. thermodenitrificans</i>	1	+	+	+	+	2.36 ±0.21	0.89 ±0.07	0.03 ±0.01	0.07 ±0.03	nd	0.02 ±0.01	nd	nd	3.37 ±0.27	0.627 ±0.106	0

ET 229	<i>B. coagulans</i>	5	+/-	+/-	+	+/-	2.83 <i>±1.61</i>	0.47 <i>±0.03</i>	0.03 <i>±0.02</i>	nd	nd	nd	nd	0.02 <i>±0.00</i>	3.35 <i>±1.67</i>	0.373 <i>±0.045</i>	0
ET 267	1+2: <i>G. thermodenitrificans</i>	5	+	+/-	+	+/-	2.19 <i>±0.90</i>	0.83 <i>±0.00</i>	0.05 <i>±0.03</i>	0.11 <i>±0.15</i>	0.03 <i>±0.04</i>	0.02 <i>±0.00</i>	nd	nd	3.22 <i>±1.13</i>	0.432 <i>±0.048</i>	1/2: 0
ET 011	1+2: <i>G. thermodenitrificans</i>	1	+	+	+	+	2.00 <i>±0.91</i>	0.79 <i>±0.22</i>	0.08 <i>±0.03</i>	0.08 <i>±0.01</i>	0.14 <i>±0.20</i>	0.02 <i>±0.00</i>	nd	nd	3.11 <i>±1.35</i>	0.539 <i>±0.117</i>	1: 0 2: bg
ET 081	<i>G. thermodenitrificans</i>	2	+	+	++	-	2.03 <i>±0.20</i>	0.93 <i>±0.11</i>	0.07 <i>±0.00</i>	0.06 <i>±0.02</i>	nd	0.01 <i>±0.00</i>	nd	nd	3.10 <i>±0.08</i>	0.740 <i>±0.091</i>	bg
ET 230	1+2: <i>B. coagulans</i>	5	+/-	+/-	++	+++	2.34 <i>±1.00</i>	0.66 <i>±0.00</i>	0.04 <i>±0.01</i>	nd	nd	nd	nd	0.02 <i>±0.00</i>	3.06 <i>±1.01</i>	0.419 <i>±0.072</i>	1/2: bg
ET 263	<i>B. thermocopriae</i>	5	+	+/-	+	+/-	1.72 <i>±0.54</i>	0.85 <i>±0.16</i>	0.03 <i>±0.01</i>	nd	0.24 <i>±0.10</i>	0.05 <i>±0.00</i>	nd	nd	2.89 <i>±0.80</i>	0.600 <i>±0.098</i>	0
ET 159	<i>G. thermodenitrificans</i>	4	+/-	+/-	+/-	+/-	0.74 <i>±0.02</i>	0.67 <i>±0.06</i>	nd	nd	nd	0.05 <i>±0.01</i>	nd	0.01 <i>±0.00</i>	1.46 <i>±0.06</i>	0.253 <i>±0.001</i>	0
ET 160	<i>G. thermodenitrificans</i>	4	+/-	+/-	-	+/-	0.69 <i>±0.03</i>	0.53 <i>±0.03</i>	nd	nd	nd	0.04 <i>±0.06</i>	nd	0.00 <i>±0.01</i>	1.27 <i>±0.11</i>	0.315 <i>±0.103</i>	bg

¹ The selection based on products was made prior to making pure cultures – the identification shown is that after making pure cultures; in cases where the pure culture was split in multiple species both are shown because in this selection round it was still a mixed culture. In the CFU-column, the identification for the subcultures is provided after the corresponding strain sub-number (e.g. 1, 2, or 3) as this was performed after making pure cultures.

² Isol: isolation condition (see Table 1).

³ Sugars used in the selection step in the isolation procedure (see Table 1), prior to making pure cultures. Abbreviations: G: glucose; S: sucrose; X: xylose; A: arabinose; na: not applicable.

⁴ Cultures were analyzed by HPLC after 48 h of growth at their isolation temperature (55 or 65°C) in 15 mL screw-capped tubes with 8 mL TMM + 0.5 g/L yeast extract + 10 g/L glucose + CaCO₃. Transformable strains are shown in bold. Two genetically accessible *Bacillus* type strains known from literature for green chemical or fuel production were taken along as reference strains, e.g. *G. thermoglucosidans* DSM 2542^T (1) and *B. coagulans* DSM 1^T (2, 3). Data are the average of duplicates; standard deviations are indicated in italics after the ‘±’. nd = not detected; nd¹ = a very high malate peak with a large shoulder was observed, indicating the presence of another product besides or instead of malate but this was not further evaluated. Abbreviations: Lac: lactate; Ace: acetate; Suc: succinate; Mal: malate; Eth: ethanol; For: formate; 2,3-BDO: 2,3-butanediol; Prop: propionate.

⁵ CFU = colony forming units per μg DNA after electrotransformation with pNW33n. Transformation was confirmed by isolating plasmid material and subsequent PCR and restriction analysis (Figure S1). Res = naturally resistant to chloramphenicol; nr = not reproducible, bg = background colonies after electroporation, also when no DNA was added, while these strains did not grow on chloramphenicol prior to transformation – colonies by these strains did not test positive in PCR. Results are shown for pure cultures, e.g. 239-1 and 239-2 are '1:0 2:bg' meaning 239-1 showed 0 colonies and 239-2 showed background colonies.

⁶ As reported by (3).

⁷ Acid production from these sugars as determined by (4).

⁸ Acid production from these sugars as determined by (5).

Table S2. Optimization of electrotransformation for *B. smithii* ET 138.

Nr	Parameter changed	Buffer ¹	Final OD ²	Hrs of growth ³	kV	μF	Ω	μg DNA	Cuvette mm	Rec. medium ⁴	Cm rec ⁵	CFU fresh ⁶	CFU -80 ⁶
1	Cm rec+sett.	SG-5	0.483	1.75	1.5	25	600	1	1	RG	N	3	nd
2	Cm rec+sett.	SG-5	0.500	1.5	1.5	25	600	1	1	RG	Y	8	nd
3	Cm rec+sett.	SG-5	0.483	1.75	2.0	25	200	1	2	RG	N	33	nd
4	Cm rec+sett.	SG-5	0.500	1.5	2.0	25	200	5	2	RG	Y	3	nd
5	Settings	SG-5	0.424	2.9	2.0	25	200	1	2	RG	N	12	nd
6	Settings	SG-5	0.424	2.9	1.5	25	200	0.7	1	RG	N	nd	113
7	Settings	SG-5	0.424	2.9	2.0	25	200	0.7	2	RG	N	nd	7
8	Settings	SG-5	0.424	2.9	2.0	25	400	0.7	2	RG	N	nd	157
9	OD	SG-5	0.452	1.4	2.0	25	400	1	2	RG	N	70	nd
10	OD	SG-5	0.678	1.4	2.0	25	400	1	2	RG	N	11	nd
11	Rec. medium	SG-5	0.574	2.1	2.0	25	400	1	2	LB2	N	712	1149
12	Rec. medium	SG-5	0.574	2.1	2.0	25	400	1	2	RG	N	0	nd
13	Rec. medium	SG-5	0.514	2.1	2.0	25	400	1	2	LB2	N	406	443
14	Rec. medium	SG-5	0.514	2.1	2.0	25	400	1	2	RG	N	1	nd
15	Glycerol, rec.	SG-10	0.446	2.5	2.0	25	400	1	2	LB2	N	796	476
16	Glycerol, rec.	SG-10	0.446	2.5	2.0	25	400	1	2	RG	N	36	nd
17	Rec. medium	SG-5	0.522	2.1	2.0	25	400	1	2	LB2	N	590	nd
18	Rec. medium	SG-5	0.522	2.1	2.0	25	400	1	2	RG	N	0	nd
19	Rec. medium	SG-5	0.519	1.75	2.0	25	400	1	2	LB2	N	960	nd
20	Rec. medium	SG-5	0.519	1.75	2.0	25	400	1	2	RG	N	5	nd
21	Buffer	SM-10	0.454	2.5	2.0	25	400	1	2	LB2	N	40	0
22	Buffer	SM-10	0.454	2.5	2.0	25	400	1	2	RG	N	0	nd
23	Fast growth ⁷	SG-5	0.539	1	2.0	25	400	2.5	2	LB2	N	1900	nd
24	Fast growth ⁷	SG-5	0.512	1.25	2.0	25	400	1	2	LB2	N	nd	1122
25	μg DNA ⁷	SG-5	0.617	1.3	2.0	25	400	0.2	2	LB2	N	2409	nd
26	μg DNA ⁷	SG-5	0.617	1.3	2.0	25	400	0.02	2	LB2	N	5118	nd

Lines indicate different experiments. Nr. 1 is the original protocol as described by Rhee *et al.*(6). Nr.3 are settings based on based on (7), which are the settings as used for screening of *Geobacillus* strains.

¹ 5 or 10 indicates the % of glycerol. SG = sucrose glycerol buffer from *Bacillus* protocol (6); SM = sorbitol mannitol buffer from *Geobacillus* protocol (8).

² Final OD₆₀₀ after the indicated number of hours (³) when growing cells prior to making them competent.

³ Number of hours cells had grown before making them competent.

⁴ RG is LB with 121 g/L sucrose and 10 g/L glucose (6).

⁵ Whether or not 1/1000 diluted chloramphenicol was added after 2h of recovery. Y = yes; N = no.

⁶ CFU: colony forming units per μg DNA. Cells were either electroporated directly after making them competent ('fresh') or after storage in -80°C ('-80'). nd = not determined.

⁷ In these experiments, after overnight growth, cells were transferred to 500 mL Erlenmeyer flasks or 1 L bottles (having a similar bottom surface) instead of to a 250 mL Erlenmeyer to allow for more aeration.

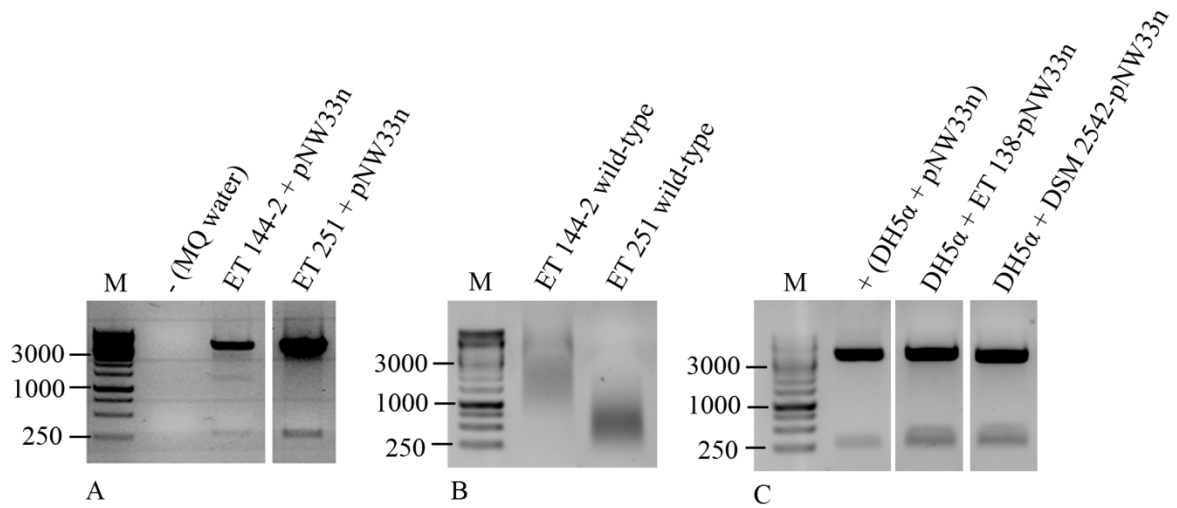


Figure S1. Restriction analysis of plasmid DNA extracted from isolates and control strain transformed with pNW33n. Restriction was performed with *Stu*I and *Hind*III; pNW33n digested with *Stu*I and *Hind*III yields bands of 282 and 3955 bp. M: Fermentas 1 kb DNA ladder. For each digestion reaction, 550-650 ng isolated plasmid DNA was used. **A:** Plasmids isolated from *G. thermodenitrificans* strains ET 144-2 and ET 251 transformed with pNW33n. **B:** Plasmids isolated from *G. thermodenitrificans* strains ET 144-2 and ET 251 prior to transformation with pNW33n (negative control) **C:** Plasmids isolated from *E. coli* DH5 α transformed pNW33n (original source) and after retransformation with plasmids isolated from *B. smithii* ET 138 transformants and *G. thermoglucosidans* DSM 2542 transformants.

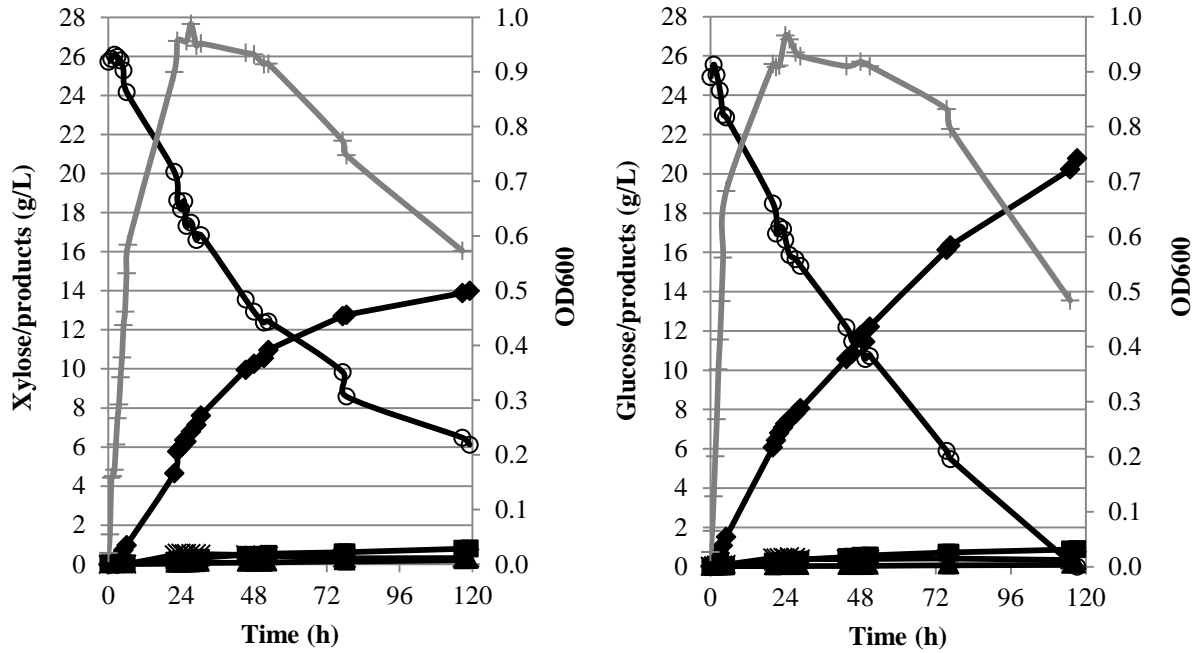


Figure S2. Fermentation of *B. smithii* ET 138 on 25 g/L xylose and glucose. Fermentation was carried out in 1 L TVMY supplemented with 25 g/L xylose or glucose, at 55°C, pH 6.5, 150 rpm and without any gas addition. Grey line with plus-sign: OD600; open circles: xylose or glucose; filled diamonds: lactate; closed squares: acetate; crosses: malate and succinate; closed triangles: pyruvate. During the xylose fermentation, severe browning of the medium was observed after approximately 70h.

SUPPLEMENTARY REFERENCES

1. **Cripps RE, Eley K, Leak DJ, Rudd B, Taylor M, Todd M, Boakes S, Martin S, Atkinson T.** 2009. Metabolic engineering of *Geobacillus thermoglucosidasius* for high yield ethanol production. *Metab. Eng.* **11**:398-408.
2. **Kovacs AT, van Hartskamp M, Kuipers OP, van Kranenburg R.** 2010. Genetic tool development for a new host for biotechnology, the thermotolerant bacterium *Bacillus coagulans*. *Appl. Environ. Microbiol.* **76**:4085-4088.
3. **Van Kranenburg R, Van Hartskamp M, Heintz E, Anthonius J, Van Mullekom E, Snelders J.** 2007. WO Patent WO/2007/085,443. Genetic modification of homolactic thermophilic *Bacilli*.
4. **Jung MY, Kim J-S, Chang Y-H.** 2009. *Bacillus acidiproducens* sp. nov., vineyard soil isolates that produce lactic acid. *Int. J. Syst. Evol. Microbiol.* **59**:2226-2231.
5. **Suzuki Y, Kishigami T, Inoue K, Mizoguchi Y, Eto N, Takagi M, Abe S.** 1983. *Bacillus thermoglucosidasius* sp. nov., a new species of obligately thermophilic *Bacilli*. *Syst. Appl. Microbiol.* **4**:487-495.
6. **Rhee MS, Kim Jw, Qian Y, Ingram LO, Shanmugam KT.** 2007. Development of plasmid vector and electroporation condition for gene transfer in sporogenic lactic acid bacterium, *Bacillus coagulans*. *Plasmid* **58**:13-22.
7. **Narumi I, Sawakami K, Nakamoto S, Nakayama N, Yanagisawa T, Takahashi N, Kihara H.** 1992. A newly isolated *Bacillus stearothermophilus* K1041 and its transformation by electroporation. *Biotechnol. Tech.* **6**:83-86.
8. **Taylor MP, Esteban CD, Leak DJ.** 2008. Development of a versatile shuttle vector for gene expression in *Geobacillus* spp. *Plasmid* **60**:45-52.