Supplementary Materials: DNA and RNA Extraction and Quantitative Real-Time PCR-Based Assays for Biogas Biocenoses in an Interlaboratory Comparison

Michael Lebuhn ^{1,*}, Jaqueline Derenkó ^{2,†}, Antje Rademacher ^{2,†}, Susanne Helbig ^{3,†}, Bernhard Munk ^{1,†}, Alexander Pechtl ^{4,†}, Yvonne Stolze ^{5,†}, Steffen Prowe ³, Wolfgang H. Schwarz ⁴, Andreas Schlüter ⁵, Wolfgang Liebl ⁴ and Michael Klocke ²



Figure S1. Effect of different MgCl₂ concentrations on the qPCR efficacy. DNA from cattle manure spiked with *E. coli* was extracted using system FSKS and quantified using the *mur*A system (see Section 2.3).

Partner/Analyst		Α	В	С	D
DNA quantification		NanoDrop 1000 + PicoGreen	IMPLEN NanoPhotometer ^{TM}	NanoDrop 1000	NanoDrop 1000 + Picogreen
qPCR reagents		Thermo Scientific DyNAmo Flash Probe qPCR Kit; Mg ²⁺ as supplied by manufacturer;	Self-made master-mix containing 2xEvaGreen, 200 μM dNTPs and 0.75 U Platinum Taq-Polymerase; Mg ²⁺ 6 mM in qPCR assays (Lebuhn <i>et al.</i> , 2003; Munk <i>et al.</i> , 2010);	Biorad SsoAdvanced™ Universal SYBR® Green Supermix; Mg²+ as supplied by manufacturer;	LC480 Roche Probes Master; Mg ²⁺ 3.2 mM in qPCR assays;
Reaction volume		18 μL master-mix + 2 μL extract	24 μ L master-mix + 1 μ L extract	15 μ L master-mix + 2 μ L extract	15 μ L master-mix + 5 μ L extract
Primers and probes	Bac fw	Bac Fw (ACT CCT ACG GGA GGCAG), [1]		Eub 338 (ACTCCTACGGGAGGCAG), [3]	Bac Fw (ACTCCTACGGGAGGCAG), [1]
	Bac probe	Bac Probe (6-FAM-TGCCA GCAGC CGCGG TAATA C-TAMRA), [1]			Bac Probe (6-FAM-TGCCA GCAGC CGCGG TAATA C-TAMRA), [1]
	Bac rev	Bac-rev (GAC TAC CAG GGT ATC TAA TCC), [1]		Eub 518 (ATTACCGCGGCTGCTGG), [3]	Bac-rev (GACTA CCAGG GTATC TAATC C), [1]
	Arc fw	Arc-fw (ATTAGA TACCCS BGTAGT CC), [1]			
	Arc Probe	Arc-Probe (6-FAM-AGGAA TTGGC GGGGG AGCAC-TAMRA), [1]			
	Arc rev	Arc-rev (GCCATG CACCWC CTCT), [1]			
	mcrA/mrtA fw		MeA-i 1046f (TAYATGWSIGGHGGIGTIGGI TTYAC), [2]		
	mcrA/mrtA rev		MeA-i 1435r (TGRTCYTGIARRTCRWAICCR WAGAAICC), [2]		
qPCR platform		CFX96 Touch (Biorad)	MX 3005P qPCR System (Agilent)	CFX96 Touch (Biorad)	LightCycler480 (Roche)

Table S1. qPCR assays used in the interlaboratory comparison.

Table S1. Cont.									
Partner/Analyst		Α	В	С	D				
Protocols	Bac	95 °C, 07:00		98 °C, 02:00	Pre-Incubation				
	protocols	95 °C, 00:15		98 °C, 00:05	95 °C, 10:00				
	(i, iv, v)	57 °C, 00:30		56 °C, 00:35	95 °C, 00:10				
		60 °C, 00:60		40x	60 °C, 00:30				
		45x			72 °C, 00:01 (acquisition)				
					45x				
	Arc	95 °C, 07:00							
	protocol (ii)	95 °C, 00:15							
		60 °C, 00:60							
		45x							
	mcrA/mrtA		95 °C, 03:00						
	protocol (iii)		95 °C, 00:15						
			60 °C, 01:00						
			82 °C, 00:10 (acquisition)						
			45x						



Figure S2. DNA concentrations calculated per mL or g of fresh sample matter of extracts produced by the five partners of the interlaboratory comparison as determined with the extinction coefficient for ds-DNA (analysts B, C and E) or the PicoGreen[®] system (analysts A, D).



Figure S3. DNA concentrations in the extracts prepared by the partners calculated per mL g of fresh sample matter using only the data from the PicoGreen[®] system. The medians over the different analysts with the SDs are presented.

Extracts 1 (Partner A)						Extracts 2 (Partner B)					Extracts 3 (Pa	artner C)		
Analyst B	A260/280	A260/230	A320	Analyst E	A260/280	A260/230	Analyst B	A260/280	A260/230	A320	Analyst B	A260/280	A260/230	A320
HR1-T1	1.767	0.03	0.011	HR1-T1	1.86	0.03	HR1-T1	<d.l.< td=""><td><d.l.< td=""><td><d.l.< td=""><td>HR1-T1</td><td>2</td><td>0.022</td><td>0.002</td></d.l.<></td></d.l.<></td></d.l.<>	<d.l.< td=""><td><d.l.< td=""><td>HR1-T1</td><td>2</td><td>0.022</td><td>0.002</td></d.l.<></td></d.l.<>	<d.l.< td=""><td>HR1-T1</td><td>2</td><td>0.022</td><td>0.002</td></d.l.<>	HR1-T1	2	0.022	0.002
HR1-T2	1.714	0.031	0.014	HR1-T2	1.5	0.04	HR1-T2	<d.l.< td=""><td><d.l.< td=""><td><d.1.< td=""><td>HR1-T2</td><td>1.667</td><td>0.024</td><td>0.003</td></d.1.<></td></d.l.<></td></d.l.<>	<d.l.< td=""><td><d.1.< td=""><td>HR1-T2</td><td>1.667</td><td>0.024</td><td>0.003</td></d.1.<></td></d.l.<>	<d.1.< td=""><td>HR1-T2</td><td>1.667</td><td>0.024</td><td>0.003</td></d.1.<>	HR1-T2	1.667	0.024	0.003
HR2-T1	1.719	0.03	0.014	HR2-T1	1.57	0.04	HR2-T1	<d.l.< td=""><td><d.l.< td=""><td><d.1.< td=""><td>HR2-T1</td><td>1.7</td><td>0.037</td><td>0.003</td></d.1.<></td></d.l.<></td></d.l.<>	<d.l.< td=""><td><d.1.< td=""><td>HR2-T1</td><td>1.7</td><td>0.037</td><td>0.003</td></d.1.<></td></d.l.<>	<d.1.< td=""><td>HR2-T1</td><td>1.7</td><td>0.037</td><td>0.003</td></d.1.<>	HR2-T1	1.7	0.037	0.003
HR2-T2	1.647	0.042	0.014	HR2-T2	1.63	0.05	HR2-T2	<d.l.< td=""><td><d.l.< td=""><td><d.1.< td=""><td>HR2-T2</td><td>2</td><td>0.034</td><td>0.004</td></d.1.<></td></d.l.<></td></d.l.<>	<d.l.< td=""><td><d.1.< td=""><td>HR2-T2</td><td>2</td><td>0.034</td><td>0.004</td></d.1.<></td></d.l.<>	<d.1.< td=""><td>HR2-T2</td><td>2</td><td>0.034</td><td>0.004</td></d.1.<>	HR2-T2	2	0.034	0.004
HR1-M1	1.686	0.042	0.021	HR1-M1	1.54	0.04	HR1-M1	<d.l.< td=""><td><d.l.< td=""><td><d.l.< td=""><td>HR1-M1</td><td>1.4</td><td>0.025</td><td>0.003</td></d.l.<></td></d.l.<></td></d.l.<>	<d.l.< td=""><td><d.l.< td=""><td>HR1-M1</td><td>1.4</td><td>0.025</td><td>0.003</td></d.l.<></td></d.l.<>	<d.l.< td=""><td>HR1-M1</td><td>1.4</td><td>0.025</td><td>0.003</td></d.l.<>	HR1-M1	1.4	0.025	0.003
HR1-M2	2.75	0.044	-0.002	HR1-M2	1.42	0.09	HR1-M2	<d.l.< td=""><td><d.l.< td=""><td><d.1.< td=""><td>HR1-M2</td><td>2.4</td><td>0.017</td><td>0.002</td></d.1.<></td></d.l.<></td></d.l.<>	<d.l.< td=""><td><d.1.< td=""><td>HR1-M2</td><td>2.4</td><td>0.017</td><td>0.002</td></d.1.<></td></d.l.<>	<d.1.< td=""><td>HR1-M2</td><td>2.4</td><td>0.017</td><td>0.002</td></d.1.<>	HR1-M2	2.4	0.017	0.002
HR2-M1	1.756	0.044	0.015	HR2-M1	1.63	0.05	HR2-M1	<d.1.< td=""><td><d.l.< td=""><td><d.1.< td=""><td>HR2-M1</td><td>1.667</td><td>0.032</td><td>0.002</td></d.1.<></td></d.l.<></td></d.1.<>	<d.l.< td=""><td><d.1.< td=""><td>HR2-M1</td><td>1.667</td><td>0.032</td><td>0.002</td></d.1.<></td></d.l.<>	<d.1.< td=""><td>HR2-M1</td><td>1.667</td><td>0.032</td><td>0.002</td></d.1.<>	HR2-M1	1.667	0.032	0.002
HR2-M2	1.826	0.04	0.024	HR2-M2	1.62	0.04	HR2-M2	<d.l.< td=""><td><d.l.< td=""><td><d.1.< td=""><td>HR2-M2</td><td>1.8</td><td>0.033</td><td>0.002</td></d.1.<></td></d.l.<></td></d.l.<>	<d.l.< td=""><td><d.1.< td=""><td>HR2-M2</td><td>1.8</td><td>0.033</td><td>0.002</td></d.1.<></td></d.l.<>	<d.1.< td=""><td>HR2-M2</td><td>1.8</td><td>0.033</td><td>0.002</td></d.1.<>	HR2-M2	1.8	0.033	0.002
CSTR1-T1	1.957	0.021	0.012	CSTR1-T1	1.67	0.02	CSTR1-T1	1.727	0.052	0.005	CSTR1-T1	0.8	0.016	0.002
CSTR1-T2	1.676	0.032	0.044	CSTR1-T2	1.67	0.03	CSTR1-T2	1.615	0.041	0.004	CSTR1-T2	1.556	0.037	0.003
CSTR2-T1	1.742	0.031	0.015	CSTR2-T1	1.5	0.03	CSTR2-T1	1.6	0.045	0.005	CSTR2-T1	2	0.031	0.003
CSTR2-T2	1.848	0.031	0.015	CSTR2-T2	1.63	0.03	CSTR2-T2	1.5	0.146	0.004	CSTR2-T2	1.625	0.038	0.002
CSTR1-M1	1.793	0.025	0.014	CSTR1-M1	1.38	0.02	CSTR1-M1	1.692	0.14	0.003	CSTR1-M1	1.667	0.033	0.001
CSTR1-M2 *	1.929	0.036	0.028	CSTR1-M2	1.73	0.02	CSTR1-M2	1.632	0.091	0.009	CSTR1-M2	1.75	0.044	0.002
CSTR2-M1	1.926	0.028	0.019	CSTR2-M1	1.38	0.03	CSTR2-M1	1.733	0.058	0.004	CSTR2-M1	1.8	0.1	0.002
CSTR2-M2	1.71	0.028	0.017	CSTR2-M2	1.67	0.02	CSTR2-M2	1.786	0.139	0.005	CSTR2-M2	1.714	0.041	0.001
E	Extracts 4 (Par	rtner D)			Extracts 5 (Partner E)									
Analyst B	A260/280	A260/230	A320	Analyst B	A260/280	A260/230	A320	Analyst E	A260/280	A260/230				
HR1-T1	1.75	0.019	0.008	HR1-T1	1.765	0.435	0.002	HR1-T1	1.85	1.26				
HR1-T2	1.647	0.019	0.008	HR1-T2	1.806	0.699	0.002	HR1-T2	1.76	0.79				
HR2-T1	1.833	0.019	0.009	HR2-T1	1.933	0.426	0.001	HR2-T1	1.74	0.66				
HR2-T2	1.5	0.061	0.01	HR2-T2	1.905	0.506	0.002	HR2-T2	1.77	1.05				
HR1-M1	1.8	0.017	0.007	HR1-M1	1.812	0.403	0.001	HR1-M1	1.78	0.51				
HR1-M2	1.684	0.022	0.01	HR1-M2	1.818	0.525	0.001	HR1-M2	1.77	0.67				
HR2-M1	1.591	0.031	0.01	HR2-M1	1.759	0.646	0.001	HR2-M1	1.8	1.24				
HR2-M2	1.706	0.022	0.008	HR2-M2	1.795	0.814	0.004	HR2-M2	1.82	1.17				
CSTR1-T1	2.077	0.036	0.016	CSTR1-T1	n.d.	n.d.	n.d.	CSTR1-T1	1.345	0.34				
CSTR1-T2	1.909	0.023	0.008	CSTR1-T2	1.783	1.019	0.004	CSTR1-T2	2.5	1.12				
CSTR2-T1	2.14	0.042	0.021	CSTR2-T1	n.d.	n.d.	n.d.	CSTR2-T1	n.d.	n.d.				
CSTR2-T2	2.293	<d.l.< td=""><td>0.019</td><td>CSTR2-T2</td><td>n.d.</td><td>n.d.</td><td>n.d.</td><td>CSTR2-T2</td><td>n.d.</td><td>n.d.</td><td></td><td></td><td></td><td></td></d.l.<>	0.019	CSTR2-T2	n.d.	n.d.	n.d.	CSTR2-T2	n.d.	n.d.				
CSTR1-M1*	2.654	0.03	0.017	CSTR1-M1	1.763	1.305	0.005	CSTR1-M1	2.04	1.34				
CSTR1-M2	2.306	0.036	0.025	CSTR1-M2	1.773	1.376	0.005	CSTR1-M2	1.92	1.89				
CSTR2-M1*	2.552	0.033	0.019	CSTR2-M1	1.775	1.153	0.004	CSTR2-M1	1.93	1.79				
CSTR2-M2*	3 235	<d 1<="" td=""><td>0.022</td><td>CSTR2-M2</td><td>18</td><td>0.462</td><td>0.002</td><td>CSTR2-M2</td><td>2.06</td><td>1 41</td><td></td><td></td><td></td><td></td></d>	0.022	CSTR2-M2	18	0.462	0.002	CSTR2-M2	2.06	1 41				

Table S2. Absorbance ratios A260/280 and A260/230, and absorbance at 320 nm of extracts produced by the five and analyzed by two partners/analysts of the interlaboratory comparison.

*: qPCR inhibition by undiluted extract; n.d.: not determined/not delivered; <d.l.: below detection limit; red color denotes values below and pink color values above conventional thresholds.

References

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