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2 **Supplementary material**

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4 **Unexpected diversity and high abundance of putative nitric oxide dismutase (Nod)**
5 **genes in contaminated aquifers and wastewater treatment systems**

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7 Baoli Zhu^a, Lauren Bradford^a, Sichao Huang^a, Anna Szalay^a, Carmen Leix^b, Max Weissbach^b,
8 András Táncsics^c, Jörg E. Drewes^b, Tillmann Lueders^a

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10 ^a Institute of Groundwater Ecology, Helmholtz-Zentrum Muenchen, Germany

11 ^b Chair of Urban Water Systems Engineering, Technical University of Munich, Germany

12 ^c Szent István University, Gödöllő, Hungary

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17 **Supplementary Fig. S1.** Multiple sequence alignment of selected putative Nod and qNor
18 enzymes around the quinol-binding site and the catalytic site of qNor. Representative
19 environmental Nods were deduced from the gene sequences generated in this study (in
20 bold). The alignment is extended by further Nod sequences generated in this study
21 compared to Fig. 5. The conserved residuals for quinol-binding (A) and catalytic functioning
22 (B) in qNor are highlighted in red, whereas substitutions at these sites in putative Nod and
23 putative Nor are shown in green. Note that the number and order of Nod sequences included
24 in (A) and (B) are different. Accession numbers are the same as in Fig. 2. The alignment
25 was generated with ClustalW in MEGA 6.

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		328	332	336	746	
qNor	<i>Geobacillus stearothermophilus</i>	ALLAH	HYYTE	PDS	FFGI	PDI
	<i>Staphylococcus aureus</i>	ELLAH	YVENK	-	FFGI	WDI
	<i>Neisseria gonorrhoeae</i>	GLTAH	YTV	EGQG	FFGI	PDL
	<i>M. oxyfera</i> DAMO_1889	AAVAH	YRAE	PGKF	FFGL	GDA
	HdN1	GFTA	HYT	EGQT	FFGI	GDV
unknown Nor-related	<i>Bacteroidetes bacterium</i>	VLTVH	DFV	GFVN	FFGF	GGS
	<i>Sediminibacterium</i> sp.	ILTVH	DFV	GFVH	FFGV	GGA
	<i>Chlorobi bacterium</i>	ILTVH	DFV	GFVN	FFGY	GGA
	<i>Algoriphagus mannitolivorans</i>	VLTVH	DFV	GFTS	FFGL	GGS
	<i>Mariniradius saccharolyticus</i>	VLTVH	DFV	GFTN	FFGV	GGS
	<i>Cecembia lonarensis</i>	VLTVH	DFV	GFTK	FFGW	GGA
	<i>Indibacter alkaliphilus</i>	VLTVH	DFV	GFTK	FFGW	GGA
	<i>Flaviumibacter</i> sp.	ILTVH	DFV	GFVN	FFGF	GGS
	Siklós Sik2DC09 KX364417	VLTVH	DFVN	FTV	FFGF	· · ·
	<i>Muricauda ruestringensis</i>	FITINE	FIDY	LG	YFGI	GAC
	<i>Arenibacter algicola</i>	FVTINE	FVDY	LG	FFGV	GAC
	<i>M. oxyfera</i> DAMO_2437	ILGAED	FVGG	PG	EAI	GEV
	<i>M. oxyfera</i> DAMO_2434	ILSAED	FVGG	PG	SAI	GGA
	<i>Methylomirabilis</i> sp.	ILSAED	FVGG	PG	SAL	GGA
	<i>Methylomirabilis</i> sp.	ILGAED	FVGG	PG	GESI	GGA
HdN1 Nod	IAAAW	DFVK	P	----	GIA	
Nod	Siklós Sik2DC15 KX364445	ILGAED	FVGG	PG	GETI	
	Siklós Sik2DC08 KX364446	IIGAED	FIGG	PP	VDAM	
	Siklós Sik2DC03 KX364447	ILSAEN	FVK	SG	PGTVI	
	Siklós Sik2DA06 KX364449	ILGAED	FVGG	PG	EAI	
	Siklós Sik2DA05 KX364450	ILSAED	FVGG	PG	SAL	
	Siklós Sik1DB16 KX364451	ILGAED	FVGG	PG	GETI	
	Siklós Sik1DB14 KX364452	ILSAED	FVGG	PG	GNAI	
	WWTP-TUM 8-2-M12 KX364419	VIAAM	DFV	LP		
	WWTP-TUM 1-4 KX364420	IISAT	D	FLRP		
	WWTP-TUM 1-11 KX364422	IVGAS	D	FIKP		
	WWTP-Kempton R2-3 KX364423	VIAAW	DFI	KP		
	WWTP-Kempton R2-2 KX364424	MAAAW	DFV	KP		
	WWTP-Kempton R2 KX364428	VIAAM	DFV	LP		
	2-stage-nitritation 2-4 KX364429	IAAAW	DFV	KP		
	2-stage-nitritation 2-1 KX364430	IAAAW	DFV	KP		
	2-stage-AMX 3-5 KX364431	ILCAT	D	FVRP		
	2-stage-AMX 3-3 KX364432	IAAAW	DFV	KP		
	Swing-redox 1-5 KX364433	ILAST	D	FVRP		
	Suspension R3-2 KX364435	VVSAW	D	FIRP		
	Suspension R3-1 KX364436	ILCAN	D	FVRT		
	CANDO R5-3 KX364441	MAAAW	DFV	KP		
	CANDO R4-1 KX364444	VIAAW	DFI	KP		
	Siklós Sik2DA12 KX364448	IVGA	A	DFIKP		
	Flingern Fln8DB07 KX364453	VISAT	D	FLRP		
	WWTP-TUM 1-12 KX364421	· · ·	FVRT	DRL		
	Suspension R1-3 KX36439	· · · · ·	D	FIKP		

Fig S1 A

		508	512	559 560	581
qNor	<i>Geobacillus stearothermophilus</i>	I I H L W V E G		G H H Y	L E V
	<i>Staphylococcus aureus</i>	I V H L W V E G		G H H Y	L E V
	<i>Neisseria gonorrhoeae</i>	V V H L W V E G		L H H L	L E V
	<i>M. oxyfera</i> DAMO_1889	I V H L W V E G		G H H W	M E V
	HdN1	V V H L W V E G		F H H L	L E V
unknown Nor-related	<i>Bacteroidetes bacterium</i>	V I H M W A E A		S H N F	L Q V
	<i>Sediminibacterium</i> sp.	V I H M W A E A		S H N F	L Q V
	<i>Chlorobi bacterium</i>	V I H M W A E A		S H N F	L Q V
	<i>Algoriphagus mannitolivorans</i>	V V H M W V E A		S H N F	L Q V
	<i>Mariniradius saccharolyticus</i>	V V H M W V E A		S H N F	L Q V
	<i>Cecembia lonarensis</i>	V V H M W V E A		S H N F	L Q V
	<i>Indibacter alkaliphilus</i>	V V H M W V E A		S H N F	L Q V
	<i>Flavihumbacter</i> sp.	V V H M W A E A		S H N F	L Q V
	WWTP-Kempton R2-7 KX364416	V I H M W V E A		A H N F	. . .
	Siklós Sik2DC09 KX364417	V I H M W A E A		S H N F	. . .
	<i>Muricauda ruestringensis</i>	V V H M W V E A		S H N F	L Q F
	<i>Arenibacter algicola</i>	V V H M W V E A		S H N F	L Q F
	<i>M. oxyfera</i> DAMO_2437	N I H M W V E V		S H N F	M Q V
	<i>M. oxyfera</i> DAMO_2434	N I H M W V E V		S H N F	M Q V
	<i>Methylomirabilis</i> sp.	N I H M W V E V		S H N F	M Q V
<i>Methylomirabilis</i> sp.	N I H M W V E V		S H N F	M Q V	
HdN1 Nod	V V H M W V E V		S H N F	L Q V	
Nod	WWTP-Kempton R2-3 KX364423	T V H M W V E V		S H N F	L Q V
	WWTP-Kempton R2-2 KX364424	V V H M W V E V		S H N F	L Q V
	Siklós Sik2DA12 KX364448	V V H M W V E V		S H N F	L Q V
	Siklós Sik2DA06 KX364449	N I H M W V E V		S H N F	M Q V
	Siklós Sik2DA05 KX364450	N I H M W V E V		S H N F	M Q V
	WWTP-TUM M16-4 KX364418	V A H M W V E V		A H N F	
	WWTP-TUM M12-8-2 KX364419	V V H M W V E V		S H N F	
	WWTP-TUM 1-12 KX364421	V V H M W V E A		S H N F	
	WWTP-Kempton R2-8 KX364425	V V H M W V E A		S H N F	
	WWTP-Kempton R2-2 KX364426	T V H M W V E V		S H N F	
	WWTP-Kempton R2-1 KX364427	V V H M W V E V		A H N F	
	2-stage-nitrification 2-4 KX364429	V V H M W V E V		S H N F	
	Suspension R3-22 KX364434	V V H M W V E V		S H N F	
	Suspension R3-2 KX364435	T V H M W V E V		S H N F	
	Suspension R1-7 KX364437	V V H M W V E V		A H N F	
	Suspension R1-5 KX364438	V V H M W V E V		S H N F	
	Suspension R1-3 KX364439	V V H M W V E V		S H N F	
	CANDO R5-6 KX364440	V V H M W V E V		S H N F	
	CANDO R5-2 KX364442	V V H M W V E V		S H N F	
	CANDO R4-12 KX364443	V V H M W V E V		A H N F	
CANDO R4-1 KX364444	T V H M W V E V		S H N F		
Siklós Sik2DC15 KX364445	N I H M W V E V		S H N F		
Siklós Sik2DC08 KX364446	N I H M W V E V		S H N F		
Siklós Sik2DC03 KX364447	T V H M W V E V		S H N F		

Fig S1 B