Supplementary Material

Carbon Steel Corrosion by Bacteria from Failed Seal Rings at an Offshore Facility

Silvia J. Salgar-Chaparro^a, Adam Darwin^b, Anna H. Kaksonen^c, Laura L. Machuca^{a*}

^a Curtin Corrosion Centre, WA School of Mines: Minerals, Energy and Chemical Engineering, Curtin

University, Kent Street, Bentley, WA 6102, Australia

^b Woodside Energy Ltd., Perth, 6000 WA, Australia

^c Commonwealth Scientific and Industrial Research Organization (CSIRO), Land and Water, 147 Underwood Avenue, Floreat WA 6014, Australia

* Correspondence:

Laura L. Machuca

I.machuca2@curtin.edu.au

Content

Table S1. Microbial community composition in failed seal rings based on partial 16S rRNA gene sequencing

Table S2. Morphological characterisation of bacteria isolated from corroded seal rings

Table S3. Weight loss of coupons exposed to abiotic and biotic conditions for 7 days.

Table S4. *p*-value calculated from the Kruskal-Wallis test applied to evaluate differences in the corrosion rates and pit depths among coupons exposed to the isolated bacteria and the control for 7 days.

Table S5. Classification of corrosion rates triggered by isolated bacteria

Table S6. Weight loss of coupons exposed to abiotic and biotic conditions for 21 days.

Table S7. *p*-value calculated from the Kruskal-Wallis test applied to evaluate differences in the corrosion rates and pit depths among coupons exposed to the isolated bacteria and the control for 21 days.

Fig S1. Images of the carbon steel coupons exposed for 7 days to the bacteria isolated from corroded seal rings.

Figure S2. Test solution pH (A) and planktonic cell numbers (B) during the long term study. Replenishment of media is indicated by the dashed line.

Table S1. Microbial community composition in failed seal rings based on partial 16S rRNA gene sequencing

Taxa	Relative abundance (%)			
Taxa	S1	S2	\$3	
Actinobacteria				
Micrococcus	0.01			
<u>Bacteroidetes</u>				
Marinifilum	0.01			
<u>Cyanobacteria</u>				
Synechococcus CC9902			0.02	
<u>Epsilonbacteraeota</u>				
Arcobacter	2.22	0.19	0.09	
Firmicutes				
Bacillus	0.21	0.27	1.82	
Oceanobacillus		0.19	0.03	
Sporosarcina			0.37	
Unclassified Carnobacteriaceae			0.36	
Vallitalea	0.01			
Proteobacteria				
Brevundimonas		0.15		
Martelella	0.20	5.66	0.05	
Labrenzia	1.48	24.67	0.07	
Bradyrhizobium		0.05		
Nautella	0.02			
Unclassified Rhodobacteraceae	0.01			
Tistrella	0.35			
Desulfofaba	0.02			
Halodesulfovibrio	0.05			
Desulfuromusa	0.01			
Geoalkalibacter	0.11	0.17		
Idiomarina	0.03			
Shewanella	10.91			
Achromobacter	0.06			
Ralstonia		0.08		
Enterobacter	3.39	0.47		
Escherichia-Shigella	0.57	2.67	0.08	
Alcanivorax	6.46			
Chromohalobacter			0.73	
Halomonas	1.91			
Salinicola	0.02			
Marinomonas	0.11	18.49		
Acinetobacter	0.01	0.16	0.11	
Alkanindiges	0.04			
Psychrobacter			1.36	
Pseudomonas	65.89	46.73	94.91	
Vibrio	5.87			
Stenotrophomonas		0.05		

Table S2. Morphological characterisation of bacteria isolated from corroded seal rings

Colony ID	Colony Colour	Colony form	Colony elevation	Colony margin	Gram	Morphology
CCC-APB1	white	circular	raised	entire	negative	rod
CCC-APB3	pale yellow	circular	raised	entire	negative	rod
CCC-APB5	pale orange	circular	raised	entire	negative	rod
CCC-SPP14	white	circular	raised	entire	negative	rod
CCC-SPP15	black	punctiform	raised	entire	negative	rod
CCC-IOB1	brown	circular	raised	entire	negative	rod
CCC-IOB3	metallic brown	circular	flat	entire	negative	rod
CCC-IOB9	orange	punctiform	raised	entire	negative	rod
CCC-IOB10	dark orange	circular	raised	entire	negative	rod

 Table S3. Weight loss of coupons exposed to abiotic and biotic conditions for 7 days.

	Weight loss (mg)			
Test	Coupon 1	Coupon 2	Coupon 3	
Control	4.6	4.5	4.3	
CCC-APB1	4.4	6.5	5.8	
CCC-APB3	16.1	5.1	13.8	
CCC-APB5	10.6	11.8	10.6	
CCC-SPP14	27.4	44.1	20.1	
CCC-SPP15	3.9	4.6	3.1	
CCC-IOB1	14.1	22.1	17.2	
CCC-IOB3	2.4	1.6	2.8	
CCC-IOB9	9.7	5.7	11.8	
CCC-IOB10	14.0	5.1	7.9	

Table S4. *p*-value calculated from the Kruskal-Wallis test applied to evaluate differences in the corrosion rates and pit depths among coupons exposed to the isolated bacteria and the control for 7 days.

		<i>p</i> value		
Test 1	Test 2	Corrosion	Pit depth	
		Tale		
Control	CCC-APB1	0.275	0.007	
Control	CCC-APB3	0.049	0.007	
Control	CCC-APB5	0.049	0.007	
Control	CCC-SPP14	0.049	0.007	
Control	CCC-SPP15	0.275	0.008	
Control	CCC-IOB1	0.049	0.007	
Control	CCC-IOB3	0.049	0.212	
Control	CCC-IOB9	0.049	0.007	
Control	CCC-IOB10	0.049	0.008	

p > 0.05 = not significant

	Corrosion rate	Maximum pitting rate
	categorisation*	Categorisation*
CCC-APB1	Moderate	High
CCC-APB3	Severe	Severe
CCC-APB5	Severe	High
CCC-SPP14	Severe	Severe
CCC-SPP15	Low	Severe
CCC-IOB1	Severe	Severe
CCC-IOB3	Low	Low
CCC-IOB9	Severe	Severe
CCC-IOB10	Severe	Severe

*Categorization corresponds to the corrosion and pitting rate calculated after subtracting the corrosion and pitting rate measured in the abiotic control. Corrosion rate categories (mm/y) Low (< 0.025), Moderate (0.025-0.12), High (0.13-0.25), Severe (> 0.25). Pitting rate categories (mm/y) Low (< 0.13), Moderate (0.13-0.20), High (0.21-0.38), and Severe (> 0.38).

 Table S6. Weight loss of coupons exposed to abiotic and biotic conditions for 21 days.

Weight loss (mg) Coupon 1 Coupon 2 Coupon 3			
7.1	6.2	4.8	
11.8	8.8	10.6	
7.6	7.7	6.3	
	Coupon 1 1.9 7.1 11.8 7.6	Coupon 1 Coupon 2 1.9 2.2 7.1 6.2 11.8 8.8 7.6 7.7	

Table S7. *p*-value calculated from the Kruskal-Wallis test applied to evaluate differences in the corrosion rates and pit depths among coupons exposed to the isolated bacteria and the control for 21 days.

		<i>p</i> value	
Test 1	Test 2	Corrosion rate	Pit depth
Control	CCC-IOB3	0.049	0.004
Control	CCC-SPP14	0.049	0.004
Control	CCC-APB5	0.049	0.004



Figure S1. Images of the carbon steel coupons exposed for 7 days to the bacteria isolated from corroded seal rings.



Figure S2. Test solution pH (A) and planktonic cell numbers (B) during the long term study. Replenishment of media is indicated by the dashed line. Figures generated using OriginPro version 2020 (OriginLab Corporation). https://www.originlab.com/