# Genomics 101: How genomics can assist in MIC management

### **Lisa Gieg** Biological Sciences





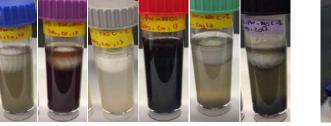
- Environmental samples can be monitored for the presence/activity of microorganisms in several ways – most common ways in O&G industry:
  - Enzyme tests e.g., ATP assay
  - 'Bug bottles' e.g., BART tests, MPNs
    - SRB
    - APB
    - HAB
  - Molecular microbiological methods (MMM)
    - = Genomics

- ATP assay
- 'Bug bottles' e.g., BART tests, MPNs
  - SRB
    APB
    HAB

**KPI** – 'number of bacteria' for system monitoring

- 'High' numbers don't necessarily mean MIC will be a problem
- 'Low' numbers don't necessarily mean MIC won't be a problem

- 'Bug bottles' e.g., BART tests, MPNs
  - SRB
  - APB
  - HAB
  - IRB, etc







### **Major Limitation:**

• Growth based, targets <1% of microbes in a sample

-e.g., captures only some SRB but not S<sup>0</sup> or thiosulfate-reducers

Most microbes will not be captured in growth-based tests

### Molecular microbiological methods (MMM) a.k.a. Genomics

- Newer approach, but now being widely used to identify microbes in O&G systems
- Not growth based



## Based on DNA – life's blueprint!



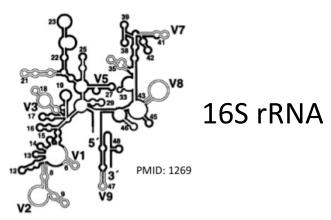
### Genomics

Based on DNA – life's blueprint!

- Uses genetic information contained in microbial cells to determine what types of microbes and/or their potential metabolism (e.g., sulfide production)
- Not based on growth therefore better at capturing the diveristy of microbes in a sample

## Genomics as a Monitoring Tool – 3 ways

- 1. 16S rRNA gene survey commonly-used method
- Biological molecule present in all microbial life (involved in protein synthesis) conserved and variable regions



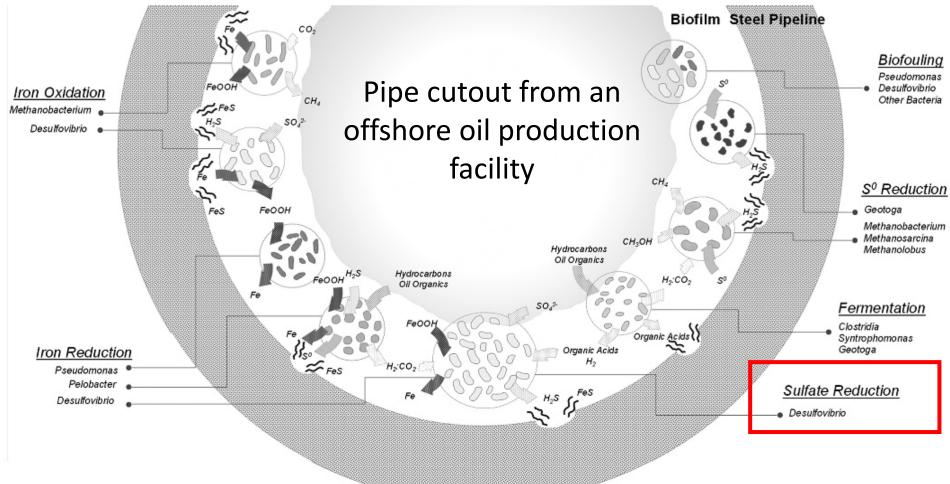
- Extract DNA
- AMPLIFY using primers in a PCR reaction make millions and millions of copies of a part of the 16S rRNA gene
- SEQUENCE on an instrument
- LIST of microbial names not quantitative but get a 'relative abundance'

## Genomics as a Monitoring Tool – 16S

#Taxonomy (Class; Order; Family; Genus)	Relative abundance (% of total sequencing reads)					
	9FW 08/15	10TW 08/15	9FW 01/15	10TW 01/15	4PW 01/15	
Gammaproteobacteria;Alteromonadales; Alteromonadaceae;Marinobacter;	2.7	1.0	8.1	1.4	1.8	
Clostridia; Halanaerobiales; Halanaerobiaceae; Halanaerobium;	31.1	29.3	31.1	66.5	36.0	
Gammaproteobacteria; Thiotrichales; Piscirickettsiaceae; Thiomicrospira;	0.1	0.2	0.0	0.0	0.0	
Gammaproteobacteria; Oceanospirillales; Halomonadaceae; Modicisalibacter;	28.0	35.8	0.0	0.0	0.0	
Deltaproteobacteria; Desulfovibrionales; Desulfohalobiaceae; Desulfovermiculus;	1.0	1.2	1.0	6.4	40.5	
Gammaproteobacteria; Oceanospirillales; Halomonadaceae; Halomonas;	1.4	0.9	0.0	0.9	0.0	
Methanomicrobia; Methanosarcinales; Methanosarcinaceae; Methanohalophilus;	16.4	16.0	5.1	3.2	2.9	
Gammaproteobacteria;	0.0	0.0	0.4	0.3	0.0	
Betaproteobacteria; Burkholderiales; Burkholderiaceae; Ralstonia;	0.0	0.0	25.6	0.1	0.0	
Methanococci; Methanococcales; Methanococcaceae; Methanothermococcus;	1.2	1.3	0.0	1.4	1.4	
Gammaproteobacteria; Oceanospirillales; Halomonadaceae; Chromohalobacter;	9.8	5.5	0.0	0.0	1.3	

### Genomics as a Monitoring Tool – 16S

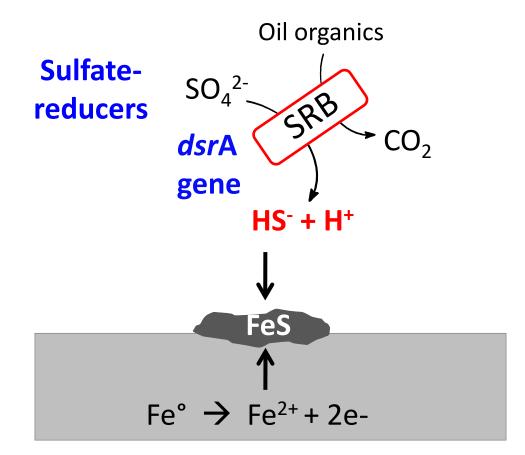
### 'List of microorganisms' – infer metabolism!



From Vigneron et al., 2016

## Genomics as a Monitoring Tool – 3 ways

- 2. 'Functional' gene analysis commonly-used method
- Targets gene specific to a certain kind of microbe or metabolism
- Can quantify these to obtain numbers of specific microbes
  - qPCR
  - Primers specific to genes for sulfate-reducers, methanogens, etc.
  - Total Bacteria, Archaea



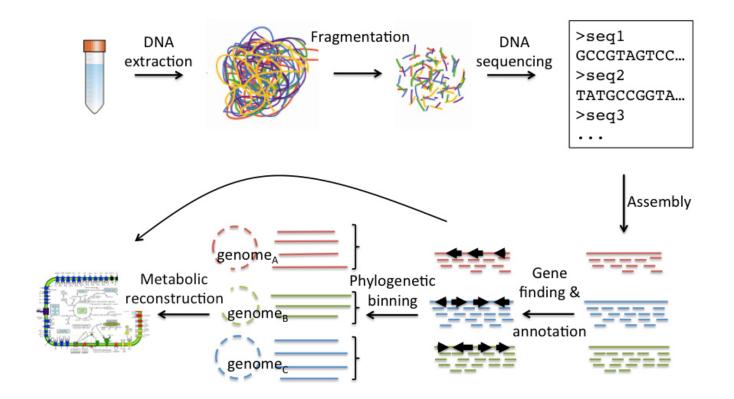
**Chemical MIC** 

### Genomics as a Monitoring Tool – qPCR

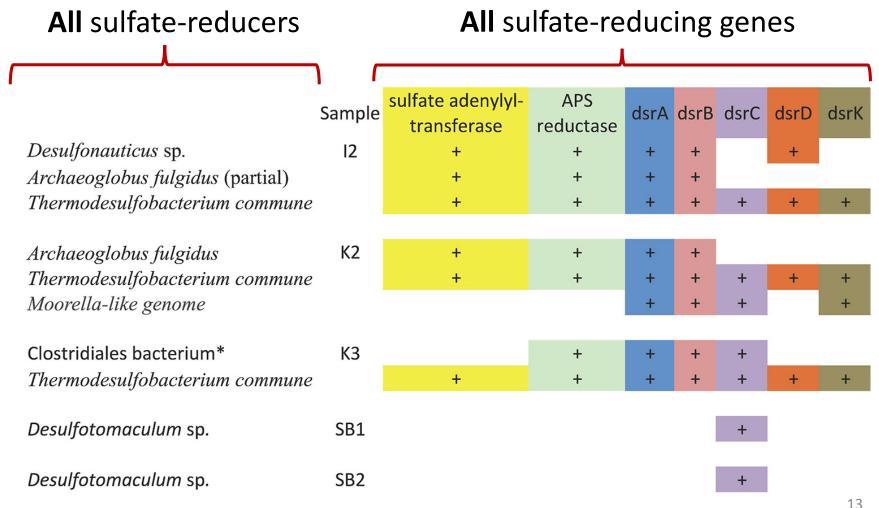
Analysis	Location <sup>a</sup>			
qPCR	Eider Production Manifold	Otter Production Pipeline	Inlet to Crude Crude Oil Oil Coalescer Coalescer PV V-1100 outlet V-110	
qPCR (gene abundance per cm <sup>2</sup> ) <sup>c</sup> Sessile samples from corrosion coupons				
Total bacteria	$< 4.0 \times 10^{2}$	$4.6 \times 10^{5}$	$< 4.0 \times 10^{2}$	$2.9 \times 10^3$
SRB	$< 4.0 \times 10^{2}$	$2.9 \times 10^4$	$< 4.0 \times 10^{2}$	$3.0 \times 10^3$
SRA	$< 4.0 \times 10^{2}$	$1.9  imes 10^4$	$< 4.0 \times 10^{2}$	$1.1 \times 10^4$
Methanogens	$1.4  imes 10^5$	$4.2 \times 10^{11}$	$5.0  imes 10^5$	$4.2 \times 10^5$
MPN (cells per cm <sup>2</sup> ) <sup>b</sup> Sessile samples from corrosion coupons				
mSRB (30 °C)	$5.9 \times 10^{\circ}$	$3.3 \times 10^1$	$0.5  imes 10^{0}$	$0.5  imes 10^{0}$

## Genomics as a Monitoring Tool – 3 ways

- 3. Metagenomics less commonly used
  - All DNA sequenced in a sample
    - All the genetic potential of all microorganisms



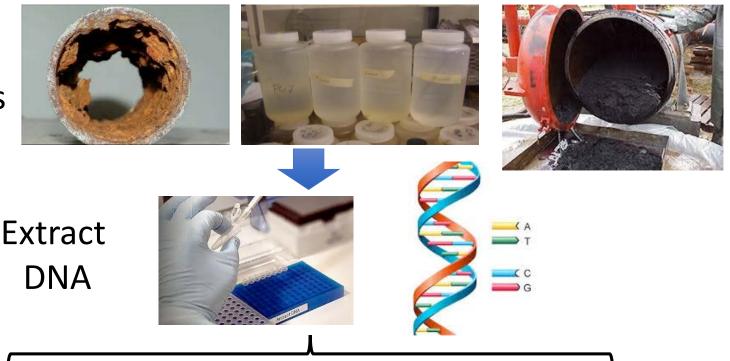
# Genomics as a Monitoring Tool – metagenomics



From Hu et al. 2016

## Genomics as a Monitoring Tool

Field Samples



### **Amplicon sequencing**

-16S rRNA gene

- -Identifies all microbes
- -Other 'functional' genes

-Identifies specific microbes/functions

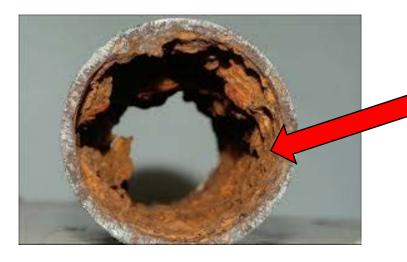
### **Metagenomic Sequencing**

-All DNA sequenced

-Complete genetic potential

# Genomics as a Monitoring Tool

• Sampling & sample preservation very important!



# Surface solids samples – where the key microbes are

- Very important to measure physical & chemical parameters
- Know operating conditions!

\*Genomics data must be interpreted in context of a given operation and associated chemistry!

### Microbiologically-influenced corrosion = microbiology + corrosion

\*holds true for data from all microbial tests

## Genomics in MIC Management

Corrosion Management Process Step	Benefits of applying Genomics/MMM	
MIC Threat Assessment	<ul> <li>Characterize baseline microbiological conditions</li> <li>Look for associations between biofilm, biofilm composition and corrosion</li> <li>Relate biofilm characteristics with operating conditions</li> </ul>	
MIC Mitigation Selection		
MIC Barrier Monitoring	<ul> <li>Monitor long term and short term effects of mitigation on biofilm</li> <li>Monitor changes in chemical effectiveness</li> <li>Watch for shifts in microbiological populations</li> </ul>	

## A 'Genomics' Case Study

APPLIED AND ENVIRONMENTAL MICROBIOLOGY, Oct. 2011, p. 6908–6917 0099-2240/11/\$12.00 doi:10.1128/AEM.05891-11 Copyright © 2011, American Society for Microbiology. All Rights Reserved. Vol. 77, No. 19

#### Effect of Sodium Bisulfite Injection on the Microbial Community Composition in a Brackish-Water-Transporting Pipeline<sup>∀</sup>†

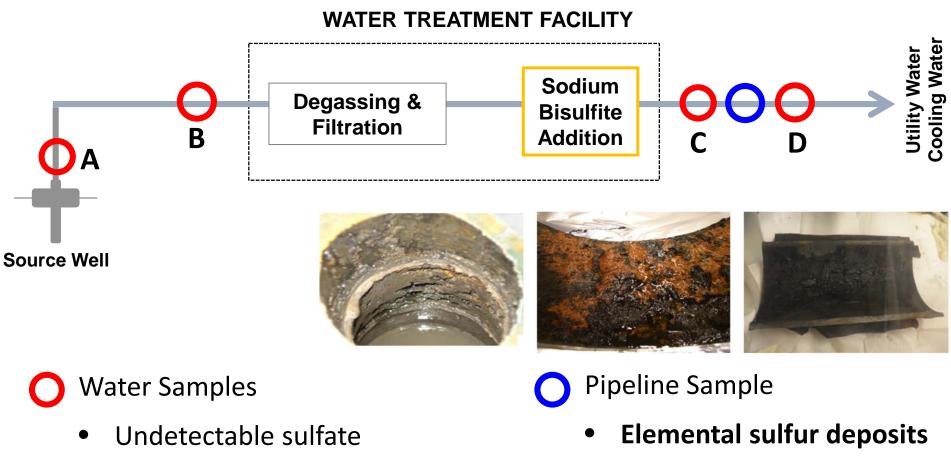
Hyung Soo Park,<sup>1</sup>‡ Indranil Chatterjee,<sup>1</sup>‡ Xiaoli Dong,<sup>2</sup> Sheng-Hung Wang,<sup>2</sup> Christoph W. Sensen,<sup>2</sup> Sean M. Caffrey,<sup>1</sup> Thomas R. Jack,<sup>1</sup> Joe Boivin,<sup>3</sup> and Gerrit Voordouw<sup>1</sup>\*

Petroleum Microbiology Research Group, Department of Biological Sciences, University of Calgary, Calgary, Alberta T2N 1N4, Canada<sup>1</sup>; Visual Genomics Centre, Faculty of Medicine, University of Calgary, Calgary, Alberta T2N 4N1, Canada<sup>2</sup>; and Cormetrics Limited, 56 Hawkwood Place NW, Calgary, Alberta T3G 1X6, Canada<sup>3</sup>

Received 18 June 2011/Accepted 5 August 2011

## Effect of Bisulfite on Biocorrosion

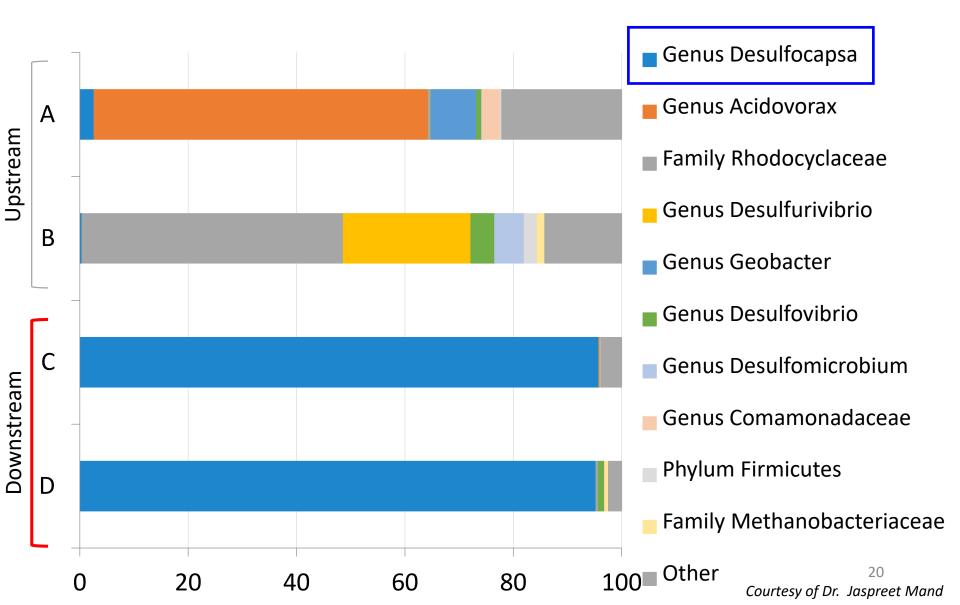
### 'Who is there?' 16S rRNA gene sequencing...



• Low sulfide concentrations

**Iron sulfide deposits** 

## Microbial Community Composition



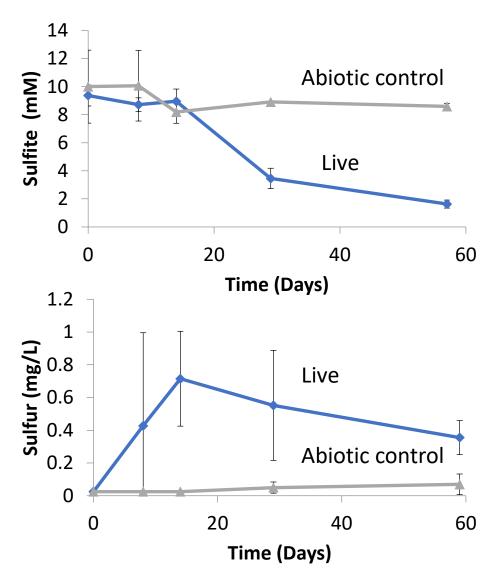
# Sulfite and MIC

- It was assumed that sulfur, found downstream of bisulfite addition, was due to oxygen ingress
- However, sulfur may be result of bisulfite metabolism (disproportionation) by *Desulfocapsa*:

 $3HSO_3^- \rightarrow S^0 + 2 SO_4^{2-} + H_2O + H^+$ 

Resulting byproducts may be used by other microorganisms in the system

## Microbial Enrichment on Sulfite



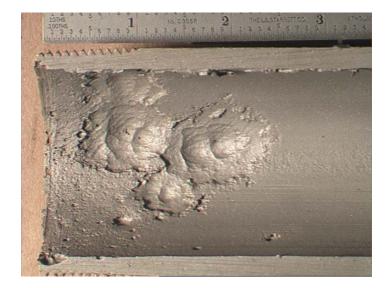
- Sulfite degradation is seen in incubations with the field sample relative to the abiotic control
- Elemental sulfur is produced
  - Genomics helped to pinpoint the problem
  - Altered KPI for bisulfite treatment

22 Courtesy of Dr. Jaspreet Mand

## Genomics in MIC Management

- Genomics is a powerful microbiological monitoring tool that captures broadest diversity in a field sample
- contributes a 'piece of the corrosion puzzle' to help guide corrosion management and mitigation
  - monitoring programs (KPI)
  - effectiveness of treatments
  - failure analysis
  - risk management

Thank you! Questions?



R. Eckert, Materials Performance