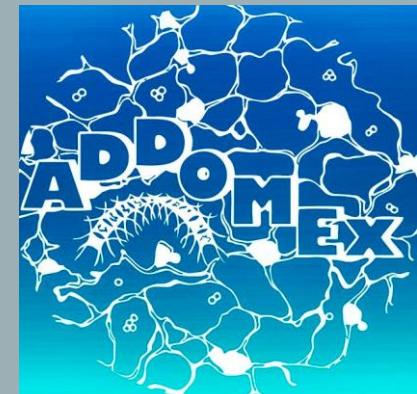


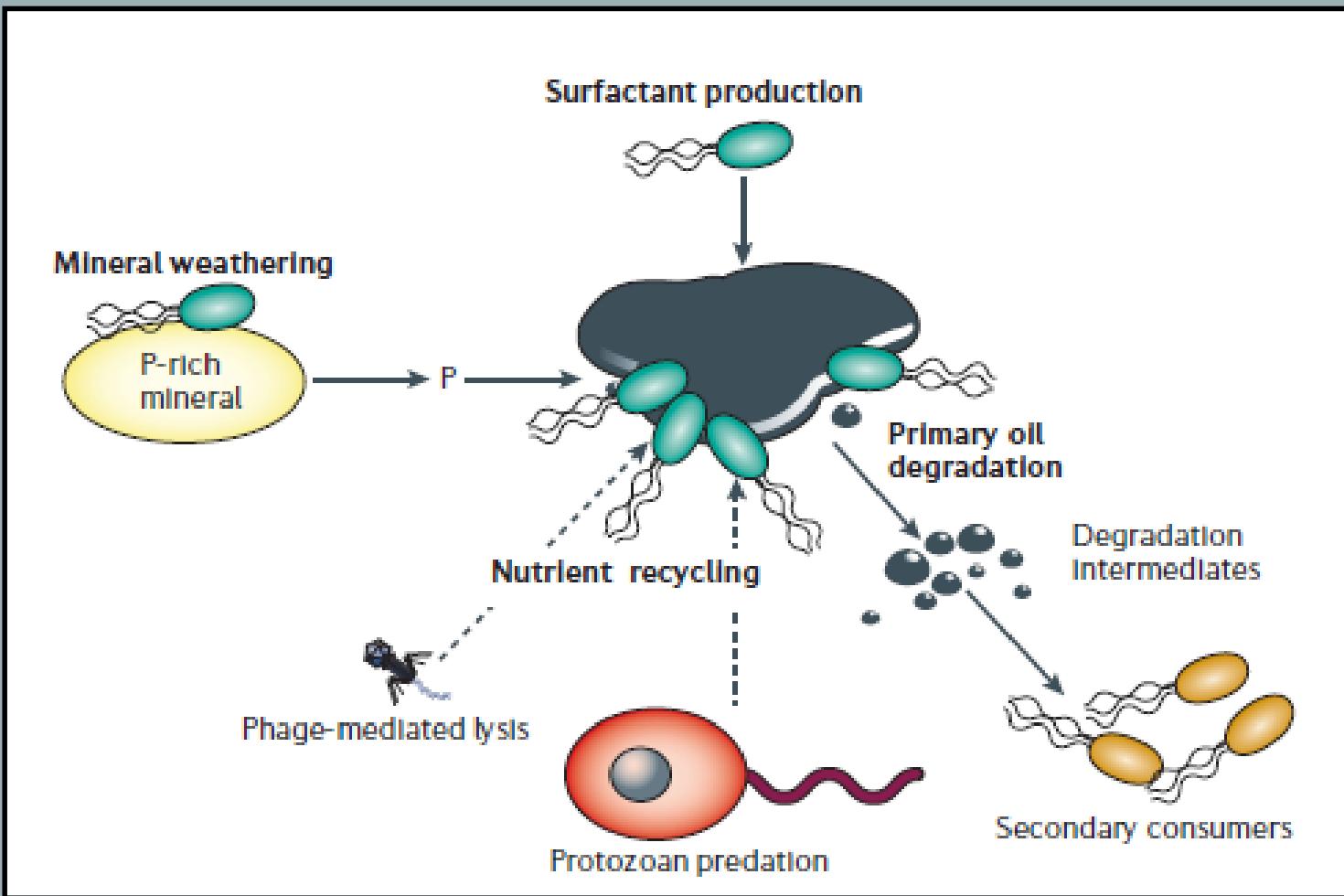
MICROBIAL INTERACTIONS ON MOS

Kai Zier vogel

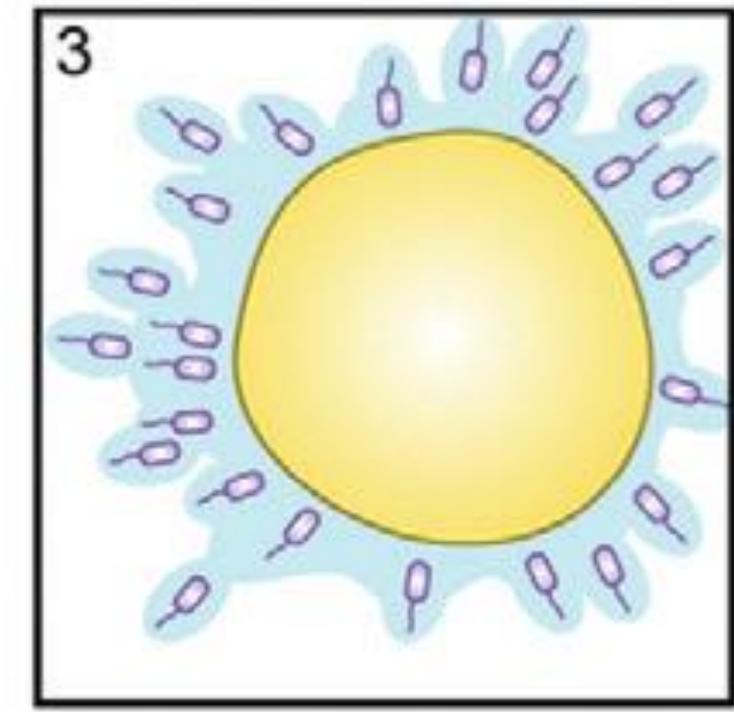
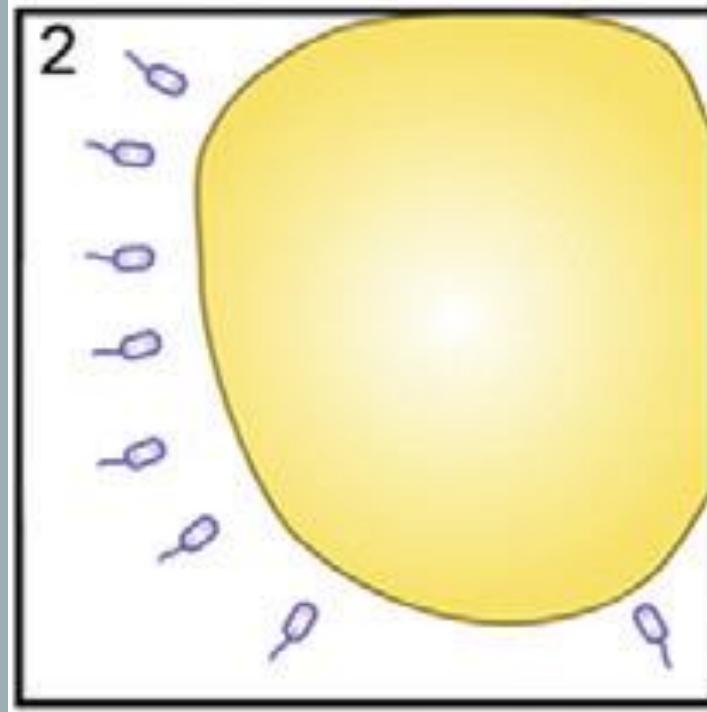
University of New Hampshire,
University of California, Santa Barbara



MICROBES AND OIL



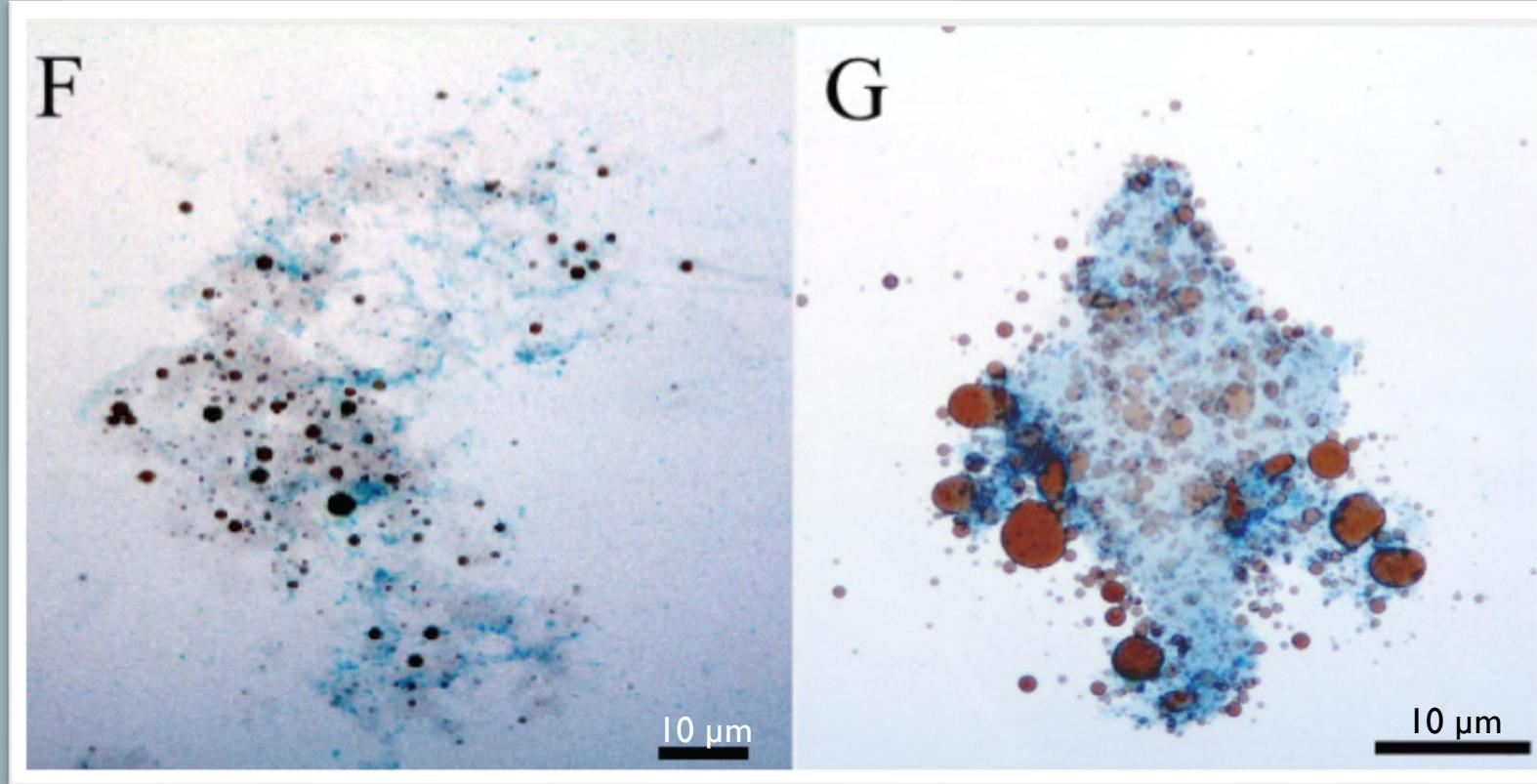
EPS – EXOPOLYMERIC SUBSTANCES



Bælum et al. (2012)

Yellow – oil, blue – EPS, with associated microbial cells

EPS AND OIL: (F) POLYSACCHARIDES AND (G) PROTEINS

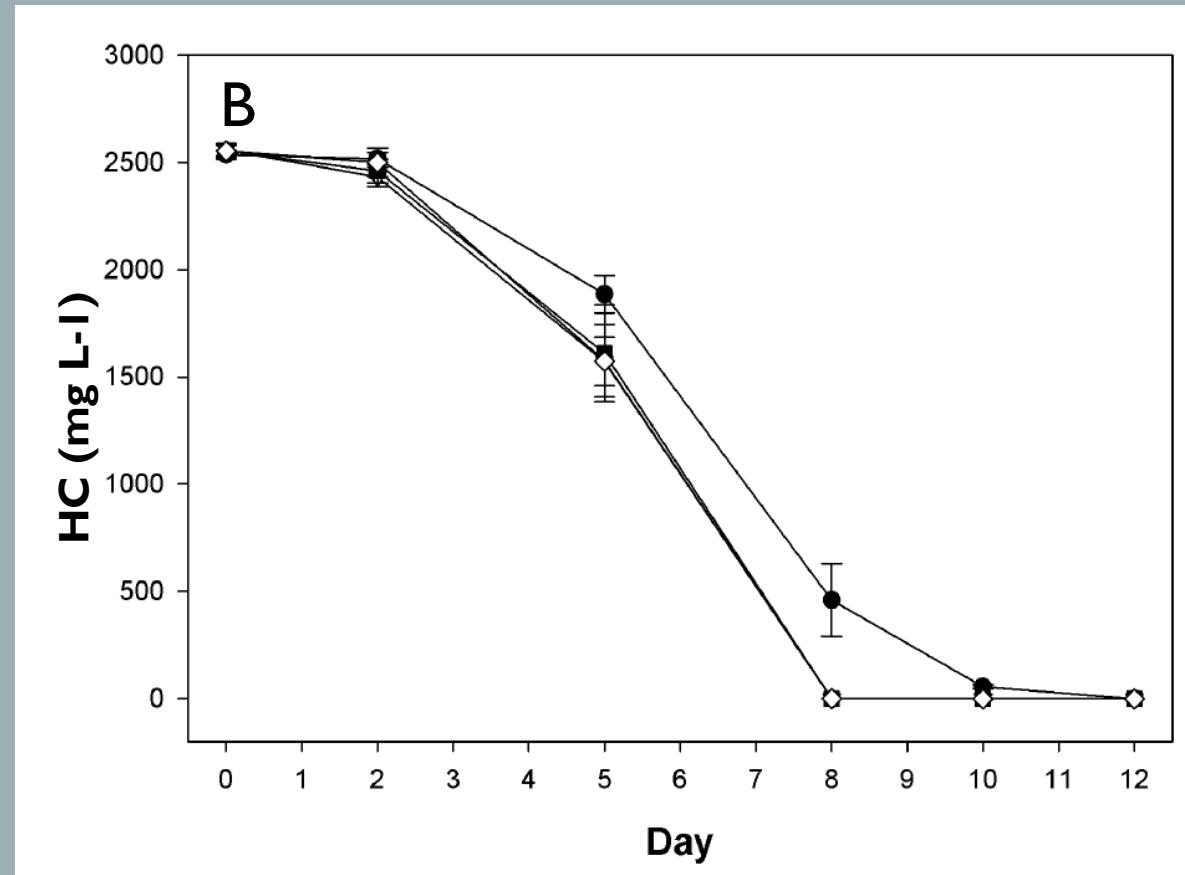
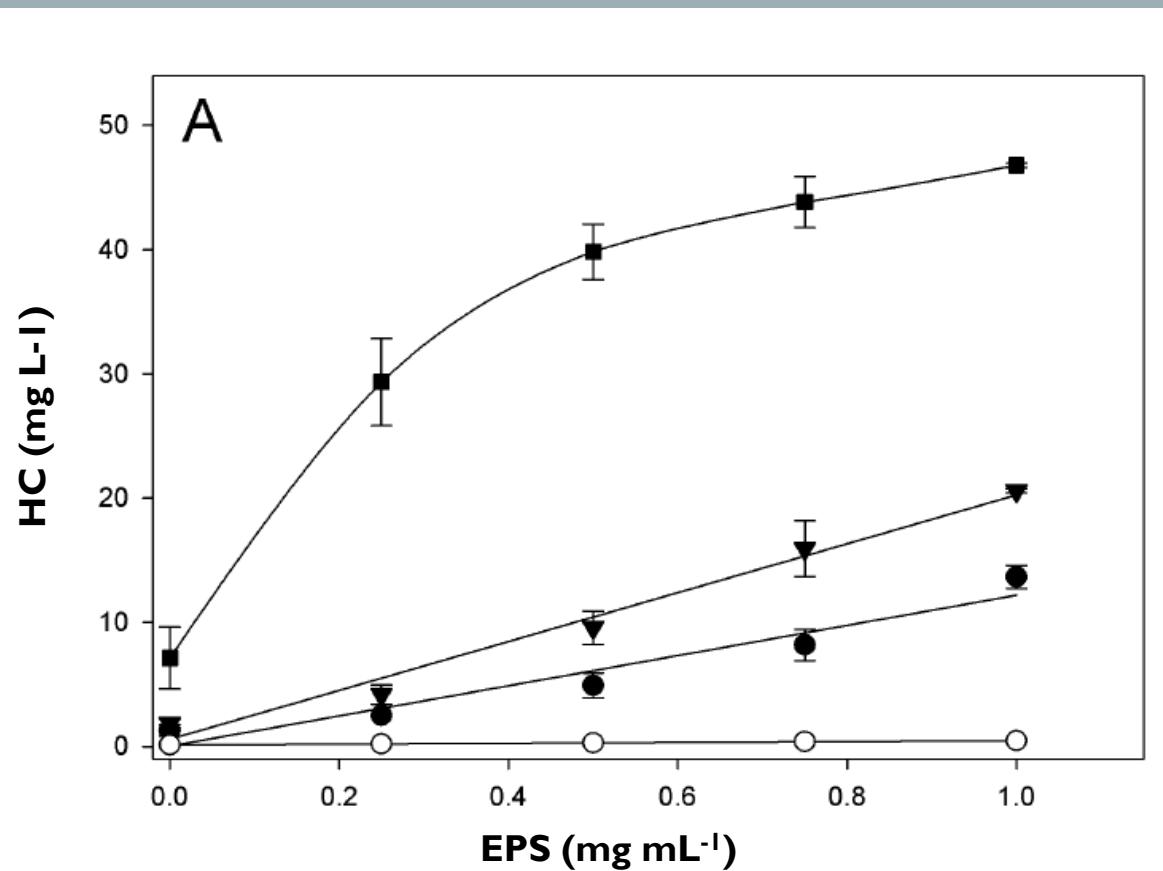


Gutierrez et al. (2013)

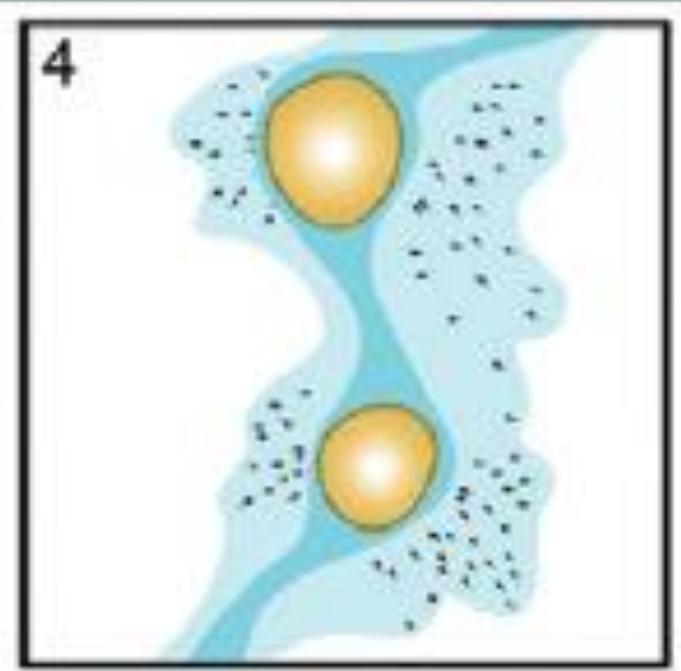
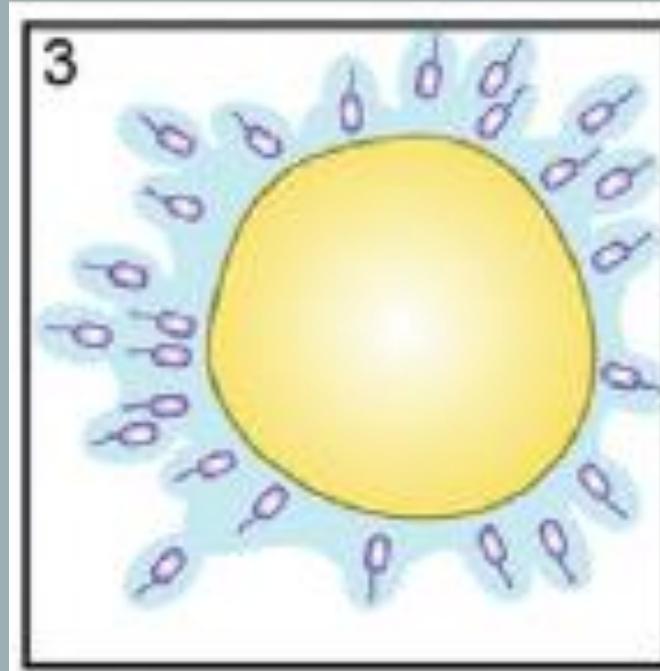
**Black/reddish dots – oil droplets, blue – EPS, microbial cells not visible
(*Halomonas* sp.)**

EPS AND OIL:

(A) ENHANCED SOLUBILITY OF HYDROCARBONS (HC)
(B) ENHANCED HC DEGRADATION



EPS AS MATRIX FOR MICROBIAL OIL FLOCS



Bælum et al. (2012)

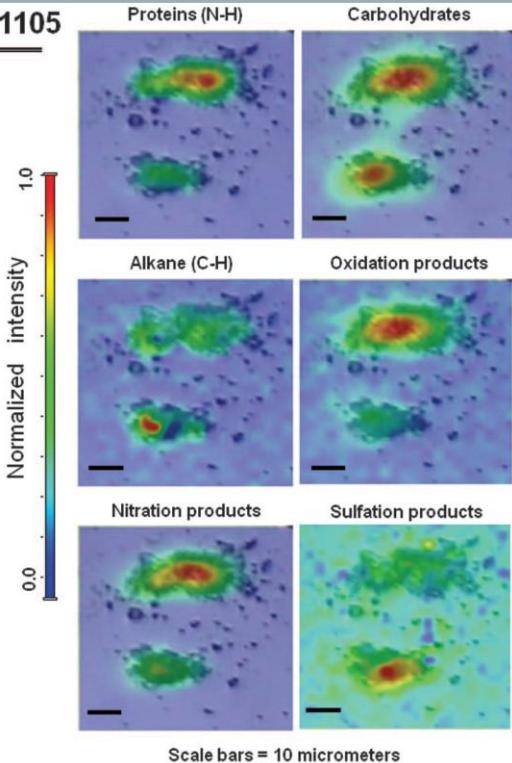
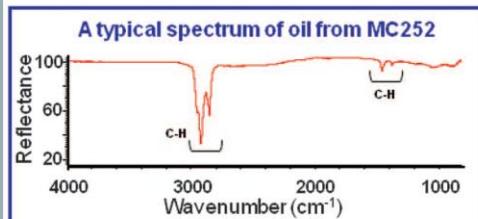
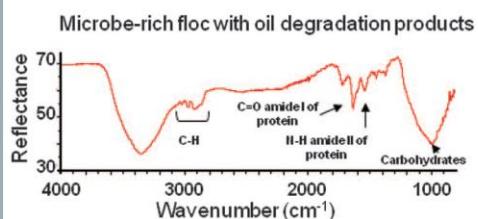
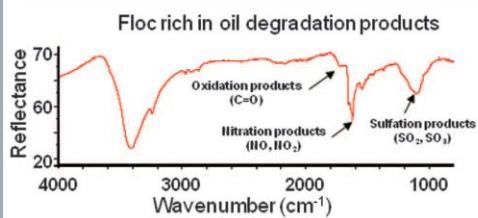
Yellow – oil, blue – EPS, with associated microbial cells

MICROBIAL OIL FLOCS

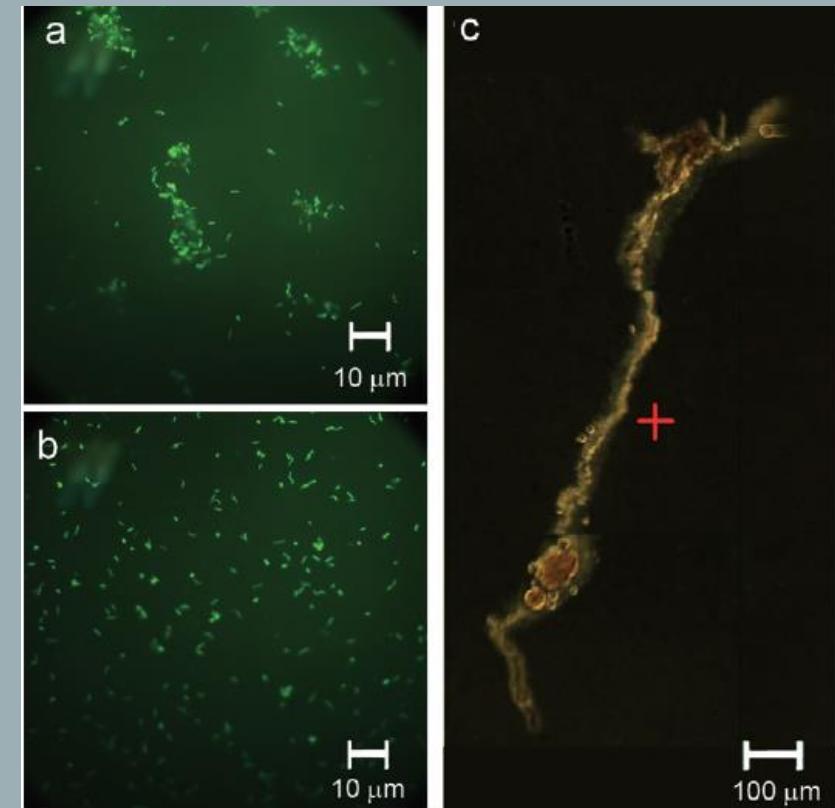
‘Cellular flocs’ – bacteria, EPS, oil, and oil degradation products (non-sinking flocs)

Flocs in deep plume water (Hazen et al., 2010)

SR-FTIR analysis of Sample # OV01105

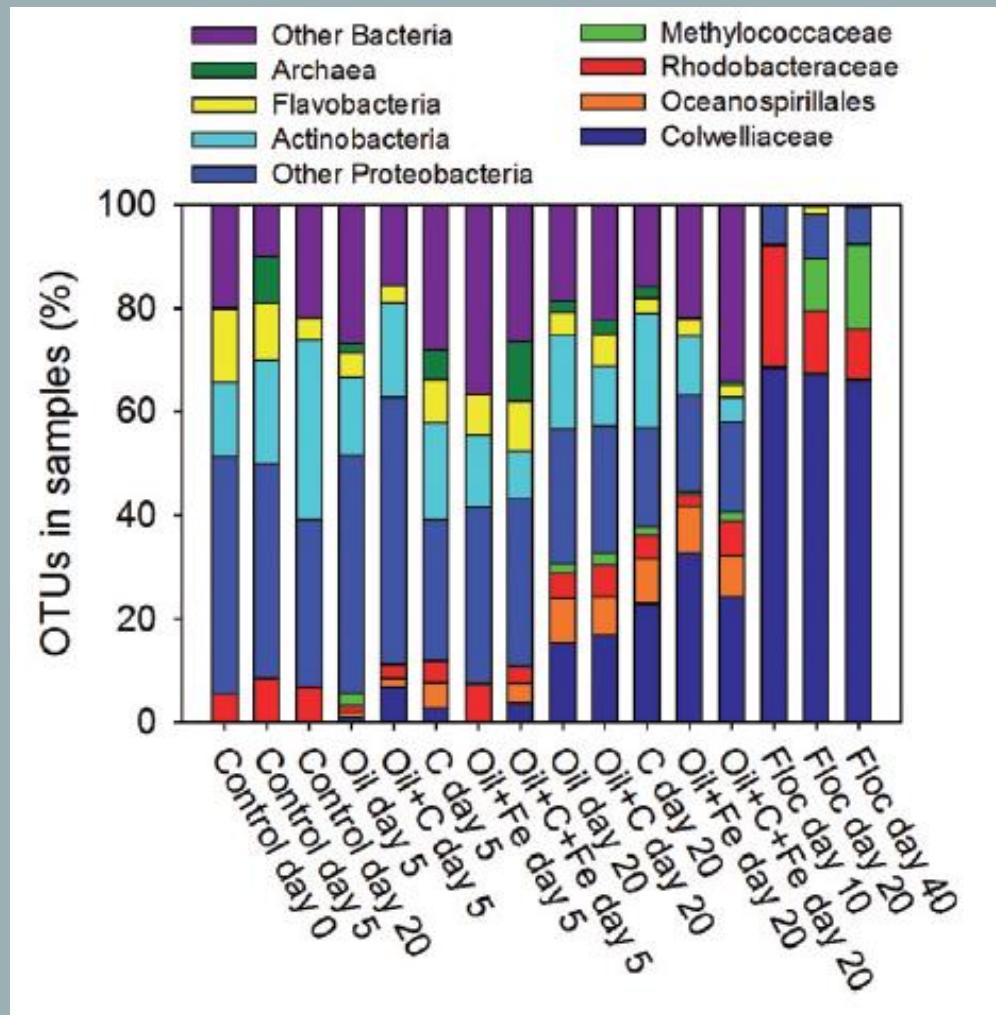


Floc formation *ex situ* (Bælum et al., 2012)

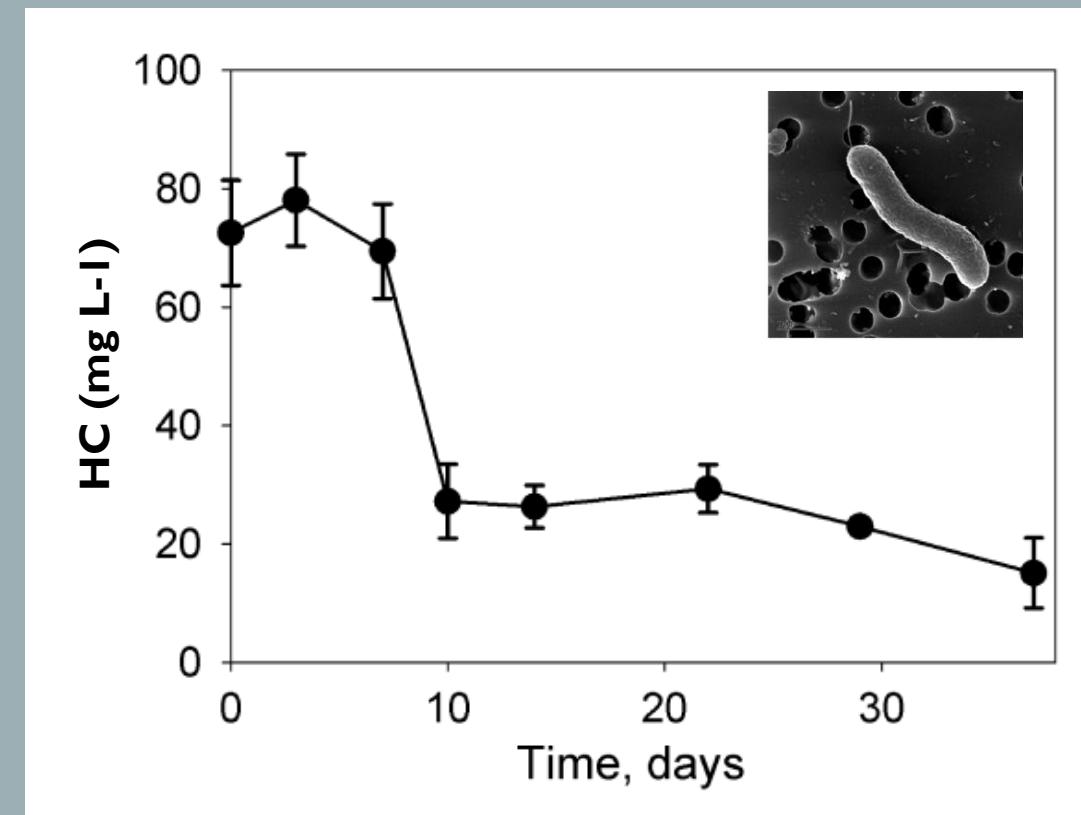


→ see also Doyle et al. (2018): Cells clumped around oil droplets (10 – 200 μm in diameter)

MICROBIAL OIL FLOCS DOMINATED BY COLWELLIA



HC degradation by *Colwellia* sp. isolated from floc



Bælum et al. (2012)

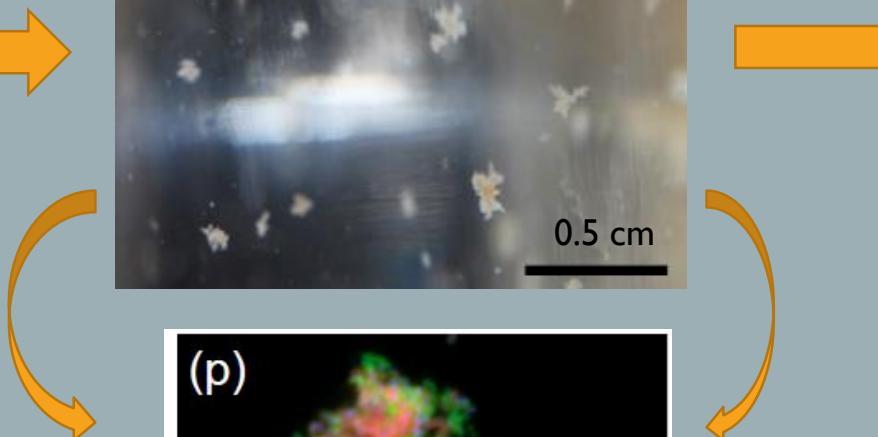
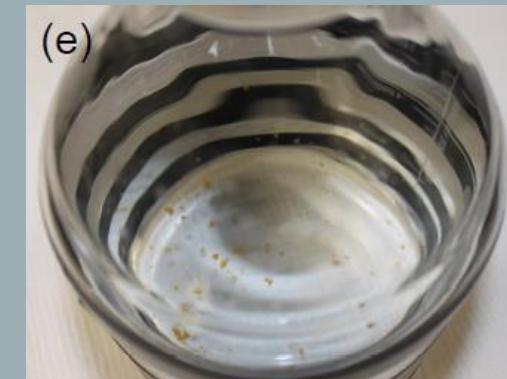
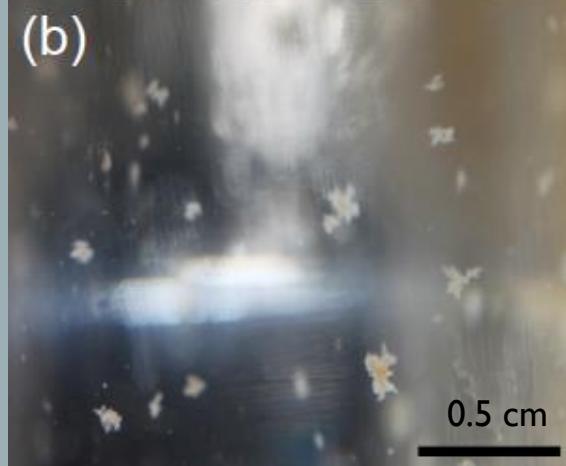
COLWELLIA IN MOS FORMED IN ROLLER BOTTLES

Enrichment of oil-degrading bacteria from oil seep (~1100m)

Stage I: EPS-oil matrix (15 d)



Stage II: EPS-oil + ballast (50 d)



Dual CARD-FISH hybridization of MOS:
Red - *Colwellia*; green – all Bacteria

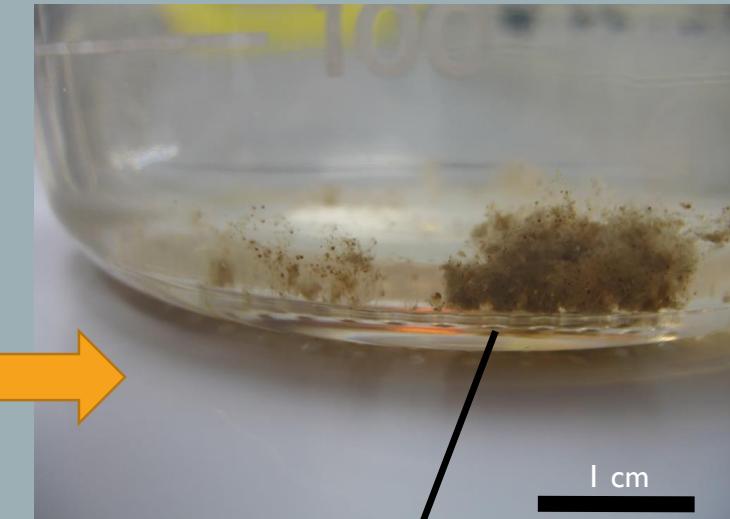
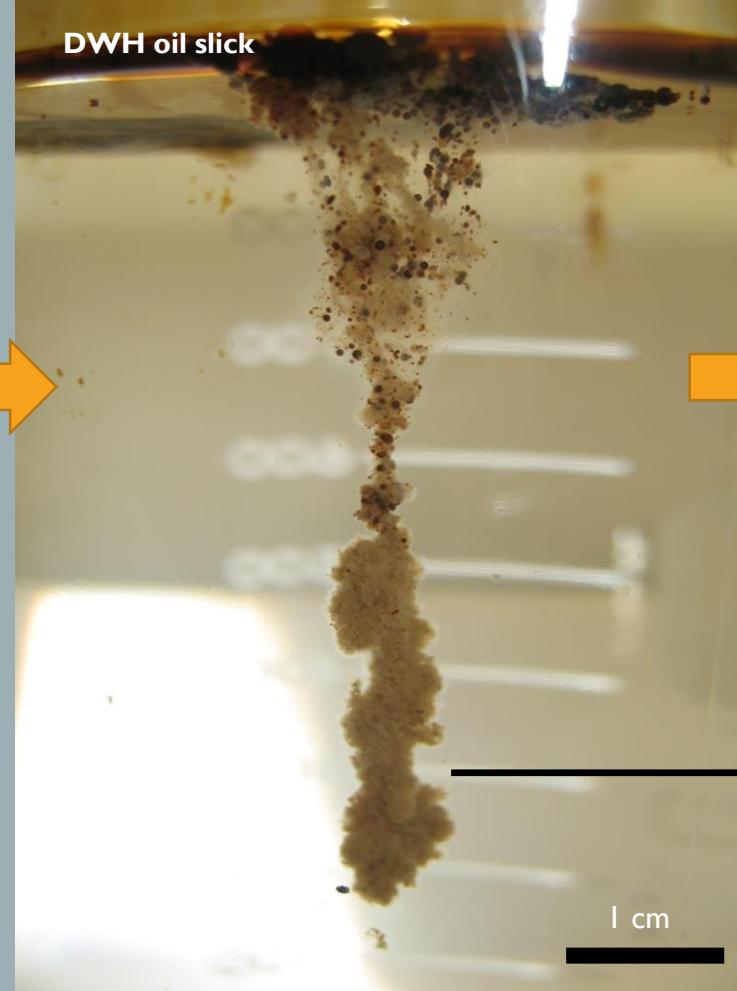
MICROBIAL MOS IN ROLLER BOTTLES

Enrichment of oil-degrading bacteria from oil slick (DWH, May 2010)

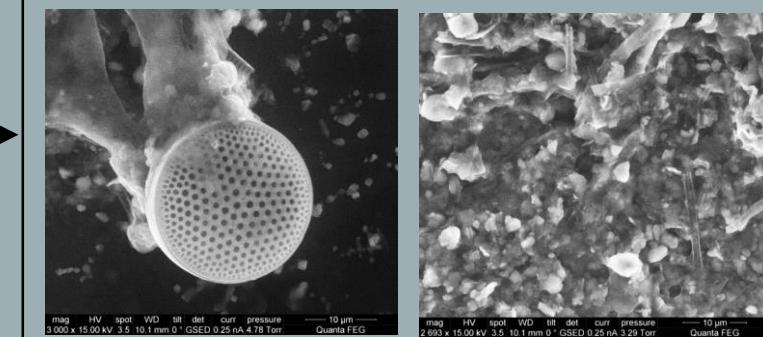
Stage 1: EPS-oil matrix (7 d)



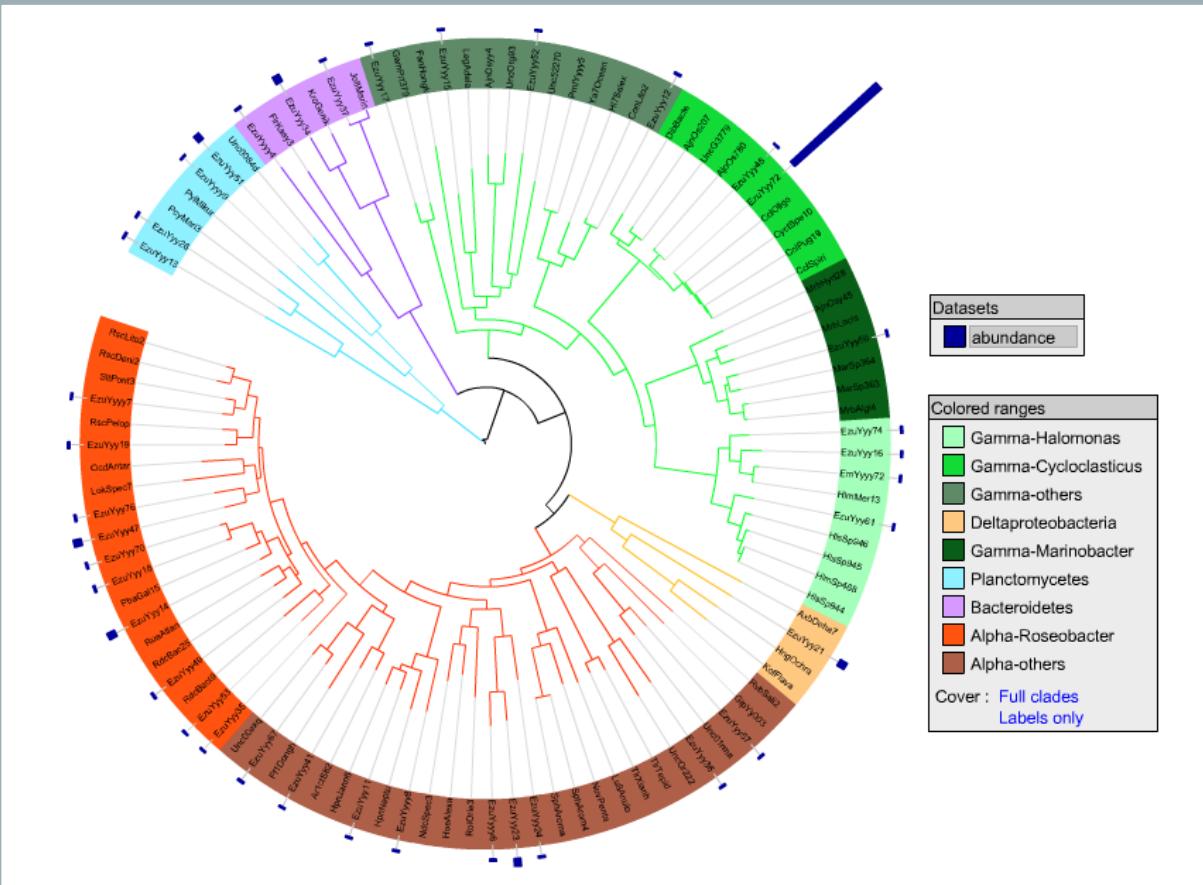
Stage 2: EPS-oil + ballast (21 d)



Ballast: Planktonic cells and minerals



MOS-ASSOCIATED MICROBIAL COMMUNITY: MORE DIVERSE COMPARED TO DEEP FLOCS/MOS



❖ Gammaproteobacteria *Cycloclasticus* – PAH-degraders

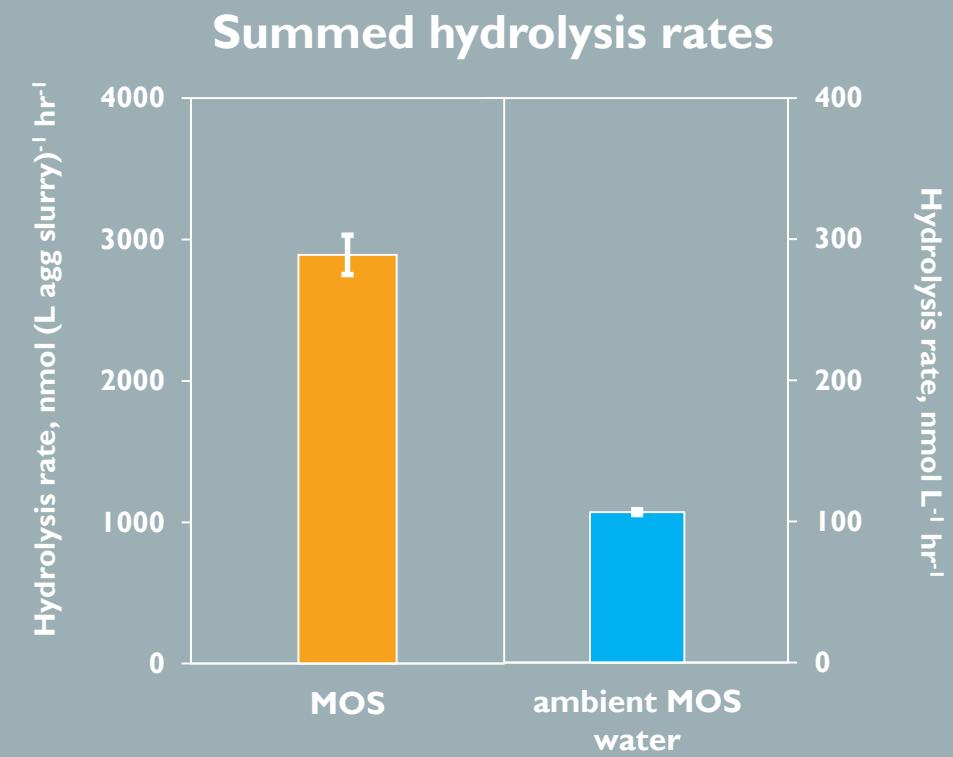
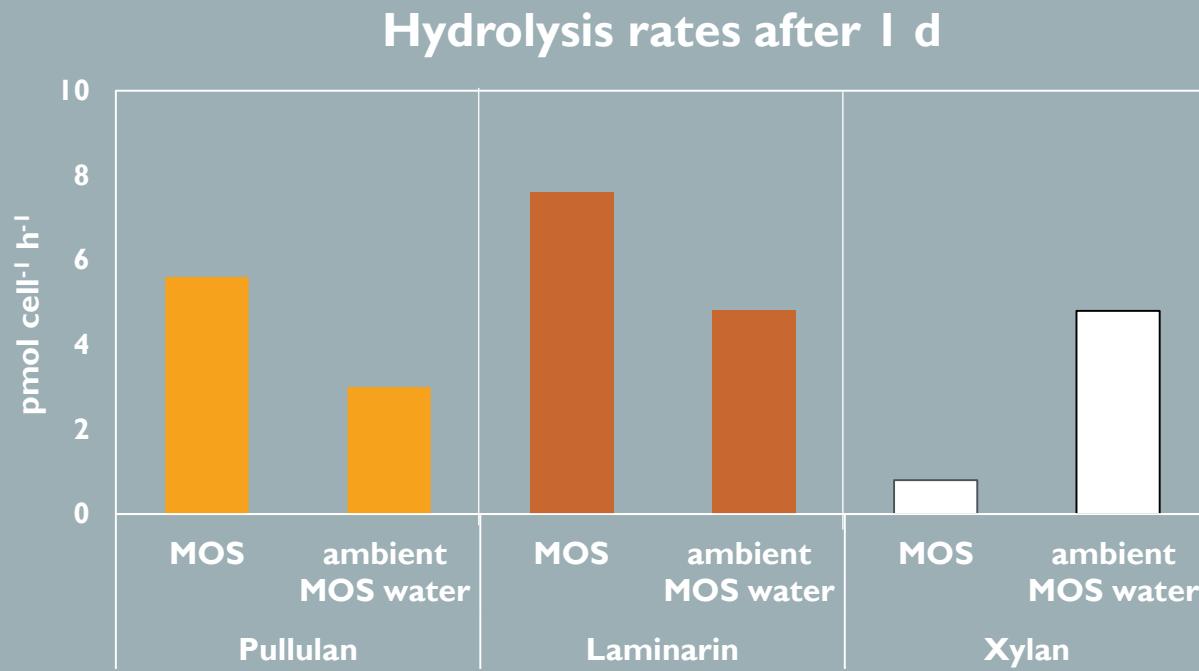
Marinobacter, Halomonas – Alkane-degraders and EPS-producers (see also Gutierrez et al., 2013)

❖ Alphaproteobacteria, Deltaproteobacteria, Bacteriodetes

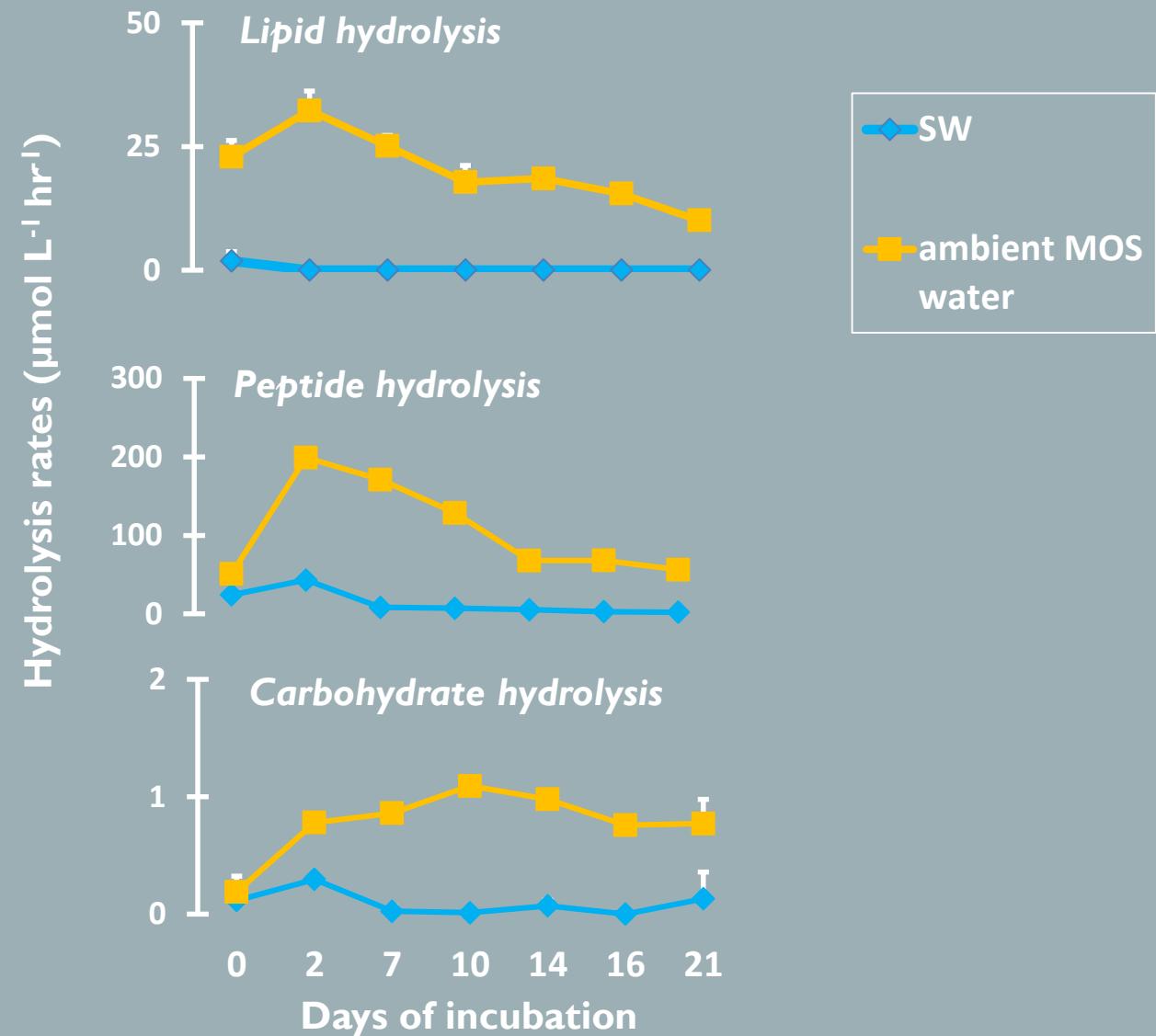
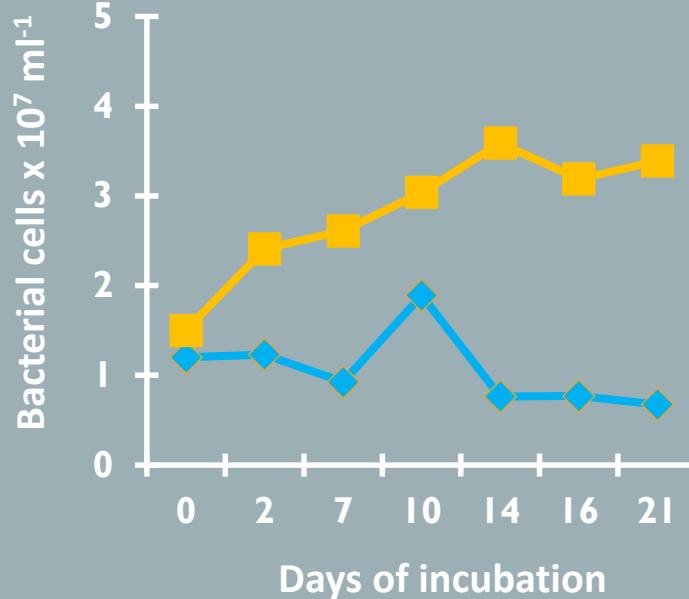
Broad metabolic capabilities → Secondary consumers (Head et al., 2006)

MICROBIAL ACTIVITIES IN MOS

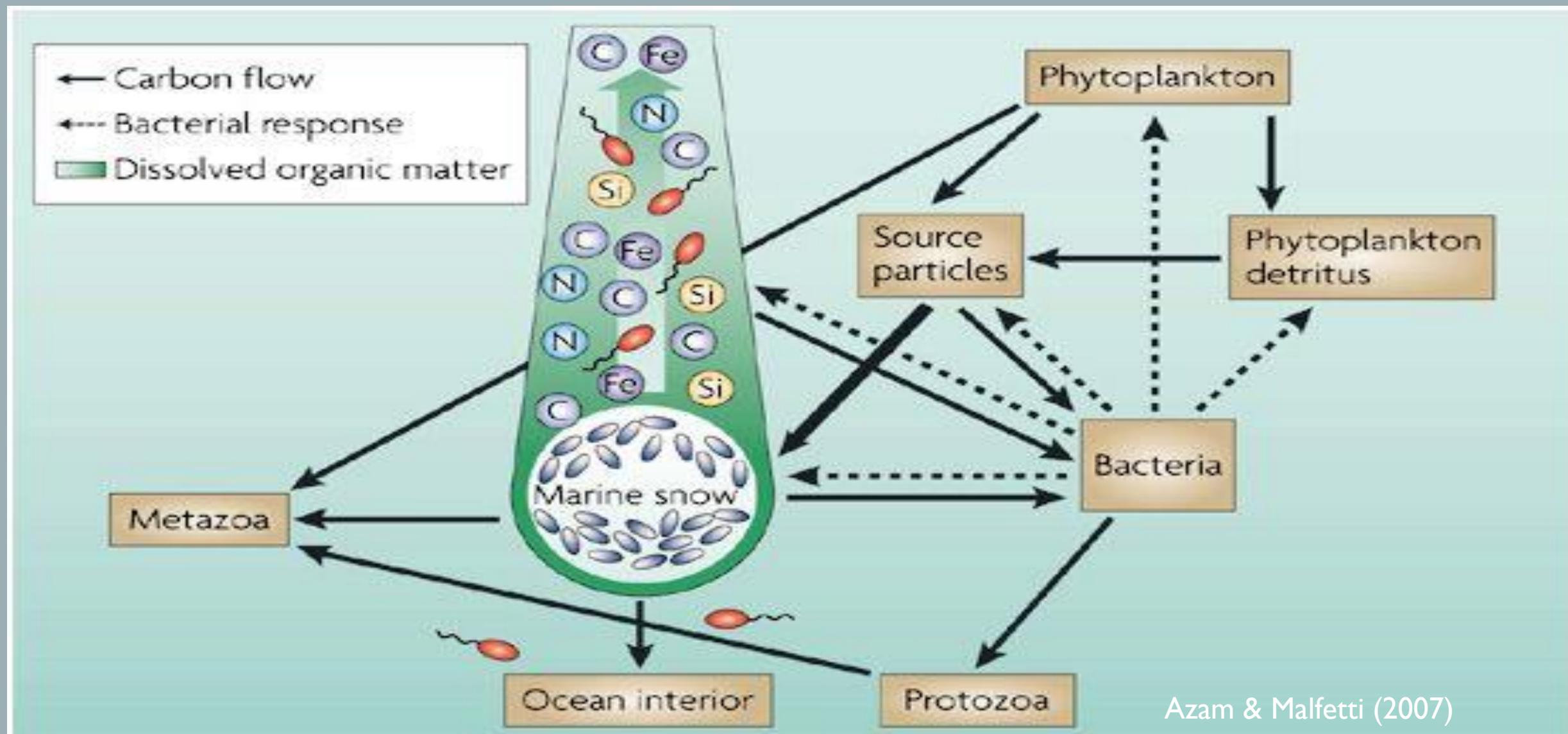
Polysaccharide degradation in MOS → Secondary consumers



MOS STIMULATES MICROBIAL ACTIVITIES IN AMBIENT WATER



MOS STIMULATES MICROBIAL ACTIVITIES IN AMBIENT WATER



Azam & Malfetti (2007)

SUMMARY

- ❖ Oil-degrading microbes trigger MOS formation

Step I: Formation of ‘cellular flocs’ (cells, EPS, oil, oil degradation products)

→ physiological mechanism for oil-degrading microbes (e.g. Bælum et al., 2012)?

Step II: MOS formation

→ ‘Cellular flocs’ grow and sink (e.g. Kleindienst et al., 2015)

→ Ballast material trapped in mucus matrix (e.g. Ziervogel et al., 2012)

- ❖ Microbial communities on MOS in deep water less diverse compared to surface water MOS

- ❖ Microbial activities on MOS (including Secondary consumers) affects elemental fluxes in the water column

WHAT ABOUT COREXIT????

- ❖ ‘Cellular flocs’ formed with and w/o COREXIT (Bælum et al., 2012; Doyle et al., 2018)
- ❖ Microbial MOS in deep waters with and w/o COREXIT and nutrients (Kleindienst et al., 2015)
- ❖ Microbial MOS in surface waters during spill ?COREXIT? (Ziervogel et al., 2012)
- ❖ Microbial MOS with COREXIT and nutrients only (Suja et al., 2017):

