Marine Oil Snow, a Microbial Perspective

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Under certain conditions, dispersed crude oil in the sea combines with organisms, organic matter, and minerals to form marine oil snow (MOS), thereby contributing to the sinking of oil to the seafloor. Marine microbes are the main players in MOS formation, particularly via the production of extracellular polymeric substances. Distinct groups of microbes also consume the majority of the hydrocarbons during descent, leading to enrichment of the less bioavailable hydrocarbons and asphaltenes in the residue. Here we discuss the dynamics of microbial communities in MOS together with their impacts on MOS evolution. We explore the effects of dispersant application on MOS formation, and consider ways in which laboratory experiments investigating MOS formation can be more representative of the situation in the marine environment, which in turn will improve our understanding of the contribution of MOS to the fate of spilled oil.

Keywords: marine oil snow, marine snow, hydrocarbon biodegradation, hydrocarbonoclastic bacteria, extracellular polymeric substances, oil-spill response

INTRODUCTION AND SCOPE

In order to respond effectively to crude oil released to the marine environment it is imperative to understand the interplay between the physical, chemical, and biological factors that determine petroleum fate, transport, and impact on the biota and surrounding environment. Marine snow (MS) came to prominence as a vehicle for transporting oil to the benthos during the 2010 Macondo spill (hereafter referred to as the MC-252 spill, because the Macondo Prospect is located in Mississippi Canyon Block 252 of the Gulf of Mexico), and has stimulated a lot of research to determine its quantitative importance, resulting in several reviews on the topic (e.g., Daly et al., 2016; Passow and Ziervogel, 2016; Quigg et al., 2016, 2020; Decho and Gutierrez, 2017; Brakstad et al., 2018b; Burd et al., 2020; Kujawinski et al., 2020; Passow and Overton, 2021).

Here we consider the microbial communities associated with marine oil snow (MOS), including producers and degraders of extracellular polymeric substances (EPS). We discuss the evidence for biodegradation of different hydrocarbon components during the descent of MOS, the impacts of MOS on the benthos, and the potential influence of oil on the formation of MOS and on its associated microbiome. We discuss the efficacy of the methods used to investigate aspects of MOS, and conclude by suggesting ways to advance research into the formation and fate of MOS.
MARINE SNOW (MS) FORMATION

The ocean contains ~600 gigatonnes of carbon in the form of dissolved organic matter (DOM; Hansell et al., 2009), the vast majority in the nanometre size range (Benner and Amon, 2015). Around 10% of this DOM can self-assemble to form marine microgels, three-dimensional polymer networks transiently suspended in seawater, which can aggregate into particulate organic matter (POM; Chin et al., 1998; Verdugo, 2012; Orellana and Leck, 2015; Figure 1). This transition from DOM to POM plays a critical role in marine snow (MS) formation (Passow et al., 2012). MS is defined as aggregates ≥500 µm in diameter that descend from the euphotic layer of the epipelagic zone (0–200 m depth) into the mesopelagic zone and beyond, ultimately reaching the sea floor (Turner, 2015; Figure 1). MS is made up of various organic and inorganic particles, eukaryotic microalgae, and Bacteria, along with Archaea, microzooplankton, fungi, viruses, fecal material, detritus, and feeding structures, all bound together by biopolymers (Shanks and Trent, 1980; Allredge and Silver, 1988; Allredge et al., 1993; Simon et al., 2002; Volkman and Tanoue, 2002; Bochdansky et al., 2017). Bacteria are abundant in MS (Mestre et al., 2018), where their large surface-area-to-volume ratio makes them particularly effective in transporting and transforming organic matter and cycling nutrients (Azam and Long, 2001). Analysis of large, slowly sinking MS particles in the bathypelagic zone showed that, while bacteria were extremely numerically dominant, two distinct eukaryotic groups, the fungi and the labyrinthulomycetes, together contributed about twice the bacterial biomass on the MS (Bochdansky et al., 2017). Marine snow transports microbes from the surface (Mestre et al., 2018), provides an important food source for organisms living in the deep water column, and drives the organic-carbon pump that exports carbon and energy to the aphotic zone, sequestering atmospheric CO2 into the deep ocean and sediment (DeVries et al., 2012). MS sinking velocities range from 1 to 368 m/day (median 95 m/day) (Allredge and Silver, 1988), dependent on particle size, density, and porosity (De La Rocha and Passow, 2007; Brakstad et al., 2018b), and can change while sinking.

Extracellular polymeric substances (EPS) are a major component of the total DOM pool in the ocean. They consist primarily of polysaccharides (up to 90%), but also proteins, glycoproteins, nucleic acids, and lipids, and are secreted or released by phytoplankton and heterotrophic microbes (Mühlenerbruch et al., 2018). Polysaccharide-rich EPS are transparent, but can be visualized with Alcian Blue (Allredge et al., 1993), and hereafter, this form of EPS will be referred to by its widely used descriptor of “transparent exopolymer particles” (TEP; Allredge et al., 1993; Passow and Allredge, 1995). Protein-rich EPS (which are also transparent in nature but not in name), can be distinguished by staining with Coomassie Blue (Long and Azam, 1996). TEP are “sticky,” largely due to acidic sugar units, and so they are often considered to be most significant in the formation of MS (Daly et al., 2016).

Thus, EPS can enhance the aggregation of particles, enabling the formation of MS (Quigg et al., 2016; Decho and Gutierrez, 2017), although not all EPS assemblies to form these discrete particles (Passow, 2002a). Firstly, EPS nanofibers (∼5–50 nm) assemble, forming nanogels (100–200 nm) that are stabilized by entanglement and, due to their polyanionic nature, cross-linking with divalent Ca2+ (Verdugo, 2012; Wakeham and Lee, 2019). These DOM polymer networks diffuse into the bulk seawater and collide with neighboring nanogels, forming microgels (∼3–6 µm) in a reversible process (Verdugo, 2012). Microgels are ubiquitous throughout the oceans, generally decreasing in abundance below the photic zone (Engel et al., 2020), and continue to collide and aggregate, eventually forming stable macrogels of several hundred micrometers, operationally defined as particles retained on 0.4 µm polycarbonate filters (though some researchers use 0.2 µm filters).

Aggregate formation depends on the number of particle collisions, and is a function of the abundance of particles, including cells, present in the medium, and their cohesiveness, which enhances the probability that such particles will stay attached after collision (Jackson, 1990; Burd and Jackson, 2009; Cruz and Neuer, 2019). TEP aggregate with other particles and act as a biological glue to form sinking MS (Engel et al., 2004; Mari et al., 2017). Several studies have established a link between increases in TEP concentration and the enhanced aggregation and sinking of MS (Passow et al., 1994, 2001; Logan et al., 1995; Gärdes et al., 2011). By contrast, Coomassie-blue stainable particles are not considered to be as important as TEP in MS formation (Prieto et al., 2002; Cisternas-Novoa et al., 2015). However, protein-rich EPS, which are as common as TEP in the ocean (Busch et al., 2017), consist of both hydrophilic and hydrophobic domains, which result in surface processes that are less predictable and more dependent on the context (e.g., types of associated minerals, presence of hydrocarbons), and may be important for MS formation (Santschi et al., 2020). Actually, the world of hydrated gels is less binary than portrayed here, with the existence of hybrid and intermediate forms, and understanding the varied nature of these gels and their roles in, inter alia, the microbial loop and marine snow formation remains work in progress (Thornton, 2018).

In addition to these biochemical processes, hydrodynamics influence the formation and transport of MS and marine oil snow (MOS), indirectly (e.g., by transport of limiting nutrients to microbiota), and directly (e.g., by turbulence-enhanced aggregation of particles due to increased collision rate or, when excessive, disaggregation of MS) (Burd and Jackson, 2009; Daly et al., 2016). Currents, eddies, subduction, upwelling, river discharge, benthic resuspension, and other physical processes can all alter the formation and distribution of MS/MOS (see Daly et al., 2016, 2020). Also, density gradients in the water column can decrease the settling rate of MS/MOS particles (Prairie et al., 2015).

Abbreviations: CEWAF, Chemically Enhanced Water Accommodated Fraction; DOM, Dissolved Organic Matter; EPS, Extracellular Polymeric Substances; MC-252, Mississippi Canyon Block 252; MOS, Marine Oil Snow; MS, Marine Snow; OHCB, Obligate Hydrocarbonoclastic Bacteria; POM, Particulate Organic Matter; PAH, Polycyclic Aromatic Hydrocarbons; TEP, Transparent Exopolymer Particles; WAF, Water Accommodated Fraction.
**MARINE OIL SNOW (MOS) FORMATION AND ITS CONTRIBUTION TO TRANSPORTING WEATHERED CRUDE OIL TO DEEP-SEA SEDIMENTS**

**Composition of MOS and Potential Impact on the Benthos and Associated Biota**

Crude oil spilled on the sea is rapidly weathered by processes such as spreading, evaporation, natural dispersion, emulsification, and photooxidation. The most soluble hydrocarbons, such as benzene and toluene dissolve into the underlying water, and hydrocarbons with up to 15 carbon atoms evaporate in the first day or so. Floating oil undergoes photooxidation and emulsification, and any wave action will disperse some of the oil into the water column. Oil-spill dispersants lower the interfacial tension between oil and water, and thereby allow dispersion with much less wave energy, and the generation of much smaller droplets. Crude oil spilled from a deep-sea well blowout will rise to the surface, losing most of its soluble components during the ascent (Ryerson et al., 2012), and at least partially dispersing as small droplets. Again, this process can be stimulated with dispersants (Gros et al., 2017).

Dispersed oil droplets are subject to rapid biodegradation, and can also interact with minerals and organic matter. In coastal areas oil-mineral aggregates are common (Owens, 1999; Loh and Yim, 2016; Li et al., 2020) and result in sinking or suspension in the water column depending on the nature of the aggregate (Quigg et al., 2020). Suspended sediment particles from the Mississippi River outflow generated oil mineral aggregates during the MC-252 spill (Figure 2), although their quantitative influence is not clear (Daly et al., 2016). A substantial settling of MS during the early days of the MC-252 spill led to the suggestion that MS could also transport oil to the sediment (Passow et al., 2012; Passow, 2016; Passow and Ziervogel, 2016; Larson et al., 2018). There are many examples from mesocosm studies of the formation of oil-algal-bacterial complexes (e.g., Lee et al., 1985; Macnaughton et al., 2003; Coulon et al., 2012). In the open sea the resultant oil-containing flocs have been called marine oil snow (MOS) or oil-related marine snow (Brakstad et al., 2018b) among other names (Quigg et al., 2020), shown schematically in Figure 3. From analysis of MOS captured in sediment traps (Stout and German, 2018) it has been estimated that the extensively biodegraded residue of 217,700 to 229,900 barrels of oil was deposited in the form of MOS over an area of at least 7,600 km² following the MC-252 spill, representing around 7% of the estimated 3.19 million barrels of spilled oil (Barbier, 2015). MOS particles consist of a mixture of organic matter (e.g., phytoplankton, heterotrophic microbes, fecal material, dead, or decaying material), sediment particles and weathered crude oil.
FIGURE 2 | A mind-map showing the factors that may influence the likelihood of a marine oil snow sedimentation and flocculent accumulation (MOSSFA) event, (A) in the event of freshwater influence from rivers, and (B) and (C) after common oil-spill-response strategies, such as (B) in-situ burning, and (C) dispersant application. Abbreviations include: dissolved organic matter (DOM), extracellular polymeric substances (EPS), oil-mineral aggregates (OMA), protein/carbohydrate ratio (P/C ratio) and transparent exopolymer particles (TEP). Up-arrows denote an increase of the process and down-arrows denote a decrease.
within a matrix of EPS and especially TEP (Daly et al., 2016; Passow and Ziervogel, 2016; Brakstad et al., 2018b). Thus, TEP can be considered as the glue, while the cells and other particles act as ballast, providing the density to sink the less dense oil and TEP. In addition, the presence of oil may lead to a tighter packaging of microbial cells causing MOS particles to sink faster than MS (Passow et al., 2019). However, as with MS, MOS particles are hot-spots of microbial activity (Ziervogel et al., 2012), and constantly changing as colonization, consumption, clumping, and fragmentation alter the rate of their descent. The sinking of these MOS particles and their accumulation on the seafloor of the Gulf of Mexico was referred to as a marine oil snow sedimentation and flocculent accumulation event (MOSSF&A; Valentine et al., 2014; Brooks et al., 2015; Chanton et al., 2015; Romero et al., 2015, 2017; Yan et al., 2016; Schwing et al., 2017). Oil is also transported to deep-water sediments by other mechanisms, such as incorporation into invertebrate fecal pellets (Lee et al., 2012; Almeda et al., 2016), direct sinking of very degraded oil (Murray et al., 2020), sinking of degraded oil attached to mineral particles (Li et al., 2020), settling of oil-mud complexes formed from the injection of drilling fluids (Stout and Payne, 2017), and sinking of heavy residue from burning oil (Stout and German, 2018). For example, in-situ burning removed 220,000–310,000 barrels of surfaced oil following the MC-252 spill, generating some 33,800–54,700 barrels of heavy viscous residue that sank to the seafloor (Stout and Payne, 2016b). Also, 4% of the burned oil was emitted as soot particles (Middlebrook et al., 2012), which may have contributed to MOS formation due to aerosol deposition and adsorption of DOM on the surfaces of soot particles, thereby stimulating aggregation (Mari et al., 2014; Figure 2).

Oil and sediment deposition following the MC-252 spill impacted the benthic macrofauna and meiofauna (including foraminifera whose carbonate shells provide a temporal sedimentary record), reducing their abundance and biodiversity (Montagna et al., 2013; Baguley et al., 2015; Schwing et al., 2015, 2017, 2018; Washburn et al., 2017). The main drivers of these changes have been assigned to the introduction of toxic oil compounds (van Eenennaam et al., 2018, 2019), an increase in anoxic conditions due to aerobic consumption of the added organic carbon (Hastings et al., 2016, 2020), or a combination of both (Schwing et al., 2015). However, the composition of the oil assumed in these predictions did not take into account the biodegradation that occurs during descent through the water column prior to deposition on the seafloor (Stout and Payne, 2016a; Bagby et al., 2017; Supplementary Table 1). For example, van Eenennaam et al. (2018, 2019) applied 3 or 10 g per m² of fresh Macondo surrogate oil in their mesocosm experiments. The use of fresh oil is unrepresentative of what was actually deposited on the seafloor, which was a wax-rich, severely biodegraded residue lacking the more bioavailable and toxic components of the oil (Stout and Payne, 2016a). Any future toxicity studies should use weathered oil in order to fully assess the impact of MOS on the benthos.

MOS may have formed after other major spills. For example, MOS is thought to have formed during the 1979 Ixtoc I blowout in the Bay of Campeche based on the presence of river-derived POM (Vonk et al., 2015). Two lines of evidence support a significant depositional pulse of oil following the spill: (1) petroleum biomarkers and molecular compound classes indicative of the Ixtoc I oil were detected in seafloor sediments in the Bay of Campeche from deep (>500 m) waters (Lincoln et al., 2020); and (2) similar laminations and visible redox boundaries in sediment cores to those seen following the MC-252 spill (Brooks et al., 2015; Schwing et al., 2020). However, it cannot be determined what proportion of oil was deposited due to MOS sedimentation compared to the alternative mechanisms discussed above.

**Mechanisms of MOS Formation**

To our knowledge, controlled field studies to investigate MOS formation and fate soon after an oil spill have yet to be performed as it would require researchers to be present (and divert resources) during an emergency incident. MOS formation and fate are also difficult to investigate experimentally at sea due to stringent regulations on controlled oil releases. In addition, monitoring MOS formation would be technically challenging as this needs an in-situ camera system that follows the aggregation process (Iversen and Lampitt, 2020). As an alternative, MOS experiments have been performed in microcosms with the aim of replicating in-situ conditions, often comparing MS/MOS formation in uncontaminated and crude-oil-amended seawater. Most groups use closed bottles on their sides on roller tables, where the changing direction of gravity keeps any flocs in suspension. If the bottles are full there is minimal turbulence, and if there is an air space the rotation simulates surface turbulence under mild sea conditions. The aggregates formed are generally morphologically and chemically similar to MS particles collected in situ (Unanue et al., 1998; Ziervogel et al., 2012), and rotating bottle experiments have been widely adopted to investigate MOS formation (e.g., Passow et al., 2012; Ziervogel et al., 2012). However, aggregates formed in full bottles may grow larger than those that descend through marine water layers where there is varying turbulence.

MOS may develop as: (1) pre-formed MS interacting with oil, or (2) oil acting as a nucleus for microbial biofilm growth and floc formation. The latter is referred to as a bacteria-oil aggregate (BOA; Passow and Overton, 2021). These mechanisms are not mutually exclusive, and given the temporal and spatial extent of the MC-252 spill, both mechanisms are highly likely to have been responsible for MOS formation. Mechanism 1 was probably enhanced by river outflows (e.g., diversionary channels of the Mississippi River) providing nutrients for microbial growth, as well as directly supplying microbes, clay and other particulate matter to the water column (Figure 2; Vonk et al., 2015; Daly et al., 2016; van Eenennaam et al., 2016). In addition, the transient acute toxicity of dispersed oil immediately upon dispersion may have negatively impacted grazing zooplankton, diminishing their effect on phytoplankton populations (Almeda et al., 2013a,b). Thus, the stimulated phytoplankton blooms (Hu et al., 2011; Bianchi et al., 2014), with their high levels of EPS/TEP, coupled with greater abundance of particles, increased the probability of collisions and aggregation that lead to MOS formation (Jackson, 1990, 2005; Burd and Jackson, 2009).

Furthermore, the unusual features of the MC-252 spill, i.e., its occurrence in deep water and extensive subsurface dispersant
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FIGURE 3 | Schematic representation of microbes and dispersed oil droplets in marine snow at different scales.

The microfluidic system developed in the laboratory of Jian Sheng (White et al., 2019) shows the colonization of an oil droplet by *Pseudomonas* sp. ATCC 27259 (strain P62) in real time. This coating of EPS dramatically increased drag, and would thus reduce the rising velocity of oil droplets in the water column. With further studies, applying natural communities and lower nutrient concentrations, this experimental set-up should help to explain and model various aspects of rising plumes, as well as the environmental factors influencing biodegradation, as well as dissolution, increases. It is important to understand how rapidly hydrocarbon abundance and composition change during transit and sedimentation, as well as the environmental factors influencing biodegradation, and thus the likelihood of harm to the benthic biota from MOS. Hazen et al. (2010) reported a half-life of alkanes in dispersed oil following the MC-252 spill of a few days, and several papers have confirmed this result under laboratory conditions that mimic environmentally-relevant concentrations (Prince et al., 2013, 2017; Brakstad et al., 2015a; Wang et al., 2016). A key to determining these rates is the long-term degradation of the oil, and as such it is also feasible that they reached the sediment of their own accord after extensive biodegradation (Murray et al., 2020), or by a combination of both means (Suja et al., 2017; Quigg et al., 2020).

It is noteworthy that droplets of dispersed oil persisted in the environment at low concentrations long after the MC-252 well was capped. Passow and Stout (2020) analyzed MOS captured in a sediment trap 6 weeks to 13 months later, and found low levels of very degraded oil in MOS which they believe originated in the subsurface “plume” and had never reached the surface. Four pulses of MOS were captured, demonstrating that periodic “blooms” of MS, with varying compositions of diatoms, fecal pellets, etc., occur regardless of major oil intrusion. Oil was highly degraded, and concentrations were low (the residue of 1.45 mg fresh oil/m²/d, median percentage degradation of 85%).

Changes in Hydrocarbon Composition During Sinking of MOS

In assessing the impact of MOS on the benthos it has sometimes been assumed, erroneously, that the concentration and composition of crude-oil entrained in marine snow near the surface will be the same when it reaches the seafloor. During the downward transport of MOS, weathering of the oil by evaporation and photooxidation stops, and the relative contribution of biodegradation, as well as dissolution, increases. It is important to understand how rapidly hydrocarbon abundance and composition change during transit and sedimentation, as well as the environmental factors influencing biodegradation, and the relative contribution of biodegradation, as well as dissolution, increases. It is important to understand how rapidly hydrocarbon abundance and composition change during transit and sedimentation, as well as the environmental factors influencing biodegradation, and the relative contribution of biodegradation, as well as dissolution, increases.
pentacyclic saturated hydrocarbon is known to be consumed under some conditions (Frontera-Suau et al., 2002). Therefore, estimates based on hopane’s stability may underestimate the extent of biodegradation. Alternative biomarkers can be used, such as C28 20S-triaromatic steroid (Douglas et al., 2012), or even long chain n-alkanes (Bagby et al., 2017), although it must be borne in mind that especially the latter may be biogenic (Dehmer, 1993). Oil residues in deep-sea sediments were found to be very biodegraded, with most of the resolvable alkanes and PAHs degraded >90–95%. The residues did contain measurable amounts of n-alkanes longer than about C30 (Stout and Payne, 2016a; Bagby et al., 2017; Stout and German, 2018). These studies agree that the extent of biodegradation increased with distance from the well, suggesting that the biodegradation occurred while the oil was in transit. Nevertheless, some biodegradation may also have occurred once the oil was in the sediment. The concentrations of oil were sufficiently low that there was likely enough bioavailable N and P so that biodegradation was not severely nutrient limited. The foregoing evidence highlights that biodegradation severely alters the oil composition of residues deposited on the seafloor, and so, to fully understand MOS’ impact on the benthos, we must consider the influence of oil degradation within MOS prior to deposition.

**Microbial Production and Degradation of Transparent Exopolymer Particles (TEP) and Their Impacts on MOS**

It is well-known that diverse eukaryotic phytoplankton produce copious amounts of EPS that aggregate to form TEP, with the quantity produced dependent on the species and environmental conditions (Passow, 2002a,b; Passow et al., 2019). Diatoms, dinoflagellates, and prymnesiophytes are perhaps the most relevant taxa in the context of MOS, as they tend to be the most abundant blooming microalgae in coastal areas where spills are most likely to occur. Many diatoms produce silica and some prymnesiophytes (e.g., coccolithophores) produce calcium carbonate, which cause cells to aggregate and ballast marine snow, causing it to sink (Tyrrell and Young, 2009; Lombard et al., 2013; Quigg et al., 2016). Cyanobacteria also contribute TEP, with naturally forming flocs of *Trichodesmium* trapping degrading oil in their mucus (Passow et al., 2012). The abundant *Prochlorococcus* produces TEP (Iuculano et al., 2017). Some associated heterotrophic bacteria can enhance cellular aggregation (Cruz and Neuer, 2019), but other phytoplankton blooms are accompanied by bacteria that consume TEP, such as Flavobacteria, Verrucomicrobia, and Rhodobacteriales species, as well as those feeding on other algal products, especially Gammaproteobacteria, such as Alteromonadaceae (Buchan et al., 2014; Taylor et al., 2014; Arnosti et al., 2016; Needham and Fuhrman, 2016; Bohórquez et al., 2017; Decho and Gutierrez, 2017).

Heterotrophs produce their own EPS, including polysaccharides that are often rich in uronic acids and protein-rich EPS, and the protein-to-carbohydrate ratio of bacterial EPS varies from 0.04 to 20 (see Santschi et al., 2020). Moreover, heterotrophs can influence the EPS quantity/type synthesized by proximal phytoplankton (see Quigg et al., 2020). The EPS produced by phototrophs can be used as a source of carbon and energy by heterotrophs (Abed, 2019), which release and return nutrients (e.g., N, P, and trace metals) from the EPS to the phototroph in a somewhat mutualistic system (Christie-Oleza et al., 2017). Importantly, both MS and MOS formation appears to be enhanced by these bacterial-phytoplankton interactions (Fu et al., 2014; van Eenennaam et al., 2016; Cruz and Neuer, 2019). For example, van Eenennaam et al. (2016) showed that more EPS was released by bacteria associated with phytoplankton (*Dunalieilla tertiolecta* and *Phaeocystis tricornutum*) compared to bacteria-free phytoplankton (which did not produce any EPS) or the phytoplankton-associated bacteria grown on their own (up to 2-fold less). In addition to a greater quantity of EPS being produced due to these interactions, the chemical composition was also different. The EPS produced by the bacteria associated with phytoplankton had a higher protein-to-carbohydrate ratio, which may alter how EPS aggregates and interacts with oil (Bacosa et al., 2018; Santschi et al., 2020), e.g., by influencing: (1) hydrophobicity (Xu et al., 2011); (2) surface tension (Schwehr et al., 2018); (3) light-induced chemical crosslinking (Sun et al., 2017, 2018, 2019); and, overall, affecting MS/MOS sinking rates (Xu et al., 2018a,b).

Therefore, when we talk about TEP derived from a phytoplankton bloom, the TEP will actually come from an array of microbial species—phototrophs (e.g., phytoplankton) and heterotrophs (e.g., bacteria)—plus bacteria from the gut contents of grazing microzooplankton (Turner, 2015), some of which may include ingested oil (Almeda et al., 2016). Moreover, the rapid turnover of phytoplankton and associated heterotrophic Bacteria and Archaea during blooms (Needham and Fuhrman, 2016) will add to the structural variety of TEP. Different EPS/TEP-degrading microbes will colonize sinking MS/MOS flocs (Busch et al., 2017; Duret et al., 2019), which are additionally grazed by animals (Giering et al., 2014; Figure 1), leading to structural, chemical, and biological succession during descent, and potentially altering sinking velocity (Turner, 2015; Dang and Lovell, 2016). Clearly, the presence of weathered crude oil as a substrate in the MS adds an extra microbial dimension.

**WHICH HYDROCARBON-DEGRADING MICROORGANISMS ARE ASSOCIATED WITH MOS?**

**General Considerations of Microbial Community Analysis of MOS**

The analytical work discussed above has demonstrated that the oil residues from the MC-252 well blowout in the sediments of the Gulf of Mexico are highly degraded, and at least some arrived in association with marine snow. Here we discuss the microbial communities associated with MOS, followed by an appraisal of the main taxa found, and their potential roles in MOS formation and oil degradation. The capacity to degrade hydrocarbons is found in more than 300 genera of Bacteria (Prince et al., 2018) as well as in numerous fungal taxa (Prince, 2018) and in some Archaea (Oren, 2017; Prince et al.,...
2018). However, most investigations into MOS have focused on bacteria, as, after viruses, they are the most abundant microbes in the surface ocean, are often associated with phytoplankton and typically dominate hydrocarbon degradation in the marine environment (McGenity et al., 2012; Gutierrez, 2019; Yakimov et al., 2019). We note that most of the experiments focusing on MOS have used concentrations of dispersed oil and dispersant that are substantially (sometimes >100-fold) higher than found in the marine water column after a spill (Bejarano et al., 2014; Wade et al., 2016), which may have resulted in toxic responses from some organisms as well as unnaturally high activity from others.

Some studies assessing hydrocarbon degradation have assigned this function to particular taxa by testing isolates, applying DNA-stable isotope probing (DNA-SIP) with $^{13}$C-labeled hydrocarbons (Gutierrez et al., 2013b), or drawing inferences from metatranscriptomics, metagenome-assembled genomes and single-cell amplified genomes (Mason et al., 2012), but the majority of these studies have been on bulk water samples rather than specifically focusing on MOS. In many cases the identification of a taxon’s capacity for hydrocarbon degradation is based on an increase in relative abundance of their 16S rRNA gene sequences in oiled samples vs. non-oiled controls, as well as an understanding of taxon-trait relationships. Therefore, we briefly assess the degree to which phylogenetic data can be used to infer the function of particular microbial taxa. Some genera contain species that can degrade hydrocarbons and others that cannot (e.g., *Pseudomonas* and *Marinobacter*; Duran, 2010; Palleroni et al., 2010), and so an increase in abundance in oil samples may be a direct or indirect effect of the oil, requiring complementary techniques to understand their role. In this respect, Netzer et al. (2018) provide a relevant example in the microbial community succession in MOS formed in cold (4–5°C) seawater microcosms additionally containing the obligately psychrophilic diatom *Fragilariopsis cylindrus* and 30 ppm of chemically dispersed oil (30 ppm oil and 0.3 ppm Corexit 9500). Degradation of hydrocarbons was similar with and without diatoms, but *Nonlabens*, a genus within the Bacteroidetes, was dominant in the diatom-containing aggregates, and thus proposed to be a hydrocarbon degrader. However, *Nonlabens*, clearly co-habiting with the diatom culture, was also dominant on aggregates formed in diatom-containing microcosms without hydrocarbons, and so its role needs further investigation (Netzer et al., 2018). These bacterial communities differed significantly from those in diatom-free oiled samples that were dominated by *Oleispira*, a psychrophilic alkane degrader (Yakimov et al., 2003), and members of the family Sphingomonadaceae (Netzer et al., 2018). Whilst there is no evidence for hydrocarbon utilization by most of the seawater-derived strains of Sphingomonadaceae, marine sediments have yielded isolates from this family that degrade polycyclic aromatic hydrocarbons (PAHs) (Gilewicz et al., 1997; Johnson and Hill, 2003; Kertesz et al., 2019). Thus, in this example, other approaches (e.g., cultivation or DNA-SIP) are needed to confirm hydrocarbon degradation by the identified Sphingomonadaceae and *Nonlabens*. By contrast, we can have much more confidence that *Oleispira* was degrading alkanes as they belong to a group of marine bacteria that use hydrocarbons almost exclusively as a source of carbon and energy, known as obligate hydrocarbonoclastic bacteria (OHCB). These microbes are present in non-polluted marine environments in low numbers, but often bloom and dominate the microbial community following a spill (McKew et al., 2007a,b; Yakimov et al., 2007; Teramoto et al., 2009; Vila et al., 2010). Although some species designated as OHCB will use small organic acids as a source of carbon and energy, the word “obligate” is understood in the context that OHCB cannot compete with generalists in an environment lacking hydrocarbons. OHCB include genera with species that degrade aliphatic (*Alcanivorax*, *Oleibacter*, *Oleiphilus*, *Oleispira*, *Thalassolituus*) and aromatic (*Cycloclasticus*, *Neptunomonas*) hydrocarbons (McGenity et al., 2012; Gutierrez, 2017). There are also obligate (as well as some facultative) methane-oxidizing bacteria that are termed methanotrophs (Kalyuzhnaya et al., 2019).

Baelum et al. (2012) found that cells aggregated to form bacterial-oil aggregates during incubation of enrichment cultures consisting of uncontaminated seawater from the Gulf of Mexico, 100 ppm of MC-252 oil and 60 ppm of the dispersant Corexit 9500. The aggregates were dominated by members of the family Colwelliaceae, which is discussed in detail later. Methyllococcaceae also increased in abundance from <1% at 10 days to 16% at 40 days. Methyllococcaceae are methanotrophs, meaning they can metabolize only methane or methanol as carbon and energy sources. An increase in the abundance of Methyllococcaceae following the MC-252 spill was not surprising given the unprecedented amount of methane that entered the water column (~250,000 tons) (Rogener et al., 2018). However, the increase of Methyllococcaceae in experimentally formed aggregates (Baelum et al., 2012) was surprising, as there was no obvious source of methane or methanol. The presence of these methanotrophs may suggest the existence of anoxic microenvironments in the aggregates that could allow growth of methanogenic Archaea, and although Baelum et al. (2012) adapted bacterial primers to improve archeal detection, it is not clear whether they would have amplified methanogens. Suboxic and anoxic conditions develop as a consequence of oxygen consumption during aerobic degradation of MS organic matter and insufficient oxygen diffusion into flocs (Ploug et al., 1997; Rath et al., 1998; Vojvoda et al., 2014). Reduced levels of dissolved oxygen enhance the use of alternative electron acceptors for anaerobic respiration by microorganisms (Torres-Beltrán et al., 2016) and can provide a favorable environment for methanogens (Shanks and Reeder, 1993; Vojvoda et al., 2014). The methane produced would diffuse to oxic zones and serve as a source of carbon and energy for methanotrophs.

Thus, there are numerous microbes implicated in MOS formation, oil emulsification/degradation and other activities induced indirectly by their existence on an oil-containing floc. Below, we discuss the main genera of heterotrophic bacteria involved in the formation and fate of MOS.
Role of the Generalist Genera *Alteromonas*, *Pseudoalteromonas* and *Halomonas* in the Formation and Fate of MOS

These genera are cosmopolitan marine aerobic generalists capable of consuming a wide range of organic compounds, including, in several species, hydrocarbons. Their potential involvement in hydrocarbon degradation tends to be overlooked in sequencing surveys, as their increased abundance relative to controls is not enough to suggest a direct role in degrading oil-derived hydrocarbons in that environment (Gutierrez et al., 2013a,b, 2018; Gutierrez, 2017; Gutierrez and Kleindienst, 2020).

Nevertheless, *Alteromonas*, *Pseudoalteromonas*, and *Halomonas* constitute a significant proportion of hydrocarbon-degrading communities in oil-contaminated marine environments. For example, *Alteromonas* and *Pseudoalteromonas* were enriched from surface waters and water-column samples collected following the MC-252 spill (Hazen et al., 2010; Valentine et al., 2010; Yang et al., 2016a). *Halomonas* was regularly detected in sequencing surveys of surface oil samples collected 1 month after the MC-252 spill (Gutierrez et al., 2013b) and water-column samples collected after 6 months (Yang et al., 2016a). The ability of strains within these genera to use hydrocarbons as growth substrates was confirmed with isolations of *Alteromonas* strain TK-46(2), *Pseudoalteromonas* TK-105 and three *Halomonas* strains (GOS-2, GOS-3a, and TGOS-10) from samples collected in the Gulf of Mexico (Gutierrez et al., 2013b, 2018; Bacosa et al., 2018). In addition, the similarity and co-location of these strains to those enriched in surface oil slicks suggests that they made a major contribution to the degradation of M-acondio oil (Hazen et al., 2010; Valentine et al., 2010; Gutierrez et al., 2013b, 2018). There are also many other species within these genera that degrade hydrocarbons. For example, *Alteromonas nathaleniivorans* was responsible for the in-situ biodegradation of multiple PAHs in oil-contaminated tidal-flat sediment from the Yellow Sea (Jin et al., 2012; Math et al., 2012). *Pseudoalteromonas* was enriched in ship-board mesocosms containing North Sea water and crude oil, and many isolates were able to grow on PAHs, and branched and straight-chain alkanes (Chronopoulou et al., 2015). *Halomonas* species degrade a variety of hydrocarbons including monoaromatics (Garcia et al., 2004; Abdelkafi et al., 2006), polycyclic aromatics (Melcher et al., 2002; Yang et al., 2010) and aliphatics (Pepi et al., 2005; Mnif et al., 2009).

Sequencing surveys demonstrated an increase in the relative abundance of *Alteromonas*, *Pseudoalteromonas*, and *Halomonas* in MOS particles compared to the surrounding seawater (Arnosti et al., 2016; Suja et al., 2017). Moreover, an *Alteromonas* strain isolated from a MOS particle that formed in roller-bottle incubations with sea surface water collected during the active phase of the MC-252 spill, matched the aforementioned hydrocarbon-degrading strain *Alteromonas* TK-46(2) with 100% 16S rRNA sequence identity (Gutierrez et al., 2018). Fluorescence in-situ hybridization assays also detected *Halomonas* in bacterial-oil aggregates generated in the laboratory in alkane enrichments from deep plume water (McKay et al., 2016).

Roller-bottle experiments with *Alteromonas* strain TK-46(2) (Gutierrez et al., 2018) and *Halomonas* strain TGOS-10 (Gutierrez et al., 2013a) formed MOS even when cells had been inactivated by sodium azide, suggesting that the cell surface and/or their EPS, play a role in MOS formation/colonization. Epifluorescence microscopy revealed the attachment of *Alteromonas* and *Halomonas* cells to the MOS particles. The EPS produced by these bacteria enhance aggregation of particles to form MS or MOS (Passow et al., 2012; Zierovogel et al., 2012; Gutierrez et al., 2013a), and can exhibit amphiphilic properties, leading to emulsification or dispersion of hydrophobic oil constituents, making them more amenable to biodegradation (Gutierrez et al., 2007, 2008, 2018).

**Colwelliaceae**

Numerous amplicon and metagenomic sequencing studies from the MC-252 spill (and others) have shown an increased relative abundance of Colwelliaceae in the presence of crude oil (Brakstad et al., 2015b; Kleindienst et al., 2015b; Campeão et al., 2017; Hu et al., 2017; Bacosa et al., 2018; Lofthus et al., 2018; Ribicic et al., 2018). Following the initial enrichment of Oceanospirillales (May 2010) there was a shift in dominance to *Colwellia* (until August 2010) (Redmond and Valentine, 2012). Furthermore, Mason et al. (2014) suggested that the high abundance of *Colwellia* in the sediments below the oil plume was due to its transport by MOS. Many species of *Colwellia* are psychrophilic and/or piezophilic (Bowman, 2014; Liu et al., 2020), and one strain, RC25 was able to degrade 75% of MC-252 oil at 5°C (Baerum et al., 2012). While some of this loss can be attributed to evaporation, *Colwellia* species are probably important hydrocarbon degraders in deep-sea MOS (e.g., MC-252 at 4 to 6°C and 15 MPa). As an aside, phylogenetic analysis of strain RC25 (Supplementary Figure 1) reveals that it clusters within Colwelliaceae Clade 1, which according to Liu et al. (2020) should be renamed as *Cognaticolwellia*.

Mesocosm experiments with deep-sea (1,100–1,240 m depth) Gulf of Mexico bacterial communities have indicated that Corex-9500 components provide additional carbon sources for bacteria, leading to an increase in the relative abundance of Colwelliaceae (e.g., Kleindienst et al., 2015b; Techtmann et al., 2017). *Colwellia* strain RC25, for example, degraded dioctyl sodium sulfosuccinate, a surfactant component of Corex 9500 (Chakraborty et al., 2012), which in part explains its high abundance on MOS in the study by Baerum et al. (2012). In other studies, Colwelliaceae made up <1% of the community in MOS particles formed at subarctic conditions (Suja et al., 2017) and was detected at low abundances (max. 6%) in oil-related aggregates formed in cold seawater (Netze et al., 2018). Whilst it is difficult to compare across experiments in which more than one variable has changed, this difference might be a function of differing methodologies, which include: (1) different initial microbial communities present between the Gulf of Mexico, Faroe-Shetland Channel and Norwegian Sea; (2) direct addition of oil/dispersant vs. the use of water accommodated...
fractions; (3) different dispersant concentrations (Baelum et al., 2012) used 60 ppm, Suja et al. (2017) used 17.6 ppm, and Netzer et al. (2018) used 0.3 ppm; probably more so than the type of dispersant used (Corexit 9500 and Superdispersant-25); and (4) incubations remaining static or continuously mixed (Supplementary Table 1). Again, it must also be borne in mind that the concentrations of oil and dispersants used in most of these experiments were much higher than those found in the environment (Bejarano et al., 2014; Wade et al., 2016); lower concentrations might have much smaller effects.

**Obligate Hydrocarbonoclastic Bacteria (OHCB)**

*Alcanivorax* spp. are undetectable or at low abundances in pristine marine environments, but they typically bloom to make up a significant proportion of the bacterial community after an oil spill (Coulon et al., 2012; Teramoto et al., 2013), and so play an important role in the natural bioremediation of oil spills worldwide (McKew et al., 2007a,b; Yakimov et al., 2007).

Suja et al. (2017) prepared MOS from surface waters of the Faroe-Shetland Channel and 35,000 ppm dispersed oil, enriching for *Alcanivorax*, *Alteromonas*, and *Pseudalteromonas*, which collectively represented 80% of the sequence reads at 2.5 weeks incubation. In a subsequent roller-bottle study using surface water from the same environment, and similar high concentrations of oil, a comparison was made between the bacterial communities associated with MOS aggregates and marine dispersant snow, a product of adding the dispersant Superdispersant-25 to seawater in the absence of oil (Suja et al., 2019). Both marine dispersant snow and MOS aggregates had similar bacterial communities dominated by *Alcanivorax* with a consistent but minor contribution from *Oleispira*, as well as a range of other genera (Suja et al., 2019). Whilst the exact formulation of commercial dispersants, such as Superdispersant-25 used by Suja et al. (2017, 2019), are proprietary, they typically consist of a mixture of solvents and surfactants. The solvent fraction is usually a paraffinic or synthetic hydrocarbon, while the surfactant fraction of at least some includes dioctyl sodium sulfosuccinate, Tween 80, Tween 85, and Span 80 (Place et al., 2010, 2016; Brakstad et al., 2018b). Some species of *Alcanivorax*, as many other heterotrophs, are able to grow on components of the surfactant fraction, such as Tween (Liu and Shao, 2005) and likely the solvent hydrocarbon, so the *Alcanivorax* on marine dispersant snow aggregates may have been metabolizing the solvent or surfactant fraction of the dispersant.

There are numerous biochemical and physiological reasons why *Alcanivorax* has an ecological advantage over other hydrocarbon degraders, which may explain their frequent dominance in MOS particles, including: (1) cellular hydrophobicity that allows attachment to oil (Godfrin et al., 2018); (2) production of biosurfactants to facilitate their access to hydrocarbons (Abraham et al., 1998; Yakimov et al., 1998; Qiao and Shao, 2010); (3) biosynthesis of exopolysaccharides (including alginate) and type IV pili which mediate the formation of biofilms (Schneker et al., 2006; Sabirova et al., 2011), including associations with marine plankton (Coulon et al., 2012; Yakimov et al., 2019); (4) expression of permeases and iron-scavenging siderophores that concentrate nutrients in oligotrophic marine environments (Schneker et al., 2006; Sabirova et al., 2011; Denaro et al., 2014; Kem et al., 2014); (5) the ability to use both branched and linear alkanes as growth substrates (Yakimov et al., 2007; Gregson et al., 2019). Nevertheless, *Alcanivorax* is not universally detected in studies which generated MOS, e.g., with surface (Arnosti et al., 2016) and deep-sea (Baelum et al., 2012) water from the Gulf of Mexico. Some *Alcanivorax* species are piezotolerant, so higher pressure in the deep-sea may have inhibited their growth (Scoma et al., 2016a,b); however other species of *Alcanivorax* are piezotolerant (Liu et al., 2019).

Members of the genus *Oleispira* are known to flourish in cold, oil-contaminated seawater (Yakimov et al., 2003, 2007; Coulon et al., 2007; Tremblay et al., 2019). *Oleispira* were dominant in deep-water samples following the MC-252 spill (Hazen et al., 2010; Mason et al., 2012), and on MOS formed in mesocosms using seawater from the deep-sea of the Gulf of Mexico (4.8°C; Baelum et al., 2012), the Faroe-Shetland Channel (8.7°C; Suja et al., 2017, 2019) and the Trondheim Fjord (6–8°C; Netzer et al., 2018). The type strain, *Oleispira antarctica* RB-8, is a psychrophilic alkane degrader with a broad growth optimum of 1–15°C (Yakimov et al., 2003). It uses multiple mechanisms to compete at low temperatures, such as changes in the flagella structure/output to overcome increased viscosity, switching of flagella rotation to accumulate cells in an environment where it can outcompete other bacteria, and proline metabolism to counteract oxidative stress (Gregson et al., 2020). These psychrophilic adaptations of *Oleispira* most probably explain its absence from MOS generated from 25°C surface water collected following the MC-252 spill (Arnosti et al., 2016). This single trait of temperature requirement for growth highlights the importance of understanding the succession of microbial communities, and hydrocarbon-degradation potential, in MOS that sinks from warm surface waters to cold hadal depths.

Biodegradation of PAHs in oil-contaminated marine environments is usually associated with *Cycloclasticus* spp. (Dyksterhouse et al., 1995; McKew et al., 2007a; Teira et al., 2007). Arnosti et al. (2016) demonstrated, using 1% floating oil, that bacterial communities on MOS particles were distinct from those in surrounding seawater and dominated by a *Cycloclasticus* phylotype that was nearly identical in 16S rRNA sequence to PAH-degrading *Cycloclasticus* strain TK8 (Gutierrez et al., 2013b) and *C. pugettii* (Dyksterhouse et al., 1995). This highly enriched *Cycloclasticus* phylotype in the MOS was distinct from *Cycloclasticus* found in the experimental inoculum and the water column following the MC-252 spill (Valentine et al., 2010; Redmond and Valentine, 2012; Yang et al., 2016a) [for phylogenetic analysis see Yang et al. (2016b) and Redmond and Valentine (2019)].

In the aforementioned study by Suja et al. (2017), the relative abundance of *Cycloclasticus* in the community after 4 weeks was 3.5-fold higher in MOS particles than in surrounding seawater, although it contributed only 1.6% of total sequence reads. Doyle et al. (2018) demonstrated that *Cycloclasticus* was sensitive to chemically dispersed oil at high concentrations (i.e., chemically enhanced water-accommodated fraction) but
thrived in lower concentrations of physically dispersed oil. Also, given that most *Cycloclasticus* spp. from the identified clade are able to metabolize only aromatic hydrocarbons, in some cases they are outcompeted by aliphatic hydrocarbon degraders (e.g., *Alcanivorax*) for nutrients, and do not reach high levels until after the aliphatic fraction of the oil is depleted (Röling et al., 2002). There is now evidence that some *Cycloclasticus* phytophylotypes do not degrade PAHs but are capable of short-chain alkane degradation (e.g., mussel endosymbionts; Rubin-Blum et al., 2017), and that others may degrade both types of hydrocarbon (see Gutierrez et al., 2018; Redmond and Valentine, 2019), and so interpretation of the function of *Cycloclasticus* from 16S rRNA sequences alone requires caution.

**DOES THE USE OF DISPERSANT ENHANCE OR INHIBIT MOS FORMATION?**

Dispersants lower the interfacial tension between water and oil, and so reduce the energy required to convert a slick (or plume) into small oil droplets in the water column where they undergo rapid biodegradation (Prince, 2015b; Brandvik et al., 2019). However, there is currently a debate over whether dispersants have an effect on the biodegradation of crude-oil constituents (Kleindienst et al., 2015b, 2016; Prince et al., 2016b), with conflicting reports in the literature showing it can be enhanced (Baelum et al., 2012; Techtmann et al., 2017; Tremblay et al., 2017), inhibited (Hamdan and Fulmer, 2011; Kleindienst et al., 2015a,b), or essentially unaffected (Prince et al., 2013; McFarlin et al., 2014; Louvado et al., 2019). In part this reflects a misunderstanding of the scale at which dispersants are designed to operate—they are formulated to encourage the dispersion of oil at sea that would otherwise remain as a floating slick, thereby dramatically increasing the surface area available for microbial colonization. Laboratory experiments cannot mimic this behavior at environmentally relevant concentrations unless oil is prevented from dispersing by corralling it in an appropriately scaled boom (Prince and Butler, 2014). Most of the studies listed above were conducted under conditions where the added oil dispersed regardless of the presence of dispersant, and the small stimulations and inhibitions reported are of no environmental significance. The substantial stimulation of biodegradation by dispersants (many-fold) occurs after oil goes from a floating slick to small dispersed droplets (Prince and Butler, 2014; Prince et al., 2017). Once dispersed, the different surfactants of the dispersant leave the droplets at different rates (Riehm and McCormick, 2014), and are biodegraded quite rapidly (Brakstad et al., 2018a,c; McFarlin et al., 2018).

Equally, the effect of oil on MOS formation, and any additional effects of dispersant application are a major point of contention. Some workers have found that dispersant alone generates MS (Baelum et al., 2012; van Eenennaam et al., 2016), and that adding dispersant with oil stimulates flocculation (van Eenennaam et al., 2016; Doyle et al., 2018; Bretherton et al., 2019), albeit sometimes only when extra nutrients were added (Kleindienst et al., 2015b). Wirth et al. (2018) saw larger, more stable MOS in the presence of Corexit, but other work suggests that dispersants inhibit MOS (Passow, 2016; Passow et al., 2017), while others claim they were essential for their formation (Suja et al., 2017, 2019).

A major problem seems to be that many of these studies are far-removed from the conditions found at sea. Perhaps the first thing to consider is how much dispersant and oil was available during the MC-252 oil spill response to cause an effect. Dispersants are typically applied at a nominal rate of 5 gallons (US) per acre; 47 liters per hectare or 4.7 ml/m² (a teaspoon/m²), which when diluted into the top meter of the sea becomes a concentration of 4.7 ppm [The critical micelle concentration—the concentration above which the dispersant forms micelles—for Corexit 9527 is about 400 ppm (Singer et al., 1995)]. Such dilution is generally accepted to occur within hours, with continued dilution as time progresses (Lunel, 1995; Bejarano et al., 2013, 2014; Lee et al., 2013). Claims that “the average Corexit concentration in the water after the Deepwater Horizon spill was estimated at 0.5%” (Passow, 2016) cannot be correct.

The amount of oil in dispersed oil “plumes” is similarly dilute—immediately after dispersion the concentration will be several hundred to thousands of ppm (Bejarano et al., 2013, 2014), but this mixes and dilutes to less than a ppm within a day. The vast majority of samples collected in the vicinity of MC-252 during the spill contained only background levels of total petroleum hydrocarbons, despite the 3.19 million barrels spilled (Barbier, 2015); only 5% of the 1,372 samples had more than 250 μg of total petroleum hydrocarbons per liter (250 ppb), and only 24 had above 3 ppm (Wade et al., 2016). These amounts are in line with the concentrations of oil residue found at the bottom of the Gulf of Mexico; Stout et al. (2017) estimated that the heavily biodegraded residue of ~233,000 bbl of fresh oil was deposited on ~1,470 km² of Gulf deep-sea sediment. This is the residue of an initial 25 ml fresh oil/m². If 80% of the oil is GC-detectable, and the residue is 80% degraded (Stout et al., 2017), this is 6.25 g of extensively biodegraded oil residue in the top 1–3 cm per m² in the sediment. By extension, a total of 18.75 g of hydrocarbon had been consumed in the 1 m² × 1,500 m water column, or in the sediment after deposition. Data from sediment traps is also consistent: Stout and German (2018) estimated that the biodegraded residue of 10 barrels/km² fell through the water column some 58 km northeast of the MC-252 site during the blowout—this is equivalent to the residue of 1.6 ml fresh oil per square meter of sediment, ~0.3 ml/m² of weathered oil.

As pointed out by Brakstad et al. (2018b), the majority of papers claiming to provide insight into the effects of dispersants on oil degradation and on MOS used concentrations that are substantially higher, typically by several orders of magnitude. This becomes especially important when discussing whether marine snow is generated from oil and/or dispersants, or whether the surfactants in dispersants might have disruptive effects on MOS.

Another confusing aspect is the use of Water Accommodated Fractions (WAF) instead of oil-in-water dispersions. WAF were developed by toxicologists in an attempt to distinguish true chemical toxicity from physical smothering (Widdows et al., 1982; Singer et al., 2000). Essentially the oil is stirred into seawater...
for a defined time, allowed to “settle” (i.e., for oil droplets to coalesce and float), and then WAF is drawn off from below the slick (see Prince et al., 2016a). If dispersant is also present, the liquid is known as Chemically Enhanced Water Accommodated Fraction (CEWAF), and other abbreviations abound to describe dilutions, higher energies of mixing, etc. While such methods do provide a way of reproducibly preparing samples for toxicity testing, say for comparing different species, their composition is specific to the precise conditions used; WAF and CEWAF are indeed “accommodated” fractions, including both truly soluble compounds and tiny droplets. The concentration of the latter depends on the mixing energy and the “settling time,” which in turn defines how much of a reservoir the drops provide for resupplying soluble compounds should they be consumed from the aqueous phase by biodegradation or evaporation. WAF with nominally similar concentrations have significantly different concentrations of soluble compounds if they are made directly at the nominal concentration, or by dilution from a more concentrated “stock” (Prince et al., 2016a), and one cannot readily predict the chemical composition of either. In contrast, while oil-in-water dispersions may have variable droplet sizes depending on the energy used to generate the dispersion, the chemical composition of what is in the vessel is reproducible anywhere, and is readily calculated. Another major confusion of such work is that CEWAF have much more hydrocarbon per volume than WAF; without dilution to equivalent oil concentrations it is impossible to disentangle any effects of the presence of dispersants from the presence of more oil. Nevertheless, few papers address this issue, and assume any effects are due to the dispersants rather than the extra oil.

The widespread use of WAF/CEWAF in MOS experiments may be due to the fact that crude oil adheres to the walls and lids of the roller bottles. To overcome this issue several modifications can be made to the experimental setup, including pre-treatment of glassware (Brakstad et al., 2015a), use of hydrophobic seals/lids (Prince, 2015a), and slower rotation (Brakstad et al., 2015a; Netzer et al., 2018; Henry et al., 2020, 2021).

Clearly there needs to be a concerted effort to make sense of the disparate conclusions described above. All of the papers claim to be aimed at understanding what happens to spilled oil after a major oil spill, but few are done at environmentally relevant conditions. While there is much to be learned from careful studies with well-characterized and appropriate microbial isolates, the diversity already revealed (e.g., Passow et al., 2019) indicates that extrapolating to the real world remains premature.

UNDERSTANDING MOS FORMATION AND BEHAVIOR BEYOND THE GULF OF MEXICO

Investigations into MOS formation/deposition and the microbial communities associated with these aggregates have focused on the Gulf of Mexico with only a few studies in other areas (Suja et al., 2017, 2019; Netzer et al., 2018). To be of value to oil spill response in future spills, studies need to include other locations with environmental features that may influence MOS processes, such as areas with both deep-sea oil exploration/production and periods with high biological productivity and/or inorganic matter abundance. Overlaid maps of these features or their proxies (oil platforms, chlorophyll, and suspended mineral concentration) will help to predict locations where marine oil snow sedimentation and flocculent accumulation may occur (Murk et al., 2020), even as we are still unclear whether we can limit or stimulate MOS formation. Many oil refineries operate on estuaries, and gradients for MOS formation are expected with distance from river systems as they supply nutrients/clay to promote aggregation (Figure 2; Daly et al., 2016; Quigg et al., 2020). The Arctic region has been predicted to be extremely sensitive to MOS formation (Murk et al., 2020), and although overall biodegradation of oil components may be a little slower in incubations with polar compared with temperate seawater (McFarlin et al., 2014; Garneau et al., 2016; Brakstad et al., 2018a,c), this may reflect the particularly pristine waters that were used in these studies, and the lower natural abundance of oil-degrading microbes. In support of this, Coulon et al. (2007) showed that the oil-degradation rate in North Sea microcosms at 4°C was only half that at 20°C. Slower degradation would lead to increased exposure of the (micro)biota to oil, but dilution of dispersed oil would still dominate physical processes, and Arctic species are no more sensitive to oil toxicity than temperate ones (Bejarano et al., 2017).

FURTHER EXPERIMENTAL CONSIDERATIONS

We have highlighted the overarching need to employ environmentally relevant oil and dispersant concentrations/ratios for experiments to be useful in modeling the fate of oil in the sea. Oil concentrations of 0.3 to 3 ppm should be sufficient for MOS formation, while also limiting depletion of autochthonous nutrients and oxygen. Oil in sealed roller bottles should be at least partially weathered to allow benzene and other small hydrocarbons to evaporate, as they do in the field. Also, where model microbes are used in MOS studies they must be relevant marine species, and not as occasionally seen, surrogates from terrestrial or freshwater environments. After all, no-one interested in understanding apex predators on the savannah of southern Africa would think that the best approach would be to study Polar Bears, yet several papers use terrestrial and freshwater microbes as surrogates for marine species. Here we outline a number of other important factors that must be born in mind when studying MOS.

Light

Light plays a central role in MS formation by stimulating the growth of phototrophs and thus their exudates, but it also has
a photochemical role by cross-linking EPS (Sun et al., 2017, 2018, 2019), and photooxidizing hydrocarbons (Prince et al., 2003; Bacosa et al., 2015a; Aeppli et al., 2018). Passow (2016) revealed that photooxidation of oil over a period of 3 weeks promoted MS formation, but the particles were smaller and more compact than MOS that accumulated at the sea surface after the MC-252 spill (diameters of 2.5 vs. 7 mm, respectively). Presumably oil’s interaction with marine snow depends on how the photooxidation changes specific oil compounds. Insoluble compounds (e.g., high molecular weight n-alkanes and PAHs) are incorporated into MOS within entire oil droplets, whereas more soluble compounds (e.g., low molecular weight n-alkanes (<C27) and simple aromatics) might be sorbed by cells in the MOS (Wirth et al., 2018).

Light intensity, spectrum, and photoperiod all have effects on photosynthesis, phytoplankton communities, and the metabolites produced (Schuback and Tortell, 2019), which lead to changes in the associated heterotrophic communities (Bacosa et al., 2015b). For example, incubations with sunlight (compared with dark incubations) greatly reduced the abundance of cyanobacteria, such as Synechococcus and Prochlorococcus, in seawater samples containing oil (200 ppm) or dispersant (10 ppm; Bacosa et al., 2015b). However, there was no analysis of eukaryotic phototrophs in this study, and so we do not know how the overall phytoplankton community responded. In the presence of crude oil (+/- dispersant) and sunlight, Alteromonas and Marinobacter were among the most enriched genera, and had much higher relative abundance than in the equivalent dark bottles (Bacosa et al., 2015b), suggesting an indirect effect of light, perhaps as a consequence of the death of cyanobacteria providing organic matter for the metabolically versatile copiotroph, Alteromonas (Pedler et al., 2014) Alternatively (or additionally) the growth of eukaryotic phytoplankton that potentially thrived in the absence of cyanobacteria may have changed the composition of released organic matter in favor of this genus. The microbial communities formed by different treatments (light vs. dark) may have contributed to the dissimilar size and structure of MOS particles in Passow (2016), but since community composition of the MOS particles was not determined, this cannot be addressed.

Thus, future MOS studies need to mimic natural light conditions, including light intensity, spectral composition, and light-dark cycles, in order to assess the aforementioned direct and indirect influences on MOS formation and fate. Due consideration must be shown to the light intensity used—it is very difficult to mimic noon sunlight in the laboratory, and different glass bottles absorb different fractions of the UV component of sunlight, so it may only be possible to mimic some depth into the natural photic zone in the laboratory (Ryther, 1956).

Aeration

The natural concentrations of MOS and dispersed oil in the water column are so low that aerobic seawater will remain oxygenated during bottle experiments at environmentally relevant levels. Surface seawater typically contains 200 µM dissolved oxygen, and although there is usually an oxygen minimum of about 120 µM oxygen around 500 m depth, most ocean waters are aerobic throughout (e.g., Camilli et al., 2010, but see Paulmier and Ruiz-Pino, 2009 for a discussion of truly low-oxygen water). This is enough oxygen for the complete oxidation of several ppm of oil, so sealed bottles with only a few ppm of oil will remain aerobic even in the dark. Incubation in the light would allow phototrophs to generate even more oxygen, for it is well-known that the surface layers of the ocean are supersaturated with oxygen during the day (Craig and Hayward, 1987). It is noteworthy that the initial discovery of the subsea plume of dispersed oil following the MC-252 spill included the detection of local consumption (10%) of the indigenous oxygen as microbes degraded the hydrocarbons (Hazen et al., 2010), but the plume was far from anoxic.

Ocean sediments are typically anoxic within a centimeter or two of the sediment surface even though the water itself contains typically >100 µM oxygen (Reimers et al., 1986). [The oligotrophic deep-sea sediments are an exception (Fischer et al., 2009) but are also unlikely to be impacted by oil.] Experiments aiming to study MOS and its effects once it lands on the sediment should ensure that such conditions exist in the mesocosm before adding MOS.

Temperature and Pressure

The impacts of temperature on hydrocarbon bioavailability, degradation rates and microbial communities are well-established (e.g., Coulon et al., 2007), and so in-situ temperatures are usually replicated in MOS experiments. However, there are few studies that apply a decreasing temperature regime to mimic the sinking of particles from warm surface waters to colder hadal depths. Increases in pressure can reduce the rate of hydrocarbon biodegradation (Prince et al., 2016c) and also alter microbial communities, e.g., selecting against piezo-sensitive microbes, such as some Alcanivorax (Scoma et al., 2016a,b). Although high pressure can be simulated in the laboratory, it is less straightforward to effect a gradual pressure change that would allow piezotolerant microbes time to acclimate. Moreover, in the marine environment MOS sinks into new waters, where particle association is the normal mode of existence for microbes (Zhao et al., 2020) and where they are adapted to low temperature and high pressure. Thus, new microbes would attach to the MOS during its decent, a phenomenon that would be challenging to replicate experimentally.

Nutrients

Numerous papers have shown that the natural levels of nutrients in seawater are enough for the rapid biodegradation of ppm levels of oil (Prince et al., 2013, 2017; McFarlin et al., 2014; Brakstad et al., 2015a, 2015b; Wang et al., 2016). Furthermore, regulators are unlikely to allow the addition of nutrients to the open ocean for fear of eutrophication (Altiieri and Diaz, 2019). Depending on the environment under investigation, experiments intending to inform oil-spill response measures should either not add nutrients or should supply them continuously at in-situ concentrations in order to mimic replenishment in the natural environment.
**Oil and Dispersant Type**

The majority of studies investigating MOS have focused on environmental effects (e.g., the presence of POM, the proximity of river outflow providing clay minerals and phytoplankton biomass, and the presence of EPS-producing biota), while the influence of oil and dispersant type has not received much attention.

Most MOS studies have used either oil collected directly from MC-252 (Macondo oil) or closely related surrogates that are similarly light and high in volatile hydrocarbons (Hammer et al., 2011; Wade et al., 2017), but other spilled oils can have significantly different compositions and densities. For example, the Bunker B fuel oil spilt from the *Tsesis* tanker in the brackish (6‰) Baltic was a heavy (though lighter than water), viscous, refined fluid, containing low levels of volatile hydrocarbons (Boehm et al., 1982). We have no knowledge of whether such a spill might lead to MOS. Passow et al. (2019) compared Macondo surrogate oil with a heavy California oil, and found that the latter led to a greater reduction in diatom abundance, which they predicted would negatively impact MOS formation, but the mechanisms by which this would occur are obscure. Oils are very complex mixtures, and some characterization of the oil, focusing on the progress of its biodegradation, should be part of all experiments.

By far the most studied dispersant is Corexit 9500 which was used on the surface and injected at the well-head during the MC-252 blowout (Atlas and Hazen, 2011). Other approved dispersants stockpiled in large quantities around the world include Finosol OSR 52 (Total Fluids, France) and SlickGone NS (Dasic International Ltd, UK) (Carter-Groves, 2014). Korea used HiClean II in the response to the *Hebei Spirit* spill (Jung et al., 2009), and Superdispersant-25 is licensed in the UK (Scarlett et al., 2005). The chemical composition of Corexit 9500 is known (Riehm and McCormick, 2014; Place et al., 2016), whilst limited information is available on the material safety data sheets for the others. It would certainly be interesting to see whether the different dispersants have different effects, with and without different oils, on microbial communities at environmentally-relevant concentrations. For example, several studies have shown an increase in the abundance of *Colwellia* in experimental incubations containing only Corexit 9500 (Kleindienst et al., 2015b; Techtmann et al., 2017; McFarlin et al., 2018), likely because they are degrading the Tween surfactant (Bowman, 2014) and the hydrocarbon solvent.

**CONCLUDING REMARKS AND RECOMMENDATIONS**

MOS research abounds with controversies, mostly because of extrapolations to the marine environment from experiments employing unrealistic conditions, particularly oil/dispersant concentrations/ratios. Despite the difficulties in both capturing and investigating patchily distributed MOS in the deep sea and mimicking *in-situ* conditions experimentally, our understanding of MOS formation and fate has advanced over the last decade in terms of the microbes present and their potential roles in the formation of MOS and the fate of hydrocarbons and dispersants. Nevertheless, an overarching concern for practitioners is whether particular oil-spill responses, such as dispersant application, will enhance or inhibit MOS formation (Figure 2). Dispersants can reduce the overall stickiness of particles through either destabilization of TEP precursors (e.g., microgels; Chiu et al., 2017) or direct dispersal of TEP (Passow et al., 2017). This means that particles may collide but not stick together. However, dispersants also cause microbes to alter both EPS abundance and composition (e.g., protein-carbohydrate ratio; Xu et al., 2018a,b, 2019). Protein-rich/hydrophobic EPS shows more resistance to dispersion and can facilitate MOS formation due to their high stickiness (Chiu et al., 2019; Santschi et al., 2020; Shiu et al., 2020). Therefore, MOS formation appears to be dependent on the relative strength of opposing mechanisms (e.g., microgel destabilization and TEP dispersion vs. changes in EPS release; Figure 2). However, there is currently insufficient evidence that results from these laboratory incubations will be replicated in the field.

Oil spill responders would benefit from models of the fate of hydrocarbons, from the formation of MOS to sediment deposition (Passow et al., 2019; Daly et al., 2020; Quigg et al., 2020), accounting for factors that influence MOS sinking rate, such as the degree of hydrocarbon degradation and loss. Due consideration should be given to how oil degradation will change while sinking, given that: (1) less structurally complex hydrocarbons will be degraded rapidly first (rapid degradation earlier); (2) colonization, growth and surfactant production by hydrocarbon-degrading microbes will take time (rapid degradation later); (3) temperatures are higher at the surface (rapid degradation earlier); (4) nutrient concentrations are generally higher at depth (rapid degradation later). More unpredictable features include: (1) the effects of photic processes on the alteration of hydrocarbons that could enhance their loss or diminish their biodegradability; (2) the poorly studied impact of pressure on hydrocarbon biodegradation; (3) changes in the buoyancy of MOS (e.g., due to change in hydrocarbon composition; *de-novo* EPS production as well as its consumption); (4) feeding on, and disaggregation of, MOS.

A more fundamental understanding of the interacting forces that influence MOS sinking rates and hydrocarbon biodegradation will come from applying novel techniques. For example, Fourier transform ion cyclotron resonance mass spectrometry (FTICR-MS; Wozniak et al., 2019) and solid-state 13C nuclear magnetic resonance (NMR; Hatcher et al., 2018) allow detailed analysis of polar species, hydrocarbons and their oxidation products. Synchrotron radiation-based Fourier transform infrared (SR-FTIR) spectromicroscopy is particularly valuable in illustrating the spatial organization in MOS of microorganisms, biopolymers, oil, and its degradation products (Baelum et al., 2012). Although many studies have described the bulk microbial composition of MOS, more analysis of the active microbial component is needed (e.g., via metatranscriptomics and metaproteomics). Also, new methods are being developed for studying MS that could be applied to MOS, which involve
capturing (Duret et al., 2019), preserving, embedding and imaging particles to investigate the spatial distribution of microbes and polymers (Flintrop et al., 2018; Rogge et al., 2018), and even allowing single-cell functional analysis by nanoscale secondary ion mass spectrometry (nano-SIMS; Rogge et al., 2018). However, such analyses must be coupled with realistic experiments and/or in-situ investigations if they are to aid oil-spill response.

It remains to be seen how important MOS is in the overall fate of spilled oil in light of the roles of adhesion to mineral and biological particles, and the simple loss of buoyancy as oil droplets degrade. It is clear that dispersed oil degrades promptly at sea, while undispersed and weathered oil can reach the shoreline (for example as a black tide or as tar balls), where the ecological impacts can be prolonged and the clean-up effort is often expensive and time consuming. This alternate fate must be borne in mind when addressing the environmental effects of sunken dilute degraded oil.

AUTHOR CONTRIBUTIONS

TM, BM, and RH conceived the review. BG wrote the first draft and produced the figures. TM, TN, and RP wrote sections. All authors contributed to the writing and structure of the review.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fmars.2021.619484/full#supplementary-material

SUPPLEMENTARY FIGURE 1 | Phylogenetic tree based on 16S rRNA sequences for different species of the family Colwelliaceae. The 16S rRNA sequence of Colwellia RC25 which was isolated from marine oil snow (MOS) particles in Baelum et al. (2012) is highlighted in bold. The evolutionary history was inferred using the Neighbor-Joining method. The percentage of replicate trees in which the associated taxa clustered together in the bootstrap test (1,000 replicates) are shown next to the branches. The tree is drawn to scale, with branch lengths in the same units as those of the evolutionary distances used to infer the phylogenetic tree. The evolutionary distances were computed using the Maximum Composite Likelihood method and are in the units of the number of base substitutions per site. Evolutionary analyses were conducted in MEGA X.

SUPPLEMENTARY TABLE 1 | Details of conditions used in experiments on Marine Oil Snow (MOS) and Marine Dispersant Snow (MDS) formation.

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Conflict of Interest: RH is employed by the company Oil Spill Response Limited.

TN is employed by ExxonMobil Upstream Research Company. RP is retired from ExxonMobil, and received no payments for this work.

The remaining authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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