

Marine Epibiosis: Concepts, Ecological Consequences and Host Defence

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Abstract The sessile mode of life is widespread in a variety of marine phyla. Sessile life requires a stable substratum. On the benthos, motile life stages and sessile adults compete for rigid surfaces making non-living, i.e. inanimate, hard substratum a limited resource. Epibiosis is a direct consequence of surface limitation and results in spatially close associations between two or more living organisms belonging to the same or different species. These associations can be specifically guided by host chemistry resulting in species-specific symbiotic or pathogenic assemblages. Most colonizers, however, are non-specific substratum generalists. The ecological consequences for the overgrown host (basibiont) and the colonizer (epibiont) can be positive and negative. The predominantly disadvantageous nature of epibiosis by microorganisms for the basibiont has resulted in a variety of defence mechanisms against microcolonizers, including physical and chemical modes of action. Besides antimicrobial effects of secondary metabolites emanating from the host, recent studies increasingly demonstrate that epibiotic bacteria associated with the host deter growth and attachment of co-occurring bacterial species or new epibiotic colonizers competing for the same niche.

1 Introduction

In the marine environment the sessile mode of life is dominant in the majority of phyla. Owing to their low specific weight not only microorganisms (e.g. bacteria, microalgae) but also the small motile life stages of macroorganisms (e.g. larvae and spores) behave like passive propagules in this viscous, hydrodynamically dominated environment. Clearly, for these species an attached filter-feeding mode of life is the energetically advantageous and favourable state, although the lack of locomotion necessitates a variety of new challenging survival strategies, such as reproduction and various defence forms against consumers and overgrowth.

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Sessile life requires a stable substratum. On the benthos, motile life stages and sessile adult forms compete for rigid surfaces making non-living (i.e. inanimate) hard substratum a limited resource. Epibiosis (greek *epi* “on top” and *bios* “life”) can be considered as a direct consequence of surface limitation and results in spatially close associations between two or more living organisms belonging to the same or different species. The substrate organism is considered the *basibiont*, while the organism(s) growing attached to the animate surface is referred to as the *epibiont*. Epibionts are further subdivided into *epizoans* (animals) and *epiphytes* (plants, algae). Epibiotic assemblages are rarely species-specific; on the contrary, numerous sessile organisms may live either as basibiont or as epibiont, or both simultaneously (Wahl 1997). Attachment and growth on inanimate surfaces is usually considered as *fouling*, although this term is frequently used synonymously in an epibiotic context in the literature.

Epibiosis is a typical aquatic phenomenon although many examples of terrestrial epibionts are known (e.g. algae, lichens). Depending on seasonality and location the average millilitre of seawater contains 10–100 microscopic larvae and spores, 10^3 fungal cells, 10^6 bacteria and 10^7 viruses. Thus, the colonization pressure exerted by meroplanktonic dispersal stages can be intense on submerged surfaces (Davis et al. 1989) with severe ecological consequences for basibionts and epibionts. The distinctive role of water as a food vector for sessile organisms is the main reason why surface attachment, fouling and hence epibiotic associations predominate in aquatic environments. A large variety of marine phyla have adopted the sessile mode of life for at least one ontogenetic phase. The list includes many bacteria, protozoa, diatoms, molluscs, tube-building polychaetes; most macroalgae, bryozoans, phoronids, cnidarians; some echinoderms, crustaceans; all sponges and tunicates.

2 Ecological Consequences for Epibionts

Marine fouling is an omnipresent phenomenon and the list of foulers is long. The different stages of the fouling process of dispersal stages on solid substrates have been described in successional (Davis et al. 1989) and probalistic models (Clare et al. 1992; Maki and Mitchell 2002) and are presented in detail in this volume. Irrespective of the fouling sequence, for most meroplanktonic larvae settlement is the ultimate prerequisite for successful metamorphosis into sedentary juveniles (Hadfield and Paul 2001). Also, microcolonizers such as bacteria, benthic diatoms and algal spores often proliferate more rapidly or sometimes exclusively when fixed to a solid substratum (Grossart et al. 2003). Once attached, microcolonizers are challenged by other members in the biofilm matrix (Fletcher and Callow 1992; Costerton et al. 1995). To successfully compete in biofilms, many representatives of the bacterial genus *Pseudoalteromonas* release anti-bacterial products that aid the cells in the colonization of host surfaces. Through the production of agarases, toxins, bacteriolytic substances and other enzymes, bacterial cells are assisted in their competition for nutrients and space as well as in their protection against predators grazing on surfaces (Holmström and Kjelleberg 1999).

In densely populated marine environments where competition for space is high, the advantage for colonizers in occupying empty animate surfaces is probably the main reason for epibiosis (Wahl 1989; Todd and Keough 1994) although a variety of other advantages for the colonizer may support specific host–epibiont associations. For instance, settlement on raised or elevated hosts results in a hydrodynamically favourable position of the epibiont (Keough 1986) as flow dynamics increase with distance from the benthos (Butman 1987). Increased flow ensures better supply of planktonic nutrients and more efficient removal of toxic excretory products such as ammonia. An exposed habitat supports phototrophic epibionts, especially in deeper or turbid waters where light penetration is weak (Brouns and Heijs 1986). While filter-feeding epibionts profit from nutrient currents created by the host (Laihonen and Furman 1986) deposit-feeding epibionts benefit from metabolites exuded by the basibiont (Harlin 1973). Regarding the fate of colonizers, epibionts either benefit from the host defence against consumers or other colonizers, a phenomenon termed “associational defence” (Hay 1986), or the fates of epibiont and host are closely interlinked and shared together, a phenomenon termed “shared doom” (Wahl and Hay 1995).

The predominantly advantageous associations of epibionts with host organisms indicate that the mere presence of a surface is often not the only criterion for successful colonization. Numerous colonizers are reported to be guided by specific “cues” mediating the suitability of the settlement site (Rodriguez et al. 1993; Wiczorek and Todd 1998; Steinberg et al. 2002). The recognition of appropriate cues activates the genetically scheduled sequence of behavioural and physiological processes during settlement (Morse 1990) and many larvae delay or even avoid settlement in the absence of appropriate settlement cues (Coon et al. 1990; Qian and Pechenik 1998).

3 Settlement Cues

There is clear experimental evidence for physical settlement cues, such as surface roughness (Berntsson et al. 2000) and wettability (Qian et al. 2000); environmental conditions in direct proximity to the surface, such as irradiation (Maida et al. 1994) and microhydrodynamics (Mullineaux and Butman 1991); and biogenic chemical signals emanating from the basibiont or other epibionts (e.g. bacteria) already present on the host surface (Johnson et al. 1991a,b; Krug and Manzi 1999). Several authors have presented experimental evidence for selective settlement of both generalist and specialist epibionts in response to invertebrate or plant host cues. In most of these studies, the host served as the obligate prey source for larvae or adults. This raises the question of how planktonically dispersed larvae locate their patchily distributed hosts. Given the large spatial scales that need to be screened by potential colonizers, one would expect either strong or very distinct cues that govern such host–epibiont associations. To address this question, a number of studies have focused on the selective response of sea slugs to host corals. For example, water-soluble cues from corals induce settlement and metamorphosis in larvae of the opisthobranchs *Phestilla sibogae* (Hadfield and Scheuer 1985), *Adalaria proxima* (Lambert and Todd 1997) and *Alderia modesta* (Krug and Manzi 1999). Other

well-investigated host plants are coralline algae that govern larval settlement of taxonomically distinct invertebrates such as the sea urchin *Holopneustes purpurascens* (Williamson et al. 2000), the starfish *Acanthaster planci* (Johnson et al. 1991b; Johnson and Sutton 1994) and the mollusk *Haliotis* (Morse and Morse 1984; Hahn 1989). Other well-studied systems comprise obligate associations that seemingly benefit from close proximity of conspecifics to enhance reproductive output, such as in oysters (Tamburri et al. 1992; Turner et al. 1994) and barnacles (Clare and Matsumura 2000). However, the settlement cue(s) involved in the establishment of these systems were rarely identified at the molecular level.

In contrast to the numerous partially characterized inducers, only few settlement cues isolated from natural sources were in fact chemically identified, e.g. delta-tocopherols from *Sargassum tortile* that induce settlement of the hydroid *Coryne uchidai* (Kato et al. 1975); jacarone isolated from the red alga *Delesseria sanguinea* that induces settlement of the scallop *Pecten maximus* (Yvin et al. 1985); narains and anthosamines A and B isolated from marine sponges and lumichrome isolated from conspecifics that induce settlement of ascidian larvae (Tsukamoto et al. 1994, 1995, 1999); *N*-acylhomoserine lactone quorum sensing signal molecules that aid zoospores of the green macroalgae *Ulva* to exploit a bacterial sensory system and select permanent attachment sites by responding to bacteria already present on the surface (Joint et al. 2002). In most cases, the ecological relevance of these compounds in situ is not clear, either because the source of the settlement cue is not necessarily related to the recruitment patterns of the organism (Yvin et al. 1985; Tsukamoto et al. 1994, 1995), or because the availability of the cue to settling larvae has not been demonstrated unequivocally (Tsukamoto et al. 1999).

Interestingly, there is a high similarity in host recognition by pathogens in marine and terrestrial plants (Kolattukudy et al. 1995). For example, the pathogenic filamentous green alga *Acrochaete operculata* recognized its host, the red alga *Chondrus crispus*, by cell wall polysaccharides. *C. crispus* has an isomorphic life history, in which the gametophytic and sporophytic generations differ only in minor traits, such as sulfate-ester group distribution of their matrix polysaccharides, known as κ - and λ -carrageenans. Remarkably, the sporophytic generation is highly susceptible to infection whereas the gametophytic phase is naturally resistant. The virulence of the green algal endophyte is modulated by the presence of λ -carrageenan, which stimulates protein synthesis and elicits the production of specific polypeptides in the pathogen (Bouarab et al. 2001).

Only recent years have witnessed some complete characterizations of marine invertebrate larval settlement cues. In a series of investigations Matsumura et al. (1998) identified the key molecule responsible for gregarious settlement in the fouling barnacle *Balanus amphitrite* as a settlement-inducing protein complex (SIPC). This protein complex has now been fully elucidated as a α_2 -macroglobulin-like glycoprotein (Dreanno et al. 2006). Although the SIPC is regarded as an adult cue that is recognized by the cyprid at settlement, it is also expressed in juveniles and in larvae, where it may function in larva-larva settlement interactions. In another series of investigations the structure and the different sources of coralline algae-derived settlement cues for two larval species of sea urchins, *Holopneustes purpurascens*

and *Heliocidaris erythrogramma*, have been fully elucidated. The biogenic amine histamine was isolated from the red alga *Delisea pulchra* by bioassay-guided fractionation and identified as the inducer of settlement of *H. purpurascens* (Swanson et al. 2004). The alga still evoked larval settlement after antibiotic treatments, which effectively removed epiphytic bacteria on the algal surface, demonstrating that histamine was indeed an alga-derived cue. In contrast, the coralline alga *Amphiroa anceps*, which also stimulates larval settlement of *H. purpurascens*, lacked detectable amounts of histamine. Interestingly, antibacterial treatment of *A. anceps* removed the settlement cue, suggesting a bacterial origin of the cue from this alga; indeed bacterial films of two isolates from the surface of *A. anceps* induced settlement of *H. purpurascens* in laboratory assays (Swanson et al. 2006). The role of algae-associated bacteria as producers of settlement cues has been examined in more detail for the sea urchin *H. erythrogramma* (Huggett et al. 2006). The hypothesis of a bacterially derived settlement signal was supported by the fact that a variety of bacterial isolates from the surface of coralline algae triggered larval settlement at levels comparable to those of the positive control of coralline algae. One bacterial isolate from *A. anceps*, *Thalassomonas viridans*, is a known histamine producer. Given that larvae of both urchin species settle in response to histamine, these findings demonstrate a common settlement cue in coralline algae produced by the host alga and/or by associated bacteria.

4 Ecological Consequences for Basibionts

Any potential basibiont, i.e. the majority of sessile, relatively long-lived organisms, must either tolerate epibiosis or employ some sort of defence against this phenomenon. While epibiosis entails both benefits and disadvantages for epi- and basibionts the investment into defence depends on a finely tuned and often variable energy budget of the basibiont (Wahl 1989). Epibiosis causes a variety of beneficial effects to the basibiont, such as the induction of morphogenesis in macroalgae by symbiotic bacteria (Tatewaki et al. 1983; Nakanishi 1999), the interaction between macroalgae and nitrogen-fixing bacteria (Thevanathan et al. 2000), and the protection of seaweed surfaces from bacterial colonizers by associated bacteria (Lemos et al. 1985). A well-investigated example of a symbiotic association between host and epibiotic bacteria is the embryo of the American lobster, *Homarus americanus*, which is resistant to the fungus *Lagenidium callinectes*, a pathogen of many crustaceans. The surfaces of healthy lobster embryos are covered almost exclusively by a single, Gram-negative bacterium, which produces the antifungal substance 4-hydroxyphenethyl (Gil-Turness and Fenical 1992). Testing the effects of epibiosis on herbivory and predation, research by Wahl and colleagues suggested that epibionts on the blue mussel *Mytilus edulis* affected its susceptibility to predation by the shore crab *Carcinus maenas* (Wahl et al. 1997). Similarly, epibiosis by a variety of plants and animals altered the host susceptibility of the omnivorous sea urchin *Arbacia punctulata* (Wahl and Hay 1995). Furthermore, Wahl and Mark (1999) investigated

the hypothesis that if the effects for epibiont and basibiont were predominantly beneficial then co-evolution would be expected to lead to some sort of associational specificity. However, by analyzing over 2000 patterns of epibiotic associations the authors concluded that many colonizers are non-specific substratum generalists and that epibiosis is predominantly facultative (Wahl and Mark 1999).

The adverse effects of epibiosis on the basibiont often outweigh the beneficial ones (Table 1). For instance, soft-bodied marine invertebrates and algae are susceptible to diseases and tissue necrosis induced by bacteria, fungi and microalgae (Mitchell and Chet 1975; Bouarab et al. 2001; Cooney et al. 2002). The sometimes drastic changes of pH and redox conditions created by microepibionts may attack chemically sensitive surfaces of the basibiont (Terry and Edyvean 1981). Importantly, the adverse effects of microbial epibiosis may reach beyond pathogenicity and virulence. Since microbial films are important sources of chemical cues for larval settlement in many benthic marine invertebrates (Lau et al. 2002; Harder et al. 2002), microbial epibiosis may promote subsequent colonization by rigid crustose epibiotic macroorganisms, which in turn significantly impair the basibiont's ability to exchange gases and nutrients (Jagels 1973), damage the tissue by increased weight, rigidity and drag (Dixon et al. 1981), and decrease the growth rate of photosynthetic basibionts by cutting surface irradiance levels (Sand-Jensen 1977; Silberstein et al. 1986). From a nutritional perspective it is evident that if the host and the epibiont share the same trophic requirements then planktonic nutrients reaching the basibiont may already be partially depleted after their passage through the epibiotic barrier. As epibionts may fall victim to predators of their hosts, so may basibionts suffer from

Table 1 Ecological consequences for epibiont and basibiont as a result of epibiotic associations (summarized from Wahl 1989)

	Advantages	Disadvantages
Epibiont	Colonization of new substrate	Unstable, non-durable substrate
	New surface due to growth of basibiont	Biologically variable substrate
	Nutrient flow from basibiont	Exposure to detrimental host defence
	Favourable hydrodynamic conditions	Shared doom
	Favourable exposure to light	
Basibiont	Associational resistance	
	Camouflage	Increased weight and drag
	Insulation against desiccation	Decreased elasticity
	Nutrient flow from epibiont	Increased surface roughness
	Associational resistance	Increased deposition of particulate material
		Insulation against exchange of gas and waste products
		Increased mechanical damage
	Increased chemical damage	
	Decreased nutrient flow through epibiotic filter	

“shared doom”, i.e. damage due to grazers preying on epibionts (Dixon et al. 1981). Table 1 summarizes the advantages and disadvantages of epibiosis for epi- and basibionts.

5 Defence

Many marine invertebrates and plants have evolved a variety of physical and chemical defence mechanisms to suppress epibiosis and/or remove epibionts. Epibiont removal can be physically achieved by continuous or periodic surface renewal or by means of mucus secretion (e.g. in cnidaria, algae, molluscs, echinoderms and tunicates) and periodical shedding of the cuticula or epidermis (Sieburth and Tootle 1981; Littler and Littler 1999; Nylund and Pavia 2005) (see Fig. 1).

To create unfavourable or toxic conditions at or immediately above the living surface is a wide-spread adaptation of host organisms to cope with epibionts.

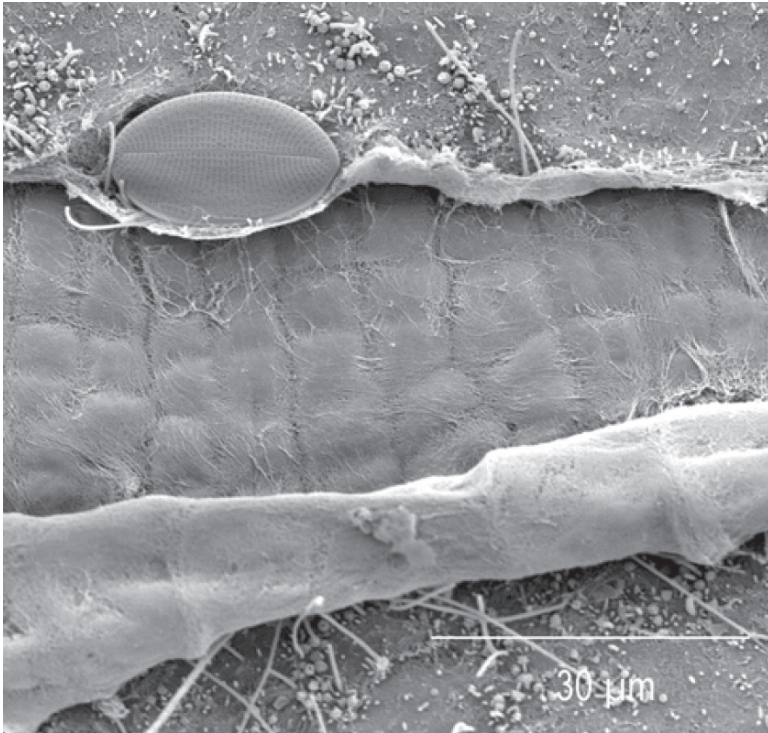


Fig. 1 Surface of the macroalga *Laminaria digitata* showing sloughing of the cuticle-containing bacteria and diatoms to reveal an uncolonized algal surface. Scale bar: 30 µm

The brown alga *Laminaria digitata* and the red alga *Gracilaria conferta* react with an oxidative burst to the presence of either alginate oligosaccharides or agar oligosaccharides, both of which are degradation products of their own cell walls (Küpper et al. 2001), resulting in the efficient elimination of bacterial epiflora (Weinberger et al. 2000). Moreover, sessile marine organisms feature a variety of chemical defence metabolites effective against different phyla of potential epibionts (reviewed by Clare 1996; Faulkner 2000). There are numerous studies on the inhibition of micro- and macroorganisms by extracts from diverse marine eukaryotes, such as corals, sponges, tunicates, ascidians and macrophytes (e.g. Michalek and Bowden 1997; Jensen et al. 1996; Wilsanand 1999; Slattery et al. 1995; Hellio et al. 2000; Dobretsov et al. 2006). Mostly, these investigations were descriptive and did not result in the purification and elucidation of inhibitory compounds. It remains unclear whether these extracts deter epibiosis at or near surfaces in situ and, if so, at what concentrations these effects are elicited. In this context, one of the better-studied models for algal secondary metabolism is the Australian red alga *Delisea pulchra*, which produces a range of structurally similar halogenated furanones (Steinberg et al. 2001). These metabolites are encapsulated in vesicles in the gland cells of *D. pulchra*, which provide a delivery mechanism to the surface of the alga at concentrations that deter a wide range of prokaryote and eukaryote epibionts (Maximilien et al. 1998). Being structurally related to acylated homoserine lactones (AHLs), halogenated furanones inhibit bacterial colonization through direct antagonism of bacterial cell-to-cell signalling. The AHL-mediated gene expression of bacteria is inhibited when halogenated furanones occupy the AHL-binding site of LuxR-like proteins, which represent the transcriptional activators in AHL regulatory systems (Manefield et al. 1999).

Information on the localization, identity and surface concentration of defence secondary metabolites is rapidly advancing (e.g. Salomon et al. 2001; Kubanek et al. 2002; Nylund et al. 2005; Paul et al. 2006) and the relevance of defence metabolites is increasingly discussed in a chemical ecological context. Moreover, recent studies on the deterrence of microbial colonization highlight that chemical antifouling defences cannot be generalized as broadly bacteriostatic or bactericidal, instead the effects are quite selective and targeted against particular microbial species (Maximilien et al. 1998; Egan et al. 2000; Kubanek et al. 2003).

Besides antimicrobial effects of secondary metabolites emanating from the host, recent studies have increasingly demonstrated that epibiotic bacteria associated with the host deter growth and attachment of co-occurring bacterial species or new epibiotic colonizers competing for the same niche (Armstrong et al. 2001; Harder et al. 2004a). A well-investigated bacterium in this context is *Pseudoalteromonas tunicata*, which has been isolated from a tunicate and a green macroalga. *P. tunicata* has been found to produce at least five extracellular compounds that inhibit other organisms from establishing themselves in an epibiotic community by inhibiting settlement of invertebrate larvae and algal spores, growth of bacteria and fungi, and surface colonization by diatoms (Holmström and Kjelleberg 1999; Holmström

et al. 1996). Thus, in terms of the chemical ecology of host–epibiont associations, it seems evident that there is a significant protective role of symbiotic microbial epibionts, which in turn release antifouling compounds. However, after more than 20 years of research there is no experimental evidence demonstrating if and how host organisms selectively attract such epibionts.

With the advancement of molecular biological tools to analyse the diversity and abundance of bacteria in biofilms (Dahllöf 2002), several studies have demonstrated that quantitative and qualitative bacterial occurrence on host organisms differs significantly from inanimate reference surfaces (Harder et al. 2003, 2004b; Lee and Qian 2004; Dobretsov et al. 2006; Rao et al. 2005). These findings firstly suggest strong host defence mechanisms against non-culturable epibiotic bacteria but also support the notion of potent effects of non-culturable epibiotic bacteria against subsequent colonizers of host organisms. It will be interesting to see follow-up studies that utilize advanced molecular biological tools, such as cloning techniques, to directly test the metabolites of non-culturable symbiotic prokaryotes on epibiotic eukaryotes.

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