

Drug development from marine natural products

Tadeusz F. Molinski*, Doralyn S. Dalisay*, Sarah L. Lievens** and Jonel P. Saludes**†

Abstract | Drug discovery from marine natural products has enjoyed a renaissance in the past few years. Ziconotide (Prialt; Elan Pharmaceuticals), a peptide originally discovered in a tropical cone snail, was the first marine-derived compound to be approved in the United States in December 2004 for the treatment of pain. Then, in October 2007, trabectedin (Yondelis; PharmaMar) became the first marine anticancer drug to be approved in the European Union. Here, we review the history of drug discovery from marine natural products, and by describing selected examples, we examine the factors that contribute to new discoveries and the difficulties associated with translating marine-derived compounds into clinical trials. Providing an outlook into the future, we also examine the advances that may further expand the promise of drugs from the sea.

In 1967, a small symposium was held in Rhode Island, USA, with the ambitious title “Drugs from the Sea”¹. The tone and theme of the meeting were somewhat hesitant, even sceptical (one paper was entitled “Dregs [sic] from the Sea”). The catchphrase of the symposium title has endured over the decades as a metaphor for drug development from marine natural products, and in time genuine drug discovery programmes quietly arose to fulfil that promise. The remarkably high hit rates of marine compounds in screening for drug leads were often cited as justification for the search and, indeed, notable discoveries were made of compounds that profoundly affect the cell cycle and cellular metabolism. If anything, marine natural-product chemistry has built a legacy of discovery of biomedical probes; for example, okadaic acid (a phosphatase inhibitor produced by dinoflagellates) and xestospingonin C (an intracellular blocker of calcium release from a marine sponge). However, the first genuine drug from the sea was a long time coming.

Natural products, especially those from terrestrial plants and microbes, have long been a traditional source of drug molecules (for example, morphine from poppies, cardiotoxic digitalis glycosides from foxgloves and penicillins from fungi). Modern pharmaceutical discovery programmes owe much to natural products. Indeed, pharmacologically active compounds from plants and microbes represent an important pipeline for new investigational drugs^{2–4}. Interest in marine natural products, however, awaited refinements in technologies (mainly scuba diving) to collect the source organisms. Even so, by the late-1950s, the concept of drugs from the sea had

attracted some interested. Beginning in 1951, Werner Bergmann published three reports^{5–7} of unusual *arabino-* and *ribo-*pentosyl nucleosides obtained from marine sponges collected in Florida, USA. The compounds eventually led to the development of the chemical derivatives ara-A (vidarabine) and ara-C (cytarabine), two nucleosides with significant anticancer properties that have been in clinical use for decades.

The role of natural products in drug discovery has undergone many changes in the past 30 years, with a noticeable decline in participation by the major pharmaceutical companies by the mid-1990s. Nevertheless, enterprising academics, mainly partnered with industry, exploited the niche left by larger research and development efforts. The field seems to have benefited from a renaissance in the past 5 years. This has partly been driven by new developments in analytical technology, spectroscopy and high-throughput screening^{8,9}, and partly driven by a broad realization that competing technologies, such as combinatorial chemistry, have failed to deliver new drug leads in significant numbers. In the meantime, basic scientific research in chemistry and pharmacology of marine natural products, and directed efforts in drug development — begun in the 1970s — have finally borne fruit for marine-based drug discovery. The first drug from the sea, ziconotide (ω -conotoxin MVIIA) — a peptide originally from a tropical marine cone snail — was approved in the United States in 2004 under the trade name Prialt for the treatment of chronic pain in spinal cord injury. A second drug — the antitumour compound trabectedin

*Department of Chemistry and Biochemistry and Skaggs School of Pharmacy and Pharmaceutical Sciences, University of California, San Diego, 9500 Gilman Drive, MC 0358, La Jolla, California 92093, USA.

†Department of Chemistry, University of California, One Shields Avenue, Davis, California 95616, USA. Correspondence to T.F.M. e-mail: tmolinski@ucsd.edu doi:10.1038/nrd2487 Published online 19 December 2008

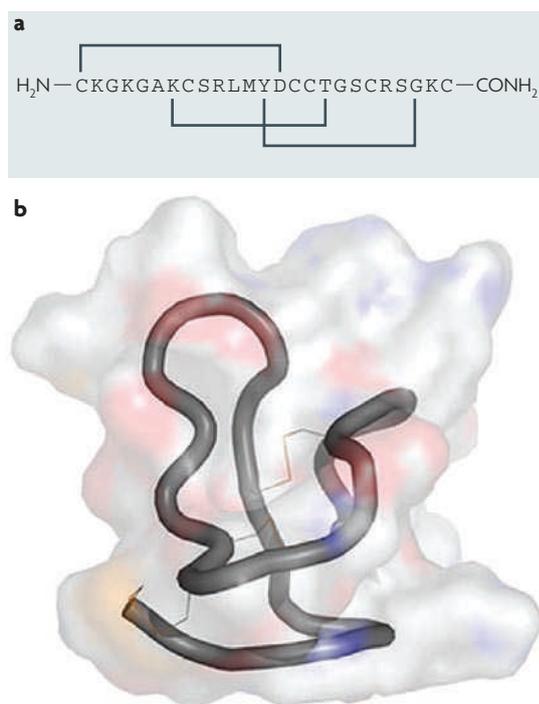


Figure 1 | ω -Conotoxin VIIA. a | Amino-acid sequence of the peptide ω -conotoxin MVIIA (ziconotide/Prialt; Elan Pharmaceuticals). **b** | Three-dimensional structure of the synthetic ω -conotoxin VIIA polypeptide. The cylinder represents the amide backbone of ω -conotoxin VIIA overlaid against an electrostatic potential surface. For a three-dimensional representation of ω -conotoxin MVIIA see the entry Ziconotide @ 3Dchem.com (see Further information).

(Yondelis/ecteinascidin-743/ET-743) from a tropical sea-squirt — was approved by the European Union in October 2007 for the treatment of soft-tissue sarcoma. Several other candidate compounds are presently in the pipeline, and marine natural products are being evaluated in Phase I–III clinical trials in the United States and in Europe for the treatment of various cancers. A recent tabulation of these compounds and their status can be found in REFS 2–4.

The difficulty of developing drugs from a marine source was obvious from the outset: the procurement or manufacture of quantities of rare compounds from marine sources to ensure a sustainable supply was a bottleneck. For example, the chemically versatile marine sponges, the source of many developmental compounds such as discodermolide and hemiasterlin, are primitive metazoans that live almost exclusively in marine habitats. Sponges and their microbial fauna are largely unculturable, and the valuable compounds they produce must be extracted and purified from specimens collected by hand using scuba diving from shallow to deep waters, or sometimes with the aid of submersibles equipped with robotic arms. Both of these techniques are expensive endeavours that are unwieldy and foreign to the modern pharmaceutical industry. Nevertheless, interest in the remarkable properties of marine natural

products remained high enough so that inspired innovative solutions to the supply problem were proposed on a case-by-case basis, ranging from aquaculture¹⁰ of marine invertebrates to semi-synthesis.

In this Review, selected examples serve to illustrate contemporary drug discovery and development of marine natural products — some successful, some less so. Each story begins with the discovery of a novel molecule from a marine organism, and their progression through pre-clinical and clinical trials (for oncology indications, with the exception of ω -conotoxin MVIIA) — mostly as the unmodified structure or as a synthetic compound modelled after the natural product. It is not intended to be comprehensive account, but to illustrate the challenges associated with moving marine-derived compounds into clinical trials. Finally, we provide an outlook to some truly revolutionary new developments in genome mining that are expected to change the way that marine-derived drugs are discovered in the future.

ω -Conotoxin MVIIA

After more than two decades of research and development, ziconotide, a synthetic form of ω -conotoxin MVIIA, became the first marine-derived drug approved by the US Food and Drug Administration (FDA). Now known under the trade name Prialt, it is approved for the specific indication of chronic pain. The venom of fish-hunting cone snails (genus *Conus*) contains a myriad of toxic peptides (conotoxins) that act synergistically to immobilize prey by targeting the neuromuscular system. The chemistry and biology of these conotoxins have been extensively reviewed^{11,12}. ω -Conotoxin MVIIA (FIG. 1a), from the Pacific piscivorous marine snail *Conus magus* (FIG. 2a) has been an exceptional lead for drug development in the management of severe and chronic pain. It was originally discovered by the group of Olivera at the University of Utah, USA, in 1979 via a bioassay that elicited a characteristic shaking behaviour in mice after the peptide was administered by intracerebral injection¹¹. ω -Conotoxin MVIIA is a linear 25 amino acid, polycationic peptide containing six cysteine residues linked by three disulphide bridges that stabilize its well-defined three-dimensional structure^{13–15}. Its complete chemical synthesis was achieved in 1987, and N-type voltage-sensitive calcium channels (NVSCCs) were subsequently identified as its target site¹⁶. It potently inhibits the conduction of nerve signals (K_i value of 0.5 μM)¹⁷ by specifically blocking the NVSCC. In the complex with NVSCC, it forms a compact folded structure with a binding loop between Cys8 and Cys15 that also contains Tyr13, an important amino-acid residue located at the binding site^{18,19}.

NVSCCs are found exclusively in presynaptic neurons where they regulate depolarization-induced calcium influx, which subsequently control a variety of calcium-dependent processes. NVSCCs are abundantly present in the superficial lamina of the spinal-cord dorsal horn, where they have an important role in the spinal processing of nociceptive afferent (pain signalling) activity^{20,21}. The potent inhibition and highly selective affinity of ω -conotoxin MVIIA to NVSCC (K_d value of 9 pM)²² attracted interest to develop this peptide into an antinociceptive agent.



Figure 2 | Marine invertebrates producing anticancer and analgesic drugs.
a | Textile cone snail *Conus magus*. **b** | The Caribbean sea-squirt *Ecteinascidia turbinata*. **c** | The Caribbean sea-squirt *Trididemnum solidum*. **d** | The sea hare *Dolabella auricularia*. **e** | The sacoglossan *Elysia rufescens* feeding on the red alga *Bryopsis* spp. **f** | The bryozoan *Bugula neritina*. Panel **a**, image reproduced with permission from Elan Pharmaceuticals, USA. Panel **b**, image courtesy of S. Lopez-Legintil, University of North Carolina, USA. Panel **d**, image reproduced with permission by W. B. Rudman, *Sea Slug Forum*, Australian Museum, Sydney, Australia. Panel **e**, image reproduced with permission by H. Flodrops, *Sea Slug Forum*, Australian Museum, Sydney, Australia. Panel **f**, image reproduced with permission by the California Academy of Sciences, USA.

Synthetic ω -conotoxin MVIIA, ziconotide (FIG. 1b), was developed and brought to the clinic by Neurex (a subsidiary of Elan Pharmaceuticals)²³. Preliminary studies demonstrated that ziconotide has a remarkable potent antinociceptive profile in animal models with acute, persistent and neuropathic pain after intrathecal administration²⁴. In a rat incisional model of post-operative pain, ziconotide demonstrated more potent (ED_{50} value of 49 pM) and longer activity than intrathecal morphine (ED_{50} value of 2.1 nM)²⁵.

The promising analgesic activity observed in animal studies enabled ziconotide to enter clinical trials in the United States and in Europe for the treatment of severe chronic pain. Its clinical development has been reviewed

recently²⁶. Ziconotide was administered to subjects by continuous infusion using an external or an implanted pump via an intrathecal catheter because of its poor tissue penetration and hypotensive effect when administered systemically.

On 22 December 2004, Elan Pharmaceuticals was finally granted FDA approval for ziconotide (an intrathecal infusion formulation) under the trade name Prialt. Two months later the European Commission approved ziconotide for the treatment of severe, chronic pain in patients who require intrathecal analgesia²⁷. The approval of this synthetic form of ω -conotoxin MVIIA offers hope for several conotoxins that are currently under investigation²⁸ for therapeutic potential and possible clinical usage.

Ecteinascidin-743

Ecteinascidin-743 (ET-743/trabectedin) (FIG. 3a) illustrates a significant milestone in the development of marine-derived drugs. Almost 40 years after its discovery and 17 years after the publication of its structure, it became the first marine-derived anticancer drug to reach the market. Extracts of the Caribbean tunicate *Ecteinascidia turbinata* (FIG. 2b) have been reported to possess anti-tumour activity as early as 1969 (REF. 29); however the minute amounts of active component obtained from laborious isolation work precluded their identification for almost two decades. Finally, in 1990, Rinehart³⁰ and Wright³¹ published the structure of the alkaloids named ecteinascidins. The structure of the most abundant active component, ET-743, and its *N*-demethyl analogue ET-729 (FIG. 3a), is comprised of three fused tetrahydroisoquinoline rings and is related to the simpler isoquinoline alkaloid saframycin A from various *Streptomyces* spp. The connection of the third tetrahydroisoquinoline ring to the base structure by a thioether bridge completes a 10-membered lactone — a distinctive structural feature of ecteinascidins. Initially, ecteinascidins were found to be cytotoxic against L1210 leukaemia cells (IC_{50} value of 0.5 ng per ml)³⁰ and were later shown to possess strong *in vivo* antitumour effects in various mice models bearing P388 lymphoma, B16 melanoma, M5076 ovarian sarcoma, Lewis and LX-1 human lung carcinoma, and MX-1 human mammary carcinoma xenografts^{30,32}. Further studies comparing ET-743 and ET-729 indicated that they have similar potency. Subsequently, ET-743 was selected for further development because it is more abundant than ET-729 in *E. turbinata*.

Still, the yield for ET-743 from the tunicate is very low (~10 parts per million). To provide more material for *in vitro* and *in vivo* animal studies, the first multi-step synthesis of the compound was completed in 1996 (0.75% yield)³³, followed by more efficient methods³⁴. Although total synthesis is not practicable to provide sufficient material for clinical use, this exercise led to the novel synthetic derivative phthalascidin, which has an antiproliferative property comparable to that of ET-743 (REF. 34). Large-scale coastal aquafarming of *E. turbinata* in Europe provided a larger supply of the drug source, but yields were variable. A breakthrough was achieved by PharmaMar, the licensee of natural

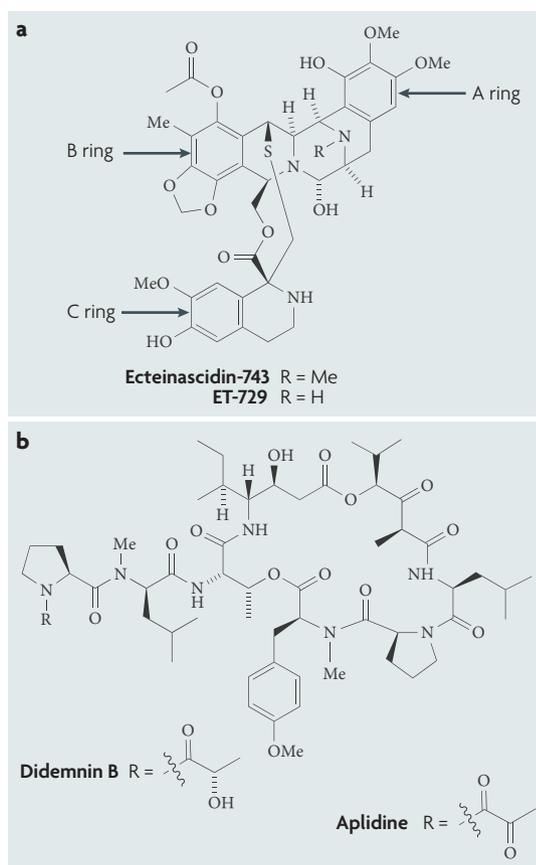


Figure 3 | Tunicate-derived anticancer drugs.
a | Ecteinascin-743 (ET-743/trabectedin; marketed under the trade name Yondelis by PharmaMar/Johnson & Johnson/OrthoBiotech), derived from *Ecteinascidia turbinata*, and its analogue ET-729. **b** | Didemnin B from *Trididemnum solidum* and aplidine (also known as dehydrodidemnin B) from *Aplidium albicans*.

ET-743, who developed a large-scale semi-synthetic method that starts with cyanosafraicin B, an antibiotic that can be produced in multi-kilogram scale by fermentation of *Pseudomonas fluorescens*³⁵. ET-743 was licensed by PharmaMar to Johnson & Johnson/OrthoBiotech for drug development in the United States³⁶.

The mechanism of action of ET-743 is ascribed to covalent modification of DNA by guanine-specific alkylation at the N2 position, a property similar to another minor-groove alkylating agent, anthramycin. In contrast to anthramycin, ET-743 is selective for GC-rich sequences and forms an adduct with duplex DNA, which is reversible upon denaturation³⁷, and induces a bend in the DNA helix directed towards the major groove³⁸. Molecular dynamics calculations shows protrusion of ring C of ET-743 into the minor groove and interference with DNA-binding factors³⁹ (FIG. 4). This partly accounts for the potency of ET-743 compared with other N2 alkylating agents such as saframycin A and anthramycin that have smaller groups corresponding to ring C. ET-743 also affects transition-coupled nucleotide excision repair and triggers cell death^{39,40}.

ET-743, under the trade name Yondelis, was approved for the treatment of refractory soft-tissue sarcomas by the European Commission in July 2007. Results of Phase II trials of ET-743 for Ewing's sarcoma and soft-tissue sarcomas⁴¹, colorectal cancer⁴², pretreated advanced breast cancer⁴³, ovarian cancer⁴⁴, and other sarcomas^{45,46} have been disclosed. Phase II studies in breast cancer and advanced prostate cancer are ongoing. Concluded Phase I and II studies demonstrate that ET-743 has remarkable antitumour activity against solid tumours, in particular breast cancer and renal carcinoma, and soft-tissue sarcomas (particularly osteosarcomas, mesothelioma, leiomyosarcoma and liposarcoma). A review of these studies has been published elsewhere⁴⁷. Gastrointestinal stromal tumours (a subclass of soft-tissue sarcomas) were found to be non-responsive to ET-743 treatment^{48,49}. Preclinical studies of ET-743 combination treatments with doxorubicin⁵⁰, paclitaxel⁵¹ or irinotecan⁵² against sarcomas revealed synergistic effects between the drugs, as have Phase II combination drug studies with doxorubicin⁵⁰. Toxicities that became dose limiting include neutropaenia and thrombocytopaenia. Hepatotoxicity, previously observed in preclinical studies, was observed in humans but could be controlled by dose adjustment. Recent findings using HepG2 human hepatocellular liver carcinoma cells revealed a cytochrome P450-mediated metabolism of ET-743 (REF. 53). It was demonstrated in rats that pretreatment with metabolism modulators such as dexamethasone and β -naphthoflavone abrogates ET-743-mediated hepatotoxicity⁵⁴, and dexamethasone-ET-743 combination drug treatment was recommended for investigation in humans^{55,56}.

Didemnin B and aplidine

ET-743 was not the only lead anticancer agent found from marine ascidians. Two closely related compounds from different organisms — didemnin B from the tropical *Trididemnum solidum* (FIG. 2c) and aplidine from *Aplidium albicans* collected in temperate Mediterranean waters — have been extensively investigated for more than 20 years; although clinical trials for didemnin B were stopped in the mid-1990s (see below). The chemical differences between the two compounds are minimal: they are both cyclic peptides in which didemnin B has a pyruvyl amide residue at the secondary amine of the proline residue and aplidine has a pyruvyl amide at this position (FIG. 3b). Nevertheless, the *in vitro* cytotoxicities and preclinical profiles seemed to be sufficiently different such that new clinical trials have been warranted for aplidine.

Didemnin B^{47,57-59} was first isolated by the Rinehart group in 1981 from the tunicate *T. solidum*⁶⁰ and displayed antiviral and *in vivo* cytotoxic activities at nanomolar concentrations⁶¹. The original structure was revised in 1987 as a result of total synthesis of the first proposed structure, which revealed inconsistencies: the correct structure replaces the non-proteinogenic amino-acid residue statine with *isostatine*⁶². Complete spectroscopic characterization of didemnin B was reported^{63,64} along with a definitive single-crystal X-ray structure published in 1988 (REF. 65).

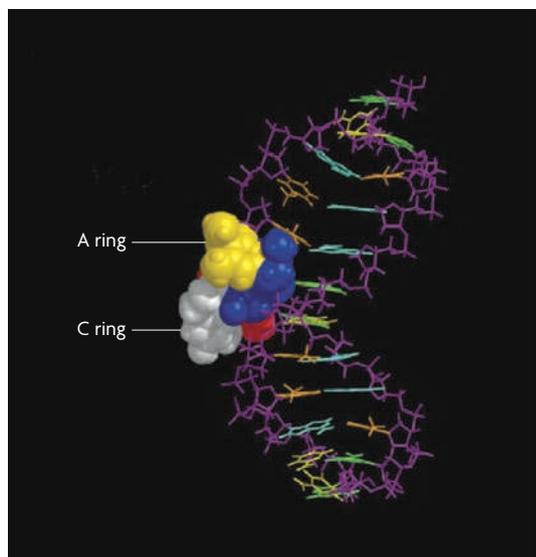


Figure 4 | **Molecular-dynamics model showing the alkylation of DNA by ET-743 at N2 of guanine in the minor groove.** The A Ring and C Ring represent the tetrahydroisoquinoline A and C rings of ET-743 (FIG. 3). Image is modified, with permission, from REF. 39 © (2001) Elsevier Science.

Precise details of the mechanism of action of didemnin B were difficult to pinpoint. Early studies showed that didemnin B inhibits palmitoyl protein thioesterase in a non-competitive manner^{66,67}; although this low-affinity target did not fully account for the nanomolar cell inhibition observed with the natural product. Didemnin B induces inhibition of protein synthesis at a concentration that is commensurate with cell growth inhibition⁶⁸. This is achieved by stabilization of aminoacyl-tRNA and prevention of EF-2-dependent translocation on the ribosome^{69–72}. However, inhibition of protein synthesis does not seem to be the primary cause of apoptosis^{73,74}. Apoptosis induced by didemnin B is dependent on protein tyrosine kinases and can be inhibited using protein tyrosine kinase inhibitors⁷⁵ or rapamycin, possibly through the interaction of rapamycin with the immunophilin FKBP25 (REF. 76).

As early as 1983, didemnin B was shown to be active against herpes simplex virus⁷⁷ and later against Ehrlich's carcinoma⁷⁸ in mice. Early cancer trials showed that didemnin B had minimal activity and demonstrated the requirement for co-treatment with anti-emetics⁷⁹. *In vitro* testing established that didemnin B was active against colorectal⁸⁰, lymphatic⁸¹ and prostate⁸² cancers. Consequently, didemnin B was submitted to several Phase I^{79,83,84} and Phase II clinical trials against previously treated non-small cell lung cancer (NSCLC)⁸⁵, breast cancer⁸⁶, small-cell lung cancer⁸⁷, non-Hodgkin's lymphoma^{88,89}, metastatic melanoma⁹⁰, glioblastoma multiforme⁹¹, and CNS tumours⁹². However, these trials resulted in significant neuromuscular toxicity and no objective responses. Nevertheless, didemnin B showed activity in patients with advanced pretreated non-Hodgkin's lymphoma, but trials were suspended owing

to onset of severe fatigue in patients⁸⁹. Other trials showed a high incidence of anaphylaxis, and were therefore terminated⁹³. All current trials of didemnin B are on hold.

While trials with didemnin B are suspended, a simple analogue of didemnin B, aplidine (dehydrodidemnin B), seems more promising. Aplidine was first reported in a 1991 patent⁹⁴ by Rinehart and is obtained from the Mediterranean tunicate *A. albicans*. Aplidine differs from didemnin B only in replacement of the *N*-lactyl side chain with a pyruvyl group. Aplidine shows similar levels of antitumour activity to didemnin B in cultured tumour cells⁹⁵, and has been shown to induce apoptosis by induction of oxidative stress⁹⁴, which triggers the pro-apoptotic receptor Fas (CD-95)⁹⁵ and induces mitochondrion-mediated apoptosis^{96,97}. Aplidine also activates p38 mitogen-activated protein kinases (MAPKs) and JNK^{98,99}, and inhibits secretion of vascular endothelial growth factor (VEGF)^{101,101}. Non-P-glycoprotein-expressing cell lines that are resistant to aplidine were shown to display temporary phosphorylation of JNK and p38 MAPKs upon exposure to aplidine, and the short duration of activation was insufficient to trigger apoptosis¹⁰². In relapsed–recalcitrant leukaemia cell lines, aplidine arrests the cell cycle at the G1 and G2/M phases, and induces p53-independent apoptosis¹⁰³.

Significant differential *in vitro* cytotoxicity of aplidine in primary cultured lymphocytes and in transformed cell lines was observed, which may explain why both didemnin B and aplidine show minimal haemotoxicity *in vivo*^{104–107}.

Aplidine has also been shown to inhibit angiogenesis in chick embryos *in vivo*¹⁰⁸, and to selectively disrupt β -sheet fibrils caused by prion protein sequence PrP 106–126 in a 1:1 molar ratio¹⁰⁹. Phase II clinical trials with aplidine are ongoing in indications that include metastatic melanoma, multiple myeloma, non-Hodgkin's lymphoma, acute lymphoblastic leukaemia, prostate cancer and bladder cancer.

Significant challenges have been encountered during the development of didemnin B and aplidine. The supply of these two peptides from nature is limited by the difficulties of collecting sparsely distributed source organisms and the lack of feasible aquaculture conditions. Consequently, supply is entirely dependent upon multi-step total synthesis of these peptides from constituent amino acids, some of which are not found in proteins. Inventive solutions to the supply problem of these and other marine natural products with therapeutic properties may ultimately determine their future use, even if clinical outcomes are favourable.

Antimitotic marine natural products

Several antitumour marine natural products, derived mainly from marine sponges or molluscs but also bryozoans and cyanobacteria, exhibit potent antimitotic properties. Several have advanced to Phase I and II clinical trials and, perhaps not coincidentally, many such compounds block progression of dividing cancer cells through M phase by targeting the same protein (tubulin) in ways similar to the clinical drugs vincristine, vinblastine and paclitaxel. Although none has yet been approved

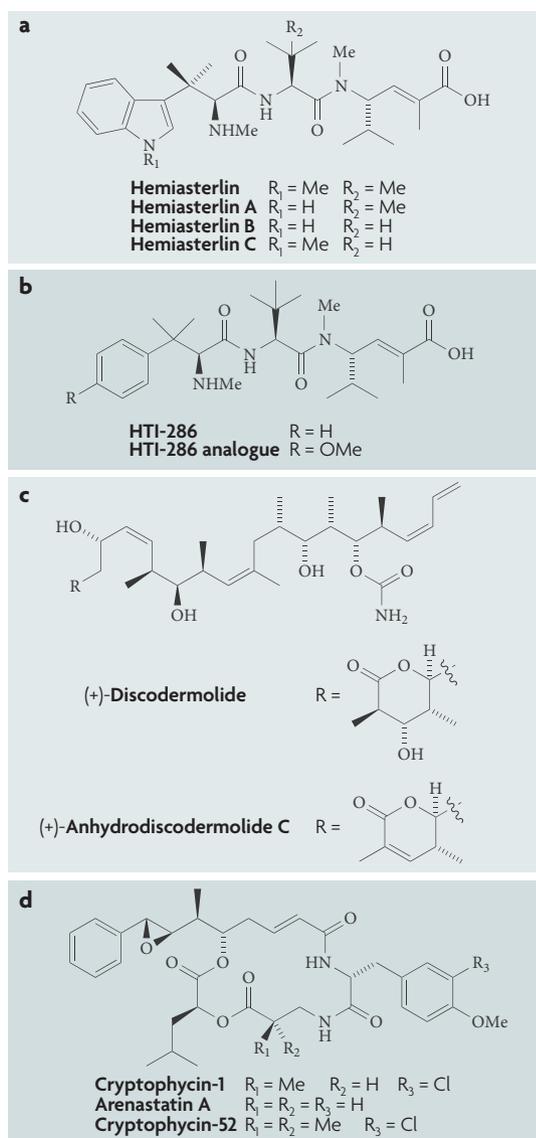


Figure 5 | Marine-derived antimitotic compounds (part 1). **a** | The hemiasterlins. All hemiasterlins showed cytotoxicity in the nanomolar range (concentrations $\sim 1 \times 10^{-9}$ M) against a variety of cultured human and murine cell lines. **b** | Hemiasterlin analogues, HTI-286 (SPA-110/taltobulin) and its analogue. **c** | The discodermolides, (+)-discodermolide and (+)-anhydrodiscodermolide. **d** | The cryptophycins.

as a drug, the following examples of natural products are illustrative of the wide range of chemotypes that modulate tubulin dynamics. Important differences have been reported in their mechanisms of action, and this in itself has been tremendously informative for understanding the complex interaction of microtubule proteins with antimitotic drugs.

Hemiasterlin/HTI-286. Hemiasterlin, an antimitotic tripeptide (FIG. 5a), was first isolated in 1994 by Kashman and co-workers from the sponge *Hemiasterella minor*¹¹⁰. Initial screening showed that it was highly cytotoxic with

an IC_{50} value of ~ 0.01 μg per ml against the P388 leukaemia cell line¹¹⁰. The related isomers hemiasterlin A and B, reported by Andersen and co-workers from sponges of the genus *Auletta* and *Cymbastella* in 1995 (REF. 111), were even more potent. All hemiasterlins showed cytotoxicity in the nanomolar range (concentrations $\sim 1 \times 10^{-9}$ M) against a variety of cultured human and murine cell lines¹¹¹. In 1996, an X-ray crystal structure analysis of the hemiasterlin methyl ester confirmed the linear structure of hemiasterlin and its unusual constituent amino acids¹¹². A fourth analogue, hemiasterlin C, was described in 1999 (REF. 113).

The potent cytotoxicity of hemiasterlins is due to the induction of mitotic arrest in metaphase with cellular dynamics similar to those of known tubulin binders, such as the chemotherapeutics paclitaxel or vinblastine, at half-maximal effective doses (ED_{50} values) that ranged from 0.5 nM (hemiasterlin) to 28 nM (hemiasterlin B)¹¹⁴. Hemiasterlin A was shown to interfere with mitotic spindle formation at low concentrations and causes tubulin depolymerization at higher concentrations (with higher efficacy than paclitaxel or vinblastine¹¹⁴). By contrast, hemiasterlin was shown to bind to the vinca peptide binding site of β -tubulin, where it competitively inhibits binding of the marine compound dolastatin 10 and non-competitively inhibits binding of vinblastine^{110,111}. Interestingly, upon binding to tubulin, hemiasterlin induces the formation of small ring oligomers¹¹⁵.

Extensive structure–activity relationship studies demonstrated that the simpler synthetic analogue of hemiasterlin, HTI-286 (SPA-110/taltobulin) (FIG. 5b), with a phenyl substituent replacing the *N*-methyltryptophan, is more potent than hemiasterlin¹¹². An analogue of HTI-286 (FIG. 5b) with a *para*-methoxyl substituent on the benzene ring was even more potent¹¹³. Other structural elements, including the geminal β,β -dimethyl group and the *N*-methyl on the first amino-acid residue (N terminus), the isopropyl and an olefin in the homologated γ -amino acid (C terminus), including a terminal carboxylic acid or methyl ester, were essential for activity. The aryl side chain on the N terminus could be replaced synthetically by alkyl groups (for example, *tert*-butyl), while still retaining potent activity^{116–120}.

The stoichiometry of HTI-286 binding to an α/β -tubulin monomer was found to be one to one¹²¹. Binding of HTI-286 to tubulin is rapid and exothermic, but formation of oligomers is slower. Disruption of tubulin dynamics by hemiasterlin and HTI-286 takes place at low concentrations, but a full equivalent of peptide is required for depolymerization of microtubules¹²². Molecular modelling suggested that binding of both compounds takes place at the vinca peptide site of β -tubulin, near the α/β interface that is also targeted by the marine-derived tubulin inhibitors cryptophycin 1, dolastatin 10 and the fungus-derived phomopsin A¹²³. Radiolabelling of protein with a photoaffinity analogue of hemiasterlin, however, indicated an exclusive binding site on α -tubulin near the α/β interface¹²⁴.

Unlike paclitaxel or vinblastine, hemiasterlin and HTI-286 are poor substrates for P-glycoproteins and maintain toxicity towards cell lines with high expression of

multidrug resistant (MDR) drug pumps. Cells resistant to either compound do not express high levels of known MDR proteins; instead it was shown that mutations on both α -tubulin and β -tubulin subunits confer resistance in most cases¹²⁵, although one study showed moderately reduced drug accumulation due to an unidentified ATP-drug pump¹²⁶. The mutations on tubulin are not localized on the vinca peptide binding site, and do not seem to affect drug binding, but instead increase microtubule stability causing cross-resistance (3-fold to 186-fold) to other depolymerizing agents and sensitization (2-fold to 14-fold) to tubulin polymerizing agents¹²⁵.

Preclinical studies showed that HTI-286 causes tumour regression and growth inhibition of human xenografts in mice. Even cell lines expressing P-glycoprotein or resistant to paclitaxel were shown to be sensitive to HTI-286 inhibition, but required higher doses than non-resistant cell lines¹²⁷. An open-label Phase I clinical trial of HTI-286 was completed in patients with advanced solid tumours; however, there were no objective responses and common toxicities observed included neutropaenia, nausea, alopecia and pain¹²⁸. Phase II trials have been halted. Nevertheless, there is still interest in HTI-286 in view of recent results including high antitumour activity in androgen-dependent and androgen-independent mouse models of refractory prostate cancer, and in a newly established *in vitro* taxane-resistant prostate PC-3 cell line¹²⁹.

(+)-Discodermolide. (+)-Discodermolide¹³⁰ (FIG. 5c) was first isolated in 1990 by Gunasekera and co-workers¹³¹ at the Harbor Branch Oceanographic Institute, Florida, USA, from the rare deep-water sponge *Discodermia dissoluta*. This sponge is found only in the Bahamas at depths of up to 300 m, although recent evidence suggests it also occurs in shallower waters. Discodermolide functions as an immunosuppressant^{132,133}, and induces G2/M phase cell-cycle arrest in lymphoid and non-lymphoid cells at nanomolar concentrations¹³⁴. The structure of (+)-discodermolide was first elucidated by analysis of nuclear magnetic resonance and mass-spectrometric data^{135,136}, and its relative stereochemistry was provided by X-ray crystal structure analysis¹³⁵. Significant differences in activity were found for the two enantiomers as shown by analysis of the unnatural antipode (–)-discodermolide prepared by total synthesis¹³⁷. (–)-Discodermolide arrests cell cycle *in vitro*, albeit at higher concentrations (72–135 nM) compared with the natural enantiomer (3–80 nM). Although both enantiomers block cell cycle at S phase, their effects are not mutually competitive¹³⁸. Subsequent synthesis produced the natural enantiomer (+)-discodermolide¹³⁹. Owing to the exceedingly limited natural supply of (+)-discodermolide, efforts focused on developing efficient methods to synthesize the compound, and several inventive multi-step, scaled-up processes for total syntheses were developed^{139–141}. These included a gram-scale synthesis^{142–144} and a heroic 60 g scale preparation by workers at Novartis that provided material for Phase I clinical trials^{145–149}.

Investigations into the mechanism of cell-cycle arrest by (+)-discodermolide showed that the compound stabilized microtubules competitively with paclitaxel,

with higher affinity for tubulin than paclitaxel^{150,151}. (+)-Discodermolide is a poor substrate for P-glycoprotein and is cytotoxic to cells with tubulin mutations that confer resistance to paclitaxel¹⁵². Interestingly, despite their mutual competitive inhibitory activity, (+)-discodermolide and paclitaxel show strongly synergistic activity^{153–155}. The cytotoxicity of (+)-discodermolide is apparent at concentrations too low to cause cell-cycle arrest, in which aberrant mitosis¹⁵⁶, altered induction of apoptosis¹⁵⁷ and a significant alteration of microtubule dynamics¹⁵⁸ can be observed. Similar to paclitaxel, (+)-discodermolide interferes with binding of tau protein to microtubules¹⁵⁹ and shows microtubule-disruption-dependent down-regulation of hypoxia-inducible transcription factor 1 α (HIF1 α)¹⁶⁰. However, as opposed to paclitaxel, (+)-discodermolide induces accelerated senescence in A549 NSCLC cells and even a strain of A549 that has higher resistance to the compound by activating the signalling molecules plasminogen activator inhibitor, type I (PAI1; also known as SERPINE1), p66Shc and extracellular signal-regulated kinases 1 and 2 (ERK1/2)¹⁶¹.

(+)-Discodermolide has further shown *in vivo* activity against HCT-116 colorectal cancer xenografts in mice. Co-treatment of mice with (+)-anhydrodiscodermolide C (FIG. 5c) and bacteriolytic treatment with *Clostridium novyi*-NT spores resulted in a rapid, complete cure of four out of five animals with a single injection¹⁶². A Phase I clinical study of (+)-discodermolide showed no neuropathy or neutropaenia, and demonstrated mild-to-moderate toxicity from 0.6 mg per m² to 19.2 mg per m². The pharmacokinetics of (+)-discodermolide were shown to be non-linear with recycling of (+)-discodermolide between tissues and the circulatory system¹⁶³. At present, Novartis has discontinued Phase I trials with (+)-discodermolide owing to lack of efficacy and toxicity problems. However, potential remains for its use in combination drug therapy.

Cryptophycins. Cryptophycins are examples of marine cyanobacteria-derived tubulin-binding compounds, which, owing to extensive academic–industry collaborations, inspired the design of several synthetic analogues. Cryptophycin A was first isolated by Schwartz and co-workers in 1990 (REF. 164) from the cyanobacteria *Nostoc* spp. strain ATCC 53789. It was found to be extremely potent in mice infected with *Cryptococcus* spp. but was not pursued because of its notable toxicity. Subsequently, it was re-isolated by Moore's group in Hawaii, USA, in 1994 along with several derivative congeners from a cultured freshwater *Nostoc* spp. strain GSV 224 (REF. 165). The name was later changed to cryptophycin-1 (Crp-1; FIG. 5d). Crp-1 showed IC₅₀ values of 3 pg per ml and 5 pg per ml against KB carcinoma and LoVo adenocarcinoma cells, and was active *in vivo* against xenografts of colon adenocarcinoma 38 and 51 cell lines, paclitaxel-sensitive and paclitaxel-resistant mammary adenocarcinoma M16 cells, and in mouse models of pancreatic ductal adenocarcinoma 03 (REF. 165). The related cytotoxic compound arenastatin A (FIG. 5d), isolated from the marine sponge *Dysidea arenaria*, was reported contemporaneously by Kobayashi¹⁶⁶, and turned

the high antimetabolic activity of the peptide (IC_{50} value of 1.2 μ M) is its prolonged intracellular retention that subsequently facilitates binding to tubulin¹⁸⁵.

Dolastatin 10 demonstrated *in vitro* inhibitory activity against various human cancer cell lines, including melanoma, sarcoma and ovarian cancer cells¹⁷⁹, and entered several Phase I and II clinical trials as a single agent. Phase I clinical trials on dolastatin 10 identified a maximum tolerated dose of 300–400 μ g per m^2 for patients with advanced solid tumours, and documented granulocytopenia as its major dose-limiting toxicity^{186,187}. Dolastatin 10 progressed to Phase II clinical trials as a single agent against prostate adenocarcinoma¹⁸⁸, NSCLC¹⁸⁹, melanoma¹⁹⁰, colorectal cancer¹⁹¹, soft-tissue sarcomas¹⁹², breast cancer¹⁹³, and pancreaticobiliary cancers¹⁹⁴. Although the toxicity profile of dolastatin 10 was acceptable, the results were disappointing as no clinically significant activity could be detected in any of the trials. The negative results of Phase II clinical trials did not justify continuation and dolastatin 10 was withdrawn from antitumour clinical trials¹⁹⁵. Nevertheless, the new synthetic derivative TZT-1027 (REF. 196) (FIG. 6a) is currently being evaluated in Phase I clinical trials in Japan, Europe and the United States^{197,198}. TZT-1027, which differs from dolastatin 10 in replacement of the terminal dolaphenine amino-acid residue with a simple phenethylamine group, is a potent microtubule and vascular blockade inhibitor¹⁹⁹, and has anticancer activity in several murine cancer models¹⁹⁶. TZT-1027 and dolastatin 10 share the same mechanism of activity: they both bind near the vinca peptide site on tubulin and inhibit tubulin polymerization with equal potency (IC_{50} values of 2.2 μ M and 2.3 μ M, respectively). Dolastatin 10 completely displaces tubulin-bound TZT-1027 (REF. 200).

Halichondrin B and eribulin mesylate. The discovery and development of the antitumour compounds halichondrin B and its analogue eribulin mesylate is an interesting account of multidisciplinary approaches that brought a promising drug candidate to patients with cancer while addressing the perennial issue of supply that limits procurement of anticancer natural products from marine invertebrates. In 1986, Uemura and co-workers discovered halichondrin B, norhalichondrin B and homohalichondrin B (FIG. 6b) from the minor fractions of extracts of *Halichondria okadai*, the same sponge that also sequesters the phosphatase inhibitor okadaic acid^{201,202}. Subsequently, halichondrin B and other halichondrins, which occur in very small concentrations (for example, halistatin 1 (REF. 203), 8.8 parts per billion; FIG. 6b) were reported from unrelated sponges: *Axinella* spp. from the western Pacific Ocean²⁰⁴ and *Phakellia carteri* from the East Indian Ocean²⁰⁴. The halichondrins were shown to have potent cell growth inhibitory activity at nanomolar concentrations ($\sim 1 \times 10^{-9}$ M)²⁰⁴. Differential cytotoxicity data indicated that halichondrin B binds tubulin at the vinca peptide binding site^{205,206}, similar to other compounds covered in this Review. The total syntheses of halichondrin B and norhalichondrin B were reported in 1992, and required approximately 90 steps from commercially available raw materials²⁰⁷. Following

favourable *in vitro* and *in vivo* data, the US National Cancer Institute advanced the compound (NCI number NSC 60935) to Phase I clinical trials in 2002.

A new source of halichondrins was discovered in a new species of deep-water sponge, *Lissodendoryx* n. sp. 1 (abbreviated here as *Lissodendoryx*), off the east coast of New Zealand's South Island in the early 1990s by the Blunt and Munro group at the University of Canterbury, New Zealand^{208,209}. This fortuitous finding enabled two developments that would greatly increase the supply of this rare natural product: trawling for the sponge in its natural habitat and transplantation-aquaculture of the sponge to in-shore artificial habitats²¹⁰. Approximately 1 tonne of *Lissodendoryx* was harvested by trawling, and subsequent isolation of halichondrin B by scaled-up isolation–purification methods provided useful quantities for preclinical studies. Meanwhile, the survival and growth of sponge explants was highly dependent upon season and water temperature, but the growing sponges did produce halichondrins, albeit at lower yields than wild-type samples of *Lissodendoryx* (halichondrin B content ~ 0.4 mg per kg; total halichondrins ~ 1.5 mg per kg)²¹⁰. Thus, the combined efforts of marine ecologists, governments and academic scientists and support from the US National Cancer Institute resulted in the delivery of 310 mg of halichondrin B and a comparable quantity of isohomohalichondrin B.

Although aquaculture could partially address the supply issue associated with halichondrin B, the structures of the halichondrins were too complex to make them feasible as targets for scaled-up production by total synthesis. A significant breakthrough was achieved by academic scientists in partnership with Eisai Company in Japan, its US subsidiary Eisai Corporation of North America, and the Eisai Institute, who showed truncated synthetic analogues of halichondrin B with approximately 70% of the molecular mass had equipotent activity with the natural product against tumour cells^{211–213}. This work culminated in the identification and scaled-up production of the antitumour halichondrin analogue, eribulin mesylate (E7389, NCI number NSC 707389; FIG. 6c), the methanesulphonate salt of a terminal amino alcohol corresponding to the C1–C35 carbon skeleton of halichondrin B²¹⁴. Eribulin mesylate, like halichondrin B, exhibits broad antiproliferative activity against tumour cells by binding to tubulin and arresting the cell cycle at mitosis²¹⁵. Clinical trials of eribulin mesylate were approved in 2002, and results from Phase II trials of eribulin mesylate as monotherapy for refractory breast cancer (88 patients) were reported in 2006 (REF. 216) and more recently in heavily pretreated patients. The median duration of confirmed response was 113 days and preliminary safety data showed that the major toxicity issue was neutropaenia and leukopaenia (31 patients) with some febrile neutropaenia (two patients). In a related study with eribulin mesylate in patients with advanced breast cancer, response rates were 15% with tolerable toxicities²¹⁷. Eribulin mesylate also showed 9.7% response rate in patients with NSCLC (10.4% in taxane-pretreated patients) and a median survival time of 9.6 months²¹⁸. Phase III trials of eribulin mesylate for breast cancer are continuing in the United States and in Europe²¹⁹.

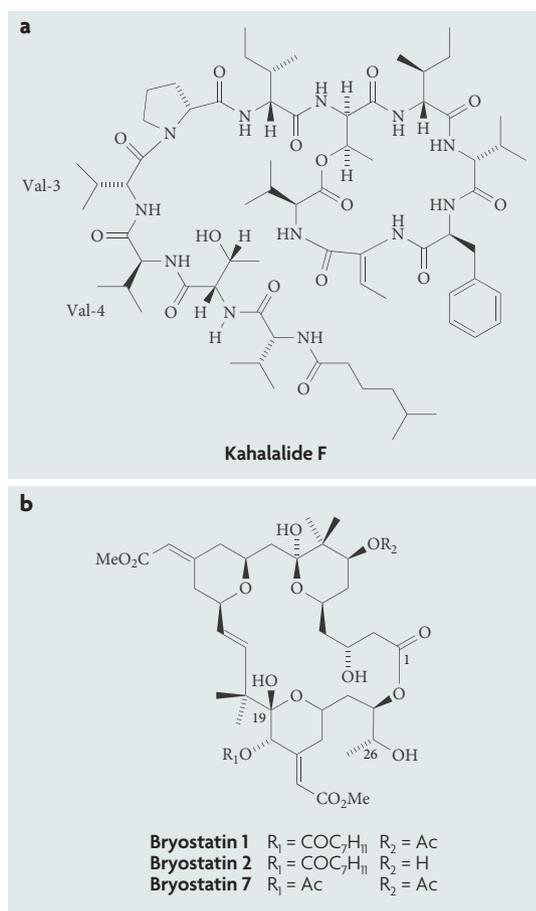


Figure 7 | **Kahalalide F and bryostatins.** **a** | Kahalalide F. **b** | The bryostatins.

Other antitumour marine natural products

Antimitotics are not the only antitumour compounds derived from marine organisms. Other natural products from various marine organisms, represented by a range of chemical structures and different cytotoxic or cytostatic modes of action, have entered clinical trials. To date, none has been approved as drugs.

Kahalalide F. Seasonal collections of the sacoglossan (sea slug) *Elysia rufescens* (FIG. 2e) by the Scheuer group from the University of Hawaii led to the isolation of the novel antitumour depsipeptide kahalalide F^{220,221} (FIG. 7a) in 1993 in addition to other analogues²²². Kahalalide F is a C₇₅ cyclic tridecapeptide that contains several unusual amino-acid residues, including the rare *Z*-dehydroaminobutyric acid found only in a few peptides including the antibiotics cypemycin²²³ and hassallidin A²²⁴. *E. rufescens* is a herbivorous Hawaiian opisthobranch that feeds on the alga *Bryopsis* spp.^{220–222}, which is the true source of kahalalide F. Accumulation of high concentrations of kahalalide F is seen in *E. rufescens* (1%) compared with the alga (0.0002%)²²², and it seems that both organisms use kahalalide F as a chemical defence against predation²²⁵. The relative configuration of kahalalide F was the subject of some revision.

A cyclic peptide with the structure originally proposed for kahalalide F was synthesized^{220,221} but it exhibited cytotoxicity an order of magnitude lower than that of the natural product. Subsequent revision of the structure — by inverting the configurations of Val3 from *L*- to *D*- and of Val4 from *D*- to *L*-, respectively — coupled with a more efficient solid-phase synthesis developed at the University of Barcelona, Spain, led to the correct structure and a product with essentially the same activity as the natural product^{226,227}. Kahalalide F was licensed by the University of Hawaii to PharmaMar in the early 1990s.

The mechanism of action of kahalalide F has not yet been fully elucidated. Kahalalide F is active mainly at the lysosomal level and induces vacuolization²²⁸, which may explain its activity on tissues that actively secrete lysosomal proteins, such as prostate cells. In human prostate cell lines and breast cancer cell lines, kahalalide F was shown to induce cell death by oncosis²²⁹ (necrosis coupled with karyolysis and swelling). In addition, a necrosis-like process was observed in several human kahalalide-F-sensitive breast, vulval, NSCLC and hepatic and colon carcinoma cell lines²³⁰, in which downregulation of the ERBB3 protein and inhibition of the phosphatidylinositol 3-kinase (PI3K)–AKT signalling pathway were identified as determinants of its cytotoxicity²³¹. Kahalalide F also induces channel formation and consequent cell-membrane permeability in sensitive HepG2 cells²³¹. It does not manifest toxicity towards murine haematopoietic progenitors and stem cells at concentrations up to 10 μM¹⁰⁷, and was shown to be metabolically stable²³².

A preclinical study of kahalalide F in rats reported renal toxicity using single doses at the maximum tolerated dose of 1,800 μg per m² administered intravenously. However, a multiple-dose regimen reduced drug-induced toxicity²³³. A Phase I study on androgen-refractory prostate cancer reported a maximum tolerated dose of 930 μg per m² per day, and determined a dose of 560 μg per m² per day for Phase II trials²³⁴. Out of 32 patients, one patient treated at a dose of 80 μg per m² had a partial response with a corresponding prostate-specific antigen reduction of at least 50%, and five patients showed stable disease. Following a Phase I trial suggesting a positive therapeutic index on advanced solid tumours²³⁵, kahalalide F is currently in Phase II clinical trials for solid tumours including melanoma, NSCLC and hepatocellular carcinoma.

Bryostatin 1. The bryozoan *Bugula neritina* (FIG. 2f) is a common ‘fouling’ organism that grows in thick colonies on pier pilings and docks. In 1968, a collaborative programme between Pettit and co-workers at the Arizona State University and the US National Cancer Institute found that extracts of *B. neritina* collected from the Gulf of Mexico exhibited remarkable activity against murine P338 lymphocytic leukaemia cells¹⁸⁰. Isolation and identification of the active principles was made difficult when follow-up collections of the bryozoan from different locations provided extracts with low titres of activity. Subsequent large-scale collections from the Gulf of California (Sea of Cortez) and locations along the coast of California, USA, were laboriously purified

to provide the first milligram of the active component, bryostatin 1 (FIG. 7b), the structure of which was finally solved by in 1982 by X-ray crystallographic and spectroscopic analyses²³⁶. Bryostatin 1 is a 26-membered macrocyclic lactone, with 11 stereocentres and a unique polyacetate carbon backbone that had not been previously encountered in natural products²³⁶. Subsequently, 19 additional bryostatin homologues and analogues were isolated from *B. neritina* (ordinal numbering up to bryostatin 20). The main differences in the molecular structures of these compounds are substitution at C₇ and C₂₀ by different acyloxy substituents¹⁸⁰. Bryostatin 1 demonstrated potent *in vitro* activity against the P388 lymphocytic leukaemia cell line with an ED₅₀ of 0.89 µg per ml²³³. However, the low abundance of compound (~0.01 parts per million) and variable occurrence in natural populations of the bryozoan precluded supplies of enough compound for early preclinical and clinical studies as an anticancer drug. In 1991, a novel process of large-scale isolation and purification afforded 18 g of bryostatin 1 from a collection of approximately 10,000 gallons of *B. neritina*²³⁷. In the meantime, the Southern California company CalBioMarine Technologies was engaged in refinement of aquaculture technology for growth and harvest, and improvement of the titre of bryostatin 1 (REF. 238). The prospects of supply of bryostatin by total synthesis were not optimistic. The unique chemical structure of bryostatin 1 posed major synthetic challenges and, as of this time, no synthesis of the more desirable bryostatin 1 has been reported, although syntheses of bryostatin 2 (REF. 239), bryostatin 3 (REFS 240,241) and bryostatin 7 (REF. 242) (FIG. 7b) have been completed together with numerous partial syntheses of its constituent molecular segments^{243–245}. Computer modelling studies at Stanford University, USA, identified that the minimum pharmacophore of bryostatin 1 required the presence of oxygen atoms at C₁, C₁₉ and C₂₆ (REF. 246), and led to the preparation of a truncated analogue with similar levels of activity to bryostatin 2 — the so-called bryopyran — in fewer synthetic steps than required for related natural products^{247,248}. Bryopyran is being considered for clinical development.

Early studies identified bryostatin 1 as a modulator of protein kinase C (PKC)²⁴⁹; however, the precise details have been difficult to define. The compound binds to the regulatory domain of PKC — the same binding site targeted by phorbol esters — at subnanomolar concentrations. Unlike phorbol esters, bryostatin 1 elicits different physiological effects and lacks tumour-promoting activity^{247,248}. Short-term exposure of tumour cells to bryostatin 1 induces PKC activation, self-phosphorylation and translocation to the membrane. Subsequent prolonged exposure inhibits PKC by causing depletion from the cell, probably due to proteolysis by a proteasome²⁴⁶. Bryostatin 1 also demonstrates various other biological activities, which include modulation of the immune system²⁵⁰, induction of cell differentiation²⁵¹, radioprotection²⁵² and synergistic interactions with other anticancer agents such as ara-C, paclitaxel, tamoxifen, auristatin PE, dolastatin, vincristine, doxorubicin and prednisone; these activities are summarized in REF. 253.

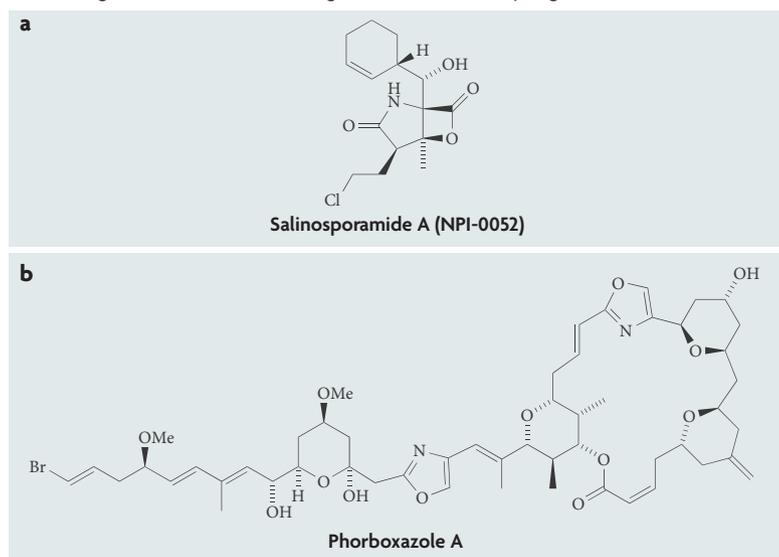
With its promising anticancer activity, bryostatin 1 has been investigated either alone or in combination with other chemotherapeutic agents in numerous clinical trials (Phase I and II) for myeloid leukaemia, lymphocytic leukaemia, melanoma, non-Hodgkin's lymphoma and NSCLC, metastatic myeloma²⁵⁴, relapsed lymphoma and chronic lymphocytic leukaemia^{255–257}, and other refractory malignancies²⁵⁸. A review of bryostatin 1 provides key information on the history of clinical development of the drug²⁵⁹. The major dose-limiting toxicity in all cases is myalgia, which explains a direct toxic effect of bryostatin 1: muscular vasoconstriction as a result of impaired oxidative metabolism and proton efflux from the muscle cells²⁶⁰. The end results of Phase II trials of bryostatin 1 as single-agent therapy against squamous cell carcinoma of the head and neck²⁶¹, colorectal cancer²⁶², melanoma²⁶³, renal cell carcinoma²⁶⁴ and ovarian carcinoma²⁶⁵ were discouraging, as no significant clinical effect against these tumour types was observed. Nonetheless, the results of a Phase I trial investigating co-drug therapy with 1-β-D-arabinofuranosylcytosine for acute leukaemia²⁶⁶ and vincristine in B-cell malignancies²⁶⁷ were promising. Patients in these trials have achieved objective remissions and prolonged stable disease. Five Phase II clinical trials were recently sponsored by the US National Cancer Institute (data from clinical trials web site; see Further information) to evaluate co-drug therapy of bryostatin 1 with rituximab, vincristine, temsirolimus, cladribine or interleukin 1 against B-cell non-Hodgkin's lymphoma, metastatic solid tumours, lymphocytic leukaemia and refractory solid tumours. Bryostatin 1 has been evaluated in combination with paclitaxel for the treatment of solid tumours and achieved superior response rates than paclitaxel alone. However, roughly one-third of patients discontinued the therapy due to myalgias²⁶⁸. Bryostatin 1 has also been reviewed in context of targeted therapies for cell-cycle checkpoint intervention²⁶⁹.

Outlook

As illustrated by the examples above, the discovery of selective and potent therapeutic activity in a rare marine natural product can drive innovative methods for its procurement, including production by aquaculture, semi-synthesis, synthesis and the development of synthetic analogues with more manageable properties. While academically inspired research has been key to marine natural product drug discovery in the past, novel approaches to translational medicine, which unite marine natural products chemists and pharmacologists with investigators in medical schools and institutes of pharmaceutical sciences, will be important in accelerating the progression of marine natural products from their discovery to the laboratory bench-top and to the clinic. Successful collaborations between academic institutes and pharmaceutical companies will continue to provide the mutual benefits that each party seeks. Academic programmes gain access capacities for sophisticated screening, pharmacological evaluation and advancement of leads to *in vivo* models, whereas

Box 1 | Marine natural product development from bacteria

Marine bacteria have recently taken the limelight as potential sources of highly novel chemical structures and potential as drug leads. For example, the p26 proteasome inhibitor salinosporamide A (NPI-0052; **a**) was discovered by Fenical, Jensen and co-workers at the Scripps Institution of Oceanography, California, USA, in a bacterium recovered from deep-sea sediment in 2003 (REF. 278). Salinosporamide A contains an unusual β -lactone and is produced by a previously undiscovered genus of actinomycetes, a newly named family of 'salt-loving' bacteria, the *Salinispora*²⁷⁹. The mechanism of action of salinosporamide A is similar to that of the proteasome inhibitor bortezomib (Velcade; Millennium/Janssen-Cilag), which is approved for the treatment of multiple myeloma, but it overcomes resistance to bortezomib in patients with relapsed or refractory multiple myeloma²⁸⁰. Salinosporamide A has completed Phase I clinical trials for multiple myeloma under the sponsorship of Nereus Pharmaceuticals in La Jolla, California^{281,282}. Furthermore, many compounds originally extracted from marine sponges are actually produced by marine bacteria. For example, the potent cytostatic agent phorboxazole A (structure 29; **b**) from a Western Australian sponge, *Phorbas* spp., that inhibits cell cycle in S-phase at subnanomolar concentrations^{283–285}, or the co-occurring phorbasisides^{286,287} may actually be produced by cyanobacteria that either live within the host, or expressed from microbial genes that have been integrated into the host sponge.



industry gains high-value leads while evading the high-risk associated with marine drug discovery. Major government funding can offset the high risk factor associated with marine natural products drug discovery, and new national programmes to exploit marine biotechnology and drug discovery have been launched by the governments of Germany, Ireland, Norway⁹ and South Korea — but, surprisingly, less so in the USA — within the past 5 years.

But why persist at all with the pursuit of natural products for new drugs in the modern age? Many reasons have been offered as to why natural products are such good sources for drug leads, but at least one study has attempted to quantify a correlation between the drug molecules and those typically found in natural products and combinatorial chemical libraries. Combinatorial libraries are purposely synthesized in large numbers, but the structures are intentionally imbued with high randomness. A multivariate comparison of the chemical space occupied by thousands of combinatorial drug compounds with that of natural

products revealed a strikingly good correlation of clinically approved drug molecules with the latter, but not the former²⁷⁰. In other words, the structures of drugs we use today more closely resemble those of natural products.

There are good reasons to be optimistic about the future. New technologies in analytical spectroscopy have pushed the limits of observation so that discovery of new molecules require only a few micrograms — a fraction of the material that was required even 10 years ago. New sources of marine natural products, such as marine bacteria, are being explored (BOX 1). Finally, there is enormous excitement and promise for drug discovery by manipulation of biosynthetic pathways in the most refractory microbes — the unculturable bacteria or metagenomic DNA from seawater and sediments. By deploying the cutting-edge tools of genetic engineering, genome mining²⁷¹ and new approaches to metagenomic mining of environmental DNA²⁷², it may be possible to unlock the genetic potential of millions of bacteria that occupy each millilitre of seawater or benthic sediment. Generally, pelagic bacteria are unculturable (~90%, or more correctly as yet, uncultured), unlike the soil-borne actinomycetes that historically have played an important role in drug discovery. Many compounds obtained from marine sponges, including those mentioned in this Review, are thought to arise from marine microbes that live in symbiosis with the larger host organisms. Recently, the prospects for sustainable production of pharmaceuticals from invertebrate-derived microbes have been reviewed²⁷³, and it is clear that many obstacles must be overcome before this approach transforms into conventional technology. The potential payoff for this approach is very high. For example, biosynthesis of the antitumour drug bryostatin 1 has been traced to an as-yet uncultivated endosymbiotic γ -proteobacterium: "*Candidatus* Endobugula sertula"²⁷⁴. Most of the genes that are responsible for the biosynthesis of the drug have been recently identified and sequenced²⁷⁵. The cloning of the biosynthetic genes for bryostatin 1 and so-called heterologous expression in a bacterium more amenable to large-scale fermentation production would overcome the limitation of procurement of the drug from the ocean (which is currently limited to expensive aquaculture or field harvesting) and ensure supply. The 'golden era' of antibiotic discovery began with microbes, and it is fitting that we now return to these origins with new tools to probe the untapped, broad molecular diversity of the bacteria that populate the oceans, benthic sediments and marine invertebrates.

In ancient poems almost 3,000 years old, Homer wrote of "the wine dark sea"^{276,277}, an evocative phrase that still rings with mystery. Did Homer choose his words to have us empathize with a grieving Achilles in his contemplation of human suffering and untimely death by marrying two familiar metaphors: wine, the velvet hued libation and tonic to health, with the fathomless complexity and mysteries of the open seas? As is clear from the drugs from the sea discussed in this Review, the coming years may provide a glimpse of hope and cures from beneath the swell and ebb of the vast oceans.

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 NCI Fact Sheets: <http://www.cancer.gov/cancertopics/factsheet>
 The Gordon Research Conference in Marine Natural Products: <http://www.grc.org/conferences.aspx?id=0000144>
 The Natural Products Branch of the Developmental Therapeutics Program of the NIH: <http://dtp.nci.nih.gov/branches/npb/repository.html>
 Ziconotide @ 3Dchem.com: <http://www.3dchem.com/moremolecules.asp?ID=260>

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