The importance of larval dispersal to the population dynamics and biogeography of marine organisms has been recognized for almost a century (Hjort, 1914; Thorson, 1950). More recently, theoretical studies have highlighted the role that connectivity may play in determining the resilience of marine populations (Hastings and Botsford, 2006). Effective spatial management of marine capture fisheries, including the design of marine reserve networks, also requires an understanding of population connectivity (Sale et al., 2005). However, remarkably few empirical estimates of larval dispersal or population connectivity in ocean environments exist.

Direct and definitive estimates of larval dispersal in the ocean require the ability to track microscopic larvae of benthic invertebrates and fishes through the pelagic environment, from spawning locations to recruitment sites. Most marine species with pelagic larvae spawn millions of propagules that are released and then subjected to significant advection, diffusion, and mortality in vast volumes of seawater, making traditional mark-recapture approaches extremely difficult (Levin, 2006). However, ecologists have embraced recent developments in probe-based mass spectrometry to examine the chemistry of calcified structures in marine invertebrates and fishes that can be used as artificial or natural tags of natal origins. These geochemical tags are revealing fascinating data on larval dispersal that are challenging widely held paradigms concerning the spatial scale of demographic connectivity in ocean ecosystems.
Natural geochemical tags are generated by variations in environmental conditions, including temperature, salinity, and seawater chemistry, that are subsequently recorded by the elemental or isotopic composition of calcified structures (see Box 1). Therefore, shells, fish ear-stones (called otoliths), and statoliths (the invertebrate analog to otoliths) may act as dated “flight recorders” because as organisms disperse across gradients in seawater composition or temperature, their travels are being constantly recorded by the chemistry of the calcified structure. Likewise, larvae developing in areas that have differing seawater characteristics build calcified structures whose elemental compositions reflect their sources (Zacherl et al., 2003a; Becker et al., 2007). Typically, geochemical tags consist of either a combination of minor and trace elements expressed as a ratio to Ca (e.g., Mg/Ca, Mn/Ca, Sr/Ca, and Ba/Ca), or stable isotope ratios such as δ18O (Killingley and Rex, 1985) and δ87Sr (Barnett-Johnson et al., 2005).

The ability of natural geochemical tags to track larval movement depends upon the existence of substantial variation in the elemental composition of those tags among locations of interest (Thorrold et al., 2002). Many studies using geochemical tags have focused on species inhabiting estuaries because of significant variation in the elemental composition of their calcified structures (e.g., Swearer et al., 2003), probably due to substantial differences in salinity, temperature, and water chemistry among estuarine waters. Variation in water chemistry among estuaries is a function of coastal geology, sources of pollution, atmospheric deposition, and variable inputs from local watersheds. In southern California, for example, because of heavily urbanized development, rainwater runoff should contribute high concentrations of trace elements associated with anthropogenic activity into the estuaries and nearshore waters. The magnitude of input of these trace elements is a function of land use, the amount of impermeable surface area within a watershed, number and size of rivers, and frequency of rain events (Ackerman and Schiff, 2003). Broadening consideration to the entire west coast of the United States, watersheds in Washington and Oregon regularly receive > 100 inches of rain annually, while watersheds in southern California rarely see > 50 inches annually (National Weather Service, 1961–1990 average annual rainfall). Because of this extreme variation in land use, geology, and runoff contributions typically associated with estuaries, these habitats would appear to provide ideal locations for generation of a natural geochemical tag (see “Progress to Date” below).

Less attention has focused on using geochemical tags in open-coast species. The limited emphasis on coastal ocean habitats is probably because gradients in oceanographic conditions can be more subtle, thus complicating the generation of unique geochemical tags. For example, Gillanders et al. (2001) found little variation in otolith chemistry of two-banded bream among locations and sites that ranged from ~ 10 to > 100 km from one another along the coast of Spain. They attributed the lack of variation in otolith chemistry to the limited number of major rivers within the study location and to low rainfall and its associated runoff. However, locations of convergence between distinct ocean currents (e.g., at Point Conception in central California and at Cape Hatteras in North Carolina, USA) are characterized by sharp gradients in temperature and, to a lesser extent, salinity, making generation of distinctive geochemical tags likely (Zacherl, 2005). Seawater elemental composition may also vary among locations at smaller spatial scales (e.g., Becker et al., 2005) due to differences in coastal geology, mesoscale oceanography, and variable inputs from local watersheds.

The tremendous advantage to using natural geochemical tags is that every larva is potentially tagged, eliminating any concerns arising from dispersion of tagged larvae and subsequent low recapture rates. However, there can be a significant uncertainty in interpreting the elemental variation in tags among locations. Unless obvious and strong broad-scale regional gradients in tag composition exist, the approach necessitates substantial sampling effort to ensure that the geochemical tags of all potential source populations have been characterized.

Artificial Tags
Calcified tissues are also ideal repositories for artificial chemical markers that are used to tag embryos or lab-reared larvae before dispersal from natal locations (Jones et al., 1999; Moran and Marko, 2005; Thorrold et al., 2006). Fluorescent compounds such as tetracycline or calcein, elemental markers (e.g., rare earth elements), and radioactive isotopes have all been used to tag calcified tissues.
Use of geochemical tags is premised on the notion that the chemistry of calcified structures in some way reflects physiochemical properties of the surrounding seawater in which the calcified structure was formed. Results from controlled laboratory experiments on gastropod larvae by Zacherl et al. (2003b) and on fish larvae by Bath et al. (2000) clearly supported this assumption and yielded predictable results (Figure A-1). In their experiments, both groups manipulated concentrations of Ba/Ca in rearing seawater, raised larvae for several weeks under experimental conditions, and then analyzed whole larval calcified structures using iCP-MS. Linear relationships between concentrations of elements in seawater and in biogenic aragonite are common and have been documented for many elements including Ba, Sr, and Mg (e.g., Lorenz and Bender, 1980; Elsdon and Gillanders, 2003).

Temperature and salinity also predictably influence element uptake into otoliths and statoliths (e.g., Martin et al., 2004; Zumholz et al., 2007). The combined results of all culturing studies suggest that calcified otoliths, statoliths, and protoconchs, like foraminifera tests (Lea et al., 1999), can indeed reliably integrate information about seawater physical and chemical properties.

However, there is still some mystery involved with how simple and predictable findings from larval culturing studies play out in a field setting, where calcified structures form under the influence of several interacting factors. Therefore, recent culturing experiments have focused on examining the interactive effects of two or more factors (e.g., Milton and Chenery 2001; Martin and Thorrold, 2005) and on ranking their relative importance (Eldon and Gillanders, 2004; Lloyd et al., in press). Further, there is very recent evidence that not only seawater physical and chemical properties but also maternal transmission of elements can influence the elemental composition of larval calcified structures. For example, there is evidence that trace elements contained in the egg can be incorporated into larval-fish otoliths (Kalish, 1990; Thorrold et al., 2006) and into larval-gastropod statoliths (Lloyd et al., in press). This knowledge complicates our ability to draw simple conclusions about the physical environment based upon the chemistry of calcified structures but provides an exciting venue for future Calcified Structure Investigations.

**BOX 1: CSI SEAWATER—DO CALCIFIED STRUCTURES INTEGRATE SEAWATER PROPERTIES?**

Use of geochemical tags is premised on the notion that the chemistry of calcified structures in some way reflects physiochemical properties of the surrounding seawater in which the calcified structure was formed. Results from controlled laboratory experiments on gastropod larvae by Zacherl et al. (2003b) and on fish larvae by Bath et al. (2000) clearly supported this assumption and yielded predictable results (Figure A-1). In their experiments, both groups manipulated concentrations of Ba/Ca in rearing seawater, raised larvae for several weeks under experimental conditions, and then analyzed whole larval calcified structures using iCP-MS. Linear relationships between concentrations of elements in seawater and in biogenic aragonite are common and have been documented for many elements including Ba, Sr, and Mg (e.g., Lorenz and Bender, 1980; Elsdon and Gillanders, 2003).

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Pacific coast. Fodrie (2006) found similar patterns in juvenile California halibut otoliths, but went a step further in documenting distinct signatures for embayments with different geomorphologies and for open-coast habitat. These findings were employed to address the movements of juveniles within bays, and allowed increasing refinement of nursery habitat determination for successful subadults as well as evaluation of the demographic consequences of using different nursery habitats.

Detecting geochemical differences imparted to pelagic larvae that disperse along the open coasts of continents has proved a greater challenge. Recent progress suggests that subtle differences in water masses and eddies, upwelling, temperature, or watershed influence may provide the needed chemical variation to allow for the application of natural geochemical signatures (e.g., Zacherl, 2005). For instance, natal sites of larval rockfish (Sebastes atrovirens) were distinguished between mainland and island sites by Zn/Ca, Sr/Ca, Ba/Ca, and Pb/Ca signatures in pre-hatch otoliths; mainland sites only 10 km apart also showed significant differences in elemental signatures (Warner et al., 2005).

**Identifying Natal Origins**

The ability to address the question, “where did newly recruited individuals come from?” is a real strength of geochemical tags, and distinguishes the method from other indirect approaches to measuring connectivity, including coupled bio-physical modeling and population genetics. Despite the potential, there are remarkably few studies that have used geochemical signatures successfully to determine larval origins, although identification of juvenile nurs-

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**Figure A-1.** The influence of ambient seawater on the elemental composition of biogenic aragonite is most easily quantified in the laboratory where water chemistry and temperature can be accurately constrained. For instance, Ba/Ca ratios (mean ± standard error) in otoliths of laboratory-reared spot, Leiostomus xanthurus, (cyan symbols) and of statoliths (green symbols) and protoconchs (red symbols) of laboratory-reared Kellet’s whelk (Kelletia kelletii) varied linearly with the Ba/Ca ratios of the rearing water. Note that the primary x-axis depicts rearing-water Ba/Ca concentrations for statoliths and protoconchs at 17.1°C and 34‰ salinity (from Zacherl et al., 2003b), while the secondary x-axis depicts rearing-water Ba/Ca concentrations for otoliths at 20.3°C and 20‰ salinity (from Bath et al., 2000).
The use of natural geochemical tags in studies of larval connectivity requires knowledge of the signatures imparted to larval structures by specific water masses at different locations. This has been a serious challenge for free-spawning species that do not brood or hold their offspring on the seabed. A solution to this problem is presented in Becker et al. (2007). The authors outplanted laboratory-spawned larvae of mytilid mussels to generate a reference map of chemical signatures at different locations in San Diego County, including open-coast and bay settings along 75 km of shoreline (Figure B-1). Larvae of two Mytilus mussel species were introduced into PVC "homes" within 12 hours of spawning (before shell formation) and were transported to nearshore moorings where they were allowed to develop for one week. During this period the larvae laid down aragonite shells that retained a chemical signature of the waters in which they developed. The elemental composition of the larval shells was then analyzed using laser ablation ICP-MS. After analyzing shell signatures, Becker and co-workers determined that the method resolution for identifying distinct water bodies was greatest at the regional (20–30 km) scale and for individual bays.

Two weeks after the outplanting, newly settled mussels (one to two weeks old) were collected from intertidal rocks near the outplant moorings. These recruits were 1–2 mm and retained their larval shell—the protoconch. Recruit mussel tissue was analyzed by a PCR-based assay to identify species (either Mytilus californianus or Mytilus galloprovincialis). The chemical composition of recruit larval shells was determined by laser ablation ICP-MS and individuals of each species at each sampling site were assigned an origin. This information was used to assess larval connectivity patterns for the California mussel and the bay mussel in San Diego County. Differences were found in larval connectivity patterns of the two species (Figure B-1); this was unexpected as their larvae are thought to mix along the San Diego coastline. M. californianus exhibited asymmetric mixing with the majority of larvae originating in the northern part of the study area; there was high self-recruitment in the north and high importation of larvae in the south. M. galloprovincialis recruits had more diverse origins from a mixture of north, south, and bay locations, but with substantial (40%) self-recruitment. This study provides information about larval connectivity for two species at one time in one region. It offers a first look at the origins of settled invertebrates using elemental signatures in shells, and supports a growing paradigm of limited dispersal, even in species with long-lived larvae. However, many questions remain about the underlying causes of species differences in connectivity, the stability of these patterns over time, and their relevance for other areas within the species’ ranges.

**BOX 2. MUSSELS IN MOTION**

The use of natural geochemical tags in studies of larval connectivity requires knowledge of the signatures imparted to larval structures by specific water masses at different locations. This has been a serious challenge for free-spawning species that do not brood or hold their offspring on the seabed. A solution to this problem is presented in Becker et al. (2007). The authors outplanted laboratory-spawned larvae of mytilid mussels to generate a reference map of chemical signatures at different locations in San Diego County, including open-coast and bay settings along 75 km of shoreline (Figure B-1). Larvae of two Mytilus mussel species were introduced into PVC "homes" within 12 hours of spawning (before shell formation) and were transported to nearshore moorings where they were allowed to develop for one week. During this period the larvae laid down aragonite shells that retained a chemical signature of the waters in which they developed. The elemental composition of the larval shells was then analyzed using laser ablation ICP-MS. After analyzing shell signatures, Becker and co-workers determined that the method resolution for identifying distinct water bodies was greatest at the regional (20–30 km) scale and for individual bays.
of crab zoea from inside and outside San Diego Bay (DiBacco and Levin, 2000). While analyses confirmed behaviorally enhanced export of larvae (DiBacco et al., 2001) and extensive mixing of larvae from different sources (DiBacco and Chadwick, 2001), the researchers did not address the question of natal origins of new recruits. Recently, natal origins of new recruits were determined in mytilid mussel populations along the coast of southern California (Becker et al., 2007). Bay mussels (*Mytilus galloprovincialis*) and California mussels (*M. californianus*) were found to exhibit differing connectivity patterns and rates of self-recruitment among study sites (see Box 2).

**EMERGING TECHNOLOGIES AND FUTURE DIRECTIONS**

The use of geochemical signatures in calcified structures of marine organisms to estimate population connectivity in marine ecosystems is still in its infancy. A number of significant challenges remain before routine estimates of population connectivity in coastal waters will be possible using these approaches. Nonetheless, exciting new results using natural and artificial geochemical tags clearly demonstrate the potential of the approach (Almany et al. 2007; Becker et al., 2007). We also envisage application of the techniques in novel environments, including hydrothermal vents and other ephemeral habitats in the deep sea, where connectivity is likely to be extremely important to population persistence and maintenance of biodiversity (e.g., Neubert et al., 2006).

Despite some progress, the application of geochemical markers to the study of larval dispersal in marine environments remains hampered by techno-
logical limitations. Elemental analysis of individual microscopic calcified structures must involve laser ablation of very small amounts of calcified material (i.e., less than 5 µg). Such small quantities of analyte provide a transient signal that is often insufficient for sequential analysis of isotopes with single collector inductively coupled plasma mass spectrometry (ICP-MS) instruments (Strasser et al., in press). However, time-of-flight mass spectrometry (TOF-MS) allows for simultaneous detection of ions over a large mass range and is therefore particularly well suited to analyses of transient signals generated by laser ablation (Vázquez et al., 2002). New developments in laser ablation also offer hope for more accurate chemical analyses of larval shells and otoliths. For instance, UV femtosecond laser systems, likely to be available commercially in the near future, will hopefully improve both the spatial resolution and accuracy of laser ablation ICP-MS analyses (Koch and Günther, 2007). Femtosecond lasers eliminate mass fractionation, at least in theory, during the ablation process and provide for stoichiometric conversion of ions that are then transported to a mass spectrometer for quantification (Russo et al., 2002). In the future we may see femtosecond lasers coupled to plasma source TOF-MS instruments producing precise and accurate elemental data at spatial resolutions less than 10 µm. This resolution could allow detailed examination of larval trajectories, reflected in the chemical composition of distinct regions of minute larval structures.

As mentioned above, studies attempting to use natural geochemical markers in open marine systems commonly encounter subtle gradients in physical and chemical properties of coastal water masses, at least compared to the stronger elemental signals imparted within river and estuarine systems. Efforts to overcome this problem include refinement of analytical techniques to increase the precision of geochemical variables. However, we may already be at a stage where physiological or ontogenetic controls on the elemental composition of biogenic carbonates may introduce more error into estimates of larval sources from geochemical signatures than the precision of the analytical technique. In the future, researchers are likely to use isotope tracers that are less subject to physiological effects and therefore more accurately reflect ambient values in the environment. The number of these isotope systems available to researchers has increased rapidly with the development of multiple collector arrays on ICP-MS instruments (Halliday et al., 1998). There may also be useful synergies with researchers attempting to develop new temperature proxies in biogenic carbonates based on Mg, Ca, and Sr isotopes that may not be subject to significant biological fractionation (e.g., Nägler et al., 2000; Fietzke and Eisenhauer, 2006).

Interest continues in artificial tagging approaches for marine larvae because few methods can provide unequivocal estimates of population connectivity in ocean ecosystems. However, the logistic difficulties associated with tagging a large proportion of the total larval production from an area has, until very recently, proved difficult to overcome. The development of a TRAnsgenerational Isotope Labeling (TRAIL) technique, based on maternal transmission of an enriched stable Ba isotope that is incorporated in the embryonic otoliths of larval fish, may help to overcome this limitation (Thorrold et al., 2006). The TRAIL approach represents a significant advance from earlier artificial tagging methods because it is possible to tag a much higher proportion of the total larval production from an area, the technique can be used on benthic and pelagic spawning fishes, and multiple tags can be applied (Figure 1). The first field test of the method demonstrated substantial (> 50%) self-recruitment of benthic and pelagic spawning fishes to

With careful targeting of specific questions to be addressed, and of focal species to be examined, studies using natural and artificial tags in calcified structures are likely to lead to significant advances in our understanding of population connectivity in ocean ecosystems.
a small coral reef reserve in Papua New Guinea (Almany et al., 2007). The success of the study was based on the ability to tag a high percentage of all larvae produced in the reserve over a period of two months. The TRAIL method is likely to be particularly useful for species that form spawning aggregations at specific locations and times, and therefore provide the opportunity for a considerable percentage of the total spawning population to be captured.

**IMPLICATIONS**

Ultimately, scientists need to provide accurate information on population connectivity, which can be used to optimize spatial management approaches, including marine protected areas (MPAs), for marine-capture fisheries. Although we know of no management plans that have specifically incorporated empirical estimates of population connectivity, such information would have been useful when choosing protected-area locations in Australia’s Great Barrier Reef Marine Park or the Channel Islands off the coast of California. However, there is also no denying that determining population connectivity using the approaches outlined here is both time-consuming and expensive, and is therefore only likely to be contemplated for a handful of species at any location. One way to maximize the generality of the results would be to use the data to test the performance of coupled biophysical models that are more easily applied to a number of different species over large spatial scales (e.g., Gilg and Hilbish, 2003; Galindo et al., 2006). Similarly, it has proved almost impossible to verify the effectiveness of MPAs without a method for quantifying larval export from the reserve to adjacent fished areas. Recent data suggest that the scale at which coral reef fish populations can be both self-sustaining and capable of providing recruitment subsidies is considerably smaller than previously imagined (Almany et al., 2007). Nonetheless, the generality of these results for other systems, and even for coral reefs, is still in question, particularly for grouper and snapper species that contribute disproportionately to artisanal and commercial fisheries on coral reefs. With careful targeting of specific questions to be addressed, and of focal species to be examined, studies using natural and artificial tags in
calciﬁed structures are likely to lead to signiﬁcant advances in our understanding of population connectivity in ocean ecosystems.

ACKNOWLEDGEMENTS

Our sincere thanks to the Division of Ocean Sciences at the National Science Foundation for funding larval connectivity research on K. kelletii (OCE 0351860 to DZ), bivalve mussels (OCE 0327209 and 0648656 to LAL), and California reef ﬁshes (OCE 0424688 to SRT). We thank B.J. Becker, J. Fodrie, and P. McMillan for sharing unpublished data on mytilid mussel and California halibut connectivity. Some of the ideas discussed here grew out of discussion among members of the Coral Reef Targeted Research (CRTR) Program Connectivity Working Group (http://www.gefcoral.org).

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