

## **Does fish larval dispersal differ between high and low latitudes?**

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## **SUMMARY**

Several factors lead to expectations that the scale of larval dispersal and population connectivity of marine animals differs with latitude. We examine this expectation for demersal shorefishes, including relevant mechanisms, assumptions, and evidence. We explore latitudinal differences in: 1) biological (e.g., species composition, spawning mode, pelagic larval duration (PLD)), 2) physical (e.g., water movement, habitat fragmentation), and 3) biophysical factors (primarily temperature, which could strongly affect development, swimming ability, or feeding). Latitudinal differences exist in taxonomic composition, habitat fragmentation, temperature, and larval swimming, and each could influence larval dispersal. Nevertheless, clear evidence for latitudinal differences in larval dispersal at the level of broad faunas is lacking. For example, PLD is strongly influenced by taxon, habitat, and geographic region, but no independent latitudinal trend is present in published PLD values. Any trends in larval dispersal may be obscured by a lack of appropriate information, or use of 'off the shelf' information that is biased with regard to the species assemblages in areas of concern. Biases may also be introduced from latitudinal differences in taxa or spawning modes, as well as limited latitudinal sampling. We suggest research to make progress on the question of latitudinal trends in larval dispersal.

**Key Words:** population connectivity, larval dispersal, pelagic larval duration, larval behaviour, genetic structure, habitat fragmentation

## INTRODUCTION

Most bottom-associated (demersal) marine animals, including fishes, spend part of their early life as larvae in open, pelagic waters before settling into a demersal life style. Pelagic larvae are subject to dispersal, and this has profound consequences for distributions, demography, genetic connectivity, and management. Several factors lead to expectations that the scale and processes of larval dispersal and population connectivity<sup>1</sup> of marine animals differ with latitude. These include contrasts in species composition and community structure, temperature influences on physiology and development, and differences in physical ocean processes. Conclusions that larval dispersal, population connectivity, or a proxy thereof, differ latitudinally have been reached by influential studies, and in each case the conclusion was that dispersal takes place over wider scales in higher latitudes. Houde [1] concluded that pelagic larval duration (PLD) is inversely associated with temperature and that fish larvae in warm seas are also more likely to starve than those in cold seas. These conclusions imply more limited larval dispersal in warm waters, because shorter PLDs are conventionally considered to lead to shorter dispersal distances (but see below), and higher mortality due to starvation should reduce effective dispersal distances [2]. Based on published studies of the influence of temperature on PLD in a range of marine taxa, O'Connor et al. [3] concluded that "maximum predicted dispersal distances for larvae in colder water are much greater than those in warmer water", and "population connectivity and effective population size should, in general, be inversely related to ocean temperature". Similarly, Bradbury et al. [4], using published genetic and PLD data, concluded that dispersal

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<sup>1</sup> For the purposes of this paper, *larval dispersal* describes the two-dimensional distribution of larval settlement originating from a single source population. *Connectivity* describes the source-destination matrix of settlers to a series of subpopulations that comprise a metapopulation connected through larval dispersal. Both terms can be spatially explicit, and are linked: short average larval dispersal distances should lead to spatially smaller metapopulations (or connectivity networks).

distance increases with latitude: for 163 marine fish species there were significant associations between maximum latitude, body size, and genetic structure ( $F_{ST}$ ). Although body size explained the most variation, this analysis revealed weaker genetic structure at latitudes above 40°, with the largest differences at the extremes of latitude (e.g., 20° vs. 60° latitude). Further, research using genetic parentage and otolith microchemistry techniques in warmer waters [5] has documented dispersal in larval reef fishes over much smaller scales than have been reported from temperate waters, leading to a perception that dispersal distance is correlated with latitude. Despite these perceptions, clear examples of latitudinal differences in larval dispersal or connectivity are rare.

This review is not a meta-analysis of past work investigating temperate-tropical differences in larval dispersal: such work does not exist. Instead, the intent of this review is to examine the hypothesis of latitudinal differences in larval-fish dispersal distance, the mechanisms and assumptions underlying the hypothesis, and evidence (including commonly-used proxies for larval dispersal) bearing upon it, to determine if it is supported. We also suggest research that will be useful in testing hypotheses of latitudinal differences in larval dispersal.

Why is it important to know if there are latitudinal differences in dispersal?

Knowledge of the spatial scale of larval dispersal in marine species, a major contributor to both evolutionarily and ecologically significant population connectivity, is critical to understanding community processes ranging from biogeography to population demography, to management of fisheries, and to biodiversity conservation. For example, space-based management of coastal oceans, including no-take marine reserves, is being implemented widely, and such management relies

on knowing the extent and patterns of connectivity [6],[7]. We know little about the fate of the increased reproduction that typically occurs inside marine reserves. This question is critical, because it addresses both the service function of reserves (e.g., export of larvae to fished areas) and the design of reserves (e.g., conservation networks connected through larval exchange, [8],[9],[10]). At present, the suggestion that connectivity among marine populations might vary geographically remains untested, thus hampering the ability of managers to apply general criteria to local problems. There is often disagreement about whether evidence gathered from one geographic area (for example, temperate coastal waters) is applicable to other geographically distinct areas (such as coral reefs).

Our focus here is on the dispersal distance of the pelagic egg and larval stage prior to settlement in demersal marine shorefishes (i.e., teleosts, the adults of which live on or near the bottom at depths <100 m). Because these species are relatively site-attached as adults, adult movement is unlikely to contribute greatly to either genetic or demographic connectivity. Even with this limited focus, many factors influence dispersal and connectivity, and the distance travelled is the result of biophysical processes involving hydrodynamics, as well as species-specific aspects of mortality, swimming, settlement behaviour, and PLD. Although post-settlement processes modify connectivity established by movement during the pre-settlement larval phase, these are beyond the scope of the present review. Note, however, that studies estimating dispersal or connectivity from settled populations (e.g., most genetic work) include influences from both larval supply and post-settlement processes, and must be interpreted with this in mind (see Supplement). It is possible that the extent to which population connectivity is maintained by pre-settlement vs. post-settlement

processes varies latitudinally (T.J. Miller, pers. com.). Even if this is true, it is appropriate to focus on the role that larval dispersal plays, as we do here.

For clarity, we divide this review into three general classes of factors that might lead to latitudinal differences in dispersal:

1. Biological differences: latitudinal differences in species composition and associated characteristics (especially spawning mode and PLD) that could affect dispersal;
2. Physical differences: latitudinal differences in water movement and habitat fragmentation that could independently affect dispersal, regardless of the underlying species composition;
3. Biophysical differences: latitudinal differences in physical factors (principally temperature) that could strongly affect biological processes (such as development, swimming ability, and feeding), that can in turn affect dispersal.

Due to space limitations, we present details of analyses in the Supplement, and confine ourselves here to overviews of results, discussion of the implications of those results, and recommendations for future research.

## **1. BIOLOGICAL DIFFERENCES**

### **(a) *Taxonomy and biogeography***

Taxonomic composition of demersal teleost shorefishes differs with latitude at all taxonomic levels, and different Orders or Suborders dominate at different latitudes (see Supplement for details). In tropical Hawaii, eastern Pacific, and Cuba, Anguilliformes, holocentroid Beryciformes, Tetraodontiformes, and perciform suborders Percoidei, Blennioidei, Gobioidae, Labroidae, and Acanthuroidei constitute 73-84% of the 430-700 demersal shorefish species. In contrast, in cold waters of northwestern Atlantic, northeastern Pacific and Antarctic, Gadiformes, perciform suborders Zoarcoidei and Notothenoidei, and scorpaeniform suborders Cottoidei and Hexagrammoidei and scorpaenid genus *Sebastes* constitute 73 to over 90% of the 55-198 species. To the extent that different taxa have different dispersal characteristics, apparent geographic differences in dispersal may simply reflect differences in faunal composition rather than differences in environments. To date, comprehensive information about taxon-specific dispersal differences is lacking, and given the non-independence of taxa and geographic distributions, it will be challenging to separate location-dependent physical and biological conditions from lineage-related factors.

### **(b) *Taxonomy and pelagic larval duration***

Longer dispersal distances are often assumed to arise from longer PLDs (e.g., [11], but see below for evaluation of this assumption). Aside from marine eels (mean PLD>100 days), available PLD data (see Supplement for sources and details) indicate that the Orders and Suborders dominating warm waters have shorter mean PLD values (23-52 days) than do taxa dominating cold waters (55-108 days: Supplement figure Sup1). The generality of latitudinal trends in PLD is questionable

because these PLD values were based only on nine Orders or Suborders from warm waters and four from cold waters. There are also possible biases due to habitats sampled – tropical data come mainly from shallow reefs, whereas temperate data come from a wider range of habitats (see Supplement discussion).

### **(c) *Spawning mode***

Spawning mode (in this case, demersal eggs vs. broadcast spawning with pelagic eggs) could have a strong effect on dispersal distance ([4],[12]). The pre-hatching period of pelagic eggs potentially increases dispersal distance, particularly in colder waters, where such periods can be weeks long [13]. This period of drift is rarely included in estimates of PLD, and it does not occur in live-bearing species or most species with demersal eggs. Further, larvae of most taxa from demersal eggs begin their pelagic larval life larger and in a more developed state than those from pelagic eggs, and the earlier acquisition of swimming ability might enable these larvae to behaviourally limit dispersal [14]. Clear latitudinal differences in spawning mode exist among taxa. In warmer locations, 60 to 80% of demersal shorefish species have pelagic eggs, whereas in colder locations (i.e.,  $>50^{\circ}$  latitude), only 15-27% of demersal species have pelagic eggs (based on faunal lists and taxon-specific spawning modes, see Supplement, figure Sup 2, and Table Sup 1). Further, in most regions, larvae from demersal eggs have shorter PLDs than those from pelagic eggs (see Supplement and section 3(c), on PLD, also [4]). Spawning mode has a strong taxonomic component, with spawning modes being mostly consistent within a family. Exceptions exist, however, and in these, the trend is for taxa from higher latitudes to shift away from broadcast spawning (Supplement). This trend toward demersal eggs in cold waters may have implications for larval dispersal and connectivity, and



highlights the need to account for spawning-mode differences in comparisons across regions.

## **2. PHYSICAL DIFFERENCES**

### **(a) *Oceanography***

Latitudinal gradients in seasonality, temperature, mixed layer depth (MLD), wind, and Coriolis force may potentially result in latitudinal differences in dispersal of fish larvae. The effects of physical oceanographic processes on latitudinal patterns in larval dispersal are not well discussed in the literature, and are included in few explicit, published hypotheses. Therefore, in the Supplement we develop hypotheses about how some aspects of physical oceanography might influence latitudinal patterns of larval dispersal.

Water movement itself varies with latitude, in part due to changes in Coriolis force. For example, Ekman coastal upwelling should be least important at low latitudes, perhaps leading to less upwelling-cell retention in the tropics (see Supplement). However, more energetic eddies should form at higher latitudes, and these can either advect larvae from their source, or retain them nearby, resulting in more variable larval dispersal. If the mixed layer depth (MLD) is shallow, larvae may be able to vertically migrate into slower-moving water below the MLD and thereby retard dispersal. Although MLD is more stable in the tropics, it may be shallower seasonally at higher latitudes, leading to differences in larval dispersal if the MLD interacts with vertical movement of larvae as outlined.

There are clear latitudinal differences in many variables that drive coastal circulation, but equally, there are large within-latitude regional and local differences in circulation

due to topography, coastal orientation, differences in tidal regimes, river input, and a variety of other factors [15]. Although certain latitudinal trends are expected, within-latitude spatial variation may frequently override those trends, thus obscuring them (see Supplement discussion).

**(b) *Habitat fragmentation***

For demersal fishes with some degree of habitat association, the strength of population connections should depend not only on spatial scales of larval dispersal, but also on the scales of patchiness of benthic habitat: clearly, larvae cannot settle successfully where there is no suitable habitat, so patchiness of habitat has a direct influence on dispersal distance [16]. At coarser scales, benthic habitats for nearshore demersal species are largely determined by the spatial distribution of coastlines, found either along continental margins or around islands. Continental margins have large areas of continuous nearshore habitat whereas islands are more isolated, with the degree of isolation depending on geographic and oceanographic distances to nearby islands or continents [17]. At finer scales, particular benthic habitats are often patchily distributed. Dispersal among patches becomes less likely as distance between suitable habitat patches increases [2].

Habitat patchiness appears to affect the scale of dispersal. A review of recent literature estimating demographic connectivity (see Supplement) shows that self-recruitment (i.e., larvae settling into the same area where they were spawned) is higher along continental coastlines compared to islands (figure 1A), but this is strongly affected by the spatial scale of the study (nearly an order of magnitude larger in continuous continental coastlines compared to patchy habitats and islands, figure 1B). Controlling for spatial extent of the study, the mean scales of connectivity

differ among contexts, with species in patchy habitats dispersing about 60 to 100 km, whereas species in continuous habitats dispersed about 900 km (figure 1C).

Combined, these data suggest that larval dispersal may be more restricted in fragmented habitats.

If habitat patchiness differs between tropical and temperate systems, then landscape context could affect dispersal. In fact, islands more than 5 km apart are 2-3 times more abundant in the tropics than in higher latitudes (figure 2; see Supplement), and this is expected to lead to more fragmented populations and shorter successful dispersal distances in tropical habitats. The degree of geographical isolation of habitat patches, however, may not be a consistent predictor of the likelihood of connectivity: oceanographic barriers (rather than simple distance, [18],[19],[20]) or larval behaviour may modify the effect of habitat fragmentation ([21],[22],[23], see Supplement).

### **3. BIOPHYSICAL DIFFERENCES**

Many variables scale with latitude including Coriolis force, seasonality, and day length, but, the most obvious and important is temperature. Many of the factors considered in this review are temperature related rather than latitude related *per se*, but other associated factors are also important.

#### **(a) *Temperature and larval swimming***

The expectation that behaviour of larvae may influence the scale of larval dispersal is based on research in three areas. First, many studies show that vertical distribution behaviour by larvae indirectly influences dispersal [14]. Second, swimming and sensory abilities of marine fish larvae are better than previously

realized [14, 23]: larvae of many species are able to swim directionally and at high speeds in the sea [22], which implies the ability to influence dispersal outcomes.

Third, larval dispersal distances can be shorter than expected from a simple combination of advection, diffusion, and PLD [5, 24],[8]. Combined with the growing perception that passive drift of larvae with currents could not account for this small scale (e.g. [25, 26]), these lines of evidence have led many to presume that behaviour by larvae may restrain dispersal.

Larval swimming is expected to be constrained by temperature due to hydrodynamic and physiological influences. For small larvae, the higher viscosity of colder water requires more swimming effort than warmer water [27], and speed is more strongly affected by viscosity than by temperature [28]. In larger larvae, effects of viscosity are reduced, but colder water should reduce metabolic rates and inhibit the motor activity required for fast swimming [27]. There is mixed support for these expectations: in the laboratory, larvae of some, but not all, species do swim faster at higher temperatures (see Supplement).

Latitudinal comparisons of swimming performance of larvae are best made with data from laboratory studies that measure "critical speed" at ambient temperatures [22, 29] because more data are available for this metric. At any size, swimming speeds differed little between tropical and warm temperate species, but speeds of cold-water species were only 25 to 50% that of warmer water species, and their ontogenetic increase in speed was slower (details in Supplement). Comparisons of larval-fish behaviour *in situ*, although hampered by the lack of data from cold temperate waters, give a somewhat different picture [22]. *In situ*, at any size, larvae of warm-temperate

species were 4–10 cm s<sup>-1</sup> slower than tropical species, and the ratio of *in situ* speed to lab-based critical speed was larger in tropical than in warm temperate species.

The only clear latitudinal pattern in behaviour of fish larvae is that, adjusted for size, tropical and warm temperate species have similar critical speeds, and these are greater than speeds of cold temperate species. However, tropical larvae may swim faster in the sea than warm temperate species (Supplement). The limited evidence indicates that larvae in warm water environments swim faster and earlier in development, and this implies that larvae in lower latitudes should have more control over their dispersal. If behavioural abilities are used to restrict advection or to find settlement habitat from greater distances, they could decrease the spatial scale of larval dispersal, a possibility supported by dispersal modelling [30], but not tested in the ocean. If so, dispersal distances in warmer waters should be smaller.

### **(b) Temperature, feeding, and mortality**

The perception exists that greater oligotrophy and higher temperatures in lower latitudes should result in more starvation of larvae [1],[31], which if true could influence larval dispersal by slowing growth or increasing mortality (see Supplement). Prey densities and feeding success may play a critical role in survival of pelagic larvae of marine fishes, and these factors can affect the degree to which subpopulations are connected via larval dispersal. This is because the numbers of larvae reaching any location - which affects the spatial extent of larval dispersal [15] - should be inversely related to mortality. If, however, larvae do not starve, but survive in poor condition, they may become more buoyant, and become concentrated near the surface [32]. In this case, passive larvae might be dispersed over greater distances because surface water typically moves faster than deeper water. If

feeding conditions in tropical waters are indeed poorer, one might expect increased dispersal in warmer water.

Are larvae in the tropics subject to poorer feeding conditions or greater mortality from starvation? Recent literature syntheses identified latitudinal differences feeding incidence, prey types, prey selectivity, and niche partitioning of fish larvae [33], [34]. Feeding rates are greater in the tropics, and fish larvae in low and high latitudes appear similarly successful at feeding (see Supplement), contrary to expectations. However, empirical estimates of starvation mortality are very limited [35] [36], and none exist for tropical demersal species. Differences in the feeding ecologies of larval fishes between low and high latitudes are present, but little empirical evidence suggests that they result in latitudinal distinctions in dispersal or systematic geographic patterns in mortality (see Supplement).

### **(c) *Temperature, development, and PLD***

Based solely on temperature-driven variation in physiological processes, larvae of tropical species are hypothesized to have reduced potential for dispersal due to faster development times and shorter PLD than temperate species (e.g., [27],[3],[37]). To test the expectation that PLD would be shorter in low latitudes, we examined PLD data for differences among latitudes (see Supplement for details).

Surprisingly, regional differences in PLDs appear to be larger than differences between warm temperate and tropical sites (figure 3). These analyses, although attempting to control for habitat, reproductive mode, and region, are still confounded by taxonomic influences (see Supplement). Therefore, for the nearshore demersal species for which PLD data are available, the expectation that warm temperate PLDs

were longer than tropical PLDs was not fulfilled. More comprehensive coverage of taxa and high-latitude PLD data are needed to relate PLD to latitude or temperature definitively. Finally, the relationships between PLD and other proxies for dispersal (such as genetic structure or species range) are not compelling (see Supplement).

## **DISCUSSION**

The widespread view that larval dispersal and the spatial scale of population connectivity of marine fish populations differ with latitude is very plausible when theoretical considerations alone are considered. Based on either limited empirical data or these same theoretical considerations, several authors have concluded that larval dispersal likely takes place over larger scales in higher latitudes. We find only partial empirical support for this view, and the existing support is based primarily on differences in spawning mode and larval-fish behaviour between tropical and warm temperate regions vs. cold temperate regions, and on habitat-fragmentation considerations.

*Biological differences:* Existing evidence indicates that species with demersal eggs have smaller scales of genetic connectivity and generally shorter PLDs than broadcast spawners, both of which are commonly assumed to be proxies for larval dispersal distance (but see above and the Supplement for a critical evaluation of the relationship between genetics, PLD, and actual dispersal distance). Most high-latitude demersal shorefish taxa are not broadcast spawners, and this should reduce the average scale of larval dispersal at high latitudes. At low to mid latitudes, most species are broadcast spawners, and this should increase the average scale of larval dispersal. This is contrary to the inferences drawn from habitat fragmentation data, some oceanographic variables, and the influence of temperature on physiology

and behaviour. Regional differences in many factors that influence larval dispersal do exist, but the ultimate net effect of these contrasting factors on larval dispersal is far from clear: direct measures of dispersal across large geographic regions are required.

Putative latitudinal differences in spawning mode, PLD, and genetic structure have been confounded by the use of data from non-representative subsets of the resident nearshore demersal fishes, biased toward pelagic spawners at high latitudes and demersal spawners at lower latitudes. However, high-latitude demersal fish assemblages are actually dominated by demersal-spawning species whereas pelagic spawners dominate warm temperate and tropical fish assemblages. Care must be taken to ensure that questions are framed and conclusions are qualified with full regard to the mix of species for which data exist.

*Physical differences:* In contrast to the biological and biophysical variables reviewed here, physical oceanographic variables have featured in few explicit hypotheses of latitudinal differences in larval dispersal. Although we develop several physical-oceanographic-based hypotheses in the Supplement, there is little relevant information available to test them. Water movement, the strength of upwelling and the mixed layer depth are factors that differ latitudinally and are likely to affect the horizontal and vertical movements of larvae. Although factors affecting coastal circulation may vary over degrees of latitude, local and regional variation can also be large. Therefore, it will be difficult to determine how and to what extent physical factors may vary with latitude in their influence on larval dispersal.



Dispersal can also be affected by the frequency and spacing of suitable settlement targets, especially islands. Island habitat relative to continuous continental habitat changes along a latitudinal gradient, with more island habitat in the tropics. Thus it is possible that tropical fishes restricted to discontinuous habitat may have shorter dispersal distances than their temperate counterparts, although empirical evidence for this is lacking.

*Biophysical differences:* Tropical waters are warmer, and it is commonly assumed this will increase development rates : more rapid development should shorten both the pre-hatching period of pelagic eggs and PLD, and hence, it is assumed, dispersal distances. Unfortunately, the correlation between PLD and dispersal distance is weak at best in the species for which there are sufficient data for testing, and data suggesting shorter PLDs in the tropics are also subject to bias because available PLD data are not representative of the taxonomic composition or spawning modes of either tropical or temperate regions. Length of PLD is influenced not only by spawning mode, but also by adult habitat and region within the same latitudinal range, as shown here, even though our analysis is confined to nearshore demersal fishes at latitudes below 50°. Importantly, even within spawning modes, clear differences between tropical and warm temperate areas are lacking. Thus, there is no simple relationship between water temperature (or latitude) and PLD, and careful partitioning of data is required for valid latitudinal comparisons. Pelagic eggs take longer to hatch in cold water [13], and drift during this time may increase dispersal distances for broadcast spawners at high latitudes.

It is important to note that although there is ample evidence of within-species temperature-dependent responses of physiological processes related to dispersal

and survival, the actual effects in nature might be minimized through adaptation of key traits. Thus, it is unclear whether well-known physiological effects of temperature actually result in geographic variation in dispersal distance or connectivity. Certainly, the strong regional and taxonomic effects on PLD (see above and Supplement) suggest that there is wide scope for adaptation.

Larval behaviour, particularly swimming and feeding, could affect realised dispersal: both strong directed swimming and increased mortality from starvation potentially can shorten average dispersal distances. Although there is some evidence that tropical larvae swim more rapidly than temperate larvae, generalizations are difficult to make, again because of taxonomic differences and limited data from cold temperate species. Further, as with genetic and PLD data, the range of species for which larval behaviour information is available is not representative of either the taxonomic composition or spawning modes of the assemblages from different latitudes. Equally, although there are differences in the feeding ecologies of larval fishes between low and high latitudes, there is little evidence that these differences result in latitudinal distinctions in feeding rates, starvation mortality, or dispersal.

## **FUTURE DIRECTIONS**

There is a clear need for more studies of larval dispersal and population connectivity across latitudinal ranges. Measuring these processes empirically remains challenging. Yet, the importance of connectivity to fisheries management, conservation, and predicting climate-driven changes to marine systems, makes a more general understanding of latitudinal and temperature effects timely and valuable.

The various oceanographic factors considered individually here will interact in the ocean, and it is difficult to predict how they will influence dispersal when combined. Biophysical modelling that incorporates many of these oceanographic factors [38] will be helpful in understanding how latitudinal changes in physical variables influence larval dispersal .

Future latitudinal comparisons will need to take into account taxonomic composition, adult habitat, and spawning mode if they are to have generality. Ideally, one would investigate a single species over large latitudinal gradients, but few species qualify. One solution is to compare species across more limited latitudinal ranges such as sub-tropical to tropical areas. In addition, there may be cases where one could control for life history and habitat difference among higher taxonomic groupings such as the Family level.

In addition, the goals for measuring connectivity must be defined clearly because these may alter the impact of any biases. For example, if the goal were fishery management or design of marine protected areas for replenishment of fished populations, a different mix of species might be appropriate to study than if the goal were biodiversity conservation or latitudinal trends in ecosystem processes. Where meta-analysis of previously published data is attempted, care must be taken to qualify interpretation and conclusions when data are biased with regard to species composition or spawning mode. Future examinations of possible latitudinal differences in larval dispersal and population connectivity will need to look beyond published data, and undertake new studies.

We must relate diet and feeding success in larvae to growth, survival, and behaviour in order to understand and model how trophic-related factors ultimately affect larval dispersal and population connectivity. Linking individual-based models of larval growth and mortality to realistic circulation models could facilitate comparisons of tropical and temperate regions [39], although many of the caveats identified here will still apply, and field-testing of model predictions is required.

Currently available estimates of PLD are largely based on few individuals from very limited locations [12]. These studies have also focused on a limited range of taxonomic groups and habitats, which makes broad latitudinal comparisons problematical. It would be valuable to broaden the taxonomic base and habitats for PLD estimates, as well as to obtain better measures of within-species variation in PLD values, especially if PLD varies with location. Most PLD estimates derive from otolith counts, and because otoliths frequently do not begin to form until some time after hatching, particularly in species with pelagic eggs, many PLD values are underestimates of the true time in the water column. Better PLD estimates might reveal relationships with latitude-based factors that are not apparent with currently available estimates.

The very limited information available on larval behaviour of temperate species is another obstacle to general comparisons among areas. It would be useful to study larvae of the same species from different latitudes within its natural range when considering behaviour or effects of temperature on physiological processes to help determine the scope for adaptation. In addition, such information is needed on a broader range of species and habitats.

At present, most of the available genetic data for high latitudes are from the northern hemisphere (particularly the Atlantic), and are from larger, often pelagic, species that are of commercial interest. More single-species studies examining trends in dispersal and gene flow along latitudinal gradients are needed. Translating the observed genetic patterns into demographic trends remains challenging [40],[41]. Better integration of genetic, demographic, and life history studies will be needed to further disentangle the patterns observed.

Managers are most often interested in direct measures of demographic connectivity [40, 41]. Advances in otolith-based approaches and genetic-parentage approaches are being applied successfully in warmer waters, where the life histories of the fishes make these approaches particularly advantageous. They have been little applied at higher latitudes, but hold great promise.

## **CONCLUSIONS**

It is important to emphasize that our conclusions apply to demersal shorefishes, and not necessarily to pelagic fishes or those from deeper waters. It is not clear that latitudinal differences in larval dispersal or associated factors exist at the level of broad faunas; certainly, they have not yet been clearly demonstrated for larvae of demersal shorefishes. This may be due to lack of appropriate data, or the use of 'off the shelf' data that are biased with regard to the species assemblages in the areas of concern. Biases may be introduced from both differences in taxa or spawning modes at different latitudes as well as limited latitudinal sampling, and as we move away from ideal study types, the uncertainty increases.

Many factors lead to expectations that larval dispersal should differ latitudinally, and although most suggest broader dispersal at higher latitudes, some do the opposite. Limited evidence is available to evaluate some of these expectations, especially for higher latitudes, and for a broad array of taxa. Some hypotheses of differences are not supported by the evidence that is available on demersal shorefishes.

Considerations of this issue have been dominated by untested assumptions, acceptance of logical, yet unsupported assertions, and limited empirical evidence.

More research on a broad array of the many factors that influence larval dispersal is required to make progress on this subject.

ALS thanks Michel Kulbicki for helpful discussion and advice. Suzanne Bullock provided editorial assistance. S.D. Simpson and T.J. Miller provided helpful criticisms. TK was supported by the Norwegian Research Council through project MENUUI #190286. JML was supported by ARC Discovery Grant DP110100695. JEC and RRW were supported by the Partnership for the Interdisciplinary Study of Coastal Oceans, funded by The David and Lucille Packard Foundation and the Gordon and Betty Moore Foundation. This is PISCO publication ###.

JML, JEC and RRW initiated and constructed the manuscript, provided overall editorial direction, and the introduction and conclusions. JML provided sections on taxonomy, biogeography, larval behaviour, and spawning modes. IRB and RDV provided genetics sections. JKL provided feeding sections. TK and CBP provided physical oceanographic sections. MJM provided sections on eels. MIO and SMS provided sections on temperature effects. ALS provided PLD sections with additions from JML. SES and EAT provided habitat fragmentation sections. RRW provided caveats sections. All contributed to future directions section.

## Figure Captions

Figure 1. Effects of the degree of habitat patchiness based on an analysis of published otolith chemistry studies on: (A) the % of self-recruitment; (B) the scale at which self-recruitment was measured; and (C) the scale over which populations were connected. Different letters above columns indicate significant pairwise differences based on post-hoc Tukey tests. Continuous refers to relatively continuous habitat on continental margins; coastal habitat patches are saltmarshes, mangroves, seagrass beds or reefs.

Figure 2. Changes in the number of: (a) all island patches and (b) non-continental island patches as a function of latitude. Each relationship was analysed by Pearson correlation. See Supplement for details of analysis.

Figure 3. Average ( $\pm$  95% confidence interval) pelagic larval durations of temperate (solid squares) and tropical (open squares) reef fishes. A) Data from all geographic locations and spawning modes combined and PLDs of demersal and broadcast spawning species plotted separately. B) Data plotted by geographic region with spawning modes combined. C) Data for demersal spawning species plotted by geographic region. D) Data for broadcast spawning species plotted by geographic region. If 95% confidence intervals overlap, means are not significantly different, but if they do not overlap they are significantly different as confirmed by t-tests.

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Figure 1

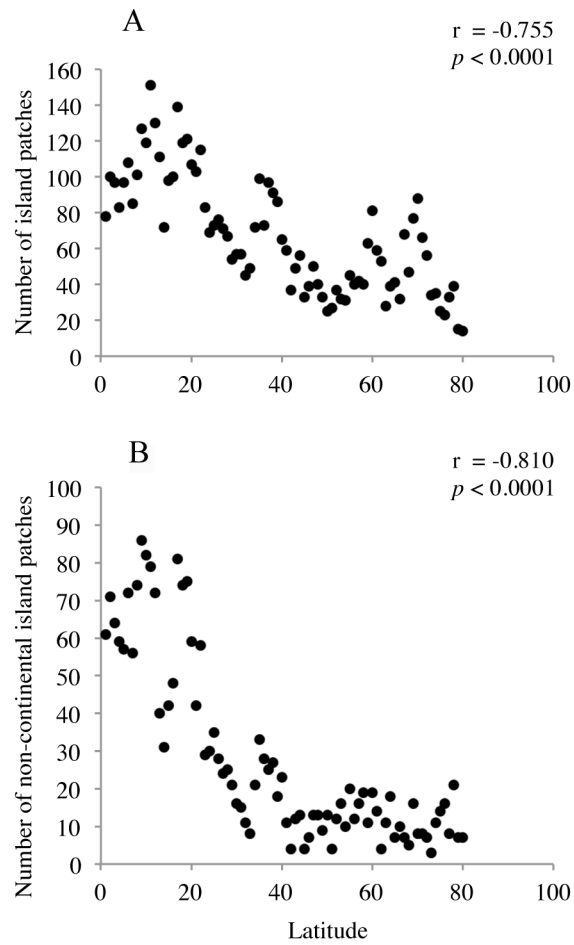


Figure 2

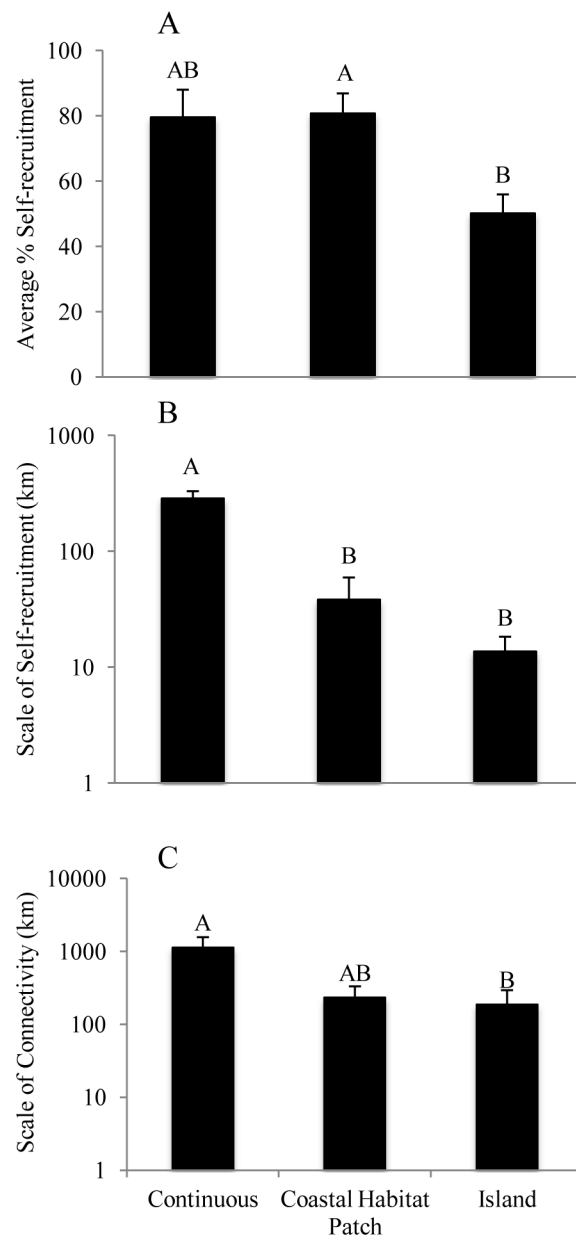
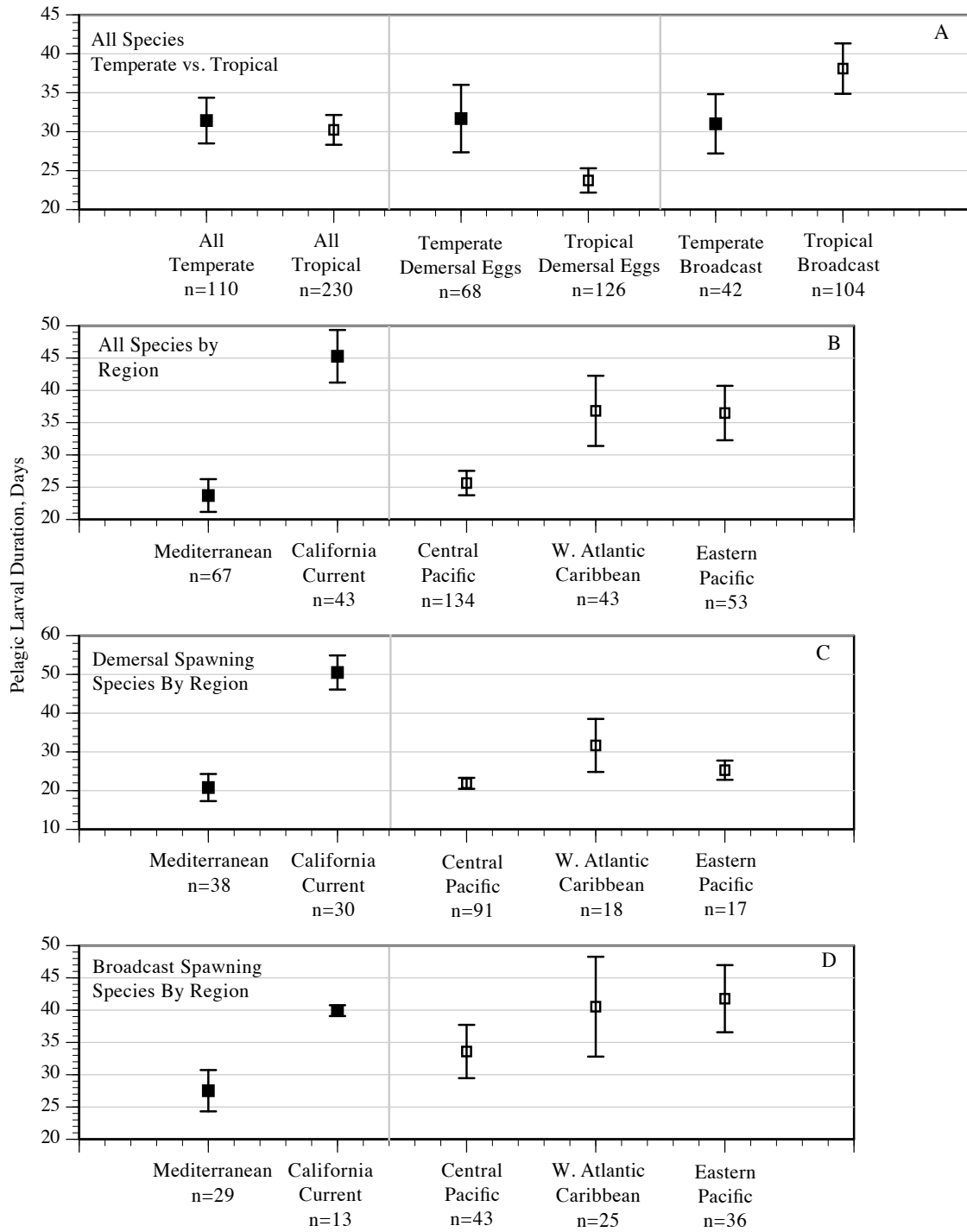


Figure 3



1 **Supplement to Leis et al. Does fish larval dispersal differ between high and**  
2 **low latitudes?**

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4 Llopiz, Michael J. Miller, Mary I. O'Connor, Claire B. Paris, Alan L. Shanks, Susan M.  
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6

7

8 **1. INTRODUCTION**

9 A large number of physical and biological variables that might affect larval dispersal  
10 vary with latitude, and many were considered in the present study. Some of these  
11 variables might theoretically have an influence, but evidence is lacking; some have  
12 been considered in previous studies to have an influence, but lack theoretical or  
13 empirical support; and for some, there is support for an influence. This supplement  
14 includes background information and analyses for each of the main factors  
15 mentioned in the text. For each category, we also provide caveats about underlying  
16 assumptions and alternative interpretations. We close with a consideration of bias  
17 introduced by differences in methodology or target species.

18

19 **2. BIOLOGICAL DIFFERENCES**

20 **(a). *Taxonomic composition, biogeography, PLD, and spawning modes***

21 *Taxonomic composition and biogeography:* Warm waters are dominated by  
22 Anguilliformes, holocentroid Beryciformes, Tetraodontiformes, and the Perciform  
23 suborders Percoidei, Blennioidei, Gobioidae, Labroidei, and Acanthuroidei. These  
24 taxa constitute 73 to 84% of the demersal, teleost shorefish species in the three  
25 tropical areas shown in Table Sup 1. In contrast, cold waters are dominated by

26 Gadiformes, Perciform suborders Zoarcoidei and Notothenoidei, and by the  
27 Scorpaeniform suborders Cottoidei and Hexagrammoidei and scorpaenid genus  
28 *Sebastes*. These taxa constitute 73 to over 90% of the demersal, teleost shorefish  
29 species in the three cold-water areas listed in Table Sup 1. Flatfishes  
30 (Pleuronectiformes) of most families are essentially warm water in distribution, but  
31 Achiropsettidae, Pleuronectidae, and to a lesser extent, Scopthalmidae have  
32 colder-water distributions. Unfortunately, there is very little information about taxon-  
33 specific dispersal differences, and given the non-independence of taxa and  
34 distributions, without this information it will be difficult to determine if any identified  
35 differences were due to adaptations to location-dependent physical and biological  
36 conditions, or due to lineage-related factors. Therefore, caution is required when  
37 attributing causes to any apparent differences in larval dispersal and connectivity  
38 between latitudes.

39  
40 *PLD*: Measurement of PLD (pelagic larval duration) is typically estimated by  
41 counting daily rings in otoliths either at settlement or inside a settlement mark [1].  
42 The PLD values used herein were from the published sources detailed below and  
43 were culled from this literature only on the basis of adult habitat, and by eliminating  
44 pelagic species. Demersal fish Orders and Suborders with primarily warm water  
45 distributions (as identified in previous section) appear to have lower mean PLD  
46 values than do taxa with primarily cold-water distributions. The PLD data in [2] and  
47 [3], which addressed 45 California Current and 727 largely tropical species,  
48 respectively, were used to determine mean values at the Ordinal and Subordinal  
49 level. These are not necessarily a representative sample of the species belonging to  
50 the Orders and Suborders identified as having either primarily warm or cold-water

51 distributions above, so these data should be interpreted cautiously. The taxa and  
52 the number of PLD estimates in each are: warm water taxa, Anguilliformes (marine  
53 eels, i.e., not including *Anguilla*) 4, Holocentroidei 10, Acanthuroidei 81, Blennioidei  
54 20, Gobioidae 36, Percoidae 87, Labroidae (Labridae) 213, Labroidae (Pomacentridae)  
55 265, Scorpaenoidei (*Scorpaena*) 2, Tetraodontiformes 9; cold water taxa,  
56 Gadiformes 9, Zoarcoidei 6, Cottoidei 13, Scorpaenoidei (*Sebastes*) 17. The mean  
57 values for each taxon are shown in figure Sup 1. The mean values for each cold-  
58 water taxon (55-108 days) are all larger than the mean values for warm water taxa  
59 (23-52 days), with the exception of non-anguillid Anguilliformes, with mean of 118  
60 days, more than twice that of other warm-water taxa. A challenge to interpretation of  
61 this analysis is that the temperate-tropical comparisons are biased by habitat  
62 differences. At least half of the temperate species in [2] were continental shelf and  
63 slope species whereas nearly all of the tropical species in [3] were from shallow  
64 reefs. Given that Shanks and Eckert [2] found large habitat-related differences in a  
65 number of life history traits (including PLD), ascribing PLD differences to a latitudinal  
66 effect without taking habitat into account is problematical.

67

68 Although these data indicate there is a taxonomic component to PLD, it is important  
69 to note that there were more than 727 PLD values for warm-water taxa, and only 45  
70 values for cold-water taxa, providing further reason to treat the data cautiously. The  
71 very high PLD values for the marine eels, a largely warm-water group, are also an  
72 indication that general statements about latitudinal differences in PLD will likely be  
73 difficult to sustain. But here, too, the fact that only four values were available for a  
74 group with about 800 species provides ample reason for caution, and indicates how  
75 understudied marine eels are.



76

77 We used an ecological-based definition of PLD (i.e., the early life-history period  
78 before settlement [4]), which clearly has meaning only for demersal species with a  
79 pelagic larval stage (it may also include juveniles if they remain pelagic, e.g., many  
80 Tetraodontiform fishes and the Scorpaeniform genus *Sebastes*). The use of an  
81 ecological criterion avoids ambiguity introduced by morphological criteria (i.e., the  
82 early-life history period before a particular set of morphological milestones are  
83 reached, for example, formation of all fins, or scales). The latter can be relevant for  
84 both demersal and pelagic species, but we avoid it precisely because we focus on  
85 the presettlement stages of demersal fishes.

86

87 *Spawning mode*: Spawning mode, in particular the lack of a pelagic egg (i.e.,  
88 demersal eggs, brooded eggs, or viviparity), has been identified as an important  
89 factor in genetic connectivity or larval dispersal in two synthetic studies that  
90 assessed a broad range of published genetic and PLD data [5],[6]. However, the  
91 study of Riginos et al. [5] was based primarily on warm-water species, with only 14 of  
92 the 148 demersal teleost species considered to have cold-water distributions.  
93 Therefore, the conclusions of [5] about the influence of spawning mode on genetic  
94 connectivity may not apply to cold-water taxa. Further, about half of the warm-water  
95 species in the Riginos et al. [5] data set spawn demersal eggs, a value considerably  
96 higher than the proportion of demersal-spawning species in warm-water faunas  
97 (figure Sup 2). Clearly, it is important to understand how spawning mode varies with  
98 latitude.

99

100 The percentage of species with pelagic eggs varies little with latitude until high  
101 latitudes (roughly 50°N, but possibly at higher latitudes in the southern hemisphere)  
102 are reached, after which a strong shift occurs to demersal eggs (figure Sup 2). High-  
103 latitude fish assemblages are dominated by species with demersal eggs (Fig Sup2),  
104 and in most cases, spawning type is consistent within families. However, several  
105 exceptions exist, and in these, there is a trend within taxa toward demersal eggs at  
106 higher latitudes. Some examples where taxa from higher latitudes are less likely to  
107 be broadcast spawners include the sebastine scorpaenids, a temperate group, which  
108 is exclusively viviparous, whereas other scorpaenid taxa, which are essentially  
109 tropical, are broadcast spawners [7]; the temperate labrid tribe Labriini, which is  
110 dominated by demersal-egg species, whereas all other labrids are broadcast  
111 spawners [8]; and the important commercial taxa, Pleuronectidae and *Gadus*, where  
112 some species are broadcast spawners, and other, more boreal species, spawn  
113 demersal eggs [9].

114

115 There is also a trend for fish eggs to be larger at higher latitudes, but this seems to  
116 apply across all latitudes only for demersal eggs. For pelagic eggs, there is little  
117 trend between the tropics and about 50° North, but poleward of this, pelagic eggs are  
118 larger [10]. It is not known if the same relationship applies in the southern  
119 hemisphere. Egg size and pre-hatching period are typically correlated, so these  
120 times should be longer for northern-hemisphere, high latitude species with pelagic  
121 eggs. Equally, dispersal distances due to passive drift of eggs should be longer than  
122 for species south of 50°N. In addition, independent of egg size, lower temperatures  
123 are expected to increase the pre-hatching period, which may lead to a latitudinal

124 gradient in pre-hatch times and therefore dispersal distances between spawning and  
125 hatching, assuming there is no adaptation.

126

127 Another potential latitudinal factor influencing larval dispersal is the timing of  
128 spawning. Many tropical demersal species have a protracted spawning season,  
129 whereas temperate species tend to have a narrow season, presumably to coincide  
130 with appropriate biological and physical oceanographic conditions [11]. For  
131 temperate species, shifts in temperature may shift the seasonal timing of spawning.  
132 The flounder *Platichthys flesus*, for example, spawns 1-2 months earlier in years with  
133 cooler than normal temperatures [12]. In other species, spawning or occurrence of  
134 larvae is delayed by cooler temperatures [13],[14],[15]. Larvae of spring-spawning  
135 species can appear in the plankton earlier in cold years whereas larvae of summer-  
136 spawning species appear later [16]. Sheaves [17] noted a latitudinal pattern within  
137 the family Sparidae, with greater variation in spawning timing for more temperate  
138 species. Such shifts in phenology will affect potential dispersal if the oceanographic  
139 environment experienced by eggs/larvae differs from that typically experienced.  
140 Spawning that is temperature-induced could result in a mismatch of spawning timing  
141 with timing of food availability or transport mechanisms. If so, this might result in  
142 more variable dispersal distances and directions, and thus a larger scale of larval  
143 dispersal in temperate than in tropical waters (also see Feeding below). In upwelling  
144 systems like the California Current, the timing of spawning has long been recognized  
145 as a critical determinant of larval success [18], and reproductive patterns of resident  
146 fishes appear to have evolved to facilitate completion of the life cycle and minimize  
147 advective losses of larvae under 'normal' oceanographic conditions [2].

148

149 **(b) Diadromy**

150 Diadromy varies with latitude, albeit differently in its different forms, but there is only  
151 limited evidence that connectivity of diadromous species varies with latitude. Life  
152 histories of diadromous fishes are split between fresh and marine waters. We  
153 consider only species with marine larvae, which excludes anadromous taxa such as  
154 salmonids, which spend their early life history in freshwater. However, anadromy is  
155 largely confined to temperate latitudes, particularly in the northern hemisphere, with  
156 few tropical anadromous species, primarily Hilsa shads [19]. This does constitute a  
157 latitudinal difference, but an unbalanced one because there are far more  
158 anadromous fish species in the northern than southern hemisphere.

159

160 Catadromous species such as anguillid eels, two species of Australian percichthyids  
161 (warm temperate), and the tropical latids, centropomids and kuhliids migrate to  
162 estuaries or the sea to reproduce and the young enter freshwater after larval growth  
163 in the ocean. There are only a few temperate catadromous species, and other than  
164 the anguillid eels, there is no clear evidence of latitudinal differences in dispersal or  
165 connectivity. Anguillid eels, however, provide a clear example: temperate anguillid  
166 species have larger scales of larval dispersal and population connectivity than do  
167 most tropical species. All five primarily temperate anguillid species display little  
168 genetic structure across their freshwater adult ranges. Their long-lived larvae are  
169 transported by currents 1000s of km from discrete tropical, oceanic spawning areas  
170 [20],[21],[22]. *Anguilla reinhartii*, with a tropical to warm temperate adult range on  
171 the western side of the Coral and Tasman Seas, is apparently similar to the  
172 temperate species in having a discrete tropical oceanic spawning area and no  
173 apparent genetic structure [23]. *Anguilla marmorata* and *A. bicolor* are primarily

174 tropical species that spawn in the Indian and Pacific Oceans and have more than  
175 one genetic lineage or spawning population [24, 25],[26],[27] within an ocean basin.  
176 The other tropical *Anguilla* species each have limited adult ranges and one or two  
177 local spawning areas [28]. They have shorter larval durations and smaller maximum  
178 larval sizes than temperate species [29], [30].

179

180 Amphidromous species such as Sicydiinae gobiids (primarily tropical), some  
181 temperate cottids, and the temperate Osmeriform families Galaxiidae and  
182 Plecoglossidae spawn in freshwater, with their larvae moving downstream to the  
183 ocean for a larval growth phase after which they return to freshwater for a juvenile  
184 growth phase [31] [32]. Amphidromous fishes are found primarily in tropical and  
185 subtropical islands, but a few extend to temperate islands such as Japan and New  
186 Zealand [33]. Most amphidromous species have long PLDs (e.g., [32],[34]) and  
187 widespread larvae compared with most marine fishes, and they have broad genetic  
188 connectivity (e.g., [35],[36],[37]). In contrast, a temperate Japanese sculpin species  
189 with a short PLD has clear population structure [38]. However, no clear latitudinal  
190 trend in larval duration, dispersal, or connectivity is evident in the species that have  
191 been studied.

192

### 193 **3. PHYSICAL DIFFERENCES**

#### 194 **(a) Oceanography**

195 Generally, stronger winds and greater eddy propagation occur in mid to high  
196 latitudes than in the tropics [39],[40]. This can affect dispersal both directly and  
197 indirectly. Latitudinal increases in both mixed layer depth (MLD) and eddy kinetic  
198 energy (EKE) tend to increase diffusion, which influences the variation in dispersal

199 distances, but an increase in MLD alone would tend to increase dispersal distance  
200 via its interaction with both diel and ontogenetic vertical migration.

201

202 Although seasonal variation in tropical MLD is generally low in any region, there are  
203 large differences in MLD among different tropical regions [41]. Temperate species  
204 that spawn when MLD decreases in spring and summer are expected to have  
205 shorter dispersal distances than those that spawn in winter, and this seasonal  
206 difference should be reinforced by the higher ML temperatures of spring and  
207 summer. This contrast may increase among-species variation in dispersal at higher  
208 latitudes.

209

210 Vertical migration behaviour of larval fish can influence dispersal differentially  
211 between tropical and temperate regions. Most fish larvae dwell within the upper 50-  
212 100 m of the water column in both tropical [42],[43],[44],[45] and temperate [46],[47]  
213 environments and are able to modify their vertical position behaviourally. Where  
214 currents vary in speed and direction with depth (e.g., [48],[49]), larval fish in the  
215 surface layer can be advected in a different direction or speed than deeper-living  
216 larvae. Therefore, vertical distribution behaviour by larvae can affect their integrated  
217 drift trajectories and geographic dispersal [50],[51],[52],[53]. Larval fish released at  
218 the same spawning ground, but that occupied different parts of the water column,  
219 become separated by hundreds of kilometers after a few months drift [54],[55],[56].  
220 In both temperate and tropical regions, warmer temperatures and low winds during  
221 the summer cause an increase in stratification and a shallowing of the MLD. The  
222 strongest currents generally occur in the ocean's surface mixed layer and decrease  
223 with depth, particularly in the pycnocline. Stratification in the upper water column and

224 MLD can influence the impact of larval vertical migration on dispersal and  
225 connectivity [42],[52],[57]. In general, where MLD is large (e.g., in many temperate  
226 areas during times other than summer) dispersal distances should be relatively long,  
227 creating larger connectivity networks.

228

229 The increase in kinetic energy, and therefore eddy generation, with latitude seems  
230 directly linked to the Coriolis-dependent geostrophic motion, which increases with  
231 latitude, and with wind, which is strongest at mid to high latitudes [58]. The size of  
232 eddies decreases with latitude because the intrinsic length scale of baroclinic  
233 instabilities is directly related to latitudinal variation of Coriolis force [58], and as a  
234 result the typical size of eddies decreases from 200 km at mid latitudes to 100 km at  
235 high latitudes. The strongest variation in eddy kinetic energy (EKE) and amplitude is  
236 associated with western boundary currents [59], resulting in within-latitude regional  
237 EKE differences. Eddies in the tropics are generated at large scales by Rossby  
238 waves (very low-frequency waves in the ocean's surface and thermocline), tropical  
239 instability waves [60], or by the interaction of currents with topography such as capes  
240 or the reef edges that can result in shedding of sub-mesoscale eddies [61],[62].

241 Thus, there are clear differences in eddy formation and size with latitude. Although  
242 slow-moving eddies can retain larvae near a source (e.g., [63]), fast-moving eddies  
243 can move larvae away from a source [64], so it is difficult to generalize about how  
244 latitudinal differences in eddies might influence larval dispersal.

245

246 Temperate regions are characterized by the widest seasonal temperature  
247 ranges([41]), and frequent storms and low-pressure systems that mix the water  
248 column, and can have major impacts on the dispersal of larvae [65],[63],[66].

249 Tropical regions have equivalent low-pressure systems, that can disperse larval fish  
250 from their source relatively quickly, but tropical cyclones are, short-term, focused  
251 storms that are much less frequent and followed by long relatively quiet periods [67].

252

253 Advection alone moves larvae away from the source whereas diffusion by itself  
254 increases variance in dispersal distance, but not the mean [68] (figure Sup 3). Zonal  
255 advection (i.e., along the east-west axis) in the surface layers increase with latitude,  
256 but the meridional advection (i.e., along the north-south axis) decreases [69].

257 Therefore, connectivity networks are expected to be more elongated zonally at mid-  
258 and high-latitudes and meridionally in the tropics. East-west spreading rate  
259 increases with time in the tropics but not at high latitudes [70]. Therefore, super-  
260 diffusive east-west spreading should be expected at low latitudes only for species  
261 with PLDs, like eels, but not for species with short PLDs.

262

263 Ekman-related coastal upwelling cells may be important for nearshore retention of  
264 larvae [63] because they allow larvae that move offshore in Ekman near-surface flow  
265 to migrate downward and be returned toward shore by deeper flow ([63],[71],[72]).

266 Coriolis force combined with long-shore wind drives Ekman coastal upwelling cells.

267 Although Coriolis force increases with latitude, wind strength is maximal in mid  
268 latitudes, so it is expected that Ekman-cell retention would be most important at mid

269 latitudes, and least important at low latitudes. If so, away from the tropics this

270 phenomenon should result in a decrease in the scale of dispersal for species that

271 utilize these cells. However, intensification of Ekman upwelling should result in

272 greater offshore transport of shallow-living larvae. Although these arguments are



273 theoretically sound [73], there is little evidence that many species do utilize Ekman-  
274 cell retention in nature [72],[74].

275

276 There is currently no empirical evidence that any of these identified oceanographic  
277 factors result in latitudinal differences in larval dispersal. By identifying their potential  
278 to influence larval dispersal, we hope to stimulate research into their effects.

279

### 280 **(b) *Habitat Fragmentation***

281 Fish larvae in warm water have good swimming abilities and can detect settlement  
282 habitat over a range of scales [75],[76],[77]), although information on larval sensory  
283 abilities in cold water is lacking. If these abilities are used to remain close to suitable  
284 benthic habitat (e.g., orientation behaviour, [78]), larvae from islands would likely  
285 remain near their origin by orientated swimming, because they are likely to receive  
286 sensory information about suitable benthic habitat only from the closest island. In  
287 contrast, a larva from coastal waters along a continental margin will receive sensory  
288 information about suitable benthic habitat as long as it minimizes offshore dispersion,  
289 even if along-shore movement is large. Hence, there is an expectation for less self-  
290 recruitment and greater population connectivity along continental margins than  
291 among islands (e.g., [79]).

292

293 The degree to which habitat patchiness influences dispersal distance in either  
294 tropical or temperate latitudes is largely untested. To explore this question, 20 recent  
295 studies of demersal fishes (Table SUP2) that used otolith geochemistry to estimate  
296 demographic connectivity were categorized based on whether the target species  
297 lived in: a) relatively continuous habitat (e.g., soft sediments in shelf waters); b)

298 coastal habitat patches; or c) offshore islands. For each study we then calculated:  
299 (1) the % self-recruitment observed (or stated); (2) the spatial scale over which self-  
300 recruitment was estimated (i.e., the scale at which the population/stock was defined);  
301 and (3) the spatial scale of observed connectivity. Whether these three metrics  
302 differed among landscapes was tested by ANOVA.

303

304 Levels of self-recruitment in a demographic context were high (mean: 63%) and  
305 were 30% greater in continental than island habitats ( $F_{2,13} = 7.045$ ,  $p = 0.009$ ; main  
306 text, figure 1A). On first inspection, this appears to contradict the reasoning that  
307 dispersal should be shorter in island areas. However, the spatial scale over which  
308 self-recruitment was estimated differed among landscape contexts ( $F_{2,15} = 9.636$ ,  $p =$   
309  $0.002$ ): self-recruitment occurred in continuous habitat at ~300 km scales whereas  
310 for both the patchily distributed landscape contexts, self-recruitment occurred at  
311 scales <40km (main text, figure 1B). Once we controlled for spatial extents of each  
312 study, the mean scales of connectivity differed among contexts, with species in  
313 patchy habitats dispersing over about 60 to 100 km, whereas species in continuous  
314 habitats dispersed about 900 km (main text, figure 1C). Combined, these data  
315 suggest that larval dispersal may be more restricted in fragmented habitats.

316

317 Are fragmented habitats more common in the tropics? To examine whether  
318 landscape context plays a role in driving differences in population connectivity  
319 between tropical and temperate species, we assembled a spatial database  
320 consisting of the Global Self-consistent, Hierarchical, High-resolution Shoreline  
321 (GSHHS; [80]) and a lattice of one-degree latitude boundaries between +/- 80°

322 latitude (Mollweide equal-area cylindrical projection). We determined the distribution  
323 of islands as a function of latitude.

324

325 To account for the observation that all suitable habitats within the sensory detection  
326 limit of larvae may not be perceived by larvae as distinct, we generalized the global  
327 shoreline data to match this scale of habitat detection. If larvae can detect habitat  
328 patches at this spatial scale, then behavioural barriers to dispersal and their impact  
329 on connectivity will be observed at scales larger than this. For this analysis we  
330 chose 5 km, which is within the range estimated for reef fishes [81, 82], although  
331 using either shorter (2.5 km) or longer (10 km) detection limits resulted in almost  
332 identical results. This generalization resulted in a grid (at 2.5 km resolution) where  
333 cell values quantify the amount of island-only habitat within the 5 km sensory radius  
334 from the centre of each cell. Island habitat cells that fell within this sensory zone  
335 were then aggregated/merged into contiguous island patches. Using the same  
336 sensory detection limit, we also reclassified island habitat patches to continental  
337 habitat if they were within the 5 km detection limit from continental shorelines. Island  
338 habitat patches were intersected with the latitude lattice to calculate the median  
339 latitude of each patch. Changes in island number with latitude were analysed by  
340 correlation. Both all-island patches (i.e., all islands regardless of distance from a  
341 continental margin) and all non-continental island patches (i.e. island patches greater  
342 than 5 km from a continental margin) declined in number with increasing latitude  
343 (main text, figure 2).

344

#### 345 **4. BIOPHYSICAL DIFFERENCES**

##### 346 **(a) *Temperature and larval behaviour***

347 Development of swimming in larvae of demersal fishes has been recently reviewed  
348 [76, 83]. Larvae of taxa from cold-temperate waters (Cottidae, Gadidae,  
349 Pleuronectidae) had some of the slowest critical speeds [84], and consistently swam  
350 at about 5 body lengths per second ( $\text{BL s}^{-1}$ ). Development of critical speed was  
351 more variable in larvae of taxa from warm-temperate waters (Moronidae,  
352 Percichthyidae, Sciaenidae, Sparidae). Larvae of these taxa were relatively slow  
353 when small ( $5\text{--}10 \text{ cm s}^{-1}$ ), but at sizes larger than 5 mm, sciaenids swam at  $10 \text{ BL s}^{-1}$ ,  
354 whereas sparids and percichthyids larger than 7–8 mm are able to swim at  $15\text{--}20$   
355  $\text{BL s}^{-1}$ . Moronid larvae, in contrast, were slow ( $<10 \text{ BL s}^{-1}$ ) until larger than 15 mm,  
356 after which they swam at about  $15 \text{ BL s}^{-1}$ . Aside from the serranids and lutjanids  
357 which were slow at smaller sizes, but eventually were among the fastest larvae,  
358 tropical taxa had larvae that were fast throughout development, with critical speeds  
359 faster than  $10 \text{ cm s}^{-1}$  and with most species swimming at  $15\text{--}20 \text{ BL s}^{-1}$  for much of  
360 their larval phase, with some reaching almost  $30 \text{ BL s}^{-1}$ . Fisher [85] suggests that  
361 most coral-reef fish families have sufficient swimming capabilities to be able to  
362 influence their dispersal outcomes substantially for over 50% of their larval phase.  
363 For temperate *Sebastes*, in contrast, which may have PLDs of up to 6 months, the  
364 potential for using directed horizontal swimming to influence dispersal may not be  
365 developed until the pelagic juvenile stage at 20-30 mm (see below).

366

367 Although the swimming data on warm temperate species were from outside tropical  
368 latitudes, the experimental temperatures ( $19\text{--}27^\circ\text{C}$ ) overlap those of the tropical  
369 species ( $26\text{--}31^\circ\text{C}$ ). The closeness of these temperature ranges could result in the  
370 similarity in swimming performance. In contrast, a large gap exists between these  
371 and the experimental temperatures for cold temperate species ( $3\text{--}10^\circ\text{C}$ ), contributing

372 to the differences in swimming performance. Measures of swimming performance  
373 for larvae of species that live at temperatures within this gap are needed.

374

375 Within species, the expected pattern of increased swimming speed with increased  
376 temperatures has been observed in larvae of several temperate fish species  
377 [86],[87],[88],[84],[89, 90], but a cold temperate species of sculpin (Cottidae) had the  
378 opposite response [84], and a tropical damselfish had no temperature-induced  
379 change in swimming speed [91]. Physiological responses to temperature are  
380 typically dome-shaped, and the apparent inconsistency in the limited empirical data  
381 might be due to differences in the portion of the temperature-response curve that  
382 was under study.

383

384 It is expected not only that the development of swimming capacity is slower in cold  
385 temperate species due to protracted morphological development, but there should  
386 also be an increasing ontogenetic divergence of swimming capabilities [76]. Larvae  
387 of live-bearing *Sebastes* rockfishes are released at a size of 4-6 mm with incomplete  
388 fins and have critical swimming speeds of  $< 2 \text{ cm s}^{-1}$  at  $12^{\circ}\text{C}$  [92]. In contrast,  
389 tropical and warm temperate larvae at this size have more advanced fin  
390 development and  $U_{\text{crit}}$  values of 2-10  $\text{cm s}^{-1}$  [93], but the extent to which these  
391 swimming speed differences are due to temperature or to fin development is unclear.  
392 The size-dependent rate of increase in swimming speed of cold temperate species  
393 lags behind that of tropical species, as expected. For each mm increase in size,  
394 tropical serranids (*Epinephelus* spp.) swam faster by 2.1-2.6  $\text{cm s}^{-1}$  [76], whereas the  
395 increase for the temperate serranid *Paralabrax clathratus* and several species of  
396 *Sebastes* was only 1.0  $\text{cm s}^{-1}$  [92]. Many tropical species settle at small sizes ( $<15$

397 mm), and have speeds equivalent to warm temperate species at similar sizes,  
398 assuming that all fins have formed. Larvae of some species attain relatively large  
399 sizes (> 20 mm) before settlement. The tropical species are fast swimmers, with  
400 critical speeds in excess of  $20 \text{ BL s}^{-1}$  by the time they settle [76, 94], but among  
401 temperate species, data are available only for species of the scorpaeniform genus  
402 *Sebastes*, which at 25-30 mm, have critical speeds of about half those of tropical  
403 species [92].

404

405 Swimming endurance of many settlement-stage tropical fish larvae is very high [95],  
406 but information on the development of endurance is limited, especially for temperate  
407 species. There was no obvious difference in swimming endurance between warm-  
408 temperate and tropical species, except that several tropical species attained greater  
409 endurance prior to settlement primarily because they settle at larger sizes [76]. There  
410 are no endurance data for larvae of cold-temperate species.

411

412 A critical aspect of behaviour-influenced dispersal is performance in relation to age  
413 (or time in the water column), not to size. The experimental data are based largely  
414 on reared larvae with growth rates that differ from those found naturally, so a simple  
415 conversion of size to age is not generally possible. However, it is generally expected  
416 that growth rates are temperature-dependent, so converting performance from a size  
417 to an age basis is expected to increase relative differences in performance between  
418 warm and cold-water species. More information is required on field-based growth  
419 rates of larvae of demersal fishes.

420

421 The identified differences in swimming performance are consistent with theoretical  
422 predictions and could arise from differences in phylogeny, sea water viscosity,  
423 temperature and associated physiological responses or to some combination. At  
424 present, none can be eliminated or shown to be relatively more important to  
425 dispersal. Although the causes for the differences are not clear, the most relevant  
426 consideration is what a larva does in conditions it typically encounters in the sea.

427

428 **(b) *Temperature, feeding, and mortality***

429 Feeding can influence larval survival in several ways. Clearly, food availability must  
430 be sufficient to prevent starvation. Spatial or temporal variability in prey levels above  
431 or below a starvation threshold will translate to concurrent variability in the  
432 proportions of larvae surviving to settle, thereby influencing larval replenishment and  
433 connectivity. Suboptimal prey levels may influence larval growth rate [96],[97], ability  
434 to avoid predators [98],[99] or swimming endurance [100], [101]. Slower growth can  
435 extend the larval period, thus increasing cumulative mortality and reducing the  
436 numbers of settlers [102],[103],[104]. Similarly, slower growth extends the duration  
437 of the more passive early stages and thus the time before larvae are able to  
438 influence dispersal.

439

440 Feeding incidence (FI; the proportion of a sample of larvae with food present in the  
441 gut) is a commonly reported proxy for larval fish feeding success. Across published  
442 studies, average FIs of larvae from coastal and offshore waters were significantly  
443 higher in taxa from lower latitudes (30°S–30°N; median FI 96%) than from middle  
444 (30–45°N/S; median FI 80%) or high (>45°N/S; median FI 72%) latitudes [105].

445 These differences also hold when comparing within taxonomic Orders. This, along

446 with observed differences in gut evacuation rates, indicates feeding rates are higher  
447 in lower latitudes, which would be expected (because of greater metabolic demands)  
448 if levels of starvation mortality were similar among latitudinal regions.

449

450 Zooplankton prey of fish larvae differ with latitude and there are indications of greater  
451 prey selectivity in lower latitudes [105]. Greater diversity of larval-fish prey and  
452 apparent finer niche partitioning in lower latitudes [105] as described in the next  
453 paragraph, could lead to species-dependent dispersal patterns if variation in  
454 abundances of the preferred prey of different taxa does not covary, as some  
455 evidence suggests (Llopiz and Cowen, unpublished data). If so, patterns of larval  
456 dispersal and degrees of connectivity may be less consistent across taxa in lower  
457 latitudes even when adult habitat, spawning behaviours, and PLDs are similar—  
458 clearly adding to the challenges of accurately modeling the successful transport of a  
459 particular species.

460

461 The frequency with which calanoid copepods and copepod nauplii dominate the diets  
462 of larval fishes decreases towards the equator, while the importance of  
463 appendicularians and several genera of cyclopoids increases. The diversity of  
464 dominant prey types also increases towards the equator. Such prey include  
465 cladocerans, bivalve larvae, pteropods, and ostracods, which rarely if ever compose  
466 the majority of a taxon's diet in high latitudes. Diets of low-latitude larvae are more  
467 often consistent in time, space, and ontogeny, and are often unique to larvae of a  
468 given taxon [105],[106],[107]. These factors, together with comparisons of diets to  
469 ambient proportions of zooplankton prey types, indicate greater degrees of prey  
470 selectivity in lower latitudes, with some evidence that highly selective larvae are



471 'hard-wired' to consume specific prey types and have a limited capacity for prey  
472 switching when preferred prey availability is low [106].

473

474 Latitudinal differences in the seasonal cycles of primary and secondary productivity  
475 may also yield trophic-related distinctions in larval dispersal and connectivity. In  
476 higher latitudes, where match-mismatch dynamics of zooplankton and fish larvae  
477 can have an important influence on larval fish growth and survival [108],[109], the  
478 timing and locations of optimal prey availability may occur over narrower spatial and  
479 temporal ranges than in lower latitudes. This could lead to greater interannual  
480 variability in growth, survival, and dispersal, and therefore connectivity at higher  
481 latitudes. In contrast, the more stable levels of productivity in the tropics, in  
482 conjunction with protracted spawning seasons, could result in more consistent  
483 among-year spatial patterns in larval dispersal.

484

485 Some studies have suggested higher instantaneous mortality rates for fish larvae in  
486 warm temperatures [110], and higher predation pressure in clear tropical waters  
487 [111]. Further, a general trend toward stronger predation in the tropics has been  
488 suggested for other taxa including marine consumers [112],[113],[114]. In contrast,  
489 expected (but not documented) longer larval durations in cold water would imply  
490 higher cumulative larval mortality [115]. Thus, physiological processes underpinning  
491 mortality rates suggest geographic variation in mortality due to temperature, but it  
492 remains unclear whether there is an overall trend that could influence connectivity.

493

494 **(c) *Temperature, development, and PLD***

495 Previous work showed that PLD differed among adult habitat types [2]; for example,  
496 average PLD of nearshore (<30 m adult depth) California Current fishes was much  
497 less than PLD of shelf/slope species regardless of spawning mode. Because  
498 differences in adult habitat can affect PLD, our analysis was based on tropical and  
499 warm temperate (<50° from the equator), nearshore reef fishes only, because  
500 sufficient data for higher latitudes or other habitats were not available. A notable  
501 limitation is the very restricted PLD data available from cold temperate nearshore  
502 fishes and from tropical shelf species.

503

504 Shanks and Eckert [2] compared a diversity of life history characteristics, including  
505 PLDs, of fish and benthic decapod species from the California Current and found  
506 large differences based on adult habitat. The data sorted into three groups:  
507 shelf/slope species, shallow water species (< 30 m), and shallow water species from  
508 the Southern California Bight. Shanks and Eckert suggested that differences in the  
509 life history traits were adaptations to the local hydrodynamics associated with each  
510 adult habitat. We attempted to investigate PLDs of temperate and tropical  
511 shelf/slope fishes, and although we could find adequate data for temperate species,  
512 we could not find enough data on tropical shelf/slope species to support an analysis.  
513 Similarly, data on cold temperate species were too limited for testing. Therefore,  
514 values of PLD for shallow water reef fishes from temperate and tropical locations  
515 were compiled for the Mediterranean and Eastern Atlantic [116],[117],[118],[119], the  
516 California Current [2], the Caribbean and Western Atlantic  
517 [120],[121],[122],[123],[124],[125],[126],[127],[128],[129],[130],[131],[132]), the  
518 tropical Eastern Pacific [131],[133],[132],[134],[135], and the central Pacific  
519 [1],[136],[136],[132]. We assembled data on 110 warm temperate and 230 tropical

520 species from 23 publications. The publications were found by a search of Aquatic  
521 Sciences and Fisheries Abstracts back to 1980 using the terms PLD and Pelagic  
522 Larval Duration. In addition, the reference section of papers that reviewed PLD were  
523 scanned for additional publications, and Science Citation Index was searched for  
524 publications that cited the more important PLD papers. The species found in this  
525 search were included in the analysis if the data were collected in a similar fashion  
526 (e.g., shallow water/reef species, PLD from otoliths) from defined geographic  
527 settings (e.g., California Current, Caribbean, Mediterranean, etc.). The mode of  
528 reproduction (i.e., live birth, broadcast spawning, or nesting) of the species that fit  
529 within the habitat and geographic range criteria was determined from a number of  
530 sources including reference books on California Current species, Fish Base and the  
531 original publications.

532

533 PLD of warm temperate and tropical reef fishes did not differ (all areas and spawning  
534 types pooled: main text, figure 3A, for statistics see main text, figure 3 caption), but  
535 when the species were compared by reproductive mode (demersal vs. broadcast  
536 spawning), PLDs of tropical demersal and broadcast spawners were significantly  
537 shorter and longer, respectively, than warm temperate species (main text figure 3A).  
538 However, when the data were analysed by geographic location, consistent  
539 differences between PLDs in warm temperate and tropical species were not found.  
540 If PLDs are separated by region rather than reproductive mode (main text figure 3B),  
541 PLDs of warm temperate Mediterranean species are significantly shorter than  
542 temperate California Current species, even though the latitudes are similar, whereas  
543 the PLDs of tropical central Pacific species are significantly shorter than those of  
544 Caribbean and tropical Eastern Pacific species, which did not differ from each other.

545 Comparing warm temperate to tropical, the average PLDs for Mediterranean and  
546 central Pacific species were similar whereas average PLD for California Current  
547 species was significantly longer than in any other region, temperate or tropical.  
548 When divided by reproductive mode, the demersal-spawning species average PLD  
549 from the Mediterranean was significantly shorter than in the California Current (main  
550 text, figure 3C), but similar to the central Pacific and eastern Pacific and only slightly  
551 shorter than PLDs in the Caribbean. The broadcast-spawning-species average PLD  
552 from the Mediterranean was again significantly shorter than in the California Current  
553 and was also shorter than in the three tropical regions (main text figure 3D). The  
554 broadcast-spawning species mean PLD for the California Current was not different  
555 from the Caribbean or Eastern Pacific, but was significantly longer than the central  
556 Pacific.

557

558 The most striking differences in PLDs were not between temperate and tropical  
559 systems, but between two temperate regions: for both demersal and broadcast  
560 spawning species, Mediterranean PLDs were much shorter than in the California  
561 Current. The Mediterranean is not particularly warm, but Mediterranean PLDs were  
562 either the shortest or not significantly different from the tropical areas. Further,  
563 Caribbean and California Current pelagic-egg PLDs did not differ, showing there is  
564 not a simple relationship between PLD and temperature.

565

566 The main text points out that the geographic patterns we found in PLD are not  
567 expected based on sea-surface temperature alone. The temperature data (annual  
568 range of monthly means of sea surface temperature, viewed May 2012) supporting  
569 this statement for each region are given here. They are from

570 <http://www.youtube.com/watch?v=4hv1a-z53Ew> (Generated from NASA Giovanni  
571 v3.0.7) for the western Mediterranean (12-26°C), a temperate area, and from  
572 <http://www.nodc.noaa.gov/dsdt/cwtg/oatl.html> for the other regions. For temperate  
573 areas, ranges were: central California (12-16°C) and southern California (14-20°C).  
574 For tropical areas: Hawaii (22-27°C), Samoa (27-29°C), and Puerto Rico (25-28°C).  
575 The Eastern Pacific is a huge and variable area but off Central America, sea surface  
576 temperatures range from 25-28°C [137].

577  
578 We re-analyzed PLD and dispersal distance data from [138] and [139] by separating  
579 the data into temperate and tropical species. There was no obvious difference in the  
580 PLD vs. dispersal distance relationship between tropical and temperate species of  
581 invertebrates or fishes (figure Sup 4). Only 19 of the 62 comparisons of PLD with  
582 dispersal distance in [138] and [139] were from tropical species, and six of the  
583 tropical data pairs (32%) were from tunicate tadpole larvae (figure Sup 4). The  
584 overall pattern of the data is a tight cluster of points above a dispersal distance of  
585 about 20 km and PLDs of > 1 week and a broad scatter of points below about 1 km  
586 dispersal distance and with PLDs from < 1 hr to several weeks. This general pattern  
587 is displayed by both the tropical and temperate data when analysed separately: there  
588 is no obvious difference between tropical and temperate invertebrate and fish  
589 species in the PLDs vs. dispersal distance relationship. However, the small sample  
590 size of the tropical data set suggests caution in this interpretation.

591  
592 An association between genetic structure and PLD is expected, but in practice the  
593 relationship is often not found [5, 6]. Although PLD is often assumed to be a strong  
594 determinant of species range, results of attempts to test this assumption have been

595 mixed at best. Further, recent studies have concluded that PLD is usually not a  
596 major determinant of range size [3],[140]. In contrast, a significant correlation was  
597 found between genetic differentiation and PLD in a study of 32 species (including 12  
598 fishes) [68]. In another study, a consistent, moderate association between genetic  
599 and PLD proxies of dispersal was found in 50 marine species (21 fishes), which was  
600 significantly better at small (<650 km) spatial scales [141]. In some cases, the  
601 association between genetic structure and predicted dispersal can be improved by  
602 using modeled dispersal trajectories based on ocean currents instead of point to  
603 point straight lines [142], perhaps explaining some of the inconsistency in other  
604 studies. In contrast, two broad assessments of marine fish populations and  
605 associated proxies [6], [5], reported that egg type (demersal vs. pelagic) was a better  
606 predictor of genetic structure than was PLD, and neither found a significant  
607 association between PLD and genetic structure. It is worth noting, however, that in  
608 the analysis of PLD data here, in each region except the California Current, the  
609 average PLD of demersal-spawning species was shorter than that of broadcast  
610 spawning species by 20 to 60% (main text figure 3), consistent with [5] and [6]. This  
611 is potentially an indication that demersal-spawning species have shorter dispersal  
612 distances, albeit with all the caveats raised above.

613

614 Although Riginos et al. [5] found little effect of PLD on genetic structure, 96 of the  
615 148 demersal study species were tropical, with only a few temperate species (mostly  
616 *Sebastes*), and only 2-3 species from cold water. These authors properly did not  
617 attempt to generalize their findings to temperate species. Bradbury et al. [6] did not  
618 observe a significant association between PLD and genetic divergence for 30  
619 species of marine fishes, but their analysis included both pelagic and demersal

620 species. Bradbury et al. [6] also found a significant, curvilinear increase in PLD with  
621 latitude at both species and family level, and in this case, the large majority of  
622 tropical and warm temperate species were demersal, site attached reef fishes,  
623 whereas the cold water species were dominated by open shelf and pelagic species.  
624 Therefore, the higher latitude species are more likely to show an influence of adult  
625 movement on genetic structure than were the tropical species. Further, as shown  
626 herein for tropical and warm temperate species, adult habitat does influence PLD.

627

## 628 **5. CAVEATS**

629 Larval dispersal is very difficult to measure directly, although recent advances in  
630 otolith tagging and genetic parentage make it more feasible, if labour intensive. As a  
631 result, the use of indirect approaches and proxies to estimate larval dispersal is  
632 common. These include some population genetic approaches, estimation of PLD  
633 and dispersal modelling. For a variety of reasons touched upon here, genetic  
634 structure in marine populations may not always derive from recent patterns of larval  
635 dispersal alone. Similarly, published PLD data often do not capture the spatial and  
636 temporal variation that intensive study reveals, and the intuitively attractive  
637 hypothesis that PLD is a strong determinant of larval dispersal distance is often  
638 rejected when tested. We have not considered dispersal modelling here, but  
639 modelling larval dispersal is challenging because of the many variables that  
640 contribute to dispersal outcomes, and our often poor understanding of larval  
641 behaviour in particular. Therefore the results of proxy-based and indirect  
642 approaches should be viewed with caution.

643

644 Genetic variation among populations results from the combined influences of  
645 selectively neutral processes such as genetic drift, mutation, and gene flow as well  
646 as adaptive processes such as natural or sexual selection. The application of  
647 genetic approaches to the study of larval dispersal in temperate and tropical marine  
648 systems can illuminate trends in migration and dispersal, when differences among  
649 populations are found. Allele frequency differences among populations and  
650 individuals can provide the means for both indirect estimation of average gene flow  
651 and the direct measurement of individual dispersal events [143],[144]. Furthermore,  
652 the study of genes experiencing natural selection allows the scale of adaptation to be  
653 resolved and resulting clines can reveal trends in successful dispersal [145].

654 Genetic approaches can overcome some of the hurdles associated with the direct  
655 tracking of many tiny propagules and resolving the tails of dispersal kernels (ie,  
656 determining the shape of the dispersal distance frequency distribution) As with all  
657 approaches, they are subject to their own sources of bias and misinterpretation [5]  
658 [143] [144]. For example, population genetics approaches reflect a combination of  
659 egg and larval dispersal, settlement, and post-settlement processes, as well as adult  
660 dispersal, and they may be sensitive to rare events over long or intermediate time  
661 periods as well as on processes that may no longer exist.

662

663 Genetic estimates of dispersal are most commonly based on either isolation by  
664 distance models or individual assignment, and more recently on parentage. The  
665 former are more commonly used in higher latitude studies and the latter in tropical  
666 studies because larger populations and larger spatial scales make assignment  
667 approaches challenging in colder regions. Estimates of average dispersal distance  
668 based on genetic isolation-by-distance relationships (i.e., [146] [147] [148]) suggest



669 local recruitment in marine species may be more prevalent than previously thought  
670 [149], [150]. Similarly, individual assignment [151] or parentage based [152]  
671 methods resolve single events and also often emphasize local recruitment.  
672 However, single case studies of limited geographic scale do not allow the full  
673 dispersal kernel to be determined and are hard to interpret in terms of the question at  
674 hand, which is whether there are tropical-temperate trends in larval transport and  
675 dispersal, because similar studies on higher-latitude fishes are lacking.

676

677 Undoubtedly, genetic estimates of larval dispersal in a single species along a  
678 latitudinal gradient are best suited to address the goal of this review, yet such studies  
679 are rare. One of the few studies to contrast genetic structure across fish species,  
680 geographic regions and life histories [6] found significant associations between  
681 maximum latitude, body size and genetic structure ( $F_{ST}$ ), and weaker genetic  
682 structure at latitudes above 40 degrees, with the largest differences observed at the  
683 extremes of latitude (i.e., 20 vs. 60 degrees latitude). That review [6] examined  
684 dispersal overall (not limited to larval dispersal), and was not limited to nearshore  
685 demersal species, examining all taxa and data available. Nonetheless, genetic  
686 structure across species of marine fish supported the hypothesis that dispersal  
687 occurs over greater spatial scales at high than at low latitudes, consistent with  
688 predictions based on the expected effect of temperature on development times [115],  
689 some oceanographic variables, and conclusions of some studies of single species of  
690 marine invertebrates [153], [154]. Although this interpretation is valid for those  
691 species included in the meta-analysis [6], the composition of the species for which  
692 genetic data exist is not reflective of the spawning modes of the species  
693 assemblages that actually occur in the areas of interest. Specifically, genetic data

694 are available for a higher proportion of demersal-egg spawners at low latitudes and  
695 broadcast spawners at high latitudes than are found in the assemblages from those  
696 areas. This is not a criticism of such synthetic studies, but it does make it difficult to  
697 apply their conclusions to broad faunas and questions like the ones of interest here.  
698 For example, some workers have concluded that larval dispersal occurs over smaller  
699 spatial scales in demersal-spawning spawners than it does in broadcast spawners  
700 [6],[5]. If true, geographic biases in the species examined limit the utility of these  
701 approaches to single species examples until the suite of species for which data exist  
702 becomes representative. However, very few direct estimates of larval dispersal are  
703 available to test the impact of any biases due to species selection.

704

705 Biases due to species selection depend on the families present at differing latitudes.  
706 For example, the families Agonidae, Ammodytidae, Anarhichadidae, Cottidae,  
707 Cyclopteridae, Pholididae, Stichaeidae and Zoarcidae constitute about half the  
708 demersal fish species in the higher latitudes of the NE Pacific and NW Atlantic  
709 (figure Sup 2), and the vast majority of the species of these families spawn demersal  
710 eggs. The available genetic data as used in [6] includes only about 5-10% of the  
711 total number of species from the listed families, and none for five of them, thus  
712 resulting in an under-representation of demersal-spawning species. The same  
713 trends were found when the data were analysed at the family level [6], but the  
714 absence of genetic data for five of these eight families limits the conclusions that can  
715 be drawn. Thus, if spawning mode is relevant to dispersal, as seems to be the case  
716 in warmer waters [5], the available genetic data are not representative of the full  
717 fauna. More study of high latitude non-commercial taxa with demersal eggs using

718 direct and indirect methods is needed to help evaluate potential biases and trends  
719 present.

720

721 Population genetic approaches (e.g., isolation-by-distance based methods) are also  
722 influenced by historical events such as glaciations, which may have long-lasting  
723 effects on genetic structure that are relevant for latitudinal comparisons. Species  
724 characterized by large populations recolonizing high-latitude areas following  
725 deglaciation may not have had sufficient time for differences to accumulate by  
726 genetic drift. In such cases the absence of genetic divergence tells us little regarding  
727 larval dispersal and connectivity [155] and it is difficult to discount the hypothesis that  
728 the observed low structure at high latitudes is not at least partly due to recent  
729 recolonization and large effective population sizes. Similarly, glaciations also  
730 influence genetic structure in the tropics, because some regions during the  
731 Quaternary, in particular areas between South-east Asia and Australia, were subject  
732 to high-frequency sea level fluctuations that alternatively flooded and dried many  
733 thousands of square km [156]. Direct genetic approaches based on individual  
734 assignment or parentage are not subject to these effects.

735

736 Might apparent geographic differences in dispersal arise from differing  
737 methodologies or research targets? Studies of dispersal and connectivity in the  
738 tropics tend to be on small, site-localized species ([43, 157], [158]; see sections on  
739 Habitat Fragmentation). Often, the study species are chosen for tractability (e.g.,  
740 small adult size, small adult home range), and if the species is not fished, all the  
741 better: this eliminates a factor that potentially complicates population dynamics.  
742 Might this approach bias estimates of dispersal distances? These studies often

743 utilize techniques that involve assigning a source to individual recruits through  
744 parentage or chemical tagging, and the geographic scale tends to be small, on the  
745 order of tens of km (e.g., [79], [159], [160], [161]). Approaches such as this do not  
746 have the capacity to identify larger scales of dispersal, although a genetic study of  
747 anemonefishes that spanned hundreds of kilometres [162] suggested short dispersal  
748 distances, at approximately the same scale as geographically limited studies of  
749 similar species.

750

751 In contrast, in temperate regions many studies of population structure are conducted  
752 on large scales on exploited stocks from deeper water, often with a focus on  
753 assessing spatial scales of management [163]. If temperate fished stocks tend to be  
754 larger and more mobile than site-attached fishes on shallow coral reefs, then  
755 estimates of what constitutes a local population within the metapopulation can be  
756 very different, and this could set a different scale for connectivity estimates.

757 However, as mentioned above, a larger spatial scale of study does not automatically  
758 result in longer estimates of dispersal. A coarse-scale genetic study of fifteen  
759 species of exploited rockfishes along the west coast of North America found little  
760 evidence of genetic structure among some species, but sharp genetic breaks among  
761 others [164]. In the more subdivided species, more detailed study could reveal fine-  
762 scale population structure and evidence of limited dispersal. Such a finding is very  
763 unlikely in the group lacking genetic structure.

764

765 **Figure Captions**

766 Figure Sup 1. Mean ( $\pm$ SE) pelagic larval duration for taxonomic groups of demersal  
767 shorefishes with primarily warm-water (solid bars) and cold-water (open bars)  
768 distributions. PLD data from [2] and [3]. See Supplement text for numbers of PLD  
769 estimates for each taxon.

770

771 Figure Sup 2. Percentages of demersal shorefish species with different spawning  
772 modes in regions of the Western Atlantic and Eastern Pacific. In each ocean, cold  
773 temperate regions have a much smaller percentage of species with pelagic eggs  
774 than do other regions. Both Argentina and Chile have a large north-south extent and  
775 their fish faunas include both warm and cold temperate components: although they  
776 have a lower percentage of species with pelagic eggs, the percentage is still about  
777 55% as compared to 15-25% in the cold temperate regions of the northern  
778 hemisphere. Data from Hawaii in the central Pacific are very similar to the other  
779 tropical values. In the Antarctic, 16% of species spawn pelagic eggs. See Table  
780 Sup1 for details and data sources.

781

782 Figure Sup 3. Larval transport by Eddy kinetic energy (EKE) and eddy-driven  
783 currents: consequences of (A) strong diffusion (dotted line), (B) anisotropic advection  
784 (dotted line), (C) advection and strong diffusion (dotted line) on the mean dispersal  
785 kernel (solid line). Larval source is located at  $x = 0$ .

786

787 Figure Sup 4. Pelagic larval duration of marine invertebrates and fishes plotted  
788 against their dispersal distance. Solid squares and open circles are data from  
789 tropical and temperate species, respectively. "T"s adjacent to values indicate data

790 from the dispersal of tadpole larvae of tunicates. Data are from [139] and [138], but  
791 with algal data removed.

792

793

794 **Table Captions**

795

796 Table Sup 1. Percentage of demersal marine teleost shorefish species with different  
797 spawning modes in selected regions. A small number of species with unknown  
798 spawning modes are omitted: these constitute at most 3% of species in any location.  
799 Species compositions from: [165] [166] [167] [168] [169] [170] [171]. Spawning  
800 modes from: [172] [169] [173].

801

802 Table Sup 2. Studies based on otolith methods to estimate demographic  
803 connectivity used in the habitat fragmentation comparison. [160], [174], [175], [176],  
804 [177], [178], [179], [180], [181], [182], [183], [184], [185], [186], [187], [188], [189],  
805 [190].

806

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1273 Table Sup 1. Percentage of demersal marine teleost shorefish species with different  
 1274 spawning modes in selected regions. A small number of species with unknown spawning  
 1275 modes are omitted: these constitute at most 3% of species in any location. Species  
 1276 compositions from: [163] [164] [165] [166] [167] [168] [169]. Spawning modes from: [170]  
 1277 [167] [171].  
 1278

Location (number of species)	Category	Broadcast spawner (pelagic egg)	Demersal spawner (demersal egg)	Brooder (egg held by adult)	Live bearer (viviparous)	No pelagic stage
<b>Atlantic Ocean</b>						
Canada Atlantic (91)	cold temperate	27.5	64.8	1.1	3.3	0
Mid Atlantic Bight (Northern Florida to Cape Hatteras) (227)	warm temperate	77.1	15.9	4.0	0.00	2.2
Cuba (535)	tropical	64.9	24.5	6.4	2.2	0.9
Argentina (124)	temperate	55.7	34.7	4.0	0.8	4.0
<b>Pacific Ocean</b>						
NE Pacific (Oregon Border to Bering Strait) (198)	cold temperate/ boreal	15.2	65.7	0.5	14.1	4.6
Californian (Cabo San Lucas to Point Conception) (306)	warm temperate	68.6	25.2	2.6	1.3	2.3
Equatorial E. Pacific (696)	tropical	60.9	30.9	2.4	1.3	4.5
Chile (115)	temperate	55.7	32.2	3.5	5.2	1.7
Hawaii (429)	tropical	75.5	20.3	3.7	0.2	0
<b>Southern Ocean</b>						
Antarctic (55)	polar	16.4	83.7	0	0	0



1280 Table Sup 2. Studies based on otolith methods to estimate demographic connectivity used in the habitat fragmentation comparison. These  
 1281 publications were found by searching the ISI Web of Science database using the search criteria ("otolith chemistry" or "otolith microchemistry"  
 1282 or "elemental fingerprints" or "environmental marker\*" or "natural tag\*") and (dispersal or connectivity or migration or "stock structur\*"). Studies  
 1283 were then filtered to include only non-philopatric species where movement among populations was due to larval dispersal. [156], [172], [173],  
 1284 [174], [175], [176], [177], [178], [179], [180], [181], [182], [183], [184], [185], [186], [187], [188]  
 1285  
 1286

Species	Egg Type	Habitat Type	Latitude	Mean % Self-recruitment	Scale of Self-recruitment (km)	Minimum Scale of Connectivity (km)	Reference
<i>Amphiprion percula</i>	Demersal	Islands	5°S	60	0.5	20	[156]
<i>Chaetodon vagabundus</i>	Pelagic	Islands	5°S	60	0.5	20	[156]
<i>Chaenocephalus aceratus</i>	Demersal	Coastal	54-63°S	No estimate	200	400	[172]
<i>Dissostichus eleginoides</i>	Pelagic	Coastal	43-55°S	88	250	1200	[172]
<i>Chromis viridis</i>	Demersal	Islands	29°N	No estimate	No estimate	15	[173]
<i>Rhinogobius giurinus</i>	Demersal	Coastal Habitat Patch	23°N	94	0.1	175	[174]
<i>Coris julis</i>	Pelagic	Islands	37-40°N	85	20	600	[175]
<i>Thalassoma bifasciatum</i>	Pelagic	Islands	17°N	45	35	No estimate	[176]
<i>Pomacentrus amboinensis</i>	Demersal	Islands	14°S	35	7	No estimate	[177]

<i>Amphiprion polymnus</i>	Demersal	Islands Coastal Habitat Patch	5°S	24	0.1	0.5	[178]
<i>Pomacentrus coelestis</i>	Demersal	Islands Coastal Habitat Patch	23°N	69	30	370	[179]
<i>Coryphaenoides rupestris</i>	Pelagic	Coastal Coastal Habitat Patch	50-58°N	71	300	2700	[180]
<i>Sebastes melanops</i>	Live bearer	Coastal Coastal Habitat Patch	42-46°N	71	120	540	[181]
<i>Pomacentrus coelestis</i>	Demersal	Islands	14-23°S	42	10	30	[182]
<i>Coris bulbifrons</i>	Pelagic	Islands Islands and Coastal Habitat Patch	31°S	50	15	600	[183]
<i>Sebastes atrovirens</i>	Live bearer	Islands Islands and Coastal Habitat Patch	34°N	97.5	30	40	[184]
<i>Merluccius merluccius</i>	Pelagic	Coastal	36-62°N	No estimate	400	1000	[185]
<i>Thalassoma bifasciatum</i>	Pelagic	Islands Coastal Habitat Patch	17°N	50	35	No estimate	[186]
<i>Forsterygion lapillum</i>	Demersal	Islands Coastal Habitat Patch	41°S	72	10	20	[187]
<i>Micropogonias undulatus</i>	Pelagic	Coastal	35-36°N	No estimate	No estimate	330	[188]

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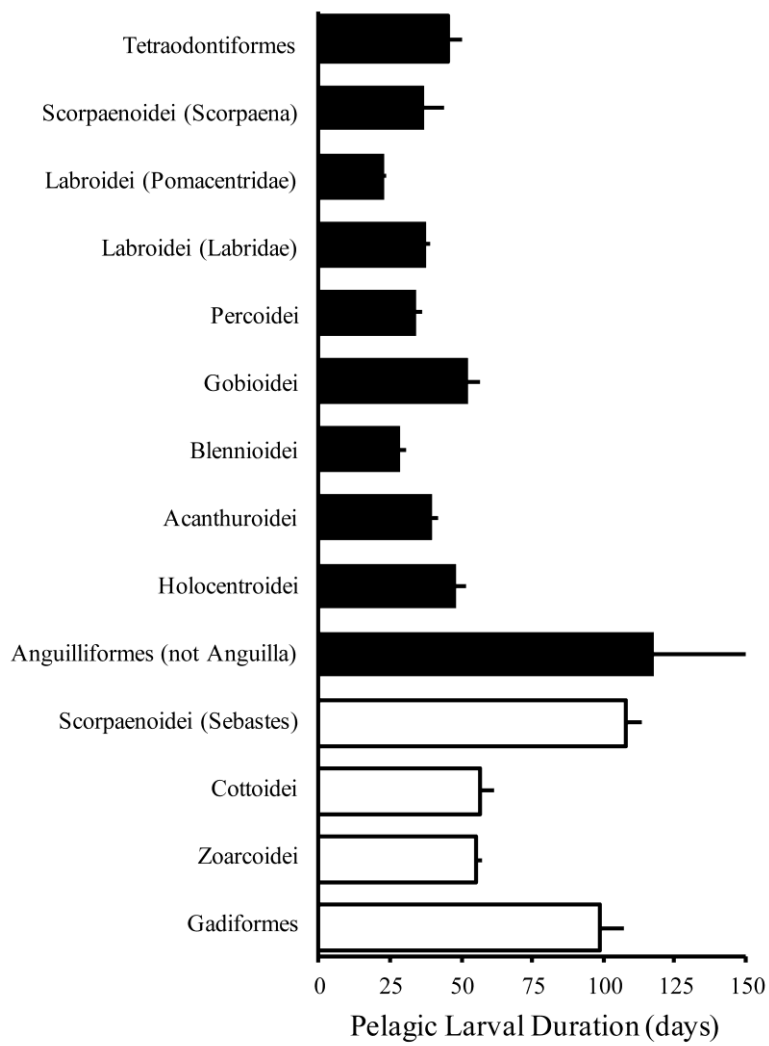


Figure Sup 1. Mean ( $\pm$ SE) pelagic larval duration for taxonomic groups of demersal shorefishes with primarily warm-water (solid bars) and cold-water (open bars) distributions. PLD data from [2] and [3]. See Supplement text for numbers of PLD estimates for each taxon.

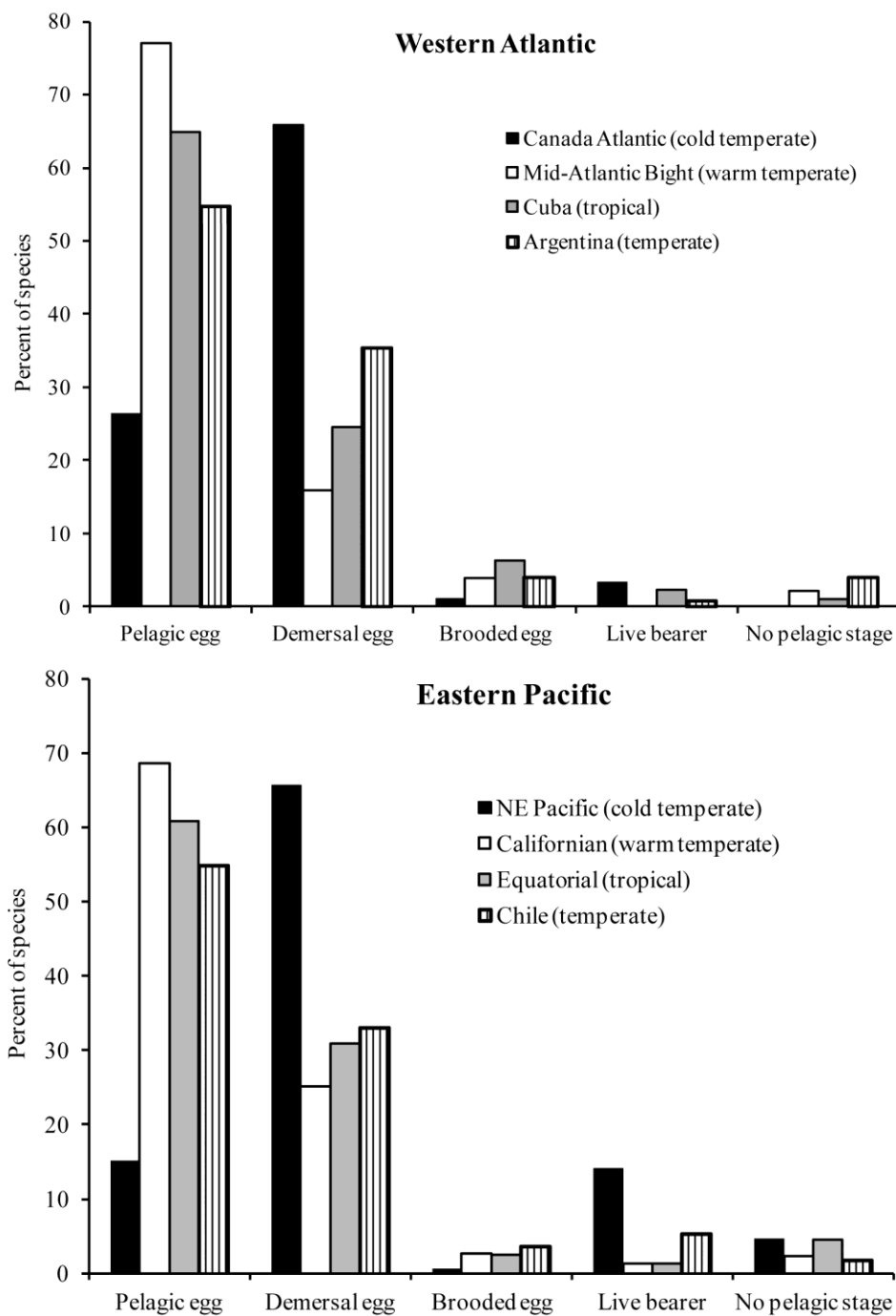


Figure Sup 2. Percentages of demersal shorefish species with different spawning modes in regions of the Western Atlantic and Eastern Pacific. In each ocean, cold temperate regions have a much smaller percentage of species with pelagic eggs than do other regions. Both Argentina and Chile have a large north-south extent and their fish faunas include both warm and cold temperate components: although they have a lower percentage of species with pelagic eggs, the percentage is still about 55% as compared to 15-25% in the cold temperate regions of the northern hemisphere. Data from Hawaii in the central Pacific are very similar to the other tropical values. In the Antarctic, 16% of species spawn pelagic eggs. See Table Sup1 for details and data sources.

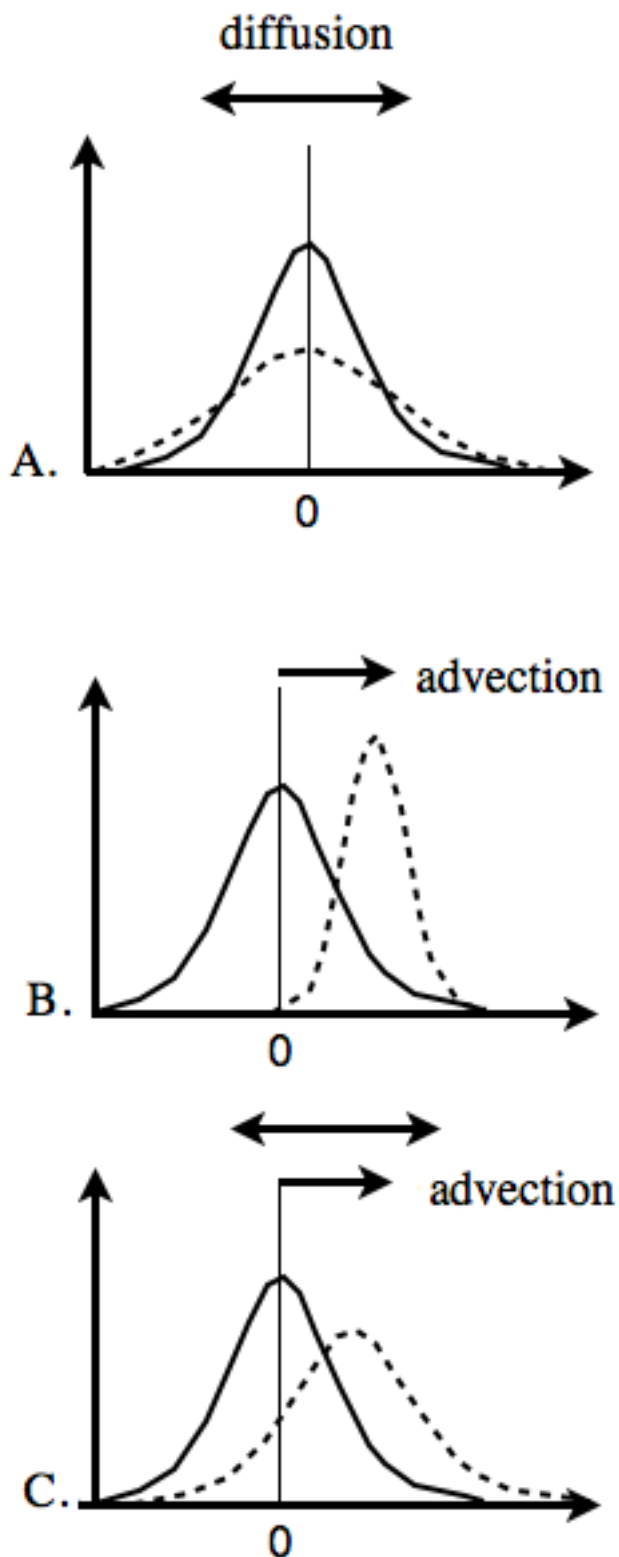


Figure Sup 3. Larval transport by Eddy kinetic energy (EKE) and eddy-driven currents: consequences of (A) strong diffusion (dotted line), (B) anisotropic advection (dotted line), (C) advection and strong diffusion (dotted line) on the mean dispersal kernel (solid line). Larval source is located at  $x = 0$ .



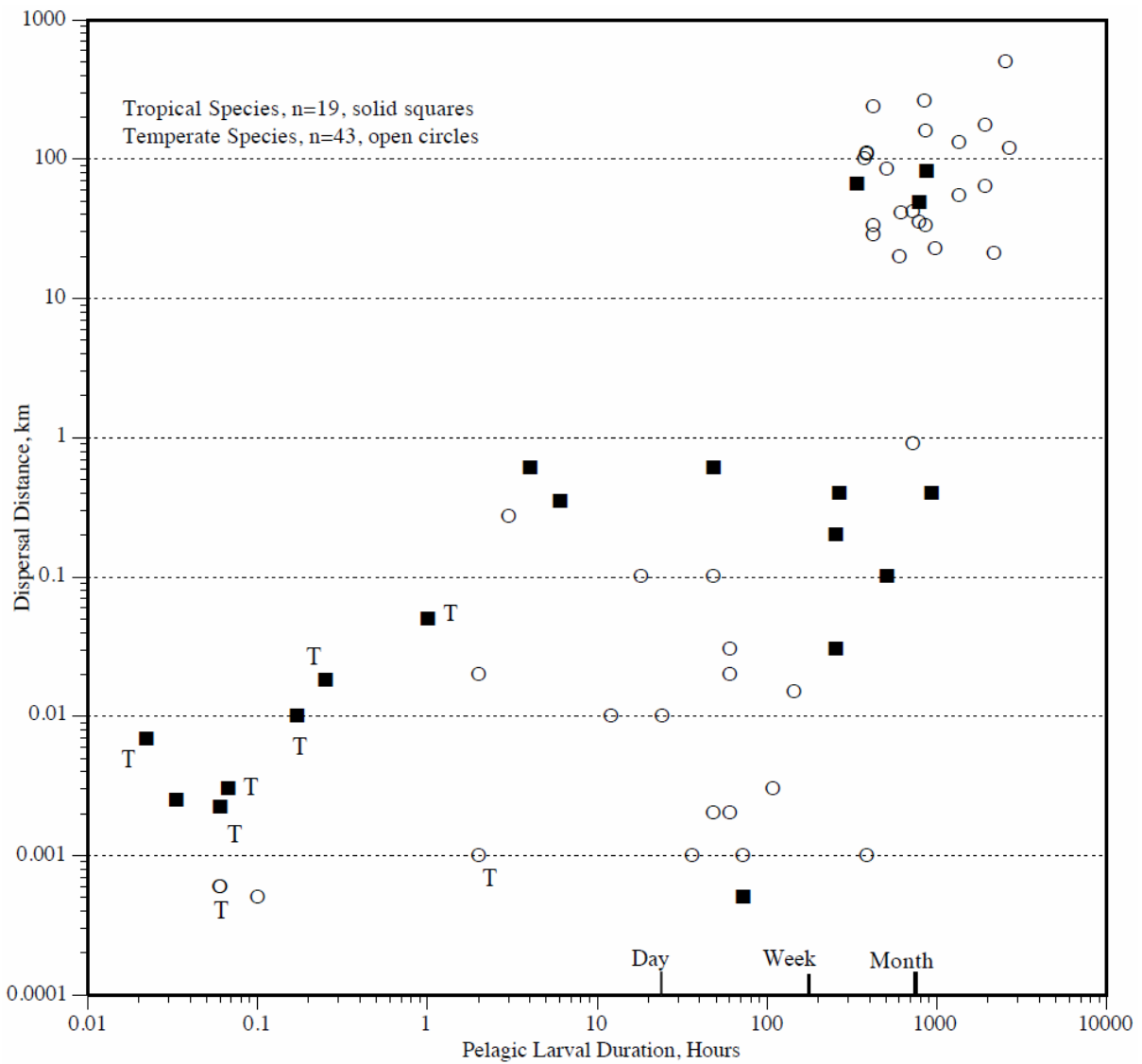


Figure Sup 4. Pelagic larval duration of marine invertebrates and fishes plotted against their dispersal distance. Solid squares and open circles are data from tropical and temperate species, respectively. "T"s adjacent to values indicate data from the dispersal of tadpole larvae of tunicates. Data are from [139] and [138], but with algal data removed.