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Guidelines for larval dispersal simulations: flow field representation versus biological traits

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14 Abstract

The natural mechanism for sessile species resilience after major population disturbance 15 is larval dispersal during reproduction. Modelling larval dispersal when processes con-16 trolling dispersal are accounted for, is a powerful tool to anticipate connectivity patterns 17 among populations. However, simulations performance might be altered considerably by 18 either flow representation in the underlying hydrodynamical simulations or the inclusion 19 of larval traits. Here, we assessed how behaviour, ocean model resolution and tempo-20 ral variability in the ocean dynamics influence connectivity patterns among coastal rocky 21 locations, some of them being designated as MPAs. We combined a fine (0.3 km) and 22 coarser (1.5 km) resolution ocean model, with or without the negatively buoyant behaviour 23 of *Paramuricea clavata* larvae, with different turbulence intensity, in three years and for 24 pelagic larval duration (PLD) ranging from 1 to 28 days. Ocean model resolution was 25 the most influential factor on the resulting connectivity levels. The results were also equally 26 influenced by buoyancy, PLD and temporal variability, in the fine resolution configura-27 tion, while turbulence could be neglected. Based on these results, guidelines for coastal 28 biophysical modelling are proposed. 29

³⁰ 1 Introduction

Anticipating the impact of the on-going global change on the already altered ocean 31 biodiversity is a major challenge for designing conservation measures in the next decade 32 (ONU SGD 14, IPBES 2030 work program, https://ipbes.net/o4-supporting-policy). Among 33 conservation measures, the Aichi Convention on Biological Diversity (CBD, 2010) de-34 fined the objective of reaching 10% of ocean surface designated as Marine Protected Ar-35 eas (MPAs) by 2020. However, designating efficient MPAs requires incorporating knowl-36 edge and understanding of mechanisms regulating species spatial distribution such as dis-37 persal (Urban et al., 2016). 38

The Mediterranean coastline is characterized by an alternance of Marine Protected Areas (MPAs) and non-protected areas (MedPAN, 2017). MPAs provide benefits to out-

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side areas and in particular, many studies have shown the efficacy of MPAs on the conservation of fishes through the spillover of juvenile and adult fishes (see Grüss et al. (2011) for a review). The effect of MPAs on the conservation of sessile species whose reproduction implies a dispersive phase mediated by ocean circulation poses the question of anticipating population connectivity within a geographically fragmented network. To this end, larval dispersal numerical models incorporating knowledge and understanding of mechanisms regulating dispersal are essential tools (Lett et al., 2008; Paris et al., 2013).

Larval dispersal simulations first require hydrodynamical models to provide a re-48 liable and validated description of the flow and its variability at the spatio-temporal scales 49 relevant for both dispersal processes at larval scale and regional connectivity estimates 50 at population scale. Connectivity studies have used a variety of ocean circulation mod-51 els, from coarse-resolution global models (10 km, Treml et al. (2008); Wood et al. (2014)) 52 to high-resolution regional models (100 m to 1 km, Guizien et al. (2012, 2014); Briton 53 et al. (2018)). Ocean circulation models have been validated against diagnostic variables 54 such as sea surface temperature and salinity, heat content, sea level, mass transport, tur-55 bulent kinetic energy or vorticity (Fox-Kemper et al., 2019). However, such validation 56 can be insufficient to ensure that simulated velocities used in the larval dispersal sim-57 ulations are representative of the advection and diffusion experienced by larvae in the 58 flow (Pineda et al., 2007). When the flow field simulations are accurate, the specificity 59 of larval dispersal simulations ultimately depends on the knowledge of biological traits 60 such as location and timing of larval release in parental populations and larval traits im-61 portant for dispersal (Scheltema, 1986). Among these, pelagic larval duration (PLD, here-62 inafter) has been extensively tested in dispersal simulations (Siegel et al., 2003; Edwards 63 et al., 2007; Guizien et al., 2012). Despite larval position in the water column likely al-64 ters dispersal patterns in three dimensional flows, two dimensional larval dispersal stud-65 ies are still frequent (Aguilar et al., 2019; Gamoyo et al., 2019; Hidalgo et al., 2019; Hock 66 et al., 2019). When vertical larval position has been accounted for in simulations, it has 67 been either prescribed (Donahue et al., 2015; Blanco et al., 2019; Criales et al., 2019) 68

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or more recently it resulted from the interaction between a larval motility behaviour and the flow dynamics (Faillettaz et al., 2018; Kvile et al., 2018; Barbut et al., 2019; Munroe et al., 2018; Bode et al., 2019; Goodwin et al., 2019; McManus et al., 2020). In contrast, the sensitivity to release timing and larval release duration have been less studied (see Guizien et al. (2012)). Moreover, the relative importance of biological traits and flow field representation in altering dispersal patterns has not been frequently evaluated together (see Huret et al. (2007)).

In the present study, we chose the rocky and coralligeneous habitat in the Ligurian 76 sea as a case study for larval dispersal simulations sensitivity analysis. We estimated the 77 sensitivity to different parameters of larval dispersal simulations. First we addressed the 78 sensitivity to the choice of hydrodynamical model and in particular to its horizontal grid 79 resolution. Second we addressed the temporal variability at different scales and finally, 80 we addressed the parameters specific of larval dispersal simulations such as turbulence, 81 larval behaviour, PLD and release timing, comparing an idealized neutrally buoyant species 82 to *Paramuricea clavata* (Risso 1826). Eventually the sensitivities to the different param-83 eters were ranked by importance to suggest guidelines for larval dispersal simulations. 84

85 2 Methods

86

2.1 Study background

The Ligurian Sea extends between Italy and the north of Corsica in the Northwest-87 ern Mediterranean Sea (Astraldi & Gasparini, 1992). It is characterized by a mean cy-88 clonic circulation (Millot, 1999; Millot & Taupier-Letage, 2005) known as the Ligurian 89 Current (LC) that flows west along the Italian coast toward France (Fig.1). Along the 90 Ligurian coast, many MPAs have been established to foster the conservation of the rich 91 biodiversity associated to coralligeneous and rocky bottom habitats (Regional Natural 92 Reserve of Bergeggi, Regional Natural Park of Portofino, Cinque Terre National Park, 93 Regional Natural Park of Porto Venere, Fig.1). Coralligeneous and rocky bottom habi-94

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tats are patchily distributed in the Ligurian sea (http://srvcarto.regione.liguria

.it/geoviewer2/pages/apps/geoportale/index.html?id=2121).

In these MPAs, the conspicuous octocoral species *Paramuricea clavata* has been 97 used as a flag species for the conservation of the Mediterranean coralligeneous assem-98 blage (Carpine & GRASSHOFF, 1975), due to its role of ecosystem engineer (sensu, Jones 99 et al. (1994)), and its patrimonial value linked to the beauty it gives to seascapes. P. clavata 100 is one of the tallest sessile Mediterranean species with erected colonies as high as 1.3 m 101 (Harmelin & Marinopoulos, 1994) and is found on hard substrate from 10 to 200 m wa-102 ter depth (Gori et al., 2017). During reproduction, P. clavata releases eggs that get fe-103 cundated and maturate during a few days at the colony surface. Once the maternal mu-104 cus is dissolved lecithotrophic planula larvae are released in the ocean currents. Off the 105 Catalan coast, larval release happens during short events of a few days separated by 15 106 days (Coma et al., 1995; Gori et al., 2007) in late June/early July. No information on 107 the exact timing and occurrence of *P. clavata* larval release in the Ligurian Sea are avail-108 able in the scientific literature but several observations by recreational divers report a 109 similar phenomenon in the month of July. 110

111

2.2 Fine resolution hydrodynamical simulations

The dynamic of the Ligurian sea is simulated using the Ligurian Integrated Mod-112 elling Effort, gcm (LIMEgcm, hereinafter Fig.1 blue dashed rectangle). LIMEgcm is based 113 on the the Massachusetts Institute of Technology general circulation model (MITgcm,Marshall 114 et al. (1997)). The MITgcm solves the Boussinesq form of the NavierStokes equations 115 either on a generalized curvilinear grid or an orthogonal grid. The finite-volume discretiza-116 tion is rendered on a horizontal Arakawa C-grid with vertical z-levels. In this configu-117 ration the equation of state is due to (Jackett & McDougall, 1995), whereas advection 118 for tracers is computed via a third-order direct spacetime flux limited scheme with zero 119



Figure 1. The Ligurian Sea with the main circulation branch and the release locations used in larval dispersal simulations, black circles (Supp. Inf. Tab. 1). Blue dashed rectangle shows the LIMEgcm domain, red rectangle the LIME-ROMS domain. Blue circles and red crosses indicate the numerical grids of LIMEgcm and LIME-ROMS, respectively. For visualization purposes the LIMEgcm mesh is plotted every 6 grid point while the LIME-ROMS mesh is plotted every three grid point.

explicit diffusivity. The K-profile parameterization (KPP, Large et al. (1994)) is used with a background vertical viscosity of $n_{\nu} = 10^{-5} \text{ m}^2 \text{s}^{-1}$.

The numerical domain was discretized with an orthogonal horizontal grid of 560 122 x 160 points. The horizontal mesh had a resolution of 300 m (Fig.1 blue circles) while 123 the vertical was discretized with 100 levels. The vertical resolution increased linearly from 124 1m to 20m in the first 20 levels, was kept constant at 20m for the following 50 levels, and 125 finally increased to 25m for the 30 deepest layers. The bathymetry is interpolated from 126 the EMODNET 1/8 arcminute product (Consortium, 2016). Realistic surface forcings 127 came from the hourly, Weather Research and Forecasting (WRF) model setup by the 128 University of Genoa (Cassola et al., 2015), while 6-hourly open boundary conditions were 129 derived from LIME-ROMS model outputs (Sciascia et al., 2019). Model outputs were 130 saved every three hours. 131

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2.3 Coarse resolution hydrodynamical simulations

The circulation of the Ligurian Sea was simulated using the Ligurian Integrated 133 Modelling Effort (LIME-ROMS, Sciascia et al. (2019)). LIME-ROMS was based on the 134 Rutgers University kernel of the Regional Ocean Modelling System (ROMS, Haidvogel 135 et al. (2000); Shchepetkin and McWilliams (2005)). The numerical domain was discretized 136 with an horizontal grid of $590 \ge 314$ points. The mesh was unevenly spaced and the most 137 resolved area (Fig.1 red rectangle), which includes the area of the Ligurian-Northern Tyrrhe-138 nian shelf, had a horizontal resolution of $1/64^{\circ} \ge 1/64^{\circ}$ (~ 1.5 km, Fig.1 red crosses). 139 The vertical dimension was discretized with 50 sigma layers. Realistic surface forcings 140 came from the 3-hourly, 0.25° horizontal resolution ERA-Interim reanalysis fields (Dee 141 et al., 2011) available at the http://apps.ecmwf.int/datasets/data/interim-full 142 -daily link, while the daily open boundary values were derived from the Mediterranean 143 Forecasting System (MFS) SYS4a3 product (Delrosso et al., 2016). Model outputs were 144

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saved every six hours. The reader is referred to Sciascia et al. (2019) for a detailed de scription and validation of the model.

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2.4 Larval dispersal simulations

Dispersal simulations of virtual larvae were performed independently using the LIME-ROMS (hereinafter, coarse) and LIMEgcm (hereinafter, fine) resolutions flow fields simulated as described above.

The offline Lagrangian Transport model (LTRANS, Schlag and North (2012)) was selected to compute particle advection in the coarse resolution hydrodynamical fields for its capability to cope with sigma layers. The offline Connectivity Modelling System (CMS, Paris et al. (2013)) was selected to compute particle advection in the fine resolution hydrodynamical fields for its capability to cope with z layers. Both Lagrangian models compute 3D trajectories using a 4th order Runge-Kutta for both neutrally buoyant and active (e.g. sinking, swimming) particles, eventually including turbulent diffusion.

In both larval dispersal simulation set ups, particles were released in five locations 158 where P. clavata has been reported (Bergeggi (BER), Portofino (POR), Sestri Levante 159 (SES), Punta Mesco (PTM), Porto Venere (PRV), Fig. 1, Supp. Inf. Tab. 1, Padrón et 160 al. (2018)). Each population was described by ten release points, separated by 100 m. 161 Particles were released ~ 3 km away from the coast line, which represented two grid points 162 of the coarse resolution flow field in order to avoid any retention bias due to the numer-163 ical boundary layer of this simulation. Particles were released 4 m above the bottom in 164 the coarse resolution simulations and 2 m above the bottom in the fine resolution sim-165 ulations (Supp. Inf., Tab. 1). Particles were released every hour from June 15 at mid-166 night until July 15 at 11pm and tracked during 28 days after release. Tacking time step 167 was 120 s in the coarse resolution flow field and 600 s in the fine resolution flow field. P. 168 calvata populations suffered two mass mortality events in 1999 and 2003 (Cerrano et al., 169 2000; Cupido et al., 2008) and get to sexual maturation after at least four years (Coma 170

et al., 1995). Hence the reproductive events that more likely contribute to the popula-171 tions found now at sea took place from 2004 onward. For the aforementioned reasons, 172 larval dispersal simulations were performed during three successive reproductive years 173 in the fine resolution flow field forcing (2004, 2005 and 2006) but only in two reproduc-174 tive years in the coarse resolution flow field forcing (2004 and 2006 only, 2005 being hy-175 drodynamically very close to 2004). In order to explore the sensitivity of larval disper-176 sal simulations to the flow field resolution, to larval behaviour and flow turbulence in-177 tensity, various simulations were performed as summarized in Tab. 1. 178

The dispersal of neutrally buoyant particles was simulated using both the coarse and fine resolution simulations. The dispersal of particles with a specific buoyancy was carried out only with the fine resolution simulations in order to simulate the dispersal of non-motile *P. clavata* larvae (Coma et al., 1995; Linares et al., 2008). Particle density was set to vary from 1.0282 to 1.0295, with an equivalent diameter of 761 10^{-6} m, to achieve the sinking speed of 0.056 ± 0.021 cm s⁻¹ at 20°C and a salinity of 38 psu as measured experimentally (Guizien et al., 2020).

	No Turbulence	Low Turbulence	High Turbulence
	2004 LIMEgcm	2004 LIMEgcm	2004 LIMEgcm
	2005 LIMEgcm	2005 LIMEgcm	2005 LIMEgcm
Neutrally Buoyant	2006 LIMEgcm	2006 LIMEgcm	2006 LIMEgcm
	2004 LIME-ROMS		
	2006 LIME-ROMS		
	2004 LIMEgcm	2004 LIMEgcm	2004 LIMEgcm
P. clavata	2005 LIMEgcm	2005 LIMEgcm	2005 LIMEgcm
buoyancy	2006 LIMEgcm	2006 LIMEgcm	2006 LIMEgcm

Table 1. Available larval dispersal simulations and their main parameters

The effect of turbulence intensity on larval dispersal simulations was tested by re-186 peating each simulation using the fine resolution flow field forcing without turbulence, 187 with low turbulent diffusivity $(10^{-4} \text{ m}^2 \text{s}^{-1})$ and high turbulent diffusivity $(10^{-3} \text{m}^2 \text{s}^{-1})$ 188 (Tab. 1). Depending on whether turbulence was (was not) accounted for, twenty-five (one) 189 particles were released every hour during each larval release event summing up to a to-190 tal number of 180000 (7200) particles for each population. Similarly to what was done 191 in Guizien et al. (2006), the number of particles was increased by a factor twenty-five 192 to ensure the convergence of statistical properties when dispersion due to subgrid tur-193 bulence is accounted for. 194

195

2.5 Larval connectivity estimates

Larval transport values between five locations in the Ligurian basin were calculated 196 as the proportion of particles released during a larval-release event from an origin loca-197 tion that reached a destination location after a given dispersal duration (Pelagic Lar-198 val Duration, PLD). Larval transport values were gathered to build connectivity matri-199 ces, varying according to PLD, larval-release event, particle buoyancy (neutrally buoy-200 ant and *P. clavata* buoyancy), turbulence level (high, low and no) and hydrodynamical 201 grid resolution (coarse and fine). Nine values of PLD were considered, starting from one 202 day, then 3.5 days up to 28 days with 3.5 days increments. The duration of each larval-203 release event was set to three days, yielding a total of ten 3-day events spanning the 30 204 days long reproductive season of each year (2004-2006). 205

The intra-annual (VAR_{intra}) and inter-annual (VAR_{inter}) variance of any pairwise larval transport elements of the connectivity matrix T were computed as defined in equations 1 and 2 for the coarse and fine resolution simulations and all PLDs.

$$VAR_{intra} = \frac{\sum_{y=1}^{N_y} \sum_{r=1}^{N_r} (T_r^y - \overline{T^y})^2}{N_r N_y} \tag{1}$$

209

$$VAR_{inter} = \frac{\sum_{r=1}^{N_r} \sum_{y=1}^{N_y} (T_r^y - \overline{T_r})^2}{N_r N_y}$$
(2)

where N_y is the number of years, N_r is the number of release events per year and $\overline{T^y} = \sum_{r=1}^{N_r} T_r^y / N_r$ is the average connectivity across all release events of year y and $\overline{T_r} = \sum_{y=1}^{N_y} T_r^y / N_y$ is the average connectivity across all years for the release event r in the year. Ratios $\frac{VAR_{intra}}{VAR_{inter}}$ were computed for each pairwise connection among the five locations and summarized into boxplots for the five locations network.

For all the PLDs, the effect of turbulence levels (high/low/no), particle buoyancy 215 (neutrally buoyant/ P. clavata buoyancy) and the hydrodynamical grid resolution (coarse/ 216 fine) on larval transport estimates between any pairwise connection of the five locations 217 network were computed as larval transport deviations from a same reference simulation 218 set for each larval-release events and years. For each pairwise connection among the five 219 populations, the deviations across larval-release events and years was described by its 220 median, avoiding any normality assumption, to compare the effect of turbulence level, 221 particle buoyancy and hydrodynamical grid resolution. The fine resolution neutrally buoy-222 ant simulations without turbulence for the years 2004-2006 was considered as the ref-223 erence simulation set for all comparisons. All calculations and statistics were conducted 224 in Matlab (R2018). 225

226 3 Results

In the following, the effect of flow resolution, larval buoyancy and turbulence level on connectivity patterns will be presented for a PLD of 3.5 days as an example. In both fine and coarse resolution hydrodynamical simulations, the median larval transport among the five populations showed a northwestern drift (upper diagonal elements larger than lower diagonal ones) with larval export (off-diagonal elements) from PRV and SES dominating over local retention (diagonal elements) while retention rates dominated over any other connections in BER, POR and PTM (Fig. 2 a,b). However, apart this similarity,

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the two sets of simulations differed largely. Firstly, larval retention values were ten times 234 larger in the coarse resolution simulations (max $\sim 95\%$) than in the fine resolution sim-235 ulations (max $\sim 7\%$). Secondly, largest retention values were found in different loca-236 tions (BER and POR in the coarse resolution and PTM in the fine resolution; Fig. 2 a,b). 237 Thirdly, more connections between populations appeared in the fine resolution simula-238 tions compared to the coarse resolution simulations. Differences in the intensity of coastal 239 currents in the coarse and fine resolution hydrodynamical models support the differences 240 in larval transport estimates. Coastal currents were weaker in the coarse resolution hy-241 drodynamical simulation compared to the fine resolution one (Supp. Inf. Fig. 1). Inter-242 estingly, observations available in the area for the period May-June 2004 better agreed 243 with the order of magnitude of the fine resolution simulation than that of the coarse res-244 olution one (Supp. Inf. Fig. 2). 245

Intra-annual variability (between different larval release events within the same year, 246 VAR_{intra}) and inter-annual variability (between the same larval release event in differ-247 ent years, VAR_{inter}) were the same order of magnitude, whatever the PLD and hydro-248 dynamical resolution (5% maximum in the fine resolution and 30% maximum in the coarse 249 resolution, Supporting Information Fig. 3). The median of the intra-annual to inter-annual 250 ratio across all PLD was 0.93 for the fine resolution and 0.85 for the coarse resolution, 251 in the neutrally buoyant case without turbulence, considering years the 2004 and 2006 252 only (Fig. 2c). For the fine resolution simulations adding the year 2005 yielded to a me-253 dian intra-annual to inter-annual ratio across all PLDs of 0.88. When varying turbulence 254 and buoyancy for the fine resolution simulations, the intra-annual and inter-annual vari-255 ance stayed the same order of magnitude, with a median intra-annual to inter-annual 256 ratio of ~ 0.8 (Supp. In., Fig. 2). 257

The sensitivity of larval transport estimates to refining the resolution of the hydrodynamical simulations (Fig.3 a) was compared to incorporating the specific buoyancy of *P. clavata* (Fig.3 b) and to incorporating different intensity of turbulence (Fig.3 c,d)

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Figure 2. Median connectivity matrix for the larval release events of 2004 and 2006 for the neutrally buoyant simulations without turbulence and a PLD of 3.5 days. Panel (a) represents the coarse resolution simulation and panel (b) the fine resolution simulations. Box plot of the VAR_{intra}/VAR_{inter} for different PLDs for the coarse resolution (Panel (c), grey bars) and high resolution (Panel (c), black bars) simulations. VAR_{intra}/VAR_{inter} were calculated for larval release events of 2004 and 2006 for the neutrally buoyant simulations without turbulence.

261	for a PLD of 3.5 days. A clear hierarchy can be drawn with the effect of the hydrody-
262	namical resolution being an order of magnitude higher (median deviation up to 90%) than
263	that of incorporating the specific buoyancy (median deviation up to $14\%)$ and two or-
264	ders of magnitude higher than the effect of including turbulence dispersal, whatever the
265	level (median deviation up to 5 $\%$ for the two levels of turbulence tested). More specific
266	ically, the effect of incorporating the specific $P.\ clavata$ buoyancy mainly affected the re-
267	tention rates in PTM, PRV and to a lower extent in SES (Fig.3 b). The effect on all other
268	connections was similarly negligible. The effect of turbulence dispersal only affected re-
269	tention in PTM (Fig.3 c,d).

The above results were true for all simulations with short PLD (1-7 days). For longer PLD (10.5-28 days), sensitivity to the specific buoyancy of *P. clavata* and to turbulence intensity were significantly reduced, although specific buoyancy effect remained prevalent (Fig.4). While the specific buoyancy continued to affect mainly retention rates when the PLD increase from 3.5 days to 28 days (although in different locations, Fig.3b and Fig.4b), the effect of turbulence mainly affected distant connections (import for BER, Fig.4c and d).

Finally, varying the PLD from the reference value of 3.5 days to 1 or 7 days affected larval transport estimates of the same order of magnitude as those obtained by incorporating *P. clavata* buoyancy for the same PLD (Fig.5). Reducing the PLD from 3.5 days to 1 day only increased the retention rates (Fig. 5a) while extending the PLD from 3.5 to 7 days not only decreased retention rates but also distant connections from SES to POR and from PRV to PTM (Fig. 5b).

283 4 Discussion

In the present study we tested the sensitivity of larval transport simulations among coastal populations in the Ligurian Sea to different parameters. The horizontal resolution of the hydrodynamical model was found to have the strongest effect on larval trans-

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Figure 3. Median transfer deviation matrix for all the larval release events of 2004-2006 and a PLD of 3.5 days. Deviations are computed between (a) neutrally buoyant coarse resolution simulations without turbulence, (b) *P. clavata* buoyancy fine resolution simulation without turbulence, (c) neutrally buoyant fine resolution simulation with low turbulence, (d) neutrally buoyant fine resolution simulation with high turbulence and the reference fine resolution neutrally buoyant simulation without turbulence.



Figure 4. Same as 3 but for a PLD of 28 days



Figure 5. Median transfer deviation matrix for all the larval release events of the 2004-2006 fine resolution neutrally buoyant simulations without turbulence. Deviations are computed between (a) a PLD of 1 day, (b) a PLD of 7 days and a PLD of 3.5 days.

port followed by larval behaviour and PLD, while turbulence was found to have a sec-287 ond order effect. Thanks to the progresses in ocean modelling availability and perfor-288 mances, larval transport predictions are now frequently used to explore the functioning 289 of marine populations and help understanding their spatial distribution (reviewed in Swearer 290 et al. (2019)). Assessing the accuracy of larval transport predictions is more than often 291 qualitative (Barbut et al., 2019; Crosbie et al., 2019). However, quantifying the improve-292 ment of larval transport simulations thanks to the improvement in ocean modelling (i.e. 293 resolution, forcing) and larval traits knowledge (i.e. behaviour, PLD, release events) is 294 a preliminary to asses the accuracy of larval transport predictions. 295

It has been recurrently argued that including larval motility behaviour was key to 296 improve the realism of larval dispersal predictions, and led to develop tracking softwares 297 specific for larval dispersal studies (Lett et al., 2008; Schlag & North, 2012; Paris et al., 298 2013). This is particularly relevant for the dispersal of fish larvae that can orient them-299 selves (Faillettaz et al., 2018). For many benthic invertebrates species which are unable 300 to outcompete flow speeds by swimming, only larval buoyancy and motility behaviour 301 regulate their position in the water column (Leis, 2007). Yet, knowledge about larval motil-302 ity behaviour is still restricted to a limited number of species, mainly commercial species 303 such as oysters (North et al., 2008), scallop (Tremblay et al., 1994), red coral (Martínez-304 Quintana et al., 2015) or tropical reef fish species (review in Leis (2006)). Behaviour ef-305 fect on larval transport estimates was either explored in sensitivity studies confronting 306 simulations with different hypothetical behaviours (Donahue et al., 2015; Blanco et al., 307 2019) or in projections studies to inform about populations functioning or management, 308 as a replacement of observations (Munroe et al., 2018; Goodwin et al., 2019; McManus 309 et al., 2020). The present study confirmed the importance of motility behaviour in lar-310 val dispersal predictions, the inclusion of a sinking speed of about 0.5 mm/s altered lar-311 val dispersal pattern by 10% for short PLD. 312

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However, other larval traits alter dispersal patterns in simulations. Several recent 313 works performed 2D or 3D dispersal varying PLD only (Aguilar et al., 2019; Gamoyo 314 et al., 2019; Hidalgo et al., 2019). In the present study, a PLD increase from 1 to 3.5 days 315 resulted in 10% deviation in dispersal patterns, similar to the effect of *P. clavata* behaviour. 316 PLD is a very uncertain parameter as environmental mortality processes occurring dur-317 ing dispersal, such as predation or starvation for planktotrophic larvae, are underesti-318 mated in the laboratory (Rumrill, 1990). For instance, for the white gorgonian Eunicella 319 singularis, the most frequent effective PLD determined by comparing dispersal patterns 320 simulations with population genetics (Padrón et al., 2018) lies in the lower range of po-321 tential PLDs determined experimentally (Guizien et al., 2020). For the red gorgonian 322 P. clavata, population genetics indicated short distance dispersal are the most frequent, 323 but long distance migrants have been observed (Mokhtar-Jamaï et al., 2011), suggest-324 ing that the larger potential PLD observed in laboratory experiments (Guizien et al., 325 2020) may also contribute to population genetics. In conclusion, despite an accurate rep-326 resentation of motility behaviour, the output of larval dispersal studies can be signifi-327 cantly impacted by uncertainty on PLD estimates, resulting from ecosystemic interac-328 tion. 329

Nevertheless deviation among dispersal patterns of similar order of magnitude as 330 the effect of behaviour and PLD was obtained when varying another larval trait, the re-331 lease timing within the current reproductive season. Such deviation was expected from 332 the interaction of the short release event duration of *P. clavata* with the natural short 333 term variability of ocean flows in the area. A similar result was obtained in the Gulf of 334 Lion where connectivity patterns variability increased when release event duration de-335 crease (Guizien et al., 2012). In this area, such effect was even more significant as wind 336 forcing explains most of the coastal circulation (Briton et al., 2018). As a consequence, 337 despite larval behaviour and PLD might be better ascertained, uncertainty on disper-338 sal patterns will remain larger for species releasing larvae or gametes during short pulses, 339 compared to species releasing propagules regularly over long periods. 340

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The coastal circulation in the Ligurian Sea is characterized by high variability of 341 the currents at different temporal scales from seasonal (Astraldi & Gasparini, 1992) to 342 climatic (Astraldi & Gasparini, 2013) in particular the years from 2004 to 2009 were char-343 acterized by an abrupt shift in the Western Mediterranean circulation (Schroeder et al., 344 2016) starting from 2005. The present study has pinpointed that in such cases, larval 345 dispersal studies should include estimating connectivity variance with intra-annual vari-346 ability and not only with inter-annual variability. In fact the seasonal and climatic vari-347 ability equally affect larval dispersal patterns suggesting that yearly averages should be 348 avoided. 349

Aside to the temporal variability of the hydrodynamical forcing, the importance 350 of spatial resolution of the hydrodymanical ocean model in affecting the dispersal pat-351 terns of larval transport simulations was questioned in several studies (Guizien et al., 352 2006; Huret et al., 2007; Andutta et al., 2012; Putman & He, 2013) and reviewed in Swearer 353 et al. (2019). A recent sensitivity study reported after comparing two ocean model res-354 olution of 4 km and 1.6 km, with and without larval vertical behaviour in different years, 355 that inter-annual variability was the most influential factor in dispersal simulations of 356 fish larvae in the North sea (Kvile et al., 2018). The present study showed in contrast 357 that the most influential factor in the coastal Ligurian population was ocean model res-358 olution as retention increased dramatically when decreasing the horizontal cell size from 359 1.5 km to 300 m. Such bias is to be expected in coarse resolution ocean models when 360 the habitat lies within the coastal boundary layer where the model representation of the 361 flow is inaccurate due to both their coarse resolution and their lack of coastal ocean pro-362 cess parameterization (Holt et al., 2010; Griffies & Treguier, 2013; Holt et al., 2017). The 363 large deviation in connectivity fluxes found in the present study when increasing ocean 364 model resolution questions the reliability of past studies of coastal habitat connectivity 365 366 using basin or Ocean General Circulation ocean Models (OGCM) (Corell et al., 2012; Andrello et al., 2013). However, those models still have an interest for connectivity stud-367 ies to delineate well-connected oceanic regions, when care is taken of releasing particles 368

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369	outside this artefactual coastal numerical boundary layer (Rossi et al., 2014). Delineat-
370	ing such well-connected regions is essential to identify areas in which ocean circulation
371	model downscaling should be performed to study specific populations connectivity. To
372	this respect, including turbulence into dispersal simulations is expected to be less im-
373	portant when the cell size decrease as mesoscale and sub-mesoscale processes will be re-
374	solved (Guizien et al., 2006; Huret et al., 2007). In the present study, the effect of tur-
375	bulence on connectivity patterns was clearly negligible when resolving the coastal cir-
376	culation at 300 m in comparison to the effect of biological traits.
377	In conclusion, we formulate the following guidelines for larval dispersal simulations:
378	• Ensure that the grid resolution is sufficient to resolve the coastal dynamics around
379	the reproductive population where larvae are released. In such a case, multiply-
380	ing dispersal tracks to account for turbulence effect can be avoided, reducing com-
381	putational costs. Coastal dynamics may be very different from place to place but
382	this study showed that, at least for the Ligurian Sea, horizontal resolutions in the
383	order of a few hundred meters should be employed.
384	• Assess the variability of connectivity patterns arising from uncertainty on release
385	timing by simulating multiple release events. Multiple release events should at least
386	span over the reproductive season within a year or over multiple years, when inter-
387	annual climatic variability dominates over short term climatic variability. As a guide,
388	we suggest a number of release events equal to the ratio of the duration of the re-
389	productive season divided by the duration of a single release event. Such proce-
390	dure would ensure to estimate confidence level for the different connectivity pat-
391	terns.
392	• Assess the variability of connectivity patterns arising from uncertainty on PLD
393	by simulating multiple dispersal duration, spanning the range of potential PLDs.
394	Such procedure would ensure to estimate most likely PLDs by comparing simu-

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- lated connectivity patterns with observed ones, baring in mind that retention sen sitivity to PLD is the largest for short PLD.
- Include larval motility behaviour (buoyancy and swimming speed) prior to assess-
- ³⁹⁸ ing PLD effect in dispersal patterns for a specific species. This behaviour should
- be based on experimentally derived parameters, including inter-individual vari-
- ability, to reduce uncertainty on dispersal patterns.

401 Supporting Information

402 Supporting Information are available in online version of the manuscript.

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