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Coupling hydrodynamic and individual-based models to simulate longterm larval supply to coastal nursery areas

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Abstract:

For many marine fish species, recruitment is strongly related to larval survival and dispersal to nursery areas. Simulating larval drift should help assessing the sensitivity of recruitment variability to early life history. An individual-based model (IBM) coupled to a hydrodynamic model was used to simulate common sole larval supply from spawning areas to coastal and estuarine nursery grounds at the population scale in the eastern Channel on a 14-yr time series, from 1991 to 2004. The IBM allowed each particle released to be transported by currents from the hydrodynamic model, to grow with temperature, to migrate vertically giving stage development, and possibly to die according to drift duration, representing the life history from spawning to metamorphosis. Despite sensitivity to the larval mortality rate, the model provided realistic simulations of cohort decline and spatio-temporal variability of larval supply. The model outputs were analysed to explore the effects of hydrodynamics and life history on the interannual variability of settled sole larvae in coastal nurseries. Different hypotheses of the spawning spatial distribution were also tested, comparing homogeneous egg distribution to observation and potential larval survival (PLS) maps. The sensitivity analyses demonstrated that larval supply is more sensitive to the life history along larval drift than to the phenology and volume of spawning, providing explanations for the lack of significant stock-recruitment relationship. Nevertheless, larval supply is sensitive to spawning distribution. Results also suggested a very low connectivity between supposed different sub-populations in the eastern Channel.

Keywords: eastern Channel; hydrodynamic model; individual-based model; larval supply; nursery grounds; recruitment variability; *Solea solea*

1. Introduction

As marine fish species generally produce a high number of eggs with a low probability of survival (Juanes, 2007), small changes in survival rates during early life may cause large variations in recruitment (Daewel et al., 2011). A thorough exploration of survival at successive early life stages is necessary to understand fish population dynamics (Levin and Stunz, 2005). In particular, for a large proportion of commercial marine fish species, stock recruitment relationships are challenged and a better knowledge of recruitment processes would improve our understanding of population dynamics and contribute to a better stock assessment (Brunel and Boucher, 2007).

Estuaries and coastal waters are of major importance for many marine species, especially because they host nursery grounds (Beck et al., 2001). Their carrying capacity influence juvenile fish abundance thus population renewal (Rijnsdorp et al., 1992; Gibson, 1994; Iles and Beverton, 2000; Juanes, 2007). Recruitment and population dynamics of coastal and estuarine dependent species also rely on processes generating high variability in early life stages and especially larval supply driven by hydro-climatic conditions (van der Veer et al., 2000; Houde, 2008). A better understanding of larval processes would help estimating the number of larvae successfully reaching nursery grounds, identifying factors generating the associated spatio-temporal variability and their consequences for recruitment to juvenile stages, as well as population dynamics and structure.

Larval drift individual based models (IBM) have proven useful to investigate the influence of hydrodynamics on ecology of fish larvae, with promising perspectives of integration in fishery management (Hinrichsen et al., 2011). They have been used to identify spawning areas (Walsh et al., 1996), drift duration (Bonhommeau et al., 2009), vertical migration behaviour (Fox et al., 2006; Sentchev and Korotenko, 2007), dispersal kernels (Koutsikopoulos et al., 1991; Ellien et al., 2000, 2000; Sentchev and Korotenko, 2005; Huret et al., 2010) or connectivity within a meta-population (Savina et al., 2009). However, most of these studies were conducted on relatively short periods of time and limited spatial scales. The use of numerical simulations over longer time series at the population scale could provide useful tools to study spatio-temporal recruitment variability generated at larval stages for estuarine and coastal nursery dependent species (Bonhommeau et al., 2009).

The common sole of the eastern Channel (International Council for the Exploitation of the Seas, ICES, division VIId, Fig. 1) is an important commercial flatfish resource, managed as a unique stock. In the eastern Channel, the common sole spawns in early spring, pelagic larvae dwell during several weeks and newly metamorphosed individuals settle in coastal and estuarine areas, acting as exclusive nursery grounds for about 2 years, before sexual maturity (Riou et al.,

2001). Recruitment variability is high (Le Pape et al., 2003) and appears not synchronous among the different nursery areas (Riou et al., 2001), hence suggesting a possible high influence of hydro-climate on larval transport and larval supply.

<Fig. 1>

To explore the potential effects of hydrodynamics and life history from spawning to metamorphosis on the number of settled sole larvae over nurseries and the associated spatiotemporal variability, larval supply in the different nursery grounds of the eastern Channel was simulated using a three dimensional hydrodynamic model. An individual based particle-tracking module was used to simulate larval drift and survival under realistic biological processes from egg to benthic settlement on estuarine and coastal areas. The sensitivity of larval supply to spawning phenology, larval drift and mortality was assessed at both the nurseries and population scales and the consequences on population dynamics and sub-populations structure were analysed.

2. Materials and methods

2.1. Outlines of the model

The ultimate objective was to simulate larval supply in different coastal and estuarine sectors (Riou et al., 2001; Rochette et al., 2010) of the eastern Channel (Fig. 1). The approach combined a hydrodynamic model with a particle-tracking module including behaviour to simulate spawning and hatching of eggs, larval drift, growth and settlement on coastal and estuarine nursery grounds under realistic hydro-climatic conditions (Fig. 2). Particles were released following the characteristics of the spawning season and spatial distribution of eggs. The development (growth), swimming behaviour, benthic settlement and survival of each individual (Fig. 2) were modelled in parallel with its transport in the oceanographic model.

<Fig. 2>

2.2. A hydrodynamic model coupled with a particle-tracking module

The three dimensional (3D) ocean circulation model MARS (hydrodynamic Model for Application at Regional Scale; Lazure and Dumas, 2008) was used to simulate hydrodynamics in the (western and eastern) Channel during 14 years (1991 - 2004). The model used a 2 km horizontal resolution with 30 vertical sigma layers. The eastern Channel is about 35000 km² with an average depth of 35 m and a maximum of 100 m. A time step of about 2 minutes allowed the

model to solve the strong tidal currents occurring in the Channel, with respect to stability criteria. River discharges and atmospheric forcing were similar to those in Huret et al. (2010), with interannual variability at open boundaries for temperature and salinity provided by a global circulation model solution ORCA from the OPA-NEMO model (Michel et al., 2009). The simulation period 1991-2004 was the longest allowing for consistent available forcing conditions (river, open boundary and atmospheric).

A particle-tracking module was coupled on-line to the 3D hydrodynamic model with a random-walk to account for vertical turbulent mixing as in Huret et al. (2010). This module included an individual-based modelling framework simulating trajectories and life traits of released particles (Ayata et al., 2010), from spawning areas to nursery grounds (Fig. 2).

The results of the model depend upon (*i*) initial conditions, *i.e.* how particles are released (as eggs) in time and space; (*ii*) behaviour and mortality during the drift. Influences of the spatiotemporal variability of spawning and of the natural mortalities were implemented in a posttreatment analysis by applying *a posteriori* weighting factors to released particles (Fox et al., 2006). Indeed, each particle released in a specific year (y=1,...,n), date in the year (in Julian day, t_r), and point of release in the 3D grid ($s_{r(3D)}$, noted s_r) represented a certain number of eggs having the same initial conditions $N_0[y,t_r,s_r]$. Besides, a survival probability was calculated for each particle at the end of the simulated trajectory. The $N_0[y,t_r,s_r]$ individuals associated with each trajectory were limited by this survival probability during the different eggs and larval stages (Fig. 2) to estimate the number of surviving larvae settling in and outside the nursery grounds (Fig. 1).

2.3. Initial conditions: spatial and temporal variability of egg release

Initial number of particles and vertical distribution - To assess the influence of the spatiotemporal distribution of spawning, a trade-off was found between constraints of computing time and frequency and number of particles release that well represents the interannual variability. Different frequency and number of particles were tested. A patch of particles covering the entire eastern Channel was finally released every 15 days during the whole spawning season, from February to June (Anon., 1993), with particles released 1 cell out of 4 on the whole 2D model grid ($s_{r(2D)}$). Koutsikopoulos et al (1991) observed eggs over the whole water column at stage 1 but mainly just below surface for stage 2 and 3 (Fig. 3b), suggesting a buoyancy effect. Accordingly, a particle was released each meter from 0 to 5 meters, giving 5 particles at each location $s_{r(2D)}$. Around 9000 particles were released for each year simulated. Each particle represents a number of eggs $(N_0[y,t_r,s_r])$ depending on (*i*) the total number of eggs spawned each year, $N_{eggs}(y)$ and (*ii*) the spatio-temporal distribution of eggs in the eastern Channel, included as a weighting factor, $W_{time,space}[y, t_r, s_r]$:

$$N_0[y,t_r,s_r] = N_{eggs}[y] \cdot W_{time,space}[y,t_r,s_r]$$
(1)

Interannual number of eggs – The total number of eggs released each year in the eastern Channel $(N_{eggs}[y])$ was calculated as:

$$N_{eggs}[y] = \sum_{a} (N_{adults}[a, y] \cdot Pf[a] \cdot Ew[a, y] \cdot Pm[a])$$
⁽²⁾

where N_{adults} is the number of adults of age (*a*) in year (*y*) (ICES, 2010), *Pf* the female fraction, *Ew* the number of eggs per female and *Pm* the proportion of mature females (0% until age 2 and 100% after; ICES, 2010). The proportion of female was calculated from a logistic regression:

$$Pf[a] = 1 / (1 + \exp[-(a_{Pf}.a^2 + b_{Pf}.a + c_{Pf})])$$
(3)

with parameters a_{Pf} =-2.79E-3, b_{Pf} =1.58E-1 and c_{Pf} =-4.10E-1 derived from van Beek (1988).

The numbers of eggs at age (Ew) were calculated from the number of eggs-at-weight relation (Anon., 1992) and the yearly weights-at-age observed for the eastern Channel population (ICES, 2010).

Weighting factors – Each patch of particles released every 15 days along the spawning period represented a proportion of the total number of eggs spawned in a year ($W_{time}[y, t_r]$), and was distributed in space ($W_{space}[t_r, s_r]$). The spatio-temporal weighting factors $W_{time,space}[y, t_r, s_r]$ were the combination of time and space weighting factors, standardized to 1, such that the sum of weighted particles released each year equalled the number of eggs spawned on this year:

$$W_{time,space}[y,t_r,s_r] = \frac{W_{time}[y,t_r] \cdot W_{space}[t_r,s_r]}{\sum_{t_r,s_r} (W_{time}[y,t_r] \cdot W_{space}[t_r,s_r])}$$
(4)

Time weighting factor – $W_{time}[y, t_r]$ were calculated using an index of the daily number of eggs spawned varying among years, noted $\alpha_{time}[y, t_r]$:

$$W_{time}[y,t_r] = \frac{\alpha_{time}[y,t_r]}{\sum_{t_r} \alpha_{time}[y,t_r]}$$
(5)

In the eastern Channel, the sole spawning period starts in February - March when sea surface temperature reaches 7°C (Fonds, 1979) and ends in June (Fig. 2) with a spawning peak in April -

May (Anon., 1986, 1992, 1993; van Beek, 1988) correlated to sea surface temperature (van der Land, 1991). For each year, a dome shaped curve was defined from a model established on the egg survey conducted in 1991 by the UK Centre for Environment, Fisheries and Aquaculture Science (CEFAS; Anon., 1992). The daily number of eggs in 1991 ($\psi(t)$) was fitted with a parabolic equation on the log scale:

$$\log(\psi(t)) = a_{p} + b_{p} t + c_{p} t^{2}$$
(6)

where t is the Julian day and the estimated parameters a_p =-1.288, b_p =0.217and c_p =-0.001.

From Eq. (6), the spawning peak in 1991 occurs on Julian day 110 (April 20th). In the 3D numerical model, the mean sea surface temperature on the spawning zone at this date was 9°C, consistent with the estimate of Anon. (1992). Following this, for each year, the spawning peak was defined as the Julian day with maximum proportion of sea surface temperature between 8.5 and 9.5°C. $\alpha_{time}[y, t_r]$ were calculated by translating the dome-shaped curve established in Eq. (6) with regards to the estimated spawning peak.

Spatial weighting factor - The spatial weighting factors $W_{space}[t_r, s_r]$ standing for the spatial and vertical (3D) distribution of eggs were considered constant over years but varied along the spawning season. The single sole egg survey sampling the entire eastern Channel was conducted in 1991 (Anon., 1992), monthly from March to June. Other surveys (Anon., 1986; Grioche et al., 1999) sampled only the eastern half of the area. Eastwood *et al.* (2001) compiled these data and pointed out that eggs were found over sandy sediments but not on pebbles and rocks (Fig. 3a). Accordingly, no particles were released over coarse sediments. On soft bottoms, the 2D egg distribution $W_{space2D}[t_r, s_{r(2D)}]$ was derived from kriged maps based on the 1991 egg survey (one for each month). A classical kriging approach was used on log-transformed egg densities with a common exponential variogram (Matheron, 1963; Clark, 1979) and interpolations until a distance of 20 km (see the average interpolated egg distribution on Fig. 3a). As patches were released every 15 days in the model, particles of intermediate patches were weighted according to the average egg distribution of the previous and the following survey. Patches starting before mid-March were weighed from the first survey and after mid-June from the last one. Finally, the spatial weighting factors were standardized:

$$W_{space}[t_r, s_r] = \frac{W_{space2D}[t_r, s_{r(2D)}] / Nb_{Particules}[t_r, s_{r(2D)}]}{\sum_{t_r} (W_{space2D}[t_r, s_{r(2D)}] / Nb_{Particules}[t_r, s_{r(2D)}])}$$
(7)

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Further analyses were conducted to validate this spawning pattern, assuming that the spatial distribution of eggs should be close to the distribution of the adult population during the spawning season (Loots et al., 2009). Monthly spatial distributions of catches (French: ~57%, Belgian: ~26%, and English: ~17%) for years 2006 to 2008 were provided from the ICES Working group (J. Vigneau, Ifremer, com. pers.). Even if the resolution available for these catch data is large (30' of latitude; 1° of longitude) and does not show the higher egg production along the coasts (Fig. 3a), their analysis showed that monthly spatial distribution were rather constant among years. Similarly, Loots et al. (2009) pointed out the lack of interannual variations for spatial distribution of the North Sea plaice (*Pleuronectes platessa*) spawning population. Comparison with the monthly distribution of the 1991 egg survey demonstrated a good overlap in seasonal distribution at the ICES rectangle resolution and suggested the interannual steadiness of the monthly spawning distributions.

<Fig. 3>

2.4. Life traits of drifting particles

The status of particles evolves during the drift through 5 successive development stages (Fig. 2): passive transported eggs, passive transported larvae at development stage 1, vertical migrating larvae from stage 2 (Fig. 3b), larvae at metamorphosis and settled larvae (Fig. 2). Transitions between these size specific development stages and mortality rates vary depending on water temperature along trajectories:

Eggs \rightarrow Larvae 1 - Four egg development stages were distinguished (Koutsikopoulos et al., 1991; Fig. 3b). Durations of egg stages are temperature-dependent (see Fonds (1979) for explicit temperature-dependent equations). Eggs are ascending to the surface during the three first development stages using a vertical speed to converge to surface, but are also subject to vertical mixing with the model random-walk. Vertical advection is fully passive for the fourth development stage. The survival of eggs was calculated as $exp(-M_{eggs}, \Delta t_{eggs})$ where Δt_{eggs} is the duration of egg stage and M_{eggs} their mortality rate, related to egg diameter (*diam*) and temperature encountered by the associated particle (Rijnsdorp and Vingerhoed, 1994):

$$M_{eggs} = \exp(a_M + b_M \cdot diam + c_M \cdot \overline{T})$$
(8)

where \overline{T} is the mean temperature during the whole duration of the egg stage; a_M =1.615, b_M =-3.517, c_M =0.185. Egg diameter at spawning depended on the time from spawning peak as older females spawn bigger eggs later in the spawning period (size was 1,266 mm at the spawning peak; Rijnsdorp and Vingerhoed, 1994). This diameter increased by 0.002 mm.day⁻¹ (Anon., 1992). Data used to calculate egg diameter and related mortality for *Solea solea* in the eastern Channel were issued from Rijnsdorp and Vingerhoed (1994) from an egg survey conducted in 1991.

Larvae $1 \rightarrow Metamorphosis$ – Duration of larval stage 1 is temperature-dependent (see Fonds 1979). Larvae are unable to swim actively during this stage. The length of larvae at the end of stage 1 (L_1) is modelled following Eq. (9) with parameters $a_L=3.528$ and $b_L=0.044$ estimated for *Solea solea* from results of Fonds (1979) and \overline{T} the mean temperature along larval trajectory.

$$L_1 = a_L + b_L \cdot \overline{T} \tag{9}$$

At stage 2, larvae acquire a swimming capacity and go through vertical nycthemeral migration (Fig. 3b; Champalbert et al., 1989, 1992; Koutsikopoulos et al., 1991). The swimming speed is parametrised at 1 body length per second (Gibson, 2005). Such vertical migration is implemented by a depth the particles converge to, using their swimming speed, and is combined to the random-walk of the model. From stage 2 to 4, larval length at each time step is calculated from Fonds (1979):

$$L_{t+dt} = L_t . \exp(g_L . dt) \tag{10}$$

where dt is the time step and $g_L (day^{-1})$ the temperature-dependent growth coefficient, a quadratic function of the mean temperature issued from Fonds (1979):



Metamorphosis (Fig. 2) starts when larvae reaches 8.5 mm (Amara and Lagardère, 1995).

Larval mortality - The survival probability during the larval stages is calculated as $exp(-M_{larvae}, \Delta t_{larvae})$ where Δt_{larvae} is the duration of the entire larval stage until metamorphosis, specific for each trajectory. In the baseline scenario, M_{larvae} is fixed to 0.04 days⁻¹ according to previous estimates for sole (Koutsikopoulos et al., 1991; Horwood, 1993; Lund et al., 2008).

2.5. Larval settlement and end of simulations

The aim of the present model was to estimate and analyse the variability of the number of sole larvae reaching nursery grounds in the eastern Channel. From stage 2 to 4, larvae are able to use a selective tidal stream transport combined to nycthemeral migrations to reach the coasts (Rijnsdorp and Stralen, 1985; Bergman et al., 1989; Amara et al., 1998; Grioche et al., 2000; Forward and Tankersley, 2001; Fox et al., 2006). However, this behaviour only appears when larvae approach settlement zones (Boehlert and Mundy, 1988; Ramzi et al., 2001ab) and no metamorphosed larvae nor juvenile were ever found outside nursery areas (Riou et al., 2001). To avoid the spurious simulation of larval behaviour over coastal areas, the model tested if larvae reach the coastal strip before metamorphosis. Hence, the final destination over the nursery was not targeted and larval supply was only estimated at the large extent of the 6 coastal nursery sectors (Fig. 1) based on juvenile distribution (Riou et al., 2001; Rochette et al., 2010).

In the model, larvae are transported until they reach the coastal area within the 25m isobath with a soft bottom (i.e. nursery ground; Fig. 1) or until they reach metamorphosis. If they reach a nursery at stage 2 or higher before metamorphosis, they are considered to settle in this area but the larval mortality is still applied until metamorphosis (8.5 mm). If they attain metamorphosis outside these coastal sectors, a 0% survival is applied. This is consistent with the hypothesis that larvae die while reaching size of metamorphosis out of shallow areas, as distance possibly travelled by a metamorphosed larvae outside suitable habitats to reach nursery sectors is low. Indeed, the 25 m bathymetry limit indirectly accounts for a possible active travelled distance, as the observed limit for juveniles distribution is 20 m isobath in the eastern Channel (Riou et al., 2001; Rochette et al., 2010).

2.6. Model outputs: patterns in larval supply to nursery grounds and spatial structure

Model outputs were first analysed in terms of spawning dynamics, drift duration, cohort decline and number of successful larvae at the population scale. Then, the interannual variability of larval supply was estimated on the different nursery sectors.

The connectivity between spawning areas and nursery grounds was analysed. The individual-based trajectories allowed mapping the Potential Larval Survival (PLS, Daewel et al., 2011), i.e. the probability to reach a nursery for each initial location. Connectivity was also

analysed by dividing the eastern Channel into three spawning sub-areas (Fig. 3a), based on (*i*) the map of egg distributions and (*ii*) the presence of a central Channel eastward current (Hoch and Garreau, 1998) which separate English and French coasts. Three spawning sub-areas were defined: the western and eastern part of the French coasts, separated by a rocky central area, and the English coasts (Fig. 3a). Contributions of settled larvae coming from each spawning sub-area were calculated for each nursery.

2.7. Sensitivity analysis: sources of variability in spatio-temporal larval supply

Sources of interannual variability in larval supply – Three factors acting at the population scale in the model impact the interannual variability of successful larval abundance: (*i*) larval mortality, (*ii*) variability of the spawning peak and (*iii*) number of eggs released. The relative influence of the survival during transport versus the number of eggs spawned in the interannual variability of larval supply was assessed. This question is of critical importance as this determines the possible observation of the stock-recruitment relationship. The influence of the 3 factors above on the interannual fluctuations of larval supply was tested at the scales of both the whole eastern Channel population and the nursery sectors, while modifying the baseline scenario: (*i*) As the larval mortality rate M_{tarvae} (in days⁻¹) was fixed to 0.04 in the model, two other values were tested (0.03 and 0.05) with regards to literature. (*ii*) To assess the influence of the spawning rate during the spawning season was tested. (*iiii*) To assess the influence of the spawning stock biomass (SSB) a scenario with constant number of eggs released each year ($N'_{eggs}[y] = mean(N_{eggs}[1991-2004]$)) was tested.

Sensitivity to spawning distribution – The sensitivity of the larval supply to the spawning distribution was analysed from two alternative scenarios: (*i*) a homogeneous egg distribution, and (ii) an egg distribution according to the PLS map (*cf.* §2.6).

Indicators to quantify the level of sensitivity of larval supply to mortality and spawning spatio-temporal dynamics – The sensitivity analysis was performed through two statistics related to interannual recruitment success: (i) the mean bias, mean $[(N_{larvae_scenario}(y) - N_{larvae_baseline}(y)) / N_{larvae_baseline}(y)]$, to test for scale factors in larval supply, and (ii) the linear correlation coefficient, $\rho(N_{larvae_scenario}(y) \sim N_{larvae_baseline}(y))$, to test for changes in fluctuations among years. These criteria were calculated at both the population and the nursery scales.

Sensitivity of the level of spatial connectivity to model inputs – Connectivity between spawning areas and nursery grounds may also depend on larval mortality, variability in the timing of the spawning peak and spawning distribution. Hence, their influence on the PLS map and the exchanges between the 3 sub-areas (Fig. 3a) was also analysed, using the scenarios described above.

3. <u>Results</u>

3.1. General patterns of the simulated larval drift

The timing of the spawning peak showed a high interannual variability. The estimated peak ranged from mid-March for warm years to the end of April for colder ones. The drift duration, between spawning and metamorphosis, was variable between years, ranging from 51 to 68 days with a mean at 59.5 days. The average of the settlement period took place between mid-May and mid-June (Fig. 4).

<Fig. 4>

At the population scale, the mean survival, averaged over the 1991-2004 period, from spawning to settlement, estimated with the baseline mortality rate for larvae $M_{larvae}=0.04$ day⁻¹ was 10.5 settled larvae in the eastern Channel for 10000 eggs spawned (Fig. 4, Fig. 5). This low recruitment success at settlement resulted from (*i*) the high mortality from spawning to metamorphosis (98% death) and (*ii*) the high mortality at settlement due to the large proportion of particles undergoing metamorphosis outside nursery grounds (95% death). The latter consisted of particles settling in deep water in the eastern Channel (49%) or transported outside the study area, in the North Sea (48%) and in the western Channel (3%).

The settlement success varied along the season with relation to both the high mortality during the temperature-dependent drift duration and changes in current direction. Average survival increased from 6.6 settled larvae per 10000 eggs at the beginning of the spawning period (patch released \sim 30 days before spawning peak), to 11.5 at the end (\sim 30 days after).

The interannual variability of the larval supply was high (Coefficient of Variation, standard error / mean, CV=63%) during the period 1991-2004 (Fig. 5) with total larval supply (over all nursery grounds) ranging from 1.4×10^{9} in 2001 to 9.7×10^{9} in 1996. This variability was correlated to spawning peak as later spawning peak showed a higher larval supply. Although temperature influences drift duration and larval mortality, survival to drift did not explain this correlation. The correlation was mainly due to the probability to reach metamorphosis over a nursery (Fig. 5). Indeed, the loss of larvae in the North Sea was higher for warmer years (54% of metamorphosis in the North Sea on average for years 1997, 1998, 1999 and 2002) than for colder years (42% for years 1991, 1992 and 1996). This particularly affected the Somme sector, which proportion in total larval supply is especially high (46% on average; Fig. 5). The larval supply among the nursery sectors was variable among years with coefficient of variation from CV=42% (U.K. West) to CV=150% (Seine); these fluctuations were not synchronous among nurseries (Fig. 5).

<Fig 5>

3.2. Connectivity from spawning to nursery grounds

The common sole population of the eastern Channel seems to maximize its success in larval transport when spawning along the English and the eastern French coasts (Fig. 6). However, if observed egg distribution (Fig. 3a) present similarities with PLS maps (Fig. 6), it does not reflect where spawning could potentially result in high levels of early stage survival for two areas: (i) an important spawning area exists in the north of the eastern French coasts, associated with a low or null probabilities to settle on a nursery of the eastern Channel, and (ii) spawning in the Seine estuary is associated with a high probability of settling on a nursery but only a few eggs were spawned there in 1991.

PLS (Fig. 6) supported the influence of the eastward current in the middle of the eastern Channel, separating northern and southern spawning areas (Fig. 3a). The strong relation between the origin of eggs and larval settlement denoted the low connectivity (Fig. 6) between the three sub-areas from spawning to settlement. On average over the period 1991 – 2004, almost all the particles settling on a coastal nursery resulted from eggs spawned in the associated spawning area. All the percentages were over 98% of self-supply, except for one episode in 1995 at 93% and 96% for the English and western French coasts respectively, and one in 2002 at 92% for the eastern French coasts.

<Fig. 6>

3.3. Sensitivity analysis

Sensitivity of larval supply to mortalities - Estimates of larval supply were especially sensitive to the mortality rate M_{larvae} . At the scale of the whole population, the mean survival from spawning to settlement ranged from 7.1 per 10000 eggs for $M_{larvae} = 0.05 \text{ day}^{-1}$ to 15.6 per 10000 for $M_{larvae} = 0.03 \text{ day}^{-1}$. Although the mean number of larvae was changed, neither the interannual fluctuations of the larval supply at the scale of the population nor at the scale of nurseries (R²~1, Table 1) were sensitive to change in larval mortality; larval mortality appeared as a scale factor for larval settlement success.

Sensitivity of larval supply to spawning dynamics - The sensitivity analysis first showed the low influence of the spawning timing on the total number of successful larvae: when a constant spawning rate was simulated from February to June, settlement success was not biased (Table 1) and had similar fluctuations than in the baseline scenario. This was verified at the scale of nurseries, except a positive bias for Seine and Calvados nurseries and a slight change in fluctuations in U.K. West nursery (Table 1).

The scenario evaluating larval supply without interannual variations of the number of eggs spawned revealed that larval settlement success was not sensitive to the spawning biomass variability for a realistic range of variation. A constant spawning biomass did not lead to bias nor changes in fluctuations of larval supply in the eastern Channel nor at the scale of nurseries except for a positive bias for Seine and Calvados nurseries (Table 1).

At the scale of the eastern Channel, model outputs revealed poorly sensitive to the initial spatial distribution of eggs. A homogeneous spatial spawning showed no major differences in larval supply at the scale of the eastern Channel. However, this hides a different allocation of larvae among nurseries (Table 1). Although interannual fluctuations were similar to the baseline scenario, the larval supply was lower in the Somme nursery but higher elsewhere, in particular in the Seine estuary (+203%; Table 1). When the PLS map (Fig. 6) was used instead of observed eggs distribution, larval supply was higher, with similar interannual fluctuations at both the eastern Channel and the nursery scales (Table 1). Only Calvados and Veys nurseries showed slightly different interannual fluctuations.

Sensitivity of connectivity to model options - The different scenarios revealed only a marginal influence on the PLS map. Larval mortality acted as a scale factor, with a negative relation between PLS and larval mortality but without change in the relative PLS distribution.

At the scale of the 3 sub-areas, the connectivity was not modified when running the different scenarios and remained very low, except when the spatial distribution of spawning was modified. Using homogeneous and PLS map influence the origin of particles reaching the Somme nursery, with respectively 10% (with 2 episodes of 40%) and 5% mean larval supply from the western French coasts.

<Table 1>

4. Discussion

4.1. Simulating larval recruitment at large spatial and temporal scales

This modelling approach illustrated the contribution of numerical simulation models to investigate the processes underlying the variability of fish recruitment (Hinrichsen et al., 2011), with a focus on a nursery dependent flatfish species. Simulations on relatively short periods of time were previously carried out to explore the effect of the hydrodynamics and larval behaviour on spatio-temporal mortalities or location of flatfish metamorphosis (Sentchev and Korotenko,

2007; Korotenko and Sentchev, 2008; Savina et al., 2009). The specificity of the present study consisted in exploring long-term simulations to assess the interannual variability of simulated trajectories. This was possible for flatfish as their life cycle is characterized by a metamorphosis to benthic life (Gibson, 2005) occurring early in the life cycle, before significant active swimming. From metamorphosis, juveniles swim to shallow adjacent nursery grounds (Ramzi et al., 2001ab; Fox et al., 2006), using selective tidal stream transport (Rijnsdorp and Stralen, 1985; Bergman et al., 1989; Amara et al., 1998; Grioche et al., 2000; Forward and Tankersley, 2001; Fox et al., 2006). This behaviour was assumed to maintain larvae over nursery sectors once the 25 m isobath limit was reached, which drastically simplifies the end of simulations under realistic hypotheses. Indeed, estimating larval supply at the scale of nurseries did not require knowing the exact final destination of larvae but only the nursery sector where settlement occurred. This focus on large scale results (>50km²), simplifying their analysis, allowed to carry out simulations over several years.

The simulation outputs patterns first confirmed that parameters of the biological model provided realistic life traits during pelagic stages for the common sole population in the eastern Channel. For instance, duration between spawning and settlement, estimated at 59.5 days, was in accordance with previous *in situ* estimates. Anon. (1993) estimated the time before hatching at "about 8 days" (average of 7.5 days in our study) then the duration of larval pelagic phase at "about 6 weeks" (52 days here). The average survival between eggs and settled larvae estimated from our simulations were low; only 10.5 (7 – 16) settled larvae survived for 10000 eggs spawned. This rate was also consistent with previous results for other flatfish population: ~10 for sole in the Bay of Biscay (Koutsikopoulos et al., 1989, 1991) and for the North Sea plaice (Le Pape and Bonhommeau, sub.).

However, on average, 48% of sole larvae reach the size of metamorphosis in the North Sea, outside the northern limit set for the eastern Channel population (ICES, 2010). These larvae could settle on northern nursery grounds (Rijnsdorp et al., 1992) and contribute to the North Sea population (Savina et al., 2009). Accordingly, PLS map revealed discrepancies between the low probabilities of success and the high amount of spawning products in the north east of the study area. This could be explained by larvae drifting to the North Sea, not counted as successful settlers in the present studied area thus underestimating the PLS in an important realized spawning area. Moreover, the model boundaries led to overestimate the number of particles reaching metamorphosis in the North Sea. Indeed, if the western limit of the hydrodynamic model was far from the eastern Channel, its north-eastern limit was only 10 km upper the studied area. As Lagrangian residuals sea currents are eastward in the eastern Channel (Hoch and Garreau, 1998), 24% of all the particles passed through the north-eastern boundary of the model and were not further simulated. Because of the balanced tidal currents, some of these particles

would have been able to settle in the eastern Channel if they had not been considered lost with the particle-tracking system. A complementary simulation run with a hydrodynamical model having a further northern limit but a coarser resolution suggested that the number of particles reaching "Rye Bay" could have been higher for some years. Conversely, this loss barely affected other nursery sectors. Thus, simulation outputs can be estimated as reliable, except for the northeastern part of the studied area where they should be considered as a first approach.

In spite of these local underestimation of the PLS, the larval supply model provided realistic simulations of pelagic stages dispersal and realistic levels of mortality at the scale of the eastern Channel sole population. They could be used to estimate the spatial and temporal patterns in larval supply and to identify the driving factors of their variability.

4.2. The main influence of larval drift on recruitment success

The huge natural variability in the mortality of young life stages generally hampers the identification of significant stock-recruitment relationships (Brunel and Boucher, 2007; Houde, 2008). The working group in charge of stock assessment of the common sole in the eastern Channel (ICES, 2010) could not identify any clear stock-recruitment relationship and projections are carried out under the hypothesis of a constant mean recruitment, independent from the spawning biomass.

Accordingly, the larval supply model highlighted relative variations of the number of successful larvae to be dominated by larval life history without obvious relation to spawning. Indeed, the sensitivity analysis revealed the low influence of (i) SSB, (ii) spawning peak and (iii) spawning distribution on the relative fluctuations of larval supply at the population scale. Conversely, the probability to reach a nursery at metamorphosis drove larval supply variations in the eastern Channel. At the scale of local nurseries, the interannual variability of larval supply was not synchronous between the different nursery sectors. Such a spatial asynchrony largely results from the interannual variability of the hydrodynamics in the eastern Channel. At the population scale, cold winters were correlated with higher larval supply in the eastern Channel whereas warmer winters showed a more important loss in the North Sea. This may be due to North Atlantic Oscillation (NAO) being higher for warmer winters and leading to stronger currents eastward (Hurrel and van Loon, 1997; Pingree, 2005). A higher NAO and a related warmer winter could lead to higher larval loss to the North Sea. This sensitivity of larval supply to hydrodynamics partly explained the difficulty to identify stock - recruitment relationship for this population. Moreover, although the spawning distribution barely affected the fluctuations of the larval supply, it had a high influence on the average amount of successful larvae over the studied period. The importance of currents and spawning distribution on larval supply pointed out the importance of considering larval drift (Werner et al., 1997) to improve understanding of recruitment variability (Hinrichsen et al., 2011). In that perspective, ocean circulation models coupled with individual-based transport systems provide insights on different species strategies (Daewel et al., 2011) and are potential tools for fishery management (Hinrichsen et al., 2009, 2011).

4.3. Connectivity, meta-population structure

The present modeling appoach analysed the trajectories of particles from spawning grounds to nurseries and focused on the limited larval dispersion in the eastern Channel, spawning areas feeding adjacent coastal and estuarine nursery grounds without significant exchanges with other regions of the eastern Channel. This new insight about life history during pelagic stages, together with previous analyses on older life stages raised questions about the existence of quite isolated sub-populations at the scale of the eastern Channel: Coggan and Dando (1988) and Anon. (1989) showed that sole juveniles stay over nurseries during their first two years of life. Kotthaus (1963) and Anon. (1965) observed that adult soles remained close to their nursery even after seasonal spawning migration. A mark-recapture survey investigated by Burt and Millner (2008) confirmed the low mobility of adult soles in the eastern Channel: adult soles of the English coasts remained along these coasts, soles of the eastern French coasts could migrate to the English coasts and soles of the western French coasts had only very low migrations. Despite the lack of observed genetic structure within the eastern Channel (Exadactylos. et al., 2003), the low local exchanges between sub-areas at larval stage, combined with previous conclusions on juveniles and adults behaviour, suggested a high level of segregation between sub-populations in the eastern Channel.

This raised the question of the management of the common sole fishery at the scale of the eastern Channel as a single stock (ICES, 2010) and in particular the possible isolation of the western French coasts. A potentially isolated sub population in this sector will be especially sensitive to highly degraded nursery area in the Seine estuary (Gilliers et al., 2006). At the scale of the eastern Channel population, the loss in abundance related to these anthropogenic disturbance for juvenile sole was estimated to be nearly 17% (15-32%; Rochette et al., 2010). The influence of habitat degradation on the population renewal would be exacerbated in the case of an isolated sub-population in the western part of the French coast. Habitat degradation could contribute to the choice of spawning areas and potentially explain why low realized spawning production in this sector, both spawning habitat selection (Daewel et al., 2011) and additional mortality during young stages (Gilliers et al., 2006) could be induced by local disturbances and

contribute to low densities of juvenile sole observed in nursery grounds in the Seine Estuary (Rochette et al., 2010).

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7. Figures.



Fig. 1. The eastern Channel divided in 6 different coastal nursery sectors, limited by the 25m isobath (dotted lines). Light grey: continent; hatched areas: pebbles and rocks. Lower right corner: general location of the study site



Fig. 2. Chart flow for the development of early life stages during the transport



Fig. 3. Distribution of eggs and larvae. (a) Kriged distribution of means of 1-day egg densities, averaged over the spawning period, interpolated from the 1991 egg survey (Anon., 1992). The darker, the denser. Hatched areas: pebbles, no spawning. Dashed lines: limits of the three spawning sub-areas. (b) Vertical distribution of sole eggs and larvae observed in scientific campaigns in the Bay of Biscay (see Koutsikopoulos et al., 1991).



Fig. 4. Simulated cohort decline from the biophysical model. Error-bars represent the interannual variability, with minimum and maximum values (at each stage, horizontal lines represent the variability of the spawning peak and vertical lines the variability of the number of eggs / larvae).



Fig. 5. Left axis: barplot: interannual variability of the larval supply distributed among nurseries. Right axis: Dashed line: date of the spawning peak (in Julian day). Thin line: survival of particles between spawning and metamorphosis (wherever metamorphosis occured). Bold line: probability to survive to metamorphosis (*i.e.* Metamorphosis over a nursery).



Fig. 6. Connectivity from spawning to nursery areas for the entire studied period (1991-2004). For each pixel in the grid, the color represents the mean probability, averaged on all simulated years, of successfull settlement on a nursery of the eastern Channel for a particle released from this point (PLS map, Daewel et al., 2011). The darker, the higher probability. Hatched areas: pebbles, no spawning. Dashed lines: limits of the three spawning sub-areas. Pies: origin of larvae settling over a nursery (proportions of larvae from the three spawning areas).

<u>Table</u>

Table 1. Sensitivity of larval supply to spawning and drift history: comparison with the baseline scenario. Mean bias: the difference of the number of successful larvae in each scenario relative to the baseline scenario (averaged over the 1994-2001 time series). R²: the coefficient of linear correlation of the time series of larval supply in each scenario against the baseline scenario. Main differences (>20% arbitrarily) are underlined. Pop. : Population scale. Calva.: Calvados.

Scenario	Indicator	Pop.	U.K. West	Rye Bay	Somme	Seine	Calva.	Veys
M_{larvae} =0.03 day ⁻¹	Mean bias	+50%	+51%	+53%	+50%	<u>+42%</u>	<u>+51%</u>	+52%
	R ²	~1	~1	~1	1	1	0.99	~1
M_{larvae} =0.05 day ⁻¹	Mean bias	<u>-32%</u>	<u>-33%</u>	<u>-34%</u>	<u>-32%</u>	<u>-29%</u>	<u>-33%</u>	<u>-33%</u>
	R ²	~1	~1	~1	~1	1	0.99	0.99
No peak of spawning: $W_{time}[y, t_r] = \text{constant}$	Mean bias	-4%	1%	0%	+3%	<u>+78%</u>	<u>+23%</u>	+7%
	R ²	0.96	<u>0.76</u>	0.85	0.99	0.85	0.91	0.80
Constant SSB: $N_{eggs}[y] = \text{constant}$	Mean bias	+3%	+1%	-2%	0%	<u>+73%</u>	<u>+21%</u>	+9%
	R ²	0.94	0.84	0.84	0.97	0.99	0.89	0.93
Homogeneous spatial spawning <i>Wspace[t_r, s_r]</i> = constant	Mean bias	-1%	+31%	<u>+61%</u>	<u>-52%</u>	<u>+203%</u>	<u>+69%</u>	<u>+61%</u>
	R ²	0.86	0.97	0.94	0.98	0.93	0.90	0.93
PLS map used for spatial spawning	Mean bias	<u>+242%</u>	<u>+400%</u>	<u>+323%</u>	<u>+148%</u>	<u>+312%</u>	<u>+48%</u>	<u>+25%</u>
	R ²	0.88	0.90	0.81	0.91	0.89	<u>0.69</u>	<u>0.79</u>