Movements of grey mullet *Liza aurata* and *Chelon labrosus* associated with coastal fish farms in the western Mediterranean Sea

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**ABSTRACT:** Grey mullet occur in abundance around sea bream and sea bass farms where they forage on waste fish feed, a behaviour that could modify their natural movement pattern and distribution. In this study, we used visual census to record grey mullet aggregations at fish farms in the western Mediterranean Sea. We also mapped the movements of 2 species (*Liza aurata* and *Chelon labrosus*) between farms and adjacent coastal fishing areas, using acoustic telemetry. Grey mullet were frequently observed in the vicinity of the aquaculture cages and represented an important abundance and biomass at the farms. The presence and swimming depth of the tagged mugilids at any of the farms were neither significantly related to the time of the day nor the feeding period, except for *C. labrosus*, which showed a tendency towards deeper waters (~15 m) during feeding periods. Some of the tagged fish stayed in the vicinity of the farms for longer periods and also moved frequently to other farms and nearby commercial fishing areas. Other tagged fish remained at the release location for shorter periods, before they moved out of the study area or possibly were caught by local fishermen. This is the first study using acoustic tagging in wild fish around Mediterranean fish farms that demonstrates that offshore aquaculture farms and local fishing grounds in the western Mediterranean Sea are connected through movements of wild fish. These farms attract and affect large numbers of commercially important fish species, probably causing ecological changes not only in the immediate proximity of farms but also several kilometres away from the farms.

**KEY WORDS:** Aggregation · Aquaculture · Behaviour · Connectivity · Fisheries · Telemetry · Wild fish

INTRODUCTION

A wide range of wild fish species are attracted to coastal aquaculture farms, where they shelter and/or feed on organic matter from cages (faeces and pellets) and organically enriched sediments (e.g. Carss 1990, Dempster et al. 2002, Tuya et al. 2006). The changed feeding habits of aggregated wild fish affect their biological and physiological condition in a way that, for example, their reproductive ability could be affected (Fernandez-Jover et al. 2007, Coz-Rakovac et al. 2008, Arechavala-Lopez et al. in press). Moreover, attracted marine fishes could move frequently and quickly among farms, visiting different facilities in the same area, as has been demonstrated for saithe *Pollachius virens* in Norwegian fjords (Uglem et al. 2009). This behavioural pattern could make wild fish potential — and hitherto not recognised — vectors for transmission of diseases and parasites among farms or to wild fish stocks.

It is believed that the biomass of wild fish in the areas around farms increases due to the fact that the high abundance of waste food increases growth rates (Dempster et al. 2002). Further, it is also possible that fish farms are connected to fishing areas through movements of wild fish. In the Mediterranean, artisanal fisheries exploit wild fish species aggregated at fish farms...
and may therefore benefit from this ‘biomass export’ at a local scale (Machias et al. 2006, Akyol & Ertosluk 2010, Arechavala-Lopez et al. in press). Knowledge about the connectivity of fish farms and other marine areas through wild fish movements is crucial to understand how fish farming and wild fish stocks interact, which, in turn, is essential for managing aquaculture and fisheries in coastal zones.

Grey mullet (Osteichthyes, family Mugilidae) are commonly found in the coastal regions and support artisanal and recreational fisheries in the Mediterranean Sea (McDowall 1988, Blaber 1997, Coll et al. 2004). Grey mullet are also frequently found around sea bream and sea bass farms (Dempster et al. 2002, Fernandez-Jover et al. 2008), where they play an important ecological role, consuming considerable amounts of waste feed from the farms and thus also diminishing the environmental impact of the fish farming activity (Porter et al. 1996, Katz et al. 2002, Lupatsch et al. 2003, Fernandez-Jover et al. 2008). In the present study, we determined the abundance and movement patterns of grey mullet around European sea bass and gilthead sea bream farms on a typical farm area in southeast Spain. Visual census beneath sea cages was used to assess abundances, and individual mullet equipped with acoustic transmitters were tracked in an array of automatic receivers positioned at and around several fish farms. Specifically, we sought to (1) document grey mullet aggregations in close proximity to sea cages, (2) study the movement of mugilids among fish farms and to adjacent coastal fishing areas and (3) assess the presence at and pattern of association to the farms in relation to time of day and feeding time for 2 mugilid species, *Liza aurata* (Risso, 1810) and *Chelon labrosus* (Risso, 1827).

**MATERIALS AND METHODS**

**Study area.** The study was carried out from 13 October 2008 to 15 March 2009 in Guardamar Bay (UTM: 30S 0710736 4219249), Alicante, southeast Spain (Fig. 1). In this bay, 6 fish farm facilities, belonging to 3 different aquaculture companies, grow European sea bass *Dicentrarchus labrax*, gilthead sea bream *Sparus aurata* and meagre *Argyrosomus regius*. The farms are located 3 to 4 km from the shore and the distances between farms vary from 1 to 5 km. The farms are located on soft muddy bottoms at depths ranging from 23 to 30 m. The area supports small-scale traditional and recreational fisheries (Forcada et al. 2009).

**Estimation of abundances of grey mullet around farms.** The abundance and the approximate size distributions of grey mullet around farms in the Guardamar Bay were estimated by conducting rapid visual counts (RVCs; Kingsford & Battershill 1998) to reduce count bias (Ribeiro et al. 2005), following the method out-
lined by Dempster et al. (2002). Fish were counted at 5 different times from October 2008 to March 2009, at 6 randomly selected sites in immediate proximity to the farms F1 and F2 (Fig. 1). Each count covered a volume of approximately 11 250 m³ (15 m wide × 15 m deep × 50 m long) (Dempster et al. 2002) and was conducted simultaneously by 2 divers. The first diver estimated the abundance of the dominant species present. Fish were counted in groups of 1, 2–5, 6–10, 11–30, 31–50, 51–100, 101–200, 201–500 and 500+ to minimise error, based on the method of Harmelin-Vivien et al. (1985). The average total length (TL) of each group was also recorded. The second diver followed slightly behind the first and specifically looked for both highly mobile species and smaller, less obvious fish that may have been missed by the first diver. As grey mullet commonly occur in mixed species shoals, it was difficult to distinguish different mullet species during the visual counts. *Chelon labrosus*, *Liza aurata*, *L. saliens*, *L. ramada* and *Mugil cephalus* were therefore pooled as Mugilidae. At least 1 individual of each of these species was observed during the counts. Biomass conversions from visual census abundance were made on the basis of published length–weight relationship of wild fish (Verdiell-Cubedo et al. 2006), and all raw data were arranged using ecoCEN software (Bayle-Sempere et al. 2001).

**Tagging and tracking of grey mullet.** Movement patterns of mugilids at farms were studied by tagging 14 golden grey mullet *Liza aurata* and 8 thicklip grey mullet *Chelon labrosus* with acoustic transmitters and by monitoring their spatiotemporal distribution using an array of automatic receivers positioned at different farms and adjacent coastal fishing areas in Guardamar Bay (Fig. 1). The mugilids were captured in immediate proximity to farm F1 (distance from farm < 50 m, see Fig. 1) by a baited fish trap (*L. aurata*) or gill nets (*C. labrosus*). Only fish showing no signs of damage after capture were tagged. *L. aurata* were tagged on 15 October 2008 and released on 17 October 2008 after being kept in a sea cage for 2 d. *C. labrosus* were tagged and released on 4 February 2009. Both species were released close (<10 m) to the cages of farm F1. The size of the tagged fish corresponded to the size of mugilids that aggregated at the farms in the study area.

Before tagging, the fish were anaesthetized by immersion in an aqueous solution of MS222 (0.1 to 0.12 g MS222 l⁻¹, anaesthetic volume 40 l, immersion period 3 ± 1 min, temperature in solution: 15 to 17°C). Once anaesthetized, the fish were placed ventral side up onto a V-shaped surgical table. An incision (~1 cm) was made on the ventral surface posterior to the pelvic girdle using a scalpel. The transmitter (Vemco, model V9P-6L-69KHz-S256, 9 × 39 mm, weight in air/water = 4.6/2.2 g, depth range = 200 m) was inserted through the incision and pushed into the body cavity above the pelvic girdle. The transmitters were equipped with a pressure sensor. The incision was closed with 2 or 3 independent silk sutures (3/0 Ethicon). The fish were regularly sprayed with water during the surgery (mean ± SD handling time = 3 ± 1 min, mean recovery time = 2 ± 1 min). Before each incision, the surgical equipment was rinsed in 70% ethanol and allowed to dry. In addition, both species were tagged with external streamer tags near the dorsal fin base (Hallprint) to enable individual identification in case of recapture. Tagged *Liza aurata* were allowed to recover for 2 d in a storage pen before being released by slowly lowering one side of the net wall. After tagging, the *Chelon labrosus* were allowed to recover for 3 to 7 min in a large container (2250 l) on board the fishing vessel. They were released after they showed normal swimming behaviour. All handling and tagging was conducted according to the Spanish regulations for the treatment and welfare of animals (Real Decreto 1201/2005, published in BOE no. 252, 21 October 2005).

The movements and distribution of the tagged grey mullet were recorded by 9 receivers (Vemco, model VR2) positioned to monitor fish farms and coastal fishing areas (Fig. 1). The receivers at the 6 study farms and at the 3 coastal sites were attached on anchored ropes at a water depth of 10 to 12 m. Range tests indicated that the average receiver detection range varied between 300 and 400 m. The transmitters emitted unique coded signals such that each fish could be individually recognized. When a tagged fish was present within a receiver range, the transmitter identification code, date, time and depth of detection were recorded. All receivers were deployed for 23 wk following 13 October 2008.

**Data analyses.** The receivers occasionally recorded acoustic noise that was interpreted as a single reception of a transmitter ID code. To exclude such false signals, single detections within a 30 min period were considered erroneous. Presence within the detection range of a receiver was thus defined as when a fish was detected twice or more within a 30 min period. Likewise, a fish was considered as having left a receiver site if the period between detections was more than 30 min. The 30 min interval was determined on the basis of observations of detection intervals during the recovery period for the 14 *Liza aurata*, i.e. the maximum number of fish that could be expected to be in the detection range of a receiver simultaneously. During this period, the observed average individual detection intervals were approximately 12 to 15 min.

Mortality was determined by analysing variations in swimming depth. When swimming depths corresponded to the depth at the specific receiver location and remained the same until the end of the study pe-
period, the fish was defined as being dead. Movements among receivers were defined as 1-way movements, i.e. if a fish moved from one receiver area to another and then returned, this was recorded as 2 separate movements. As the recovery method varied between the 2 species, data from the first 48 h after release was not used for statistical analyses, to avoid possible tagging effects aimed at comparing the 2 species. Variation in diurnal presence of grey mullet at fish farms was examined by comparing the total daily numbers of detections with the number of detections within 24 × 1 h intervals at any of the 6 farms in the bay for each fish (12 individuals of *Liza aurata* and 6 individuals of *Chelon labrosus*). Only days where a fish had been detected more than twice at a farm were included in the analyses of diurnal presence. Variation in vertical distribution of grey mullet at farms in relation to time of day and farm feeding schedule was examined by comparing the mean swimming depths of each fish (12 individuals of *L. aurata* and 6 individuals of *C. labrosus*) at any of the 6 farms among different times of the day (same data set as was used to examine variation in diurnal presence). To analyse diurnal variation in swimming depth, each day was divided into 2 × 12 h periods with the period from 08:00 to 19:59 h being defined as day (from dawn to dusk) and the remaining period as night (from dusk to dawn). Furthermore, to analyse swimming depth in relation to feeding at farms, each day was divided into a feeding period (09:00 to 14:59 h) and a non-feeding period (00:00 to 08:59 and 15:00 to 23:59 h). Levene’s tests were used to analyse heterogeneity of variances. Generalized linear model (GLM) repeated measurements analyses were used to test for differences in presence and daily detections (dependent variable) between species (fixed factor) and individuals (random factor) among time of the day (24 h, covariate); and for differences in swimming depth (dependent variable) between species (fixed factor) and individuals (random factor) among 2 different periods of a day (day vs. night periods, and feeding vs. non-feeding periods; covariates). The GLM repeated measures procedure provides analysis of variance when the same measurement (detections) is made several times on each subject or case (individuals). All data were analysed using commercially available statistical software (SPSS 15.0).

### RESULTS

Altogether, 12 groups of fishes were observed in the vicinity of the study farms in Guardamar Bay during the visual census (Table 1). Grey mullet (*Mugilidae*) were the most frequently observed fish in the visual counts (found in 63% of the counts) at the farms, followed by bogue *Boops boops* (57%) and horse mackerel *Trachurus mediterraneus* (47%) (Table 1). *Derbio Trachinotus ovatus* and round sardinella *Sardinella aurita* were the most abundant fish species occurring at farms in terms of number of fish (22.4 ± 9.1 and 17.6 ± 7.4 ind. 1000 m–3, respectively), while grey mullet were the third most abundant group of fish at farms (15.5 ± 5.2 ind. 1000 m–3, 19.4% of the total number of fish, Table 1). Furthermore, grey mullet represented 38.9% of the total biomass present at the fish farms (14 537.5 ± 7206.7 kg 1000 m–3), being the second most abundant fish with respect to biomass, after *derbio T. ovatus* (21 381.8 ± 10945.5 kg 1000 m–3). The estimated size distributions of grey mullet captured at the farms are shown in Table 2. Length and weight of observed (visual counts) and tagged/released grey mullet at the fish farms in Guardamar Bay.

<table>
<thead>
<tr>
<th>Taxon</th>
<th>N</th>
<th>Capture type</th>
<th>Length (cm)</th>
<th>Weight (kg)</th>
<th>Release date (d/mo/yr)</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>Liza aurata</em></td>
<td>14</td>
<td>Baited trap</td>
<td>47.9 ± 7.2</td>
<td>1.9 ± 1.0</td>
<td>17/10/08</td>
</tr>
<tr>
<td><em>Chelon labrosus</em></td>
<td>8</td>
<td>Gill net</td>
<td>60.1 ± 4.8</td>
<td>2.4 ± 0.5</td>
<td>04/02/09</td>
</tr>
<tr>
<td><em>Mugilidae</em></td>
<td>4956</td>
<td>Visual census</td>
<td>47.7 ± 10.8</td>
<td>2.0 ± 1.5</td>
<td>N/A</td>
</tr>
</tbody>
</table>
fish farms indicated that the size of the tagged mullet were representative of the mugilids aggregated in immediate proximity to the farms (Table 2).

The tagged *Liza aurata* were detected by all receivers (Fig. 2). During the first 2 wk after release, a relatively high proportion of the *L. aurata* moved from the release farm to other farms and to the area covered by the receivers positioned between the farms and the shoreline (i.e. the coastal fishing area). Two of the 14 tagged *L. aurata* died during the first 2 wk after release. Nine *L. aurata* (64.3%) were not observed after Week 4, possibly because they had left the study area, had been fished or were outside the detection ranges of the receivers (Fig. 2). Three *L. aurata* (21.4%) remained around the fish farms for the entire 22 wk study period (Fig. 2). During this period, these fish made repeated movements among the release farm, the southern farms and the area covered by the receivers positioned between the farms and the shoreline (Fig. 2, Table 3). Four *Chelon labrosus* (50%) were observed in the areas around the release farm the first week after release. These fish were detected during the 6 wk study period (Fig. 2). During the first 4 wk

These fish were detected during the 6 wk study period (Fig. 2). During the first 4 wk

### Table 3. Recorded time and number of movements (mov.) of the 18 tagged grey mullet (12 individuals of *Liza aurata* and 6 individuals of *Chelon labrosus*) that moved from the release farm to other fish farms (F) or coastal fishing areas (C) during the study period. Num. F: number of farms where the fish were detected. Fish defined as being dead were not included.

<table>
<thead>
<tr>
<th>Code</th>
<th>Recorded time (day)</th>
<th>Total mov.</th>
<th>F–F mov. (%)</th>
<th>F–C mov. (%)</th>
<th>F–F mov. (d&lt;sup&gt;–1&lt;/sup&gt;)</th>
<th>F–C mov. (d&lt;sup&gt;–1&lt;/sup&gt;)</th>
<th>Num. F</th>
</tr>
</thead>
<tbody>
<tr>
<td><em>L. aurata</em></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>L1</td>
<td>2</td>
<td>2</td>
<td>0</td>
<td>100</td>
<td>0</td>
<td>1</td>
<td>1</td>
</tr>
<tr>
<td>L2</td>
<td>21</td>
<td>7</td>
<td>100</td>
<td>0</td>
<td>0.33</td>
<td>0</td>
<td>3</td>
</tr>
<tr>
<td>L3</td>
<td>2</td>
<td>2</td>
<td>50</td>
<td>50</td>
<td>0.5</td>
<td>0.5</td>
<td>0</td>
</tr>
<tr>
<td>L4</td>
<td>146</td>
<td>101</td>
<td>100</td>
<td>0</td>
<td>0.69</td>
<td>0</td>
<td>5</td>
</tr>
<tr>
<td>L5</td>
<td>9</td>
<td>42</td>
<td>85.7</td>
<td>14.3</td>
<td>4</td>
<td>0.66</td>
<td>5</td>
</tr>
<tr>
<td>L7</td>
<td>5</td>
<td>7</td>
<td>71.4</td>
<td>28.6</td>
<td>1</td>
<td>0.4</td>
<td>3</td>
</tr>
<tr>
<td>L9</td>
<td>140</td>
<td>5</td>
<td>100</td>
<td>0</td>
<td>0.03</td>
<td>0</td>
<td>3</td>
</tr>
<tr>
<td>L10</td>
<td>5</td>
<td>6</td>
<td>100</td>
<td>0</td>
<td>1.2</td>
<td>0</td>
<td>2</td>
</tr>
<tr>
<td>L11</td>
<td>7</td>
<td>22</td>
<td>100</td>
<td>0</td>
<td>3.14</td>
<td>0</td>
<td>4</td>
</tr>
<tr>
<td>L12</td>
<td>2</td>
<td>2</td>
<td>50</td>
<td>50</td>
<td>0.5</td>
<td>0.5</td>
<td>1</td>
</tr>
<tr>
<td>L13</td>
<td>4</td>
<td>3</td>
<td>66.7</td>
<td>33.3</td>
<td>0.5</td>
<td>0.25</td>
<td>2</td>
</tr>
<tr>
<td>L14</td>
<td>150</td>
<td>152</td>
<td>100</td>
<td>0</td>
<td>1.01</td>
<td>0</td>
<td>4</td>
</tr>
<tr>
<td>Average</td>
<td>41.1</td>
<td>29.3</td>
<td>77</td>
<td>23</td>
<td>1.08</td>
<td>0.28</td>
<td>2.8</td>
</tr>
<tr>
<td><em>C. labrosus</em></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Ch2</td>
<td>20</td>
<td>30</td>
<td>63.3</td>
<td>36.7</td>
<td>0.95</td>
<td>0.55</td>
<td>2</td>
</tr>
<tr>
<td>Ch3</td>
<td>3</td>
<td>1</td>
<td>100</td>
<td>0</td>
<td>0.33</td>
<td>0</td>
<td>1</td>
</tr>
<tr>
<td>Ch4</td>
<td>2</td>
<td>2</td>
<td>50</td>
<td>50</td>
<td>0.5</td>
<td>0.5</td>
<td>0</td>
</tr>
<tr>
<td>Ch5</td>
<td>6</td>
<td>7</td>
<td>57.1</td>
<td>42.9</td>
<td>0.66</td>
<td>0.5</td>
<td>1</td>
</tr>
<tr>
<td>Ch6</td>
<td>30</td>
<td>38</td>
<td>47.4</td>
<td>52.6</td>
<td>0.6</td>
<td>0.66</td>
<td>5</td>
</tr>
<tr>
<td>Ch7</td>
<td>23</td>
<td>37</td>
<td>73</td>
<td>27</td>
<td>1.17</td>
<td>0.43</td>
<td>4</td>
</tr>
<tr>
<td>Average</td>
<td>14</td>
<td>19.2</td>
<td>65.1</td>
<td>34.9</td>
<td>0.70</td>
<td>0.44</td>
<td>2.2</td>
</tr>
</tbody>
</table>
week after release, but never again thereafter in the whole study area (Fig. 2, Table 3). One *C. labrosus* died a few days after tagging (Fig. 2).

The number of movements among farms of the *Liza aurata* and the *Cheilon labrosus* that were detected at farms other than the release farm, ranged from 1 to 152 and 1 to 38 movements, respectively (Table 3). A high proportion of the total movements of *L. aurata* occurred among farms (77%) and the rest between farms and coastal areas (23%). The proportion of the total movements among farms of *C. labrosus* (65.1%) was lower than *L. aurata*, while the movements from farms to coastal areas were higher (34.9%; Table 3). For *L. aurata*, the average numbers of movements among farms and from farms to the coastal area in relation to time (number of days from release to last detection) were 1.08 and 0.28 movements d⁻¹, respectively (Table 3). For *C. labrosus* the average numbers of movements among farms and from farms to the coastal area in relation to time within the study area were 0.70 and 0.44 movements d⁻¹, respectively (Table 3). On average, the *L. aurata* that were detected at farms other than the release farm, were observed at 2.8 farms, while the *C. labrosus* that moved among farms were observed at 2.2 farms (Table 3).

The presence of tagged mugilids at any of the farms was not significantly related to the time of the day (Fig. 3; $F = 0.9$, df = 23, $p = 0.6$) and significant differences were not observed among different individuals throughout the 24 h of a day, regardless of the species or period ($F = 3.4$, df = 17, $p = 0.09$). The daily detection rate of *Liza aurata* appeared to be higher than for *Cheilon labrosus*, but significant differences were not found (Fig. 3; $F = 1.1$, df = 1, $p = 0.3$). However, the detection rate at farms tended to vary among individual *L. aurata* (Fig. 4; $F = 4.278$, df = 11, $p = 0.06$), but not among individual *C. labrosus* (Fig. 4; $F = 3.4$, df = 5, $p = 0.12$).

During daytime, the estimated marginal mean swimming depth of *Liza aurata* was 4.46 ± 1.21 m, while mean swimming depth during nighttime was 4.11 ± 0.86 m (Fig. 5A; $F = 0.3$, df = 11, $p = 0.6$). The estimated marginal mean of swimming depth of *Cheilon labrosus* was 7.44 ± 2.21 m, while swimming depth during nighttime was 5.05 ± 1.59 m (Fig. 5B; $F = 0.5$, df = 5, $p = 0.5$). No overall differences in swimming depth were found between the 2 species ($F = 1.2$, df = 1, $p = 0.3$),
apart from that the swimming depth of the 2 species differed during feeding time ($F = 12.8$, $p = 0.003$). The swimming depths of *L. aurata* did not differ between feeding and non-feeding periods (4.99 ± 1.02 m vs. 3.94 ± 1.02 m; $F = 13.6$, df = 11, $p = 0.09$; Fig. 5C), while the swimming depth of *C. labrosus* tended to be greater during the feeding period compared to the non-feeding period (9.696 ± 2.7 m vs. 4.85 ± 2.08 m; $F = 5.931$, df = 5, $p = 0.06$; Fig. 5D). Furthermore, the individual variation in swimming depth differed significantly within both species between day- and nighttime (*L. aurata*: $F = 18.1$, df = 11, $p < 0.001$; *C. labrosus*: $F = 44.0$, df = 5, $p < 0.001$), as well as between feeding and non-feeding periods (*L. aurata*: $F = 13.6$, df = 11, $p = 0.004$; *C. labrosus*: $F = 11.2$, df = 5, $p = 0.02$).

**DISCUSSION**

In agreement with previous studies (Dempster et al. 2002, Smith et al. 2003, Fernandez-Jover et al. 2008), our results show that grey mullet are abundant around fish farms in the western Mediterranean Sea. In Guardamar Bay, grey mullet represented 19.4% and 38.9% of the total wild fish number and biomass at the farms, respectively. Both *Liza aurata* and *Chelon labrosus* were frequently detected in deeper waters (around 15 m depth) in the vicinity of the farms, occupying the mid-water and cage substratum at the farms. This corresponds to the depths where waste feed from the cages is available. These results support results from other studies regarding attraction of wild fish to farms,

Fig. 5. Estimated marginal means of swimming depth (GLM results) at which *Liza aurata* (n = 12) and *Chelon labrosus* (n = 6) were recorded in the vicinity of the farms during (A,B) day- and nighttime, and (C,D) periods of feeding during daytime.
which suggest that waste feed abundance is an important causal mechanism for attraction of wild fish to farms (Dempster et al. 2005, Tuya et al. 2006, Fernandez-Jover et al. 2008, Uglem et al. 2009). Gut content analyses have shown that waste fish feed is a major part of the diet of wild fish aggregating at fish farms (Fernandez-Jover et al. 2008). However, apart from a tendency towards detecting C. labrosus in deeper waters during the feeding periods, the variation in presence at farms and swimming depth of the tagged mullet was not significantly associated with time of the day or the feeding period at the farms. High variability existed among the behaviour of individual mullet; combined with the small sample size of tracked individuals, this may have masked behavioural patterns in depth distributions related to both time of day or feeding periods.

Our results also show that grey mullet attracted to fish farms may move rapidly and repeatedly both among farms and also to nearby coastal fishing areas. Similar behavioural patterns have been previously detected for other species in a completely different fish farming system (Uglem et al. 2009). Some tagged grey mullet stayed for prolonged periods around fish farms and moved frequently among several of the farms. This might indicate that some of the grey mullet, at least periodically, might exhibit a specialized behavioural pattern where they predominantly stay at or in the proximity of fish farms, most likely to forage on waste fish pellets. The proximate mechanism for attraction of wild fish to farms is unknown. However, the chemosensory system in fish is well developed (Sorensen & Caprio 1998, Vickers 2000), and attraction to water-soluble odorants from food pellets and the large numbers of cultured fish in the cages may be one explanation (Bjørn et al. 2009). Alternatively, aquaculture activity produces significant amounts of noise (Santulli et al. 1999), which also may attract wild fish (Popper et al. 2003). Our results indicate that the presence of Liza aurata at farms tended to be higher than for Chelon labrosus, while the C. labrosus moved more often between farms and coastal areas. This is most likely a result of the varying habitat use of the 2 species, as L. aurata exhibit a more pelagic behaviour compared to C. labrosus, which are more euryhaline and usually frequent estuarine and coastal areas (de Sostoa et al. 1990). The finding that the swimming depths varied within the different species might be related to the fact that grey mullet usually appear in shoals of different sized individuals comprising different species, which often break up and re-form (P. Arechavala-Lopez pers. obs.). The majority of the tagged grey mullet remained within the study area for shorter periods; usually less than a week. These fish most likely moved out of the study area, but they may also have been caught by local fishermen. Nonetheless, no recaptures of tagged fish were reported in Guardamar Bay following the releases.

The findings that large numbers of mullet are attracted to fish farms, most likely to forage on waste fish feed, and that these fish can move quickly and repeatedly among farms or to local fishing areas, may have several implications for integrated coastal management of fisheries and aquaculture. Firstly, the fact that farms and fishing areas are connected through movements of wild fish supports the hypothesis that farm-attracted wild fish may be vectors of diseases or parasites between farmed and wild fish stocks (Uglem et al. 2009). This does, however, assume that farmed and wild stocks share pathogens, which in turn are transferred among these stocks. Sea bream, sea bass and several species of mugilids are similarly susceptible to infection by many different viruses (e.g. family Nodaviridae), bacteria (e.g. Listonella anguillarum, Photobacterium damsela ssp. piscicida, Chlamydia spp., Mycobacterium marinum) and parasites (e.g. Enteromyxum leei, Polysporoplasma sparis, Myxobolus spp., Ceratothoa oestroides, Caligus spp.) (Raynard et al. 2007), but whether transmission is possible between reared fish and farm-associated mugilids is unknown. Compared with other geographical regions, there are few epidemiological studies and little evidence of pathogen exchange between wild and cultured fish in the Mediterranean Sea. Further research is thus required to verify if and to what extent shared diseases are transferred between reared and wild fish stocks in this area.

The movements of the farm-associated grey mullet to adjacent fishing areas also indicate that these fish are available for the local fisheries. If fish farms act as settlement sites for wild juvenile fish, including grey mullet (Fernandez-Jover et al. 2009), and if farm activity increases wild fish biomass and fish condition (Arechavala-Lopez et al. in press), the movement of farm-associated mullet to local fishing areas might be beneficial for local fisheries. In this case, the farms might actually represent small-scale marine protected areas that ‘export biomass’ of exploited species (Dempster et al. 2006). However, attraction of mullet to fish farms might also make them more available for the fisheries by concentrating them in a confined area around fish farms. Indeed, increased fishing pressure around the farms has been noticed in Guardamar Bay, particularly due to the extensive use of gill nets and purse seines close to the farms (Arechavala-Lopez et al. in press). It is also possible that fish farms could disrupt the natural movement patterns and behaviour of grey mullet, and in this way affect their recruitment. Therefore, fish farms could act as artificial reefs that may increase local fish production and in turn also the
export of biomass. However, increased aggregation of fish at farms might also lead to increased fishing effort and thus result in overexploitation of stocks by increasing access to previously unexploited stock segments and/or concentrating previously exploited segments of the stock (Grossman et al. 1997).

In conclusion, the present study supports results from earlier studies in both warm-water and cold-water ecosystems, and indicates that fish farms are connected through wild fish movements and that the spatial distribution of wild fish aggregations at farms is most likely related to specific habitat preferences or feeding requirements for different species. In addition, through the use of acoustic tagging in wild fish around fish farms, this study has demonstrated that sea-cage fish farms and local fishing grounds in the western Mediterranean Sea are connected through the movements of wild fish. Exactly how fish farms in the western Mediterranean affect wild stocks at population level is still unclear and further research is required to reveal the nature of the interaction between fish farms and wild fish stocks. In particular, research aimed at determining how farms affect the reproductive potential and spawning migrations of attracted fish, and how this, in turn, may affect the recruitment and population dynamics of wild fish stocks would be of great significance. Furthermore, our results illustrate that an integrated management of aquaculture and fisheries is imperative in order to ensure a sustainable maintenance and development of both industries according to an ecosystem approach.

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