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Short communication

Size-related variation in fecundity of European eel (Anguilla anguilla)

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Declining European eel (Anguilla anguilla) recruitment has focused attention on conservation of potential spawners leaving continental waters. Fecundity of wild, seaward-migrating silver-phase eels was shown to be size-related and higher than previously reported from artificial maturation experiments. Reliable information on fecundity is essential for stock modelling and future development of eel management policies.

Keywords: Anguilla, fecundity, female size, silver-phase eel, spawner stock management.

Introduction

Since the 1980s, the European eel has undergone a serious population collapse throughout its range, and a variety of causes have been proposed (Stone, 2003). In response, the Council of the European Union (2007) has established a legislative framework (Council Regulation No. 1100/2007) to restore spawner escapement biomass from river basins to levels comparable (at least >40%) to those that occurred when pristine environmental conditions existed. Accordingly, European eel spawning dynamics have become a priority research area. Recent satellite tracking of the oceanic migration route (Aarestrup et al., 2009) and swim trial experiments (e.g. Palstra and van den Thillart, 2010) have contributed to our understanding of the reproductive migration. The artificial completion of the Japanese eel Anguilla japonica lifecycle (Ijiri et al., 2011) is encouraging for European researchers (PRO-EEL, 2011), although a complete understanding of the lifecycle and causes of the collapse of A. anguilla are necessary before artificial propagation will become a viable conservation action.

Current European eel stock recovery plans are almost entirely focused on increasing European eel escapement biomass. However, to determine what proportion of eels successfully migrate and reproduce, information on the health and quality status of potential spawners (e.g. Belpaire *et al.*, 2009; Clevestam *et al.*, 2011) is essential. In particular, knowledge of the reproductive ecology, including fecundity, would enable estimation of the egg numbers required to maintain the standing stock, and could also facilitate future development of eel management policies.

A small number of published fecundity estimates of wild eels exist: for American eel *A. rostrata* (Wenner and Musick, 1974; Barbin and McCleave, 1997; Tremblay, 2009); New Zealand shortfin eel *A. australis*, and longfin eel *A. dieffenbachii* (Todd, 1981); *A. japonica* (Matsui, 1952); and the tropical giant mottled eel *A. marmorata* (Aoyama and Miller, 2003). It appears that European eel fecundity estimates are exclusively of artificiallymatured eels (Kokhnenko *et al.*, 1977; Boetius and Boetius, 1980; van Ginneken *et al.*, 2005). Therefore, the aim of the present study was to estimate the fecundity of wild silver-phase European eels, captured undergoing their seaward spawning migration, and to relate this to body size.

Material and methods Sampling area and collection

The Shannon International River Basin District, defined according to the Water Framework Directive (European Commission, 2010), occupies an area of 18 000 km². The River Shannon predominantly drains the central lowlands of Ireland and discharges through a 97 km estuary into the North Atlantic (Figure 1). It is regulated

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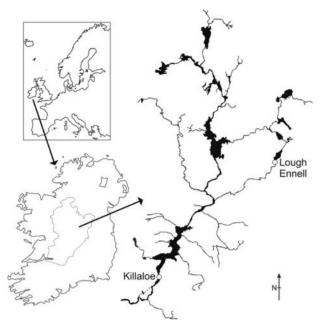


Figure 1. Silver eel sampling locations (Killaloe eel weir and Lough Ennell outlet) in the River Shannon catchment, Ireland.

for hydroelectric generation, and the mean annual discharge is $186 \text{ m}^3 \text{ s}^{-1}$ (McCarthy and Cullen, 2000). During the 2007– 2008 migration season, 25 silver-phase eels were randomly subsampled from the catch of a commercial fishing crew operating a winged stow net at the outlet of Lough Ennell (53°27'N 7°23'W). This 14.3 km² mesotrophic lowland lake in the upper Shannon catchment forms part of the River Brosna tributary. The limited size range of Lough Ennell eels (84% were 630-750 mm) precluded analysis of the complete River Shannon female size range (McCarthy and Cullen, 2000) from this location. Therefore, during the 2008-2009 migration season, supplementary silver-phase eels were obtained at Killaloe eel weir (52°48'N 8°27'W) on the lower River Shannon. Thirteen eels were selected from the catch, to represent the entire River Shannon female size range. The fishing gears at both locations captured all sizes of female silver-phase eels (McCarthy and Cullen, 2000; Tesch, 2003).

Treatment and analysis

Eels were sacrificed by immersion in a solution of clove oil and ethanol in water. The body length (to the nearest 1 mm) and body weight $(\pm 1 \text{ g})$ of each eel was recorded. Horizontal and vertical eye diameters were measured to the nearest 0.1 mm for Killaloe eels only, and eye index was calculated as:

{[(horizontal eye diameter + vertical eye diamater)/4]²
$$\times \pi$$
/totallength} $\times 100$

(Pankhurst, 1982). Sex was determined by macroscopic examination of the gonads (Tesch, 2003) and all eels were confirmed to be females. Both ovaries were removed from the body cavity and weighed to the nearest 0.01 g. Gonadosomatic index (GSI) was calculated as:

$$(gonad weight/body weight) \times 100$$

(Durif et al., 2005). Eels were classified as silver-phase by external appearance, eye index >6.5 (Pankhurst, 1982; Aoyama and Miller, 2003) (Killaloe eels only) and GSI >1.2% (Durif et al., 2005) (all eels). Treatment with 250 ml 2% acetic acid was carried out on fresh ovaries following the protocol described by Barbin and McCleave (1997). Each solution was agitated daily, and all eggs/ ovarian tissue were separated within 7 days. The solutions were then diluted using distilled water. Most (76.3%) eels were diluted to 2 l, but the larger eels were diluted to 6-10 l (Barbin and McCleave, 1997). Egg counts were made on 1 ml volumetric subsamples examined at ×40 magnification. Four subsamples were counted and an estimate of fecundity was calculated by reference to the mean egg count and the dilution factor (Barbin and McCleave, 1997; Tremblay, 2009). Body length, body weight, gonad weight, and number of eggs (fecundity) were log₁₀transformed to meet the requirements of parametric analysis (i.e. normality and equality of variances). Pearson correlation coefficients (r) were calculated for the relationships between fecunditylength, fecundity-body weight and fecundity-gonad weight. Simple linear regression analysis of length on body weight and fecundity on length were undertaken following the form:

$$\log Y = a + b \log X$$

Differences between the intercept and slope of the Killaloe and Lough Ennell fecundity–length regression equations were tested using the General Linear Test Method (Neter *et al.*, 1996).

Results

A within-river comparison of the fecundity-size relationship showed no difference between sampling location [General Linear Test: F = 0.313; *d.f.* (degrees of freedom) = 2, 34; p = 0.73]. Therefore, all data were pooled and analysed as a single River Shannon sample (n = 38). The length-body weight relationship is given by the equation:

$$log_{10} length = 1.991 \ (\pm 0.033) + 0.302 \ (\pm 0.012)$$
$$\times log_{10} lody weight \ (r^2 = 0.949, p < 0.001)$$

Values in parentheses are standard error of mean (S.E.M.). Fecundity was positively correlated with length (r = 0.943; p < 0.001), body weight (r = 0.955; p < 0.001) and gonad weight(r = 0.936; p < 0.001), and increased exponentially with length according to the following regression equation:

$$log_{10} \text{ fecundity} = -2.992 (\pm 0.550) + 3.293 (\pm 0.193)$$
$$\times log_{10} \text{ length } (r^2 = 0.889; p < 0.001)$$

The log₁₀-transformed fecundity–length regression (and associated 95% confidence intervals) is illustrated (Figure 2). Fecundity estimates for the eels examined ranged from 626 000 to 8 006 667 for individuals of 465 mm (211 g) to 1003 mm (2472 g). Based on the fecundity–length regression equation, these eels would have estimated fecundities of 619 331 to 7 785 455. The relative fecundity (eggs/kg) was estimated to be 3 591 699. The morphological characteristics and fecundity of the silver-phase eels examined (by location and pooled data) are presented in Table 1.

Discussion

Fecundity in the temperate eels *A. rostrata* (Wenner and Musick, 1974; Barbin and McCleave, 1997; Tremblay, 2009), *A. australis* and *A. dieffenbachii* (Todd, 1981) has been shown to increase exponentially with increasing body size (Table 2). In the present study, which provides the first fecundity estimates of wild *A. anguilla*, fecundity was also shown to be size-related in this species. No difference in fecundity–size relationship was observed between the upper and lower River Shannon sampling locations. Tremblay (2009) did find differences in fecundity between five subpopulations of *A. rostrata* in a large North American catchment

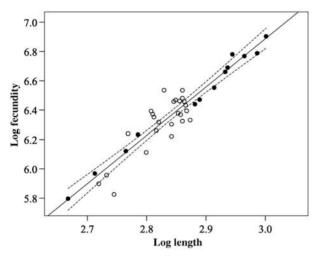


Figure 2. Log_{10} -transformed fecundity–length regression of pooled River Shannon (Killaloe; closed circles, Lough Ennell; open circles) silver-phase eels, with 95% confidence intervals.

(Saint Lawrence River), but concluded that this was not related to migration distance. However, considerable variation in *A. rostrata* fecundity estimates from Chesapeake ($37^{\circ}N$) (Wenner and Musick, 1974), Maine ($45^{\circ}N$) (Barbin and McCleave, 1997) and the Saint Lawrence ($44-49^{\circ}N$) (Tremblay, 2009) suggest differences may exist on a larger spatial (or temporal) scale (Table 2). No such data is available for *A. anguilla* at present, but possible geographical variation in reproductive potential, reflecting energy requirements and migration distance, has been suggested by Belpaire *et al.* (2009).

Factors other than spatial variation may also affect fecundity. The accumulation of lipophilic compounds in gonads may reduce egg production and development (e.g. Belpaire et al., 2009). The water quality of the River Shannon is mainly classified as unpolluted (Lucey, 2009) and in general, contaminant levels in Irish eels are low compared to other European countries (McHugh et al., 2010). Habitat use and migratory type may also affect fecundity, as observed in relation to other biological characteristics (e.g. Svedäng et al., 1996; Arai et al., 2006). Otolith microchemical analysis of upper and lower River Shannon eels indicates that the eel populations are composed almost entirely of freshwater residents (Arai et al., 2006). It seems that further eel fecundity studies from a range of locations may be necessary, particularly when significant differences in biological characteristics can occur even at the scale of neighbouring catchments (Acou et al., 2009). Furthermore, silver-phase eel quality and maturation status within a river system may also change during the downstream migration season, and this should be incorporated in a more comprehensive sampling programme.

Differences between wild and artificially-matured *A. anguilla* fecundity estimates (Table 2) may be due to the methodology used, or may reflect changes in the gonads accompanying the

Table 1. Summary of the morphological characteristics and fecundity of the silver-phase eels examined.							
L,	Ennell ($n = 25$)	Killaloe ($n = 13$)	Pooled $(n = 38)$				

L. Ennell $(n = 25)$		Killaloe $(n = 13)$		Pooled $(n = 38)$				
Mean	<u>+</u> S.E.	Min – max	Mean	<u>+</u> S.E.	Min – max	Mean	<u>+</u> S.E.	Min- max
673	13	524-747	771	49	465 - 1003	707	20	465 - 1003
587	31	267 - 798	1121	188	211-2472	770	78	211-2472
1.75	0.06	1.24 - 2.57	1.66	0.05	1.44 – 1.99	1.72	0.04	1.24 - 2.57
			7.13	0.30	6.54 - 10.57			
2.23	0.15	0.67 - 3.43	3.80	0.65	0.63 - 8.01	2.76	0.27	0.63 - 8.01
	Mean 673 587 1.75	Mean ± S.E. 673 13 587 31 1.75 0.06	Mean ± S.E. Min-max 673 13 524-747 587 31 267-798 1.75 0.06 1.24-2.57	Mean ± S.E. Min-max Mean 673 13 524-747 771 587 31 267-798 1121 1.75 0.06 1.24-2.57 1.66 7.13 7.13 7.13	Mean \pm S.E. Min-max Mean \pm S.E. 673 13 524-747 771 49 587 31 267-798 1121 188 1.75 0.06 1.24-2.57 1.66 0.05 7.13 0.30	Mean \pm S.E.Min-maxMean \pm S.E.Min-max67313524-74777149465-100358731267-7981121188211-24721.750.061.24-2.571.660.051.44-1.997.130.306.54-10.57	Mean \pm S.E.Min-maxMean \pm S.E.Min-maxMean67313524-74777149465-100370758731267-7981121188211-24727701.750.061.24-2.571.660.051.44-1.991.727.130.306.54-10.57700	Mean \pm S.E.Min-maxMean \pm S.E.Min-maxMean \pm S.E.67313524-74777149465-10037072058731267-7981121188211-2472770781.750.061.24-2.571.660.051.44-1.991.720.047.130.306.54-10.571.651.651.651.65

 Table 2.
 Summary of anguillid fecundity studies.

Species	Study	Size range	Min–max fecundity (millions of eggs)	Relative fecundity (millions of eggs/kg)
A. anguilla	Kokhnenko <i>et al</i> . (1977) [§]	N/a	N/a	3.0
	Boetius and Boetius (1980) [§]	640–920 mm	0.7 – 2.6	1.6
	van Ginneken <i>et al.</i> (2005) [§]	690–870 mm	0.8-4.0	1.8
	This study	465 – 1003 mm	0.6-8.0	3.6
A. rostrata	Wenner and Musick (1974)	490–724 mm	0.5 - 2.6	3.8 [†]
	Barbin and McCleave (1997)	452 – 1133 mm	1.7 – 20.7	8.1 [†]
	Tremblay (2009)	532 – 1159 mm	3.4-22.0	6.5 – 10.0
A. australis	Todd (1981)	516–933 mm	0.5 - 3.1	1.9
A. dieffenbachii	Todd (1981)	711 – 1452 mm	1.1 – 20.8	2.0
A. japonica	Matsui (1952)	357–924 mm	7.2 – 12.7	N/a
A. marmorata	Aoyama and Miller (2003)	2400 g	34.8	N/a

[§]Eels were artificially-matured.

[†]Estimated from the regression equation for a 1 kg eel.

maturation process. The treatment of ovaries with 2% acetic acid and subsequent volumetric subsampling has successfully been used in previous anguillid fecundity studies (Barbin and McCleave, 1997; Tremblay, 2009). However, A. anguilla fecundity estimates obtained by Boetius and Boetius (1980) by counting eggs retained by a 0.224 mm mesh, were underestimated, as only eggs which had responded to hormonal treatment were counted, and the relative fecundity (eggs/kg) reported (1.6 million) is less than half that obtained in the present study (3.6 million). Likewise, the relative fecundity of artificially-matured A. anguilla by van Ginneken *et al.* (2005) is considerably lower (1.8 million) than in the present study, possibly reflecting the counting method (gravimetric subsampling) or an effect of the artificial maturation process. Russian maturation experiments conducted in the 1970s quote a relative fecundity of 3 million (Kokhnenko et al., 1977), although no details of counting method are described.

The application of fecundity estimates derived from artificial maturation experiments to wild eel populations may not be appropriate, given the issues discussed above. Similarly, fecundity estimates derived from wild eels captured in continental waters (i.e. early vitellogenic stage) may differ from the actual fecundity at the spawning grounds. Ideally, eels in advanced spawning condition (mid-/late-vitellogenic stage) should be examined. However, to date only female eels of A. japonica and A. marmorata have been captured at their spawning grounds (Iriji et al., 2011). Analysis of changes in the number of eggs in the gonads during the maturation process may be possible using artificially-matured eels (e.g. Durif et al., 2006), although the extent to which hormonal treatment reflects the natural maturation of eels at sea will need to be verified. The limited reproductive success of artificial maturation experiments with A. anguilla (Pedersen, 2004; Palstra et al., 2005) suggests that hormonal treatment results in certain artifactual outcomes (e.g. poor egg quality, delayed hatching and abnormal morphology), which may bias fecundity estimates.

Different growth strategies have been proposed for female eels during the continental phase of the lifecycle i.e. size-maximizing, with its associated higher pre-reproductive mortality rates, to achieve maximum fecundity (e.g. Davey and Jellyman, 2005) or time-minimizing (Svedäng et al., 1996). Laboratory observations of the spawning behaviour of artificially-matured eels suggest batch spawning by females (Boetius and Boetius, 1980), and that a single male may be capable of fertilizing several egg batches (van Ginneken et al., 2005). If this is typical of natural spawning events, female eels could be considered of greater reproductive value than males. Therefore, stock recovery plans should prioritize the protection of large, highly fecund, female eels. Clevestam et al. (2011) proposed similar protection measures for large females in the Baltic Sea, as their large size and high lipid content would make them most likely to successfully migrate and spawn (Belpaire et al. 2009; Palstra and van den Thillart, 2010).

Modelling of *A. anguilla* population dynamics has been attempted using fisheries data (Dekker, 2000), demographics (van der Meer *et al.*, 2011), and integrated genetic and demographic models (Pujolar *et al.*, 2010; Andrello *et al.*, 2011). Such models often involve numerous input variables, including fecundity. However, the use of differing fecundity estimates [e.g. Boetius and Boetius (1980) in Andrello *et al.* (2011); Barbin and McCleave (1997) in Pujolar *et al.* (2010)] has a knock-on effect on subsequent calculations, and it would seem that neither estimate is appropriate, as respectively, they relate to artificially-matured eels (underestimate) and *A. rostrata* (overestimate) (see Table 2). Van der Meer *et al.* (2011) reject both of these estimates and instead use a notional figure of 2 million eggs for a 560 g eel, which is very similar to our presented data (1.98 million).

Andrello *et al.* (2011) divided the continental eel stock into three production units (North, north of 50°N; Atlantic, between 35°N and 50°N; Mediterranean, south of 35°N), with a sex ratio of 0.34 females in the breeding stock and mean female silver-phase eel sizes of 663 mm, 664 mm and 572 mm in each unit respectively. Applying this scenario to Dekker's (2000) estimate of silverphase eel escapement (8.8 million), we calculate an *A. anguilla* population fecundity of 5.21×10^{12} million eggs based on the mean size of eels in each production unit, and the fecundity– length relationship presented in Figure 2

The currently available models do not take into account the complexities associated with, for example, the impact of pollution on gonadal development and egg production (e.g. Belpaire *et al.*, 2009), or the possibility of multiple spawning events (Boetius and Boetius, 1980; Ijiri *et al.*, 2011). However, as illustrated by the present study, the need for reliable knowledge of eel fecundity and other population parameters is important for population modelling and spawner stock management. Integration of data on various aspects of European eel biology (fecundity, mating dynamics, larval survival, recruitment, escapement, spawning migration, demographics etc.) may enable estimation of total spawner numbers required in the Sargasso Sea to maintain the standing stock. If possible, this would represent a major step in the conservation of this endangered species.

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