

A stage-structured model to predict the effect of temperature and salinity on glass eel *Anguilla anguilla* pigmentation development

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(Received 2 August 2004, Accepted 7 April 2005)

The pigmentation development process of glass eels *Anguilla anguilla* from stage V_B to VI_{A3} was modelled by gamma cumulative functions. These functions varied with respect to the factors temperature and salinity whose effects were adjusted by beta functions. Temperature was shown to accelerate pigmentation, while salinity acted as a secondary factor slowing down the pigmentation. The model fits the development of 15 samples kept at various temperatures and salinities in the Vilaine River, as well as samples monitored at other dates and places in Europe. It allows the prediction of the duration of estuarine residency for glass eels, in winter and spring, in the Atlantic estuaries.

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Key words: *Anguilla anguilla*; European eel; glass eel; pigment stage; salinity; temperature.

INTRODUCTION

Metamorphosis from the larval to the juvenile stage in fishes involves important morphological, ecological and physiological changes, more evident in those teleosts that show pelagic larvae *v.* benthic adults. Such is the case of the anguillids, whose larva is the leptocephalus, a distinctive long-lived larval stage well adapted to the pelagic life in the ocean (Hulet, 1978; Hulet & Robins, 1989), that shows a unique developmental strategy among teleosts, with regards to its metabolic rate, growth and osmoregulation (Bishop *et al.*, 2000; Bishop & Torres, 2001).

During metamorphosis, leptocephali decrease in size and wet mass as the internal gelatinous matrix is degraded and replaced by developing muscles and bones (Kawakami *et al.*, 1999; Pfeiler, 1999). The willow leaf-like shaped larvae begin to acquire the eel like shape and most structures such as the gut, skin, gill and kidney become more developed during this transition from the larval stage to the adult-like glass eel stage (Hulet & Robins, 1989).

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The progressive pigmentation of skin and internal tissues is also concentrated during metamorphosis from fully-grown larvae to the juvenile phase (Hulet & Robins, 1989; Tsukamoto, 1990) and the larvae progressively lose their typical fully transparent bodies (Schmidt, 1909; Robins, 1989; Smith, 1989) with only scattered melanophores (Castle, 1997). In the genus *Anguilla*, this most visible marker has, therefore, supported the stage classification of the metamorphosis process from leptocephalus (stages I–IV) to the glass eel (stages V and VI), corresponding to the reaching of the eel-like body form (Tesch, 2003). The glass eel to elver phase covers stages V_A with no pigmentation except a spot on the caudal fin, V_B with the early development of the pigmentation on the skull and VI_{A0} with development of pigmentation along the base of dorsal fin, and with the first pigment cells beyond the skull spot (Bertin, 1951; Elie *et al.*, 1982). The next stages VI_{A1}, VI_{A2}, VI_{A3}, VI_{A4} and VI_B (Strubberg, 1913; Bertin, 1951; Elie, 1979; Elie *et al.*, 1982) are based on development of surface and branchiostegal pigmentation.

Together with these morphological changes, catadromous eels of the genus *Anguilla* perform a remarkable migration from their spawning areas in the ocean drifting to continental areas, where they recruit to freshwater habitats. In particular, *Anguilla anguilla* (L.) are thought to spawn panmictically in subtropical waters of the Atlantic Ocean, in the Sargasso Sea with their larval migration route from the centre of the spawning ground to the European and North African coasts extending for >6000 km.

It is thought today that metamorphosis onset occurs between stage I and II, the first morphological modifications (Cieri, 1999) coinciding with the moment when leptocephali reach the continental slope (Tesch, 1980), and its end occurs at stage VI_B when glass eels have achieved their estuarine migration and juvenile growth begins (Lecomte-Finiger, 1983; Jegstrup & Rosenkilde, 2003).

An important contribution to the understanding of the larval history of glass eels has been given by the interpretation of otolith microstructure, that provides a precise description of the migration timing from the ocean to the rivers (Tzeng & Tsai, 1994; Martin, 1995; Arai *et al.*, 1997; De Casamajor *et al.*, 2003), as well as an ecological interpretation of the transition between sea and estuary in the course of the ascent season (De Casamajor *et al.*, 2001a). Recruitment, besides occurring in a changing environment such as the estuary, corresponds to several behavioural changes, that include a shift from pelagic to benthic habits and a shift from vertical movements related to the use of selective tidal stream transport (McCleave & Wippelhauser, 1987) to an active counter current migration (Creutzberg, 1958). At the anatomical level, this transition brings about morphological and physiological modifications, that occur simultaneously with the development of organs associated with the achievement of metamorphosis (Hulet & Robins, 1989).

Osmoregulation compensative mechanisms occurs as consequence of the transition from a hypertonic to a hypotonic medium (Ciccotti *et al.*, 1993; Birrell *et al.*, 2000). Swimbladder activation (Hickman, 1981) is related to the change in the migration mechanism. Overall completion of metamorphosis includes pigmentation (Panu, 1929a; Pankhurst, 1983), which is the only visually obvious process allowing the discrete description of a continuous irreversible change. Therefore, if at the individual level it provides an indication of the amount of

time spent since the pigment process was triggered, at the population level the stage classification has often supported many geographical and seasonal descriptions of the coastal and estuarine recruitment process, for several species of the genus *Anguilla* (Elie, 1979; Cantrelle, 1981; Charlon & Blanc, 1982; Haro & Krueger, 1988; Boëtius & Boëtius, 1989; De Casamajor, 1998; Jellyman *et al.*, 1999).

The spatial or temporal comparisons of samples and the ecological interpretation of the time and space migration dynamics as inferred from pigment stages composition, however, are not straightforward. In fact, several uncertainties such as the individual variability of the metamorphosis, the variation in stage duration and most of all, the influence of environmental factors on the pigmentation process limits the ecological interpretations derived from pigment stages (Strubberg, 1913; Lecomte-Finiger, 1983; Tsukamoto, 1990; Ciccotti *et al.*, 1993; Dou *et al.*, 2003).

Recently, both salinity and temperature, key variables in estuaries during glass eels ascent, were confirmed to be factors influencing the development of pigmentation, and their role was described, while the experimental handling process was shown not to influence the pigmentation process (Briand *et al.*, 2004). Based on these results and some additional experiments, the objectives of this study were to: (1) propose a model describing the evolution of pigment stages with reference to temperature and salinity, (2) test the predictive model for glass eel pigment stages in other experiments and (3) discuss the applications for this model, in the general objective of providing a tool for the understanding and description of spatial and temporal dynamics of glass eels recruitment in estuarine systems.

MATERIALS AND METHODS

EXPERIMENTAL PLAN

Fifteen samples of glass eels were collected using commercial fishing boats in the Vilaine Estuary, during and after the commercial fishing season in 2000 and 2001 (see Table I for sample collection dates). Two glass eel samples were maintained *in situ* in the estuary, in fresh water upstream of the Arzal Dam (F_1 to F_6) and downstream of the dam (E_2 to E_6) respectively, in cages of 2 mm mesh-size held 1 m from the bottom with a rope (Briand *et al.*, 2004). Four samples (R_1 to R_4), collected in 2001, were maintained in laboratory conditions, *i.e.* kept in the refrigerator in the dark. These additional experiments were necessary to force the model to express positive values at low temperatures and thus describe conditions encountered in French estuaries in January and February. Temperature was recorded every 6 min with probes in each site, and varied from 4.8 to 19.1° C in fresh water, 5.4 to 20.1° C in the estuary, and from 0.4 to 4.9° C in the laboratory. The salinity was recorded close to the cages three times a week. Salinity was zero in fresh water and varied cyclically according to the discharge and tide from 0.1 to 29.2 in the estuary. The salinity was kept constant in the laboratory from 0 to 26.6 (Table I).

SAMPLE EXAMINATION

The samples were examined three times a week for pigment developmental stages. All glass eels were retrieved from each lot and anaesthetized with phenoxy ethanol at a concentration of 1‰. About 70 glass eels were collected randomly from each lot, and

TABLE I. List of samples, experimental conditions and daily mortalities

Sample	Collection date	Duration (days)	Mean salinity (range)	Mean temperature (°C)	Number of daily analysis	Daily mortality (% day ⁻¹)
Fresh water						
F ₁	28 January 2000	45	0	8.3	20	1.2 ^a
F ₂	2 February 2000	40	0	8.7	16	
F ₃	17 March 2000	36	0	10.3	14	1.5
F ₄	8 April 2000	30	0	12.3	12	1.6
F ₅	27 April 2000	21	0	16.4	9	1.1
F ₆	10 May 2000	5	0	18.1	3	0.3
Estuary						
E ₂	2 February 2000	65	7.4 (0.2–29.2)	9.7	29	1.0
E ₃	17 March 2000	36	11.7 (0.1–29.2)	10.7	16	1.3
E ₄	8 April 2000	30	8.8 (0.1–27)	12.3	14	
E ₅	27 April 2000	21	2.5 (0.1–9.5)	15.9	10	3.1
E ₆	10 May 2000	7	0.1 (0.1–0.1)	18.2	4	4.3
Refrigerator						
R ₁	26 January 2001	121	6.4	3.9	18	0.5
R ₂	2 February 2001	57	6.8	2.1	10	1.5
R ₃	27 April 2001	16	0	1.7	3	0.1
R ₄	27 April 2001	34	26.6	2.1	4	1.9
Model testing						
V ₁	3 April 1912 ^b	84	16	18	3	
V ₂	3 April 1912 ^b	133	16	12	3	
V ₃	15 December 1993 ^c	64	36	16.5	7	
V ₄	15 December 1993 ^c	64	0	16.5	7	

^aEscapee from the cages.^bStrubberg (1913).^cE. Cicotti (unpubl. data).

reintroduced after pigmentation examination. Epidermal pigmentation was assessed according to the sequence defined by Elie *et al.* (1982). This classification, that defines seven stages, was chosen in relation to the fact that it reviews preceding classifications, in particular Strubberg (1913) that considers six stages. The experiments were progressively stopped when stages VI_{A2} and VI_{A3} represented >80% of the proportion of all the stages. The pigment stages observed during the study were V_B to VI_{A3}. As the stages VI_{A4} were rare (<1%) they were included in the VI_{A3} counts.

Mortality was evaluated at the end of the trial for each experimental batch. Daily mortality was calculated as the total mortality during the experiment, divided by the duration of the experiment.

MODELLING

The modelling of pigmentation process was based on stage transition models. The series of pigment stages obtained for each sample was transformed to calculate the cumulative percentage of glass eels having reached a particular stage over time. The progressive ageing was then simulated for VI_{A0 cum} to VI_{A3 cum} by a gamma cumulated distribution function (G) whose scale parameter was set to 1 for all cumulated stages (Manly, 1990). Four parameters of shape (p_1, \dots, p_4) were thus fitted: VI_{A0 cum} = G($\phi; p_1, 1$), VI_{A1 cum} = G($\phi; p_2, 1$), VI_{A2 cum} = G($\phi; p_3, 1$) and VI_{A3 cum} = G($\phi; p_4, 1$). The duration of one pigment stage was therefore considered as a random variable following a gamma function.

The variable ϕ , 'pigmentation time', was calculated as the cumulative sum from the beginning of the experiment of the days corrected with the product of an efficient daily temperature and an efficient daily salinity. The practical meaning of this formulation can be explained as follows: daily salinities and temperatures fully relevant for the pigmentation process would correspond to efficient daily temperatures and efficient daily salinities equal to 1 and each day would contribute for 1 to the pigmentation time. When salinity or temperature conditions are less favourable, the efficient daily salinity or the efficient daily temperature are lower than 1 and one day counts for <1 for the pigmentation time. The choice of a product expresses the interacting effect of each factor on the pigmentation process. The physiological optimum for growth is found at 22–23° C for yellow eels (Sadler, 1979) and glass eels (Elie & Daguzan, 1976). This optimum in growth could also apply for other morphological development such as pigmentation. As the range of temperatures explored in the experiments (0.5–20° C) was below this optimum, beta cumulative distribution curves were chosen for both temperature and salinity to allow the expression of a wide variety of asymmetrical sigmoid responses. Limits of this beta curve were set out of the range of experimental temperatures and salinities, to ensure that the lack of flexibility of beta curves near their boundaries did not bias the model. Therefore, the 'efficient daily temperature' T_{eff}($\theta_i; p_5, p_6$) was calculated from daily temperatures with a beta cumulative distribution function of boundaries 0 and 30. The efficient daily salinity S_{eff}($\sigma_i; p_7, p_8$) was calculated from average salinity with a (1-beta) cumulative distribution function of boundaries of 5 and 40‰.

As pigment stages differed between samples at the beginning of the experiment, the pigmentation time of each sample was corrected by adding an initial pigmentation time ϕ_0 , that was calculated as the mean of inverse gamma functions for stages VI_{A0 cum} to VI_{A3 cum} calculated with parameters (p_1 to p_4) from the initial pigment stages composition of the sample.

Model fitting was performed by minimization of the square distance between experimental and modelled data (function lsqcurvefit of Matlab). The performance of the whole model, and each sample, was determined by R^2 and mean absolute errors.

MODEL VALIDATION

The model prediction was then tested against other laboratory experiments, carried out by other authors and selected because they had detailed records of salinities and

temperatures, whose range was within the range tested in the model. Experiments V_1 and V_2 were drawn from Strubberg's (1913) experiment. He kept samples of glass eels in aquaria in the laboratory and a cellar and provided detailed account of temperature, salinity and evolution of pigment stages as summarized in Table 1. As temperatures were not provided from 83 to 133 days for the laboratory experiment, a 100% VI_{A3} , the final composition of the sample, was considered as already reached after 83 days.

The model was based solely on VI_{A1} and VI_{A3} cumulated data, because the stage VI_{A1} , proposed by Elie *et al.* (1982), groups the VI_{A1} , VI_{A2-1} and VI_{A2-2} pigment stages described by Strubberg (1913).

Samples V_3 and V_4 were based on a laboratory trial carried out with glass eels collected during their ascent in December 1994 at the mouth of the River Tiber (Central Italy) and kept for 60 days at controlled temperatures 16.5 and 21° C, and at salinities 0 and 35 (E. Ciccotti, unpubl. data). Samples were examined for pigmentation at 3 and 7 days from acclimation and then at 14 days intervals up to the end of the experiment, following anaesthesia. In this trial, Boëtius's (1976) classification of five stages (A–E) was used, therefore data were converted to Elie *et al.*'s (1982) classification as following $A = V_B$, $B = VI_{A0} + VI_{A1}$, $C = VI_{A2} + VI_{A3}$ and $D = VI_{A4}$.

STAGE DURATION DERIVED FROM THE MODEL

In stage transition models, the use of a gamma distribution implies that the mean time of entry to a stage is equal to the parameter p value (Manly, 1990). As in the present model, this mean time of entry was not related to a duration but rather to the 'pigmentation time', the duration of each pigment stage was converted into days from the ratio between p values (φ) for each stage and daily pigmentation time ($\varphi \text{ day}^{-1}$) calculated from constant values of temperature and salinity.

RESULTS

IN SITU EXPERIMENTS

In fresh water and the estuary, the pigment stages evolved from 83% (range 49–99%) of glass eel stages V_B and VI_{A0} at the beginning, to 0.5% (range 0–2.9%) at the end (Table II). The duration of the experiments shortened from 40 to 65 days for the samples collected in January and February to 21–36 days for those collected in March and April and was finally reduced to 5 to 7 days for the last experiments in May. This reduction was the consequence of the shift in initial stage composition (χ^2 , d.f. = 20, $P < 0.001$), with more advanced stages at the end of the season (Table II). It was also correlated with the increase in water temperature from 8.3 to 18.2° C ($r = -0.86$, $P < 0.001$).

LABORATORY EXPERIMENTS

In the laboratory samples, the pigment stages evolved from 52% (range 14–90%) of V_B – VI_{A0} at the beginning of the experiments, to 28% (range 4–81%) at the end. Due to high mortalities, the experiments started in January and February were ended after 121 and 57 days, before stages VI_{A2} and VI_{A3} reached 80% of the total sample. The last two samples had shorter durations and were

TABLE II. Initial and final pigment stages detailed distributions for *in situ* experiments used in the model (see Table I for samples)

Sample		Stages %*				
		V _B	VI _{A0}	VI _{A1}	VI _{A2}	VI _{A3}
Fresh water						
F1	Start	34	64	0	2	0
	End	0	1	9	66	24
F2	Start	40	49	10	1	0
	End	0	0	6	63	31
F3	Start	17	72	8	3	0
	End	0	0	0	33	67
F4	Start	21	74	2	1	2
	End	0	1	5	31	63
F5	Start	11	73	12	3	1
	End	0	0	4	37	59
F6	Start	5	44	11	31	9
	End	0	0	8	46	46
Estuary						
E2	Start	40	49	10	1	0
	End	0	0	6	27	67
E3	Start	24	64	8	4	0
	End	0	0	7	36	57
E4	Start	13	82	2	1	2
	End	0	3	9	29	59
E5	Start	2	82	12	3	1
	End	0	0	3	35	62
E6	Start	0	49	11	31	9
	End	0	0	8	50	42
Refrigerator						
R1	Start	30	59	10	1	0
	End	2	15	46	37	0
R2	Start	81	9	9	1	0
	End	50	31	19	0	0
R3	Start	7	7	27	27	32
	End	3	0	35	44	18
R4	Start	7	7	27	27	32
	End	0	9	24	50	17

*Stages correspond to the pigmentation sequence defined by Elie *et al.* (1982).

composed of >90% of VI_{A1}–VI_{A3} stages in 16 days in fresh water and 34 days in the estuary (Tables I and II).

MORTALITY

The samples comprised 100 to 279 glass eels at the beginning of the experiment, and 32 to 159 (mean = 70) at the end. The daily mortality varied from 0.3 to 4.3% day⁻¹ in the cages and 0.1 to 1.9% day⁻¹ in the refrigerator. The largest mortalities occurred in estuary during the flood period and concerned the samples E₅ and E₆ (Table I).

MODELLING

A common model for all experiments provided a good fit of 728 points corresponding to the results of 11 678 glass eels distributed in 15 experiments (Fig. 1). Splitting into locations, the model was adjusted on 296, 292 and 140 points, corresponding to the repartition in four cumulated stages of 4969, 4566 and 2143 glass eels, analysed on 74, 73 and 35 occasions for the freshwater, estuary and laboratory samples respectively.

The R^2 between observed and simulated data was 0.93. It varied from 0.87 to 0.96 among samples. It was similar for fresh water (range 0.87–0.96) and the estuary (range 0.87–0.95) and was higher for the laboratory (range 0.93–0.96). The model mean absolute error was 7%, varying from 5 to 9% among samples, 6 to 9% for *in situ* samples and 5 to 7% for laboratory samples. From $VI_{A0 \text{ cum}}$ to $VI_{A3 \text{ cum}}$ mean absolute error was 5.2, 8.3, 8.1 and 6.0% respectively.

The parameters of the model allowed the description of stage durations with parameters p_1 to p_4 being the mean pigmentation time necessary for passage from stage V_B to stages VI_{A0} to VI_{A3} [Fig. 2(a)]. The parameters values and their range were fitted as following: $p_1 = 0.267$ (0.204–0.331), $p_2 = 0.835$ (0.680–0.991), $p_3 = 1.560$ (1.302–1.819) and $p_4 = 3.682$ (3.168–4.195). The notion of ‘pigmentation time’ allowed the use of the same model in experiments whose durations varied from 6 days to 3 months. The pigmentation time was fitted on both temperature and salinity, with temperature being the main factor and salinity acting as a secondary factor slowing down the pigmentation (Fig. 2(b)). Parameters p_5 and p_6 corresponding to ‘efficient daily temperature’ were calculated as $p_5 = 4.566$ (3.808–5.324) and $p_6 = 8.141$ (6.043–10.239).

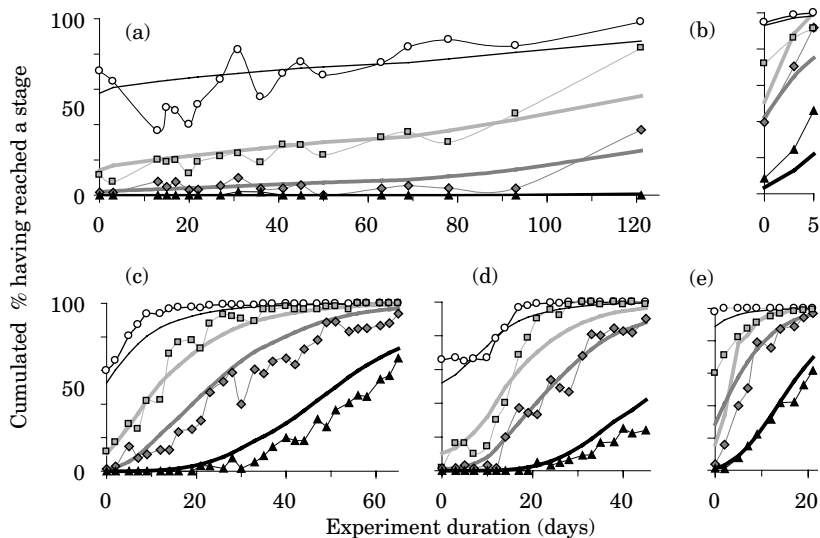


FIG. 1. Calibration of the model illustrated on five samples, for observed (—○—) and modelled (—) cumulative distributions, *i.e.* cumulative percentage of glass eels having reached particular stage over time for stages VI_{A0} (—○—), VI_{A1} (—□—), VI_{A2} (—◇—) and VI_{A3} (—▲—). Experiments in (a) refrigerator (R_1 ; 3.9° C, salinity 6.4), (b) fresh water (F_6 ; 18.1° C, salinity 0.0), (c) estuary (E_2 ; 9.7° C, salinity 7.4), (d) fresh water (F_1 ; 8.3° C, salinity 0.0) and estuary (e) (E_3 ; 15.9° C, salinity 2.5).

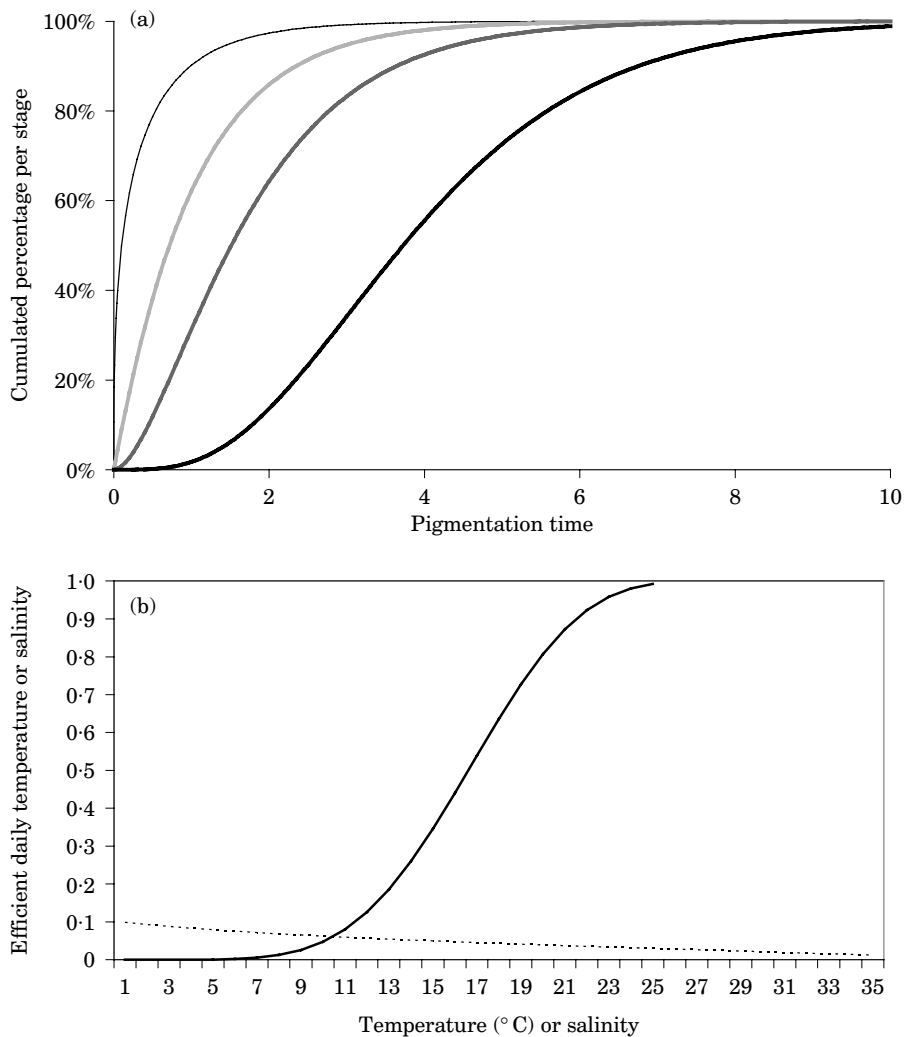


FIG. 2. (a) Modelling of stage duration by gamma functions for cumulative distributions of stages VI_{A0} (---), VI_{A1} (---), VI_{A2} (—) and VI_{A3} (—) according to pigmentation time (cumulative sum from the beginning of the experiment of the days corrected with the product of an efficient daily temperature and an efficient daily salinity). (b) Correspondence between observed daily temperature and efficient daily temperature (—) and between observed daily salinity and efficient daily salinity (---).

Parameters p_7 and p_8 corresponding to 'efficient daily salinity' were calculated as $p_7 = 0.071$ (0.050–0.092) and $p_8 = 0.456$ (0.332–0.580).

MODEL TESTING

The data provided by the two additional laboratory experiments were fitted by the model. V_1 and V_2 corresponded to 1230 and 1085 glass eels analysed on three occasions and split into nine cumulated stage values. The calculated R^2 was 0.97

and 0.74 and the mean absolute residual was 5 and 12%. V_3 and V_4 corresponded each to 300 glass eels analysed on seven occasions and split into 21 cumulated stage values. The calculated R^2 was 0.93 and 0.86 and the mean absolute residual was 5% and 10% (Fig. 3).

STAGE DURATION DERIVED FROM THE MODEL

The following result describes the model output for stage duration at constant temperature and salinity. In fresh water, the duration from stage V_B to VI_{A0} would be 20, 3 and 2 days for 5, 10 and 15° C respectively. From stage V_B to VI_{A3} , these durations would be extended to 240, 40 and 20 days. With a salinity of 20, the durations would be nearly doubled (Fig. 4).

DISCUSSION

EFFECT OF ENVIRONMENTAL FACTORS ON PIGMENTATION DEVELOPMENT

Both pigmentation evolution and modelling clearly confirm that temperature was the main factor controlling the dynamics of pigmentation during the *in situ* stay of glass eel batches, both in the estuarine and in the fluvial sites. This result is in accordance with the other experiments carried out on *A. anguilla* glass eels (Strubberg, 1913; Panu, 1929a; Lecomte-Finiger, 1983; Ciccotti *et al.*, 1993), as well as *Anguilla japonica* Temminck & Schlegel (Dou *et al.*, 2003).

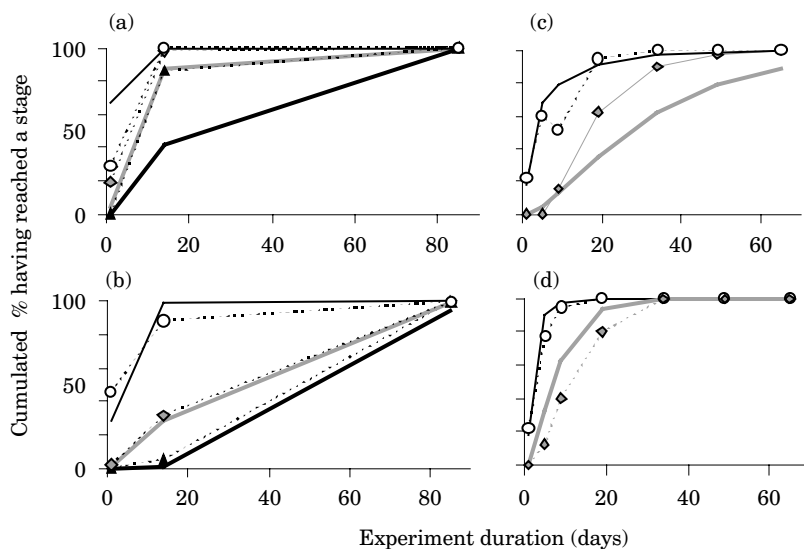


FIG. 3. Validation of the model with glass eels collected elsewhere in Europe, for observed (—○) and modelled (—) cumulative distributions of stages VI_{A0} (—; ○), VI_{A2} (—; ◆) and VI_{A3} (—; ▲). (a) Strubberg (1913) (18° C, salinity 16), (b) Strubberg (1913) (12° C, salinity 16), (c) E. Ciccotti (unpubl. data) (15° C, salinity 36) and (d) E. Ciccotti (unpubl. data) (15° C, salinity 0).

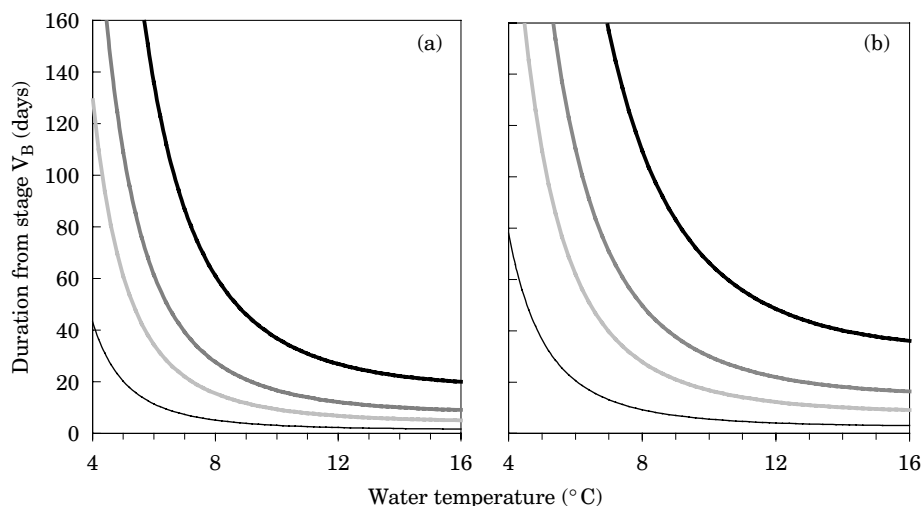


FIG. 4. Stage durations derived from the model. Durations from stage V_B to VI_{A0} (---), VI_{A1} (---), VI_{A2} (—) and VI_{A3} (—), for salinities of (a) 0 and (b) 20.

The effect of salinity was less easy to recognize in the natural conditions, because of the continuous tide-related variation of salinity, besides the possibility that the estuarine environment influences pigmentation rates by factors other than salinity. At a general level in the model, however, salinity acted as a secondary factor slowing down pigmentation, and higher salinities, related to estuarine conditions in the *in situ* experiments, were in fact related to less advanced pigmentation trends (Briand *et al.*, 2004), as also reported by other authors (Panu, 1929*b*; Lecomte-Finiger, 1983).

Environmental factors such as temperature or salinity influence eel metamorphosis, and related events *via* the pituitary-thyroid axis (Vilter, 1946; Ozaki *et al.*, 2000; Jegstrup & Rosenkilde, 2003), that controls basic physiological processes, including growth and environmental adaptation. Increased temperatures trigger metamorphosis through thyroid stimulation (Eales, 1985; Leloup & De Luze, 1985; Eales & Shostak, 1986; Mellinger, 1994; Power *et al.*, 2001). Temperatures could, however, also directly influence duration of metamorphosis shortening it by an increased metabolic rate (Youson, 1988).

The explanation of the role of salinity is less straightforward. Glass eels maintained in full salinity can develop fully and fresh water is not necessary for the completion of metamorphosis (Schmidt, 1909) as illustrated by the widespread occurrence of glass eels in fully saline conditions (Antunes & Tesch, 1997) or of yellow eels without a freshwater phase (Tsukamoto *et al.*, 1998; Tzeng *et al.*, 2001; Arai *et al.*, 2003). Acclimation by yellow eels to high salinities induces, as does temperature, significant stimulation of thyroid hormone secretion and of peripheral conversion of T4 to T3 (Leloup & De Luze, 1985). In glass eels, on the other hand, it was shown that the production of a protein containing thyroid hormone levels was twice as high in fresh water than in salt water (Monaco *et al.*, 1981). In both cases, the higher thyroid activity could be the response to the osmoregulatory challenge.

PIGMENTATION TIME

The product of the two variables 'efficient daily salinity' and 'efficient daily temperature' takes account of the interacting effect of the two factors, *i.e.* even if the salinity is favourable, a low temperature will result in a low pigmentation time. The beta function adjusting the temperature effect allows the great variability of speeds of transition between stages to be illustrated. The beta function adjusting the salinity makes it possible to modulate the effect of temperature by a light retarding effect for large salinities [Fig. 2(b)].

The pigmentation time was considered as independent from the individual level, the pigment stage and the experiment. Indeed, the model was not related to the individual level, and the times an individual spends at each stage were independently distributed (Manly, 1990). Although there might be some individual variability in the pigmentation (Schmidt, 1909), the model was adjusted on a modal value calculated on the splitting of a mean number of 63 glass eels in five stages. In practice, stage transition models seldom allow the individual level to be considered (Kempton, 1979).

The model was also considered as independent from pigment stages, in other words the influences of salinity and temperature on the ontogenetic processes were considered the same for all stages. Indeed, the enhancement of the osmoregulatory ability might not be sufficient to exert a large difference from stage V_B to stage VI_{A3} . For glass eels kept at ambient temperatures ranging from 9 to 18° C, a significant change in the drinking rate was observed, but only after 3 months in salt water, and 5 months in fresh water (Birrell *et al.*, 2000), that is, for durations longer than the present experiments. Moreover, the development of the oesophagus, an important osmoregulatory organ was shown to be partly independent from pigmentation (Ciccotti *et al.*, 1993). Ontogenetic processes are continuous and 'stage' does not imply an instantaneous state (Youson, 1988). In the case of the transition from glass eel to elver, steps are defined by pigment stages, but these do not give information on the biological or physiological state of a glass eel (Jegstrup & Rosenkilde, 2003).

Similarly the use of the same pigmentation time for all experiments implies a lesser adjustment than if separate pigmentation times were used for each experiment. This confers, however, predictive capabilities to the model under a wide range of environmental conditions.

PIGMENTATION MODELLING

In this study the fishing location was in the Vilaine Estuary, 12 km from the sea. This short distance and the heavy fishing pressure (Briand *et al.*, 2003) resulted in a majority of V_B and VI_{A0} pigment stages throughout the season (Table II). These conditions helped the building of a model using gamma distributions, because there was a single peak in numbers of glass eels entering the population at time zero (Manly, 1990). The progressive ageing of the sample collected in the estuary from February to April was corrected with the use of an initial pigmentation time calculated from the initial pigment stages composition. A sample consisting of multiple waves of glass eels having entered the estuary (*i.e.* with several peaks in number entering at time zero) would therefore not have been well described, and

this may explain the higher mean absolute residuals observed in the validation experiments V_1 and V_2 (Fig. 3).

The model should be used to describe samples collected in estuaries in winter and springtime with temperatures ranging from 0 to 20° C. Accordingly, the model should be used for salinities ranging from 0 to 27. With a salinity of 35, the validation experiment V_4 was out of the model range and had a lesser adjustment (higher mean absolute residual) with an exacerbated slowing down effect (Fig. 3). Other experiments would therefore be needed to analyse pigmentation kinetics in the sea.

Similarly, other experiments would be needed to extend the model to the first stage V_A whose importance was stressed near the mouth of the estuary (Charlon & Blanc, 1982) and to the two last and longest stages VI_{A4} and VI_B (Strubberg, 1913; Ciccotti *et al.*, 1993).

MODEL APPLICATIONS

The space and time dynamics of the glass eels migration in a given site results from three different interacting features. The first feature is inherent to the overall composition of the glass eel cohort, that depends on the developmental stage as well as on the physiological state of the single individuals (Boëtius & Boëtius, 1989; Désaunay & Guérault, 1997). De Casamajor *et al.* (2001b) did not observe any significant difference in the quality and physiological state of successive runs of unpigmented glass eels arriving at the Biscay Bay coast in the course of a season. Therefore it is possibly that the timing of arrival near the coast mostly determines the variation in the estuarine sojourn, mirrored in the developmental level and hence in the pigmentation stages of the glass eel groups collected in the estuary (Jellyman *et al.*, 1999; De Casamajor *et al.*, 2001a).

The second and strongest factor is extrinsic, and is represented by the hydrological dynamics of the water masses in the estuary, that drives the upstream transport of glass eels when they are in their passive phase and facilitate or oppose their active swimming performances when in the colonization stage (Gascuel, 1986; McCleave & Wippelhauser, 1987). As a consequence, the same glass eel cohort can evolve in different pigmentation clusters if the migratory shoal breaks up in relation to unfavourable hydrological conditions in the estuary (Elie & Rochard, 1994; De Casamajor *et al.*, 2000).

At a third level, the single estuarine abiotic conditions (*e.g.* temperature, salinity variations turbidity and current speed) influence the individuals, in that they affect their specific developmental pace but also because they might condition their individual response to environmental stimuli (Edeline *et al.*, 2004).

On the whole, most authors unanimously report a major time frequency, *i.e.* a 2 week cycle of the waves of invasion of glass eels. This periodicity of ascent is evident from abundance trends in most estuaries (Jellyman, 1979; Sorensen & Bianchini, 1986; Ciccotti *et al.*, 1995) as well as from the pigmentation frequency composition of glass eel samples (Elie, 1979; Boëtius & Boëtius, 1989). This periodicity in the peaks of abundance and in the glass eels pigmentation composition can change or level out in relation to abnormalities in the migration progress such as quantitative reductions, ecological disturbances or fishing pressure.

Within an estuary, the timing of arrival will determine the temporal variation in recruitment strength; hydrological factors will modulate the spatial and temporal dynamic of migration, while abiotic factors will determine pigment stage evolution, which could also be related to behavioural response such as settlement. Monitoring of temperatures and salinities allows the evolution of pigment stages of a sample collected at one place in the estuary and at repeated intervals to be modelled. For each stage, it is possible to add the pigmentation time necessary to reach this stage to the pigmentation time corresponding to the temperatures and salinities recorded between each sampling date, and predict a change in pigment stages. The difference between predicted and observed values provides an idea for the net increase or decrease for each stage. These variations are the consequence of recruitment, migration, mortality and settlement processes that could in turn be modelled, and fitted on the data.

The seasonal variation of pigment stages would then provide an ecological descriptor of the migration course allowing fast and affordable monitoring of the waves of glass eels entering the estuary. It could therefore be used in field studies, as the tool to trace out the migratory pattern, but also describe the anthropogenic pressures applied to glass eels. For instance, for glass eel fisheries, pigment stage collected at different space and time could be included in a model used to predict and post-evaluate the effect of fishing regulations.

We would like to thank R. Poole for correcting the manuscript.

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