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Enhanced growth without accelerated puberty in fish: A role for the melanocortin system

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ABSTRACT

In two swordtail species of the genus Xiphophorus, onset of puberty in males and females, fecundity in females, and adult size in males are modulated by sequence polymorphism and gene copy-number variation at the P locus affecting the type 4 melanocortin hormone receptor (Mc4r). The involvement of Mc4r in regulating the onset of puberty outside the genus Xiphophorus remains unclear. In this study we used a transgenic line overexpressing asip1 (asip1-Tg), an endogenous antagonist of both type 1 melanocortin hormone receptor (Mc1r) and Mc4r, to investigate the relevance of the melanocortin system on the onset of puberty and adult reproductive performance in zebrafish (Danio rerio). Comparison of growth, puberty and reproductive performance between wild-type (WT) and asip1-Tg zebrafish revealed that a decreased activity of the melanocortin system did not change the timing of puberty but significantly delayed early growth of transgenic animals. Hatching time was postponed in asip1-Tg fish and they were significantly smaller than their WT siblings at 75 dpf, despite showing enhanced linear growth after having completed puberty. asip1-Tg females produced 1.38 times more eggs but spawned less frequently, and their eggs had showed a 0.89-fold smaller diameter but a 1.04-fold increase in larvae body length at hatching. Therefore, we demonstrate that asip-tg zebrafish do not reach puberty earlier than WT counterparts as it could be expected considering the enhanced length and weight growth during early adulthood. This is so because the effects of transgene on growth are noticeable after a threshold length, when puberty has already been reached. Data show that the inhibition of melanocortin system via asip overexpression is an excellent strategy to promote growth, in absence of obesity, by enhancing food efficiency but without accelerating puberty timing. Data here presented provides a deeper characterization of the phenotype induced by the decreased activity of the melanocortin system in fish thus providing an excellent strategy for future aquaculture especially because U.S. Food and Drug Association has recently approved transgenic fish trading.

1. Introduction

The process through which an individual reaches sexual maturity and acquires reproductive capability is called puberty. Following gonadal sex differentiation and an immature, juvenile stage, genetic and environmental factors activate the brain-pituitary-gonadal axis, which promotes adult reproductive functions (Okuzawa, 2002; Chen and Ge, 2013). Some fish, such as swordtail species of the genus *Xiphophorus*, display a pronounced phenotypic diversity regarding puberty onset from early- (60–90 days) to late-maturing (200–300 days) polymorphs

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(Kallman and Schreibman, 1973). This polymorphism is also associated with adult body size and reproductive behaviour in male Xiphophorus. Because males cease to grow reaching puberty, adult male body size is correlated with the time of sexual maturation, such that early-maturing fish are smaller than late-maturing fish (McKenzie Jr. et al., 1983). Differences in body length and the timing of puberty onset are associated with different reproductive strategies, which are key for the evolutionary fitness (Lampert et al., 2010; Maderspacher, 2010). Larger latematuring males invest heavily in courtship by defending territories to be visited by gravid females and by ritualizing the pairing, whereas small fish exhibit a "sneaker behaviour". They do not court females but perform a chase behaviour and parasitically fertilize females just thrusting the gonopodium to obtain a copulation (Maderspacher, 2010; Liotta et al., 2019). Females prefer large and intermediate males suggesting sexual selection against sneaker alleles. However, smaller males evade predation better than larger males and have a larger time window for reproduction since they reach puberty earlier, so that reproductive success over the entire life cycle may be similar for small and large males (Maderspacher, 2010).

In the 70's, Kallman and Schreibman (1973) and Schreibman and Kallman (1977) demonstrated that a Mendelian locus on the sex chromosomes of the platyfish, the so-called *P locus* (Pituitary or Puberty), controls the onset of puberty in males and females, size in males and fecundity in females. The identity of P locus remained elusive for years although its position in the sex chromosomes is close to some other important loci as the sex-determining (SD) locus (SD), the Tu locus, responsible for the spontaneous melanoma (Volff et al., 2013), and some pigment genes serve as convenient gene markers of P locus (Schreibman and Kallman, 1977). Interestingly, the Plocus contains multiple copies of both functional (A allele) and non-functional versions (B1 and B2 alleles) of the melanocortin 4 receptor (mc4r) (Lampert et al., 2010). The size of males and, by extension, puberty onset correlate to the number of non-functional alleles in the Y chromosome. Thus, bigger males exhibit a higher number of non-functional alleles that delay puberty onset, presumably by diminishing the signalling of the functional alleles. On the contrary, males carrying functional alleles of mc4r are smaller and precocious (Lampert et al., 2010).

Mc4r binds the melanocyte-stimulating hormones (Mshs) and two inverse agonists, agouti-related protein (Agrp) and agouti-signalling protein (Asip). Both Agrp and Asip inhibit the constitutive activity of Mc4r and antagonistically compete with α -Msh (Tolle and Low, 2008; Sánchez et al., 2009) In mammals, Mc4r is expressed mainly in the brain but a wider expression profile is found in fish (Cerdá-Reverter et al., 2011). The hypothalamic expression is related intimately to the regulation of energy balance and growth (Cone, 2006). Therefore, the interruption of α-MSH signalling in Mc4r knockout mice induced hyperphagia, reduced energy expenditure, hyperinsulinemia, increased linear growth and maturity-onset obesity (Huszar et al., 1997). A similar metabolic syndrome is observed in transgenic mice ubiquitously overexpressing Asip or Agrp genes (Klebig et al., 1995; Ollmann et al., 1997). In zebrafish (Danio rerio), overexpression of agrp1 (Song and Cone, 2007) and asip1 (Guillot et al., 2016) also result in increased linear growth whereas morpholino-based agrp knockdown induces opposite effects (Zhang et al., 2012). The sa0149 mc4r-deficient zebrafish line also exhibited enhanced growth (Zhang et al., 2012) but recent results reported in medaka (Oryzias latipes) from de Carbio strain showed no effects on the linear growth of adult fish after TALEN-based mc4r knockout Liu et al., 2019).

The involvement of Mc4r in regulating the onset of puberty outside the genus *Xiphophorus* remains unclear. Even within this lineage, the molecular mechanism appears not to be conserved. While in *X. nigrensis* and *X. multilineatus*, puberty onset and body length are determined by *mc4r* allelic and copy number variations Lampert et al., 2010), in *X. hellerii*, a species where both large and small males also exist, only a wild-type (WT) *mc4r* allele was found (Liu et al., 2020). In *X. nigrensis*, *mc4r* expression in the brain is much higher in large than in small males. Such differential expression was also observed in X. hellerii. Hence, high expression of mc4r in large males could be related to early or late puberty onset, reflecting an ancestral scenario in the genus Xiphophorus. In medaka, the regulatory network of Mc4r signalling does not appear to be involved in the regulation of puberty, as mc4r knockout fish reach sexual maturity at a similar time as WT animals (Liu et al., 2019). Our experiments demonstrated that Asip1 work as an endogenous antagonist of both Mc1r and Mc4r (Cerdá-Reverter et al., 2005; Guillot et al., 2016). Subsequently, we generated a transgenic zebrafish strain overexpressing goldfish asip1 and demonstrated the involvement of the melanocortin system in regulating the dorsoventral pigment pattern (Ceinos et al., 2015) and growth (Guillot et al., 2016). Here, we exploit the potential of this model to study the question if the decreased activity of the melanocortin system modulates the timing of puberty in zebrafish, thus expanding studies on the melanocortinergic regulation of puberty to a key model species for vertebrate development.

2. Materials and methods

2.1. Fish and housing

Wild-type (WT) and transgenic stocks come from a background of TU (Tuebingen, Nüsslein-Volhard Lab) strain. Generation of the transgenic zebrafish line [*Tg*(*Xla.Eef1a1:Cau.Asip1*]*iim4* (*asip1*-Tg), using the Tol2 transposon system, has been previously described (Ceinos et al., 2015). Adult zebrafish were maintained at 28 ± 2 °C under a 14 h/10 h light/ dark cycle. Fish were fed three times a day until satiety with a combination of freshly-hatched brine shrimp (*Artemia* sp. nauplii) and sera Vipan flake food (Sera, Heinsberg, Germany). All experiments were performed in accordance with Spanish (Royal Decree 53/2013) and European (2010/63/EU) legislations for the protection of animals used for experimentation. The used protocols were approved by the IATS Ethics Committee (Register Number 09–0201) under the supervision of the Spanish Government.

Animals used in this study were free of any signs of disease. Approximately 600 embryos of each genotype line, WT and *asip1*-Tg, were obtained at the onset of light from natural in-tank breeding crosses. Larvae from both genotypes were raised in 15 L aquarium in strictly identical conditions. From 5 to 12 dpf fish were fed rotifers. From 13 dpf they were offered brine shrimp. At 20 dpf fish were transferred to two 45 L aquarium (n = 150 fish/aquarium) and dry food (sera Vipan flake) was gradually introduced to their diet.

2.2. Gonadal development

To increase the strength of our studies two independent experiments were conducted. In experiment 1, larvae were sampled at 30, 32, 35, 40, 46, 54, 60, and 75 dpf. In experiment 2, sampling was conducted at 30, 34, 38, 42, 46, 60 and 75 dpf. At each sampling point, at least 30 individuals were randomly collected from each tank and sacrificed by overdose of tricaine methane sulfonate (MS222, 200-300 mg/L) by prolonged immersion. Larvae were imaged by stereomicroscope (Olympus SZX16, stereo microscope, Tokyo, Japan), images captured, and their standard length recorded using ImageJ version 1.52 software. Larvae were then fixed by immersion in 1% glutaraldehyde, dehydrated, embedded in 2-hydroxyethyl methacrylate polymer resin (Technovit 7100, Heraeus Kultzer, Germany). Serial sections of 2 µm thickness were prepared and stained in toluidine-methylene blue solution for histological analysis. Gonad morphology and classification of the ontogenetic gonad differentiation into ovary or testis was done according to Maack and Segner (2003). Gonads with proliferating germ cells that could not be identified as female or male germ cells were classified as undifferentiated, gonads with both germ cell types in transition from a bipotential 'juvenile ovary' were classified as transitioning ovary (Supplementary Fig. S1). The maturation stages were categorized using a

numerical staging system based on the most mature germ cells present in the gonads. Identification of germ cells at different stages of gametogenesis was done according to the described above (Selman et al., 1993; Leal et al., 2009). Ovarian development was classified in six stages (Supplementary Fig. S2) based on their size and vitellogenic state (Wang and Ge, 2004): I, primary growth (~0.1 mm); II, previtellogenic (~0.25 mm); III, early vitellogenic (~0.35 mm); IV, mid vitellogenic (~0.45 mm); V, late vitellogenic (~0.55 mm) and VI, full grown (~0.65 mm). Testicular development was adjusted to four stages (Supplementary Fig. S3) based on Begtashi et al. (2004) stage 1, immature (type A spermatogonia); stage 2, early maturation (type B spermatogonia); stage 3, mid maturation (spermatogonia to spermatocytes) and stage 4, late maturation (spermatocytes, spermatids, and spermatozoa).

2.3. Maturation curve

According to the histology based criteria proposed by Vazzoler (1996), a scale of three maturity stages was stablished as follow: A, immature; B, maturing; C, spawning. Fish in stage B and C were considered to be initiating or completing puberty. In females, puberty onset is characterized by some follicles entering the previtellogenic stage (appearance of cortical alveoli in the oocytes), and in males by the presence of cysts containing type B spermatogonia. Fish were grouped by age (dpf) and by intervals of 2.5 mm in body length. The fraction of mature fish by age, length and sex was estimated through the logistic equation described by O'Brien et al. (1993):

$$P = \frac{1}{1 + e^{-(a+bX)}}$$

where *P* corresponds to the proportion of maturing fish, *X* the length or age and *a* and *b* are the equation estimated coefficients. The parameters were estimated by a lineal regression analysis using Graphpad Prism version 8.3. The length (L_{50}) and age (A_{50}) at which 50% of the male and female population initiate puberty was estimated as a ratio of *a/b*.

2.4. Reproductive performance

Five males and five females of *asip1*-Tg or WT fish were individually placed into 2-l tanks and fed three times a day until satiety with a combination of freshly-hatched brine shrimp and dry food. After 1-week acclimation, one female with bulging abdomens were randomly placed into individual spawning tanks with one male and left overnight. Zebrafish spawn within the first few hours after sunrise (Hisaoka and Firlit, 1962) and to ensure complete spawning, the assessment of egg production took place between 8 and 10 am. The occurrence of a spawning event and the total number of eggs spawned per female were assessed during seven spawn events. A total of thirty-five spawning couples were tested for each line (five pairs per event).

For assessing the egg fertilization, we distinguished fertilized eggs by the presence of a multi-cellular blastodisc (Kimmel et al., 1995). Fertilized eggs were collected and incubated at 28 °C. Mortality at 24 hpf and number of embryos hatched at 48 and 72 hpf was recorded. At least 50 eggs from 3 spawning events were photographed. The egg size and diameter of the yolk at the gastrulation stage were measured using ImageJ version 1.52 software. In addition, at least 30 larvae of age 4 dpf were photographed individually to compare the larval standard lengthat- hatch and larval yolk sac-volume between the strain fish. The yolksac volume was estimated using the following formula (Chambers et al., 1989):

$V = \pi \left(6LH^2 \right)^{-1}$

where L represents the length (horizontal measurement; mm) and H the height (vertical measurement; mm) of the yolk-sac.

2.5. Statistical analysis

Statistical treatment of the data was done with both GraphPad Prism version 8.3 The Mann-Whitney and the Kolmogorov-Smirnov nonparametric tests were used to compare the linear length between WT and *asip1*-Tg lines. To compare differences in the linear length between days, for each strain performed Dunn's test of multiple comparisons following a significant Kruskal-Wallis test. The Fisher's exact-test was used for comparisons of gonadal development proportions. The strength of the association between the pair of parameters linear length and gonadal development was evaluated by calculating the correlation coefficient, r, using the Spearman rank order correlation nonparametric test. Differences in rate success spawning were analyzed by Fisher's exact-test. Number of eggs and fertilized eggs, mortality, number of embryos hatched, size egg, diameter of the yolk, standard length- at- hatch and yolk sac-volume were analyzed by unpaired *t* test with Welch's correction. For all performed tests, the significance level was set at 0.05.

3. Results

3.1. Linear growth

The effect of overexpression of asip1 might have on the growth of juvenile and adult zebrafish was determined by measuring standard length in both asip1-Tg and WT fish from 30 to 75 dpf (days postfertilization). Two independent experiments were conducted. In experiment 1, WT zebrafish grew from 8.9 mm to 22.1 mm while asip1-Tg fish grew from 10.5 mm to 20.9 mm (Fig. 1A). Over the course of experiment 2, the length of WT zebrafish increased from 6.12 mm to 22.48 mm while in asip1-Tg it ranged from 6.5 mm to 22.4 mm (Fig. 1B). In general, for both WT and asip1-Tg fish, we found no significant differences in the body length during the sampling period. Exceptions include length increase in WT fish from 40 to 46 dpf and from 60 to 75 dpf in experiment 1 (Fig. 1A) and 38-42 dpf in experiment 2 (Fig. 1B). Length increase of asip1-Tg fish was significantly different within 35-40 dpf in experiment 1 and 46-60 dpf in experiment 2. The distribution of the standard length was significantly different among fish lines. At 30 dpf, in both experiment 1 and 2, standard length of WT was significantly lower than in asip1-Tg fish. However, from 42 dpf in experiment 2, 46 dpf in experiment 1, the standard length of WT was in general significantly higher than that of transgenic fish (Fig. 1). Body length data were in addition classified according to sex. For both WT and asip1-Tg female fish, we found no significant differences in the growth during the sampling period. Exceptions include growth increase in WT fish within 38-42 dpf in experiment 2. Regarding male fish, growth of WT was found to be significantly different but only after they attained the adult stage (Fig. 3). No differences were found in male asip1-Tg fish (Fig. 3). At 30 dpf (experiment 1) and 38 dpf (experiment 2), length of female WT was significantly lower than that of asip1-Tg (Fig. 2B). Nevertheless, at 42 dpf in experiment 2 and 46 dpf in experiment 1, this trait was reversed with WT female having a higher standard length than asip1-Tg females. Standard length of WT males was significantly higher than that of asip1-Tg males from 46 dpf until 75 dpf in experiment 2. The same trend was seen in experiment 1, although differences were statistically significant only at 75 dpf.

3.2. Gonadal differentiation

To investigate if a decreased activity of the melanocortin system induced by an overexpression of *asip1* might have a role on sexual differentiation, we monitored by histology the gonad development of *asip1*-Tg zebrafish between 30 and 75 dpf and compared these results with those found in WT fish. In experiment 1 (Fig. 4A), gonads from 226 WT fish and 237 *asip1*-Tg fish were analyzed. A female biased sex ratio was observed, with 56.6% of WT and 55.7% of *asip1*-Tg fish being identified as females. At 30 dpf (Fig. 4A2), the fraction of undifferentiated gonads



Fig. 1. Standard length of WT and *asip1*-Tg fish in (A) experiment 1 and (B) experiment 2. Data are represented as mean \pm SD. Numbers inside the bars indicate sample size (n). Statistical significance is indicated as asterisks (*). For all statistics: *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001.



Fig. 2. Standard length of WT and asip1-Tg female fish in (A) experiment 1 and (B) experiment 2. Data are represented as mean \pm SD. Statistical significance is indicated as asterisks (*).



Fig. 3. Standard length of WT and asip1-Tg male fish in (A) experiment 1 and (B) experiment 2. Data are represented as mean \pm SD. Statistical significance is indicated as asterisks (*).

was significantly higher in *asip1*-Tg than in WT fish (p = 0.0122), and 96.5% of WT gonads were identified as females (p = 0.0122) (Fig. 4A1). In both fish lines, signs of overt sexual differentiation (transitioning phase of testis development) began at 32 dpf. However, the number of transitioning gonads was significantly higher in WT fish at 32 dpf (p = 0.0475) and 46 dpf (p = 0.0237). Male sex was revealed in gonads of 35 dpf fish. In experiment 2 (Fig. 4B), 292 gonads from WT fish were analyzed and as in experiment 1, a female skewed sex ratio was observed with female representing 53.8% of the population cohort. From a total of 316 *asip1*-Tg fish, 44% were female and 51.3% male. At 30 dpf, there was no difference in the fraction of undifferentiated gonads. However, the percentage of ovaries (Fig. 4B1) was significantly higher in WT than in *asip1*-Tg fish (p = 0.0095) and the same was registered at 38 dpf (p =

0.0099). Gonads in the transitioning phase of testis development could be observed from 30 dpf onwards. The fraction of animals in the transitional phase was significantly higher in the WT line at 34 dpf (p =0.0136) and 42 dpf (p = 0.0048). In the transgenic *asip1*-Tg line (Fig. 4B2) histological examination revealed a male fate of the gonad at 30 dpf. This sex proportion rate was significantly overrepresented at 34 (p = 0.0007), 38 (p = 0.0002) and 42 dpf (p = 0.0010).

3.3. Gonadal maturation

In the zebrafish, the transition from primary growth (stage I) to previtellogenesis (stage II) in the ovary is considered the sign of puberty onset in females (Ge, 2005). From 30 to 34 (experiment 1) / 35



Fig. 4. Gonadal differentiation of WT and *asip1*-Tg fish in (A) experiment 1 (A1: WT; A2: *asip1*-Tg) and (B) experiment 2 (B1: WT; B2: *asip1*-Tg). Data are represented as a percentage of the total number of fish analyzed for each genotype. Statistical significance after Fisher's exact test is indicated as asterisks (*).



Fig. 5. Ovary development of WT and *asip1*-Tg fish in (A) experiment 1 (A1: WT; A2: *asip1*-Tg) and (B) experiment 2 (B1: WT; B2: *asip1*-Tg). Data are represented as a percentage of the total number of fish analyzed for each genotype. Statistical significance after Fisher's exact test is indicated as asterisks (*).

(experiment 2) dpf, oocytes of all females, regardless of the line and trial, were at the primary growth stage (Fig. 5A, B). In both experiments 1 and 2, towards the end of the sampling period, the percentage of ovarian follicles in primary growth stage was higher in *asip1*-Tg females than in WT females. In experiment 1, ovarian previtellogenic follicles that are characterized by the presence of cortical alveoli oocytes were

first seen at 40 dpf in the leading wave of developing oocytes. Even thought they were found initially in a similar proportion in both WT and *asip1*-Tg lines, at 46 dpf they were found in a higher percentage in WT than in *asip1*-Tg females (Fig. 5A1: 64.3%, p = 0.0236). In experiment 2, stage II ovaries could be recognized at 34 dpf but only in WT female (12%) and at a level not significantly different from *asip1*-Tg fish



Fig. 6. Testis development of WT and *asip1*-Tg fish in (A) experiment 1 (A1: WT; A2: *asip1*-Tg) and (B) experiment 2 (B1: WT; B2: *asip1*-Tg). Data are represented as a percentage of the total number of fish analyzed for each genotype. Statistical significance after Fisher's exact test is indicated as asterisks (*).

(Fig. 5B1). After the vitellogenesis stages III-V, when oocytes grow fast due to the accumulation of yolk in the cytoplasm, follicles in the ovaries of both WT and *asip1*-Tg fish entered the maturation stage (VI). In experiment 2, at 75 dpf there was a higher proportion (36.4%) of WT females in stage VI, although this value was not significantly different from *asip1*-Tg female (5.6%). Despite puberty completion (first egg laying) was not followed in this study, the *asip1*-Tg line can be propagated in a standard propagation scheme.

In experiment 2, gonad histology analysis revealed the presence of immature testis already at 30 dpf in transgenic *asip1*-Tg fish (Fig. 6B2). In contrast, WT fish sampled at this age were still undergoing sexual differentiation (Fig. 6B1). In both experiments 1 and 2, at day 35 or 34, a higher proportion of males in stage 1 was recorded in the *asip1*-Tg line, reaching statistical significance in experiment 1 (Fig. 6). In *asip1*-Tg fish, the fraction of stage 1 testis steadily decreased until 46 dpf (when it could not be any longer recognized), when in WT at stage 1 testis, would still account for a 30.4%. The most advanced stage of testicular maturation, stage 4, could be histologically identified in *asip1*-Tg fish already at 34 dpf (20%, experiment 2) but only at 46 dpf (experiment 1: 42%, p = 0.0239) its proportion is significantly higher than in WT males (Fig. 6A2).

3.4. Interaction between growth and gonad development

The interaction between standard length and gonad development is illustrated in Figs. 7 and 8. In experiment 1 and 2, a strong positive association was found between these variables, irrespective of sex and genotype.

3.5. Length and age at maturity

The body length of the analyzed females varies similarly in both genotypes (experiment 1: WT = 14.9 and 22 mm; asip1-Tg = 14.8 y 20.9 mm and experiment 2: WT = 8.8 and 21.6 mm; asip1-Tg = 10.9 and 22.4

mm). The logistic function applied on female gonad maturation data shows that the body length of 15 mm in experiment 1 and 12.5 mm in experiment 2 seemed to be a threshold for reaching maturity (Fig. 9). In both experiments and for both WT and asip1-Tg fish, once body exceeds these lengths, and until a size of 20 mm, cortical alveoli appear and the oocytes started to accumulate in the oocytes, as a sign of transition from primary-growth to previtellogenic stages. In 22.5 mm fish, all females had reached maturity stage with full-grown follicles present in the ovary. In experiment 1, the maturity ogive estimated that the length at 50% of maturity (L₅₀) in WT and asip1-Tg lines was 18 mm and 17.4 mm, respectively (Fig. 9A). In experiment 2, L₅₀ was estimated as 16.2 mm for WT females and 16.7 mm for asip1-Tg females (Fig. 9B). As observed in females, the body length data varied similarly in both genotypes (experiment 1: WT = 11.7 and 21.3 mm; *asip1*-Tg = 13.5 and 18.9 mm and experiment 2: WT = 9.9 and 22.5 mm; *asip1*-Tg = 9.9 and 21.7 mm). The logistic function for data in experiment 1, shows that males start to mature at a smaller length than females. Between 20 and 22.5 mm, the percentage of mature males reaches 89 and 100%, respectively (Fig. 10A). On the other hand, data collected from asip1-Tg males, indicates that 74 and 100% of the mature male, are between 17.5 and 20 mm body length. In experiment 2, mature males from both genotypes could be observed as early as 10 mm of body length (Fig. 10B). A successive increase in the proportion of testis presenting the hallmarks of maturation was then observed. 44% of asip1-Tg males reached maturity with a body length of 12.5 mm while 41% of the WT males were mature with a body length of 15 mm. All males of 22.5 mm were classified as mature. In experiment 1, the maturity ogive estimated that the L₅₀ for males of WT and asip1-Tg genotypes was 17.3 mm and 16.6 mm, respectively (Fig. 10A). In experiment 2, L₅₀ was estimated as 16.5 mm for WT males and 14.4 mm for asip1-Tg males (Fig. 10B).

In experiment 1, the age of the sampled mature female ranged from 40 to 75 dpf. Sampling in experiment 2 allowed to identify females of the WT genotype maturing at a younger age than *asip1*-Tg females (34 and 38 dpf, respectively). The logistic function indicates that below 46 dpf



Fig. 7. Correlation analysis to evaluate the strength of relationship between ovary development and standard length of WT and asip1-Tg fish in (**A**) experiment 1 (**A**1: WT; **A2**: asip1-Tg) and (**B**) experiment 2 (**B1**: WT; **B2**: asip1-Tg). The degree of association between variables was measured with the Spearman's rank correlation test with a statistical significance of p < 0.05.

(experiment 1, Fig. 11A) and 42 dpf (experiment 2, Fig. 11B), the proportion of mature females decreases and from 60 dpf onwards, more than half of the females were mature (experiment 1: WT = 79%; *asip1*-Tg = 56% and experiment 2: WT = 73%; *asip1*-Tg = 53%). At 75 dpf, 100% of the females were mature. The maturity ogives from experiment 1 and 2 estimated that the age at 50% maturity (A₅₀) for females of the WT line was 53–54 dpf and for the *asip1*-Tg genotype was 58–59 dpf (Fig. 11A).

Regarding the males, the age of WT mature fish was similar in both experiment 1 and 2, ranging from 34 to 75 dpf. On the contrary, *asip1*-Tg males sampled during experiment 1 were found to be mature from 40 to 75 dpf, while in experiment 2, mature fish could already be identified at 30 dpf. The logistic function indicates that below 40 dpf (experiment 1, Fig. 12A) and 38 dpf (experiment 2, Fig. 12B), the proportion of mature males decreases and, as seen for female fish, the majority of the males were mature from 60 dpf onwards, reaching 100% of maturity at 75 dpf. The maturity ogives from experiment 1 and 2 estimated that the A_{50} for males of the WT line was 52–53 dpf and for the *asip1*-Tg genotype, 53 dpf in experiment 1 and 49 dpf in experiment 2 (Fig. 12A, B).

3.6. Reproductive performance

The rate of spawning success (that is, spawning resulting in 1 or more ova) was 97.14% for WT and 77.14% for *asip* 1-Tg (p = 0.0275; Fig. 13A).

The total number of eggs per female (mean \pm SEM) was significantly (p = 0.0113) higher for *asip*1-Tg (493.6 \pm 45.24 eggs) than for the WT line (357.8 \pm 23.73 eggs; Fig. 13B). The absolute number of eggs counted for all breeding pairs for *asip*1-Tg was 13,327 and 12,166 for WT. Similarly, the number of fertilized eggs was significantly higher (p = 0.0028) for *asip*1-Tg (463.4 \pm 48.37 eggs) than for WT fish (287.5 \pm 27.13 eggs; Fig. 13C). However, *asip*1-Tg had a significantly higher mortality (55.5 \pm 3.78%; p = 0.0014) at 24 hpf compared to WT fish (36.19 \pm 4.3%; Fig. 13D). However, at 48 hpf, we found significant differences in the proportion of hatched embryos (p = 0.0205) between WT (61.93 \pm 4.95%) and *asip*1-Tg (44.62 \pm 5.29%, Fig. 13E). The same was observed at 72 hpf (p = 0.0111), with WT having a higher proportion of hatched larvae (92.27 \pm 1.94%) than *asip*1-Tg (80.96 \pm 3.75%; Fig. 13F).

WT eggs were significantly larger $(1.294 \pm 0.0038 \text{ mm}; p < 0.0001;)$ than *asip1*-Tg line eggs $(1.164 \pm 0.0031 \text{ mm}; \text{Fig. 14A})$. Eggs from WT fish also had a significantly larger diameter of yolk $(0.6623 \pm 0.0027 \text{ mm}; p \le 0.0001)$ than *asip1*-Tg $(0.6028 \pm 0.0021; \text{Fig. 14B})$. Moreover, significant differences were also observed in the yolk-sac volume, with WT eggs presenting a bigger volume $(0.0667 \pm 0.0026 \text{ mm}^3; p < 0.0001)$ compared to *asip1*-Tg line eggs $(0.0459 \pm 0.0012 \text{ mm}3;$ Fig. 14C). On the other hand, the standard body length of freshly hatched *asip1*-Tg larvae (4 dpf) was significantly larger $(3.135 \pm 0.0090 \text{ mm}; p < 0.0001)$ than WT $(3.015 \pm 0.0074 \text{ mm}; \text{Fig. 14D})$.



Fig. 8. Correlation analysis to evaluate the strength of relationship between testis development and standard length of WT and asip1-Tg fish in (a) experiment 1 (A1: WT; A2: asip1-Tg) and (B) experiment 2 (B1: WT; B2: asip1-Tg). The degree of association between variables was measured with the Spearman's rank correlation test with a statistical significance of p < 0.05.

4. Discussion

In teleosts, a role for Mc4r in puberty onset regulation outside *Xiphophorus* has not been described yet. Recent results in medaka have shown that Mc4r does not play any role in the timing of puberty, and mechanisms observed in *Xiphophorus* may be restricted to this lineage. To study this possibility, we here took advantage of a zebrafish line that overexpresses *asip1*, an endogenous antagonist of Mc1r and Mc4r (Cerdá-Reverter et al., 2005; Guillot et al., 2016). The *asip1*-Tg zebrafish line represents an excellent model for studies exploring the relationship between melanocortin activity and the timing of puberty onset, since overexpression of *asip1* leads to a reduction of Mc4r activity (Cerdá-Reverter et al., 2005; Sánchez et al., 2009). Our results suggest that *asip1* overexpression had no effect on the timing of puberty in zebrafish but modified growth and reproductive performance.

4.1. Growth

Previous results demonstrated that asip1 overexpression in zebrafish enhanced linear growth (Guillot et al., 2016), but, independently of age, the first differences were detected only after a critical size close to 20 mm (Godino-Gimeno et al., 2020). Our current data show consistently that both male and female asip1-Tg zebrafish are significantly smaller than WT siblings at 75 dpf, when asip1-Tg zebrafish are just below 20 mm. However, asip1-Tg fish exhibit longer size at hatching despite their smaller egg and egg-yolk diameter and, by extension, smaller volume. Accordingly, knockdown of agrp1 by morpholino techniques or the chemical ablation of agrp1 neurons in zebrafish results in shorter animals when compared to their control counterparts at 8 dpf (Zhang et al., 2012; Löhr et al., 2018). Our present data confirms that the positive effects of asip1 overexpression, and by extension of the reduced signalling of melanocortin system, on zebrafish growth requires a threshold length close to 20 mm, just when full gonadal development is reached. Therefore, asip1-Tg fish hatched larger but exhibited reduced growth



Fig. 9. Female first sexual maturity ogives by length based on the histological analysis of ovaries of WT and *asip1*-Tg fish in (A) experiment 1 and (B) experiment 2. Black circle (•) represents observed data for WT female; Dash (-) represents estimates for WT females; Black square (•) represents observed data for *asip1*-Tg female; Double dash (--) represents estimates for *asip1*-Tg female.



Fig. 10. Male first sexual maturity ogives by length based on the histological analysis of testis of WT and *asip1*-Tg fish in (A) experiment 1 and (B) experiment 2. Black circle (●) represents observed data for WT male; Dash (–) represents estimates for WT males; Black square (■) represents observed data for *asip1*-Tg male; Double dash (–-) represents estimates for *asip1*-Tg male.



Fig. 11. Female first sexual maturity ogives by age based on the histological analysis of ovaries of WT and *asip1*-Tg fish in (A) experiment 1 and (B) experiment 2. Black circle (\bigcirc) represents observed data for WT female; Dash (-) represents estimates for WT females; Black square (\blacksquare) represents observed data for *asip1*-Tg female; Double dash (-) represents estimates for *asip1*-Tg female.



Fig. 12. Male first sexual maturity ogives by age based on the histological analysis of testis of WT and *asip1*-Tg fish in (A) experiment 1 and (B) experiment 2. Black circle (\bigcirc) represents observed data for WT male; Dash (-) represents estimates for WT males; Black square (\blacksquare) represents observed data for *asip1*-Tg male; Double dash (--) represents estimates for *asip1*-Tg male.



Fig. 13. Effect of asip1 overexpression on adult zebrafish reproductive function and offspring viability. Average of spawning events (**A**), number eggs per female (**B**), number fertilized eggs (**C**), egg mortality at 24 hpf (**D**), hatched at 48 hpf (**E**) and hatched at 72 hpf (**F**). All data represent the mean \pm SEM. Statistical significance is indicated as asterisks (*), *p < 0.05, **p < 0.01.

until completing gonadal development. From then on *asip1*-Tg fish grew faster than WT siblings, length differences reaching 15% (Guillot et al., 2016; Godino-Gimeno et al., 2020).

Endocrine and molecular mechanisms promoting growth under decreased melanocortin activity have been studied but are still far from being understood. In larval zebrafish, standard somatic growth requires agrp1 signalling through mc4r (Zhang et al., 2012; Godino-Gimeno et al., 2020). Therefore, *agrp1* knockdown in the morpholino zebrafish model resulted in decreased growth hormone (*gh*) expression, concomitant with increased gh-releasing hormone (*ghrh*) and decreased somatostatin I and II expression (*sstI* and *sstII*) (Zhang et al., 2012). Recently, a mechanism involving the melanocortin system in the feeding-induced growth in larval zebrafish was proposed. Overfeeding causes leptin resistance and reduced pro-opiomelanocortin (*pomc*) hypothalamic levels, leading to reduced activity of sst neurons that express Mc4r and consequently elevated *gh* expression and somatic growth (Löhr et al., 2018). It is therefore possible that *agrp1* or *asip1* over-expression in zebrafish promotes somatic growth via m sst neurons.

4.2. Timing of puberty

Gonadal development in zebrafish can be divided in three fundamental processes: sex determination, differentiation and maintenance. While it is clear the direction in which these processes occur, the timing can be strongly variable. It is now clear, that sex of zebrafish in the wild is determined by genetic factors on a polygenic basis (Bradley et al., 2011; Anderson et al., 2012; Liew et al., 2012; Liew and Orbán, 2014). However, sex ratios in domesticate lines can be greatly influenced by



Fig. 14. Morphological assessment of offspring. Average of egg size (**A**), egg yolk diameter (**B**), larval yolk-sac volume (**C**) and larval standard length (Ls)-athatch (**D**). All data are represented as the mean \pm SEM. Statistical significance is indicated as asterisks (*), *p < 0.05, ****p < 0.0001.

environmental factors such as temperature, nutrition and population density (Lawrence et al., 2008; Liew et al., 2012; Ribas et al., 2017a; Ribas et al., 2017b). In experiment 1 of our study, we observed a female skewed sex ratio in both WT and *asip1*-Tg lines that was also found in the WT population in experiment 2. The increased incidence of females in WT and *asip1*-Tg lines, indicates that equal husbandry conditions were applied to both fish populations, in particularly during the sex determination period, and that these have favor a female-biased sex ratio. Sex differentiation was first seen at 32 dpf (experiment 1) and 30 dpf (experiment 2) in both fish lines but the percentage of differentiating animals was higher in the WT line than in the *asip1*-Tg line in agreement with the delayed growth of the *asip1*-Tg line.

In the zebrafish, puberty onset depends on somatic growth rather than age (Chen and Ge, 2013; Silva et al., 2017; Hu et al., 2019). Accordingly, we found a strong positive association between growth and gonadal development in both WT and asip1-Tg lines. The primarygrowth-to previtellogenic transition in the first cohort of developing follicles is defined as indicating female puberty onset (Ge, 2005; Taranger et al., 2010). The logistic function applied on female gonad maturation data shows that the body length of 15 mm in experiment 1 and 12.5 mm in experiment 2 seemed to be a threshold for reaching maturity in both WT and *asip1*-Tg lines. These body lengths differ from the previously reported for both female Albino (Chen and Ge, 2012; Chen and Ge, 2013) and Casper (White et al., 2008) zebrafish lines, that begin sexual maturation at a critical body length of 18mm (Chen and Ge, 2013; Lessman and Brantley, 2020). The Tubingen (TU) genetic background of our WT and transgenic lines could explain these observed differences since strong genetic components are known to affect size at puberty. L₅₀ (length at which 50% of individuals were mature) is usually used as an index to compare maturation patterns between different groups of fish. Using our logistic model, we calculated an overall L₅₀ of 17 mm (mean value of experiments 1 and 2) for both the WT and asip1-

Tg lines. Although body length at puberty is not altered in female asip1-Tg, estimates of A₅₀ (age at which 50% of individuals were mature) differed between genotypes with WT fish maturing younger than transgenic fish. However, the difference is only 5 days. As expected, males start to mature at a smaller length than females but at a similar time. The logistic function applied on male gonad maturation data shows that the body length of 12.5 mm in experiment 1 and 10 mm in experiment 2 seemed to be a threshold for reaching maturity in both WT and asip1-Tg lines. Once again, these values differ from the ones reported for males of the Casper zebrafish line, that begin sexual maturation at a critical body length of 17 mm (Lessman and Brantley, 2020). Nevertheless, these differences should be assigned to the genetic background of the TU line. Based on our maturity ogives, the size of male asip1-Tg fish at maturity is slightly smaller than that of WT but the differences are minimal (0.7 mm in experiment 1 and 1.4 mm in experiment 2). Estimates of A₅₀ differed between genotypes with *asip1*-Tg fish maturing 4 days younger than WT males but only in experiment 2. Overall, the data of the present study indicates that the melanocortin system is not a critical puberty signal in zebrafish as overexpression of asip1 had no phenotypical effect on puberty timing. Our results, together with previous studies performed in medaka mc4r knockouts (Liu et al., 2019) and X. hellerii, species carrying only Mc4r functional alleles (Liu et al., 2020), suggest that the regulation of the timing of puberty onset by Mc4r signalling in species of the genus Xiphophorus may be an evolutionary adaptation only fully conserved in this lineage.

4.3. Reproductive performance

Studies on MC4R-controlled pathways that regulate reproductive performance have primarily focused mice (Sandrock et al., 2009) and chicken (Aggag and El-Sabrout, 2018). Only a few studies have brought particular attention to this aspect in teleost fish. In Xiphophorus, P locus also controls female fecundity. The female genotype P^1P^1 matures earlier and produces more eggs than late maturing P^5P^5 females. Despite their larger size, P^5P^5 females consistently spawn fewer eggs than females of any other genotype (Kallman and Borkoski, 1978). Different from Xiphophorus, we showed that asip1-Tg females produce more eggs than WT females but spawn less frequently. The number of fertilized eggs in asip1-Tg females is also higher but the hatching rate at 48 and 72 h is lower. Increases in the fecundity of asip1-Tg females is associated with a decrease of oocyte diameter. Female oviparous vertebrates have to overcome an egg size/number commitment, thus if egg size increases, egg number decrease and vice versa (Forbes et al., 2010). The reduced size of eggs is also in agreement with a reduction in egg volk diameter and volume. Good quality eggs often display low levels of mortality at fertilization, eying, hatch and first-feeding (Bromage et al., 1992). While there is no agreement as to what levels of mortality constitute a good quality egg, asip1-Tg embryonic mortality at 24 h was higher than that of WT thus suggesting asip1-Tg eggs to have lower quality.

Yolk is the main component of freshly fertilized fish eggs and is associated to nutrients stored for embryonic development (Kamler, 2008). Eggs produced by WT fish had a larger yolk diameter than asip1-Tg eggs implying greater energy resource for the developing embryos. However, the standard length of hatched asip1-Tg larvae was significantly higher than that of WT larvae. It has been reported that larval body size results not only from growth rate of embryos and yolk-sac volume but also from efficiency of yolk energy utilization by larvae for growth and yolk energy content (Kamler, 2008). Differences in yolk energy content depend, in turn, on the size and caloric value of yolk (Kamler, 2008). Because the yolk-sac volume did differ significantly between both lines, the increased standard length at hatching of asip1-Tg zebrafish may suggest improved use of yolk energy by this line, promoting higher growth rates. We also observed that the hatching time of asip1-Tg larvae was delayed compared to WT larvae. Delayed hatching time has also been observed in mc4r knockout medaka. However, the authors concluded that contrary to our results in asip1-Tg zebrafish, this delay is due to a decrease in the growth rate, rather than an increase in the body length (Liu et al., 2020). Thickening of the outer layer of the chorion makes it difficult for the embryo to break free and has been proposed to be responsible for hatching delays (Uusi-Heikkilä et al., 2010). Unfortunately, we did not measure chorion thickness in *asip1*-Tg embryos. In general, our findings suggest that *asip1*-Tg fish have a lower reproductive performance compared to WT fishes, which is reflected in lower egg quality and yolk diameter, delayed hatching time and larval growth. However, further studies are needed to investigate the mechanisms behind the observed differences in larval growth rate as well as yolk content between WT and *asip1*-Tg fish

In summary, the regulation of the timing of the onset of puberty by the reproductive axis is modulated by the growth axis. Our data further suggests an interaction of both melanocortin and reproductive systems that modulates the effects of reduced melanocortin signalling on somatic growth. A role of sex steroids in the modulation of these effects can be anticipated but more studies are required to corroborate this hypothesis.

5. Conclusion

In conclusion, we demonstrate that the decreased activity of the melanocortin system induced by asip overexpression does not accelerate the puberty timing but significantly delays early growth of transgenic animals. Once *asip-tg* animals have outperformed a threshold length, close to 2 cm, the transgene rapidly promotes linear growth in absence of obesity by increasing both food efficiency (Godino-Gimeno et al., 2020) and food intake levels (Guillot et al., 2016). Therefore, transgenic animals will become larger and heavier than the WT counterparts, but no obese, during early adulthood. These animals will be easily distinguished after potential escapes, since asip overexpression disrupts also dorsoventral pigment pattern. However, consumer perception will be not affected since transgene will not affect flank pigmentation (Ceinos et al., 2015). Therefore, faster growing will not result in early puberty, considered to be one of the major problems in farmed fish, such as in salmonids, sea bass, flatfishes, cod, tilapia, sea bream and perch. Puberty adversely affects growth, feed utilization, health, flesh quality and welfare (Taranger et al., 2010). Reproductive performance is also affected by asip overexpression since asip-tg zebrafish spawn more eggs but less frequently and their eggs show smaller diameter per contra an increase in larvae body length at hatching is observed. Altogether, results provide sound data to corroborate that the decreased activity of the melanocortin system will be a crucial point in the future of fish aquaculture particularly after the recently approved trading of transgenic fish by the U.S. Food and Drug Association. Therefore, research in transgenic technology of marine species would be potentiated in order to cope next future challenges in animal production.

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Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

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