Increase of fecundity by probiotic administration in zebrafish (*Danio rerio*)

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Abstract

It is well known established that reproduction is sensitive to the state of energy reserves, and that there is a balance between energy homeostasis and fertility. In this view, this study examined the effects of the probiotic *Lactobacillus rhamnosus*, as a feed additive, on zebrafish (*Danio rerio*) fecundity. Ten days of probiotic treatment modulate the gene expression of neuropeptide hormones and metabolic signals, such as *kiss1*, *kiss2* and *leptin* both at the CNS level and at the peripheral level. The increase in fecundity brought about by the probiotic was demonstrated by the higher number of ovulated eggs *in vivo* and by the higher germinal vesicle breakdown rate obtained with the *in vitro* maturation assay. The increase in oocyte maturation was associated with increased transcription of genes coding for signals which induce the maturation phase, such as *lhcr*, *cbr1l*, and *paqr8* genes, concomitant with a decreased transcription of genes coding for local factors which prevent oocyte maturation, such as *tgfb1*, *gdf9*, and *bmp15*. In conclusion, all these findings highlight the action of *L. rhamnosus* both on the endocrine system and at the local level by inducing oocyte maturation. The significance of the results herein obtained underlined the importance of diet in the reproductive process, supporting the hypothesis that feed additives can improve fecundity. Considering that the zebrafish has been clearly established as a vertebrate model for biomedical research, these results support the potentiality of feed additives such as probiotics, frequently used in the human diet, as a new technology to improve reproduction in all vertebrates, including humans.

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Introduction

Over the past 20 years, the zebrafish (*Danio rerio*) has attracted considerable attention as an excellent vertebrate model system for studying genetics and development (Fishman 2001), and more recently, for understanding human disease and for the screening of therapeutic drugs (Penberthy *et al.* 2002, Sumanas & Lin 2004, Keller *et al.* 2006, Shiels *et al.* 2009).

Zebrafish have asynchronous ovaries, containing follicles at all stages of development (Selman *et al.* 1993) as well as mature eggs. The growth and maturation of the oocyte occur over a period of about 10 days (Wang & Ge 2003a, 2003b), and in laboratory conditions, eggs are spawned throughout the year.

In zebrafish, as in all vertebrates, reproduction is regulated by the hypothalamus–pituitary–gonadal axis. The hypothalamus, integrating internal and external stimuli, releases GnRh (Zohar *et al.* 2010). In recent years, it has been established that GnRh transcription and secretion are gated by the state of energy reserves in the organism (Hill *et al.* 2008). The impact of energy status on the reproductive axis is conveyed through

a number of neuropeptide hormones and metabolic signals, such as *kiss1*, *kiss2*, and *leptin*, whose nature and mechanisms of action have begun to be deciphered only in recent years in mammals and, to a lesser extent, in fish (Fernandez-Fernandez *et al.* 2006, Castellano *et al.* 2009, Kitahashi *et al.* 2009). Under the influence of GnRh, the pituitary secretes Fsh and Lh, which act upon the gonads controlling follicle growth and maturation (Nagahama *et al.* 1995, Patino *et al.* 2001). In particular at ovarian level, Lh, through its receptor (*lhcr*), stimulates the production of 17 α -hydroxyprogesterone that is converted (by the action of *cbr1l*) into 17 α ,20 β -dihydroxy-4-pregnen-3-one, the maturation-inducing hormone (MIH) in zebrafish (Patino *et al.* 2001). The binding of MIH to its receptors (*paqr7b* and *paqr8*) activates the maturation processes (Hanna & Zhu 2009).

Oocyte maturation is regulated not only by the endocrine system but also by the auto/paracrine factors such as the transforming growth factor β (Tgfb) superfamily. Tgfb is a large group of oocyte maturation regulatory molecules that include positive factors such as inhibins/activins and negative factors such as *tgfb1*, bone morphogenetic protein 15 (*bmp15*), and growth

and differentiation factor 9 (*gdf9*; Liu & Ge 2007, Clelland & Peng 2009, Tan *et al.* 2009).

This study investigated for the first time the effects of probiotic *Lactobacillus rhamnosus* administration on fecundity and on the endocrine and paracrine control of follicle development in female zebrafish.

According to the WHO definition, probiotic bacteria are 'live microorganisms which when administered in adequate amounts confer a health benefit on the hosts'. *L. rhamnosus* was selected in this study since it is a common constituent of the indigenous microbiota in the intestinal and urogenital tracts of mammals and other vertebrates and is able to balance the intestinal microbiota, inducing host immunomodulation and decreasing the symptoms of a wide range of gastrointestinal disorders (Manley *et al.* 2007, Pant *et al.* 2007, Szajewska *et al.* 2007, Verdenelli *et al.* 2009).

At a physiological level, a 10-day administration of *L. rhamnosus* had an effect on the number of ovulated eggs and on the hatching rate. At a molecular level, this administration affected the expression of molecules that control the balance between energy homeostasis and fecundity as well as the expression of molecules that control follicle maturation at brain, gut, and ovarian levels. Concomitantly with these molecular changes, the *in vitro* maturation assay revealed a higher rate of germinal vesicle breakdown (GVBD) in oocytes from females administered with this probiotic.

Results

Effects of *L. rhamnosus* on metabolic control of reproduction

Figs 1 and 2 show the endocrine control of reproduction as a result of *L. rhamnosus* administration. In zebrafish, both at the gut and at the brain levels, probiotic administration induced a significant increase in the gene expression of leptin, a key hormone in energy homeostasis and neuroendocrine functions (Fig. 1A

Figure 1 (A) *lep* mRNA levels in the gut and (B) in the brain, normalized against β -actin and *gapdh* genes, in females fed on *L. rhamnosus* (PROBIO) and in control females (CTRL). Data are given as mean $\pm$ s.d. ($n=3$). Asterisks denote significant differences from the control group ($P<0.05$) analyzed using Student's *t*-test.

Figure 2 (A) *kiss1*, (B) *kiss2*, and (C) *gnhr3* mRNA levels in the brain, normalized against β -actin and *gapdh* genes, in females fed on *L. rhamnosus* (PROBIO) and in control females (CTRL). Data are given as mean $\pm$ s.d. ($n=3$). Asterisks denote significant differences from the control group ($P<0.05$) analyzed using Student's *t*-test.

and B). This increase was correlated with a significant rise in the brain gene expression of *kiss1* and *kiss2*: these newly discovered neuropeptides can control GnRh expression, representing the link between metabolic and reproductive systems (Fig. 2A and B). The increase in neuropeptides mRNA was concomitant with an increase in *gnhr3* gene expression, as reported in Fig. 2C.

Effects of *L. rhamnosus* on oocyte maturation and fecundity

In zebrafish adults, *L. rhamnosus* administration significantly increased fecundity (Fig. 3). The number of ovulated eggs per day rose significantly as a result of the probiotic treatment, starting from the second day of administration. In particular, the number of ovulated eggs by the treated females reached the maximum on day 2, with eggs production remaining high during the following days (Fig. 3). The induction of follicle maturation was concomitant with a significant decrease in the expression of genes coding for signals involved in the prevention of premature oocyte maturation such as *tgfb1*, *bmp15*, and *gdf9* (Fig. 4A–C). The decrease in *tgfb1* and *bmp15* found in the ovaries of females fed on the probiotic was concomitant with the positive control on *lhcg*, *cbr1l*, and *paqr8* gene transcription (Fig. 5A–C), all of which are important signals involved in the induction of oocyte maturation. The involvement of probiotic on follicle development was also supported by histological studies. The ovarian sections evidenced an increase of vitellogenic follicles in fish treated with probiotic (Fig. 6). The improvement in follicle maturation

Figure 3 The count of number (10 days of treatment) of eggs spawned per day by females fed on *L. rhamnosus* (PROBIO) and by control females (CTRL) was performed every day at 0900 h within 1 h after lights. Data are given as mean $\pm$ s.d. ($n=5$). Asterisks denote significant differences from the control group ($P<0.05$) analyzed using Student's *t*-test.

was also confirmed by the *in vitro* GVBD. Stage IIIb follicles isolated from the ovaries of treated females incubated with MIH showed the greatest rate of GVBD, while probiotic administration alone had no stimulatory effect (Fig. 7).

The molecular data (Figs 4 and 5) and the significant increase in the number of ovulated eggs per day (Fig. 3) observed suggest that *L. rhamnosus* administration as a feed additive promotes oocyte maturation. Moreover, this increase in follicle maturation due to probiotic administration did not negatively affect the egg quality since the embryos produced by females fed with the probiotic showed the highest hatching rate (Fig. 8) and a faster embryo development, characterized by the fact that embryo hatching was brought forward by 4 h (Table 1).

Figure 4 (A) *tgfb1*, (B) *bmp15*, and (C) *gdf9* mRNA levels in the ovary, normalized against β -actin and *gapdh* genes in females fed on *L. rhamnosus* (PROBIO) and in control females (CTRL). Data are given as mean $\pm$ s.d. ($n=3$). Asterisks denote significant differences from the control group ($P<0.05$) analyzed using Student's *t*-test.

Figure 5 (A) *lhcr*, (B) *cbr1*, (C) *paqr8* mRNA levels in the ovary, normalized against β -actin and *gapdh* genes in females fed on *L. rhamnosus* (PROBIO) and in control females (CTRL). Data are given as mean $\pm$ s.d. ($n=3$). Asterisks denote significant differences from the control group ($P<0.05$) analyzed using Student's *t*-test.

Discussion

Reproduction is gated by the state of body energy reserves and is sensitive to different metabolic cues; the neuroendocrine mechanisms responsible for the tight coupling between energy homeostasis and fertility are represented by metabolic hormones and neuropeptides that integrate the hypothalamic center governing reproduction, controlling the expression and release of GnRH (Fernandez-Fernandez *et al.* 2006, Castellano *et al.* 2009, Kitahashi *et al.* 2009, Zohar *et al.* 2010). Thus, full activation of the hypothalamic–pituitary–gonadal axis at puberty and its proper functioning in adulthood critically depend on adequate body energy stores (Hill *et al.* 2008). The identification of the adipose hormone leptin, which signals the magnitude of energy stores to the hypothalamic centers governing reproduction (Casanueva & Dieguez 1999, Goumenou *et al.* 2003), represented an important step toward understanding the mechanisms controlling this interplay.

In this study, for the first time, a possible stimulating role of *L. rhamnosus* on female zebrafish fecundity was investigated. The results suggest that the probiotic may act indirectly by activating a potent metabolic hormone such as leptin. In mammals, it was well established that leptin can regulate reproduction by acting on the hypothalamus (modulating *gnrh* mRNA transcription and release) and on the pituitary (modulating *lh* gene expression and release) as evidenced by several authors (Smith *et al.* 2002, Barb *et al.* 2005).

In this study, the expression of the *lep* gene was significantly induced after probiotic administration in the gut. Concomitantly, we also observed the induction of

Figure 6 Representative micrographs of ovaries from (A) females fed on *L. rhamnosus* and (B) control females. PV, previtellogenic follicles; V, vitellogenic follicles.

lep gene transcription in the brain. These increases were correlated with a higher expression of *kiss1*, *kiss2*, and, in turn, *gnrh3* genes found in the brain of treated females.

The stimulatory role of *L. rhamnosus* on female zebrafish fecundity may be due both to the activation of the endocrine control described above and to the direct action of leptin on the ovary. In mammals, binding of circulating or locally produced leptin to its receptor may lead to direct modulation of ovarian function by controlling the process of gonadal steroidogenesis (Moschos *et al.* 2002). Expression of the leptin receptor (*lepr*) mRNA has been identified in adult human

Figure 7 Rate (%) of GVBD in *Danio rerio* stage IIIb follicles isolated from control groups ovary incubated in L15 (CTRL) and L15+MIH (1 µg/ml, MIH) and in *D. rerio* stage IIIb follicles isolated from treated groups ovary incubated in L15 (PROBIO) and L15+MIH (PROBIO+MIH). Each data represents means of three independent observations $\pm$ s.d. Different letters indicate significant differences on the basis of the Bonferroni post-test ($P < 0.05$).

Figure 8 Hatching rate calculated as the percent of number of viable larvae after hatch divided by the number of fertilized eggs ($n=10$), in triplicate. Results are expressed as means $\pm$ s.d. Statistically significant differences between the experimental groups were tested with Student's *t*-test ($P < 0.05$).

granulosa, theca, and interstitial cells (Cioffi *et al.* 1997, Karlsson *et al.* 1997), as well as in the rat (Zamorano *et al.* 1997), in the porcine (Lin *et al.* 2000) and in the bovine (Spicer & Francisko 1998) ovaries. In zebrafish, Liu *et al.* (2010) have recently found *lepr* expression in the adult fish brain and also in several adult tissues including liver, muscle, and gonads.

The results of our study demonstrated the stimulatory role of *L. rhamnosus* administration as a feed additive in zebrafish (*D. rerio*) follicle maturation. In particular, these findings suggested *L. rhamnosus* administration can control the expression of *tgfb1* and *bmp15* mRNA levels which, in turn, can modulate *cbr11*, *lhcg*, and *paqr8* gene expression. In fact, previous studies on zebrafish have described how treatment with recombinant human *TGFB1* inhibits oocyte maturation reducing Lh activity by decreasing *lhcg* and *cbr11* mRNA levels, and have shown decreased basal and hCG-induced *paqr8* mRNA levels in zebrafish follicles (Kohli *et al.* 2005). Similar results have also been obtained with recombinant human *BMP15*, which also reduced hCG-induced oocyte maturation (Clelland *et al.* 2006). Moreover, the knockdown of *bmp15* in oocytes, by antisense oligonucleotides, increased *paqr8* expression (Tan *et al.* 2009), and the inhibition of *bmp15* expression or function in class IIIa follicles has been found to induce their sensitivity to MIH (Clelland *et al.* 2006, 2007).

The mechanisms through which *L. rhamnosus* stimulates *lep* gene expression both at central and at peripheral levels are still unknown, and further studies are needed in order to clarify this aspect.

In conclusion, the stimulatory role of *L. rhamnosus* administration as a feed additive in zebrafish follicle maturation, fecundity, and egg quality illustrated in this study indicates the great potential of this feed additive for ovarian physiology and presumably for embryo development.

Considering that the zebrafish has been established as a vertebrate model system that is useful for biomedical research, the results obtained here may also have a potential application in reproduction technology for all vertebrates, including mammals.

Table 1 Hatching time of embryos deriving from control or treated females.

	Hatching time (hpf)
Control	48 ± 1
Treated	44 ± 1

Materials and Methods

Animals and probiotic administration

Adult female and male zebrafish (*D. rerio*) were purchased from a local supplier, Acquario di Bologna, Bologna, Italy. The fish were acclimatized to the laboratory conditions and observed for clinical health for 4 weeks prior to start experiments. The animals were kept in 50 l glass tanks under controlled conditions (28 ± 0.5 °C and 14 h light:10 h darkness). The fish were fed on 1.5 g Tetramin (Tetra, Melle, Germany) granules twice a day.

Experiments were performed on two groups: control groups (CTRL) that were fed only on commercial diet and treated groups (PROBIO) that were fed on commercial diet mixed with lyophilized probiotic. The experiment was repeated three times.

The probiotic strain used was *L. rhamnosus* IMC 501, provided by Synbiotec s.r.l. (Camerino, MC, Italy) and supplied in a final concentration in the tank water of 10⁶ CFU/ml as suggested by the producer.

The count of egg spawned output was performed every day at 0900 h within 1 h after lights. The embryos were then transferred to a nursery tank and monitored till the hatching.

After 10 days of treatment, 10 females from each experimental group were killed by a lethal overdose of anesthesia (500 mg/l MS-222 (3-aminobenzoic acid ethyl ester) buffered to pH 7.4; Sigma). The brain, gut, and ovary were sampled and immediately poured in liquid nitrogen and stored at -80 °C until molecular biological analyses were performed. Five gonads from each experimental group were removed and fixed in an aqueous solution of Bouin's fixative.

Procedures were performed in accordance with the Guidelines on the handling and Training of Laboratory Animals by the Universities Federation for Animal Welfare (UFAW) and with the Italian animal welfare legislation (D.L. 116/92).

Histological study

Fixed gonads were prepared for histological examination using standard biological procedures. Gonads were embedded in paraffin wax and sectioned (7 µm) with a microtome (HM 355 Microm). The sections were stained using picroindigocarmine (Fluka no. 57 000) and mounted in Afcylene, and then examined under a light microscope (Nikon 90i) with the software NIS element AR.

RNA extraction and cDNA synthesis

Total RNA was extracted from tissues using Minikit RNeasy (Qiagen) extraction kit following the manufacturer's protocol. Total RNA extracted was eluted in 25 µl RNase-free water. Final RNA concentrations were determined by spectrophotometer, and the RNA integrity was verified by ethidium bromide staining of 28S and 18S ribosomal RNA bands on 1% agarose gel. RNA was stored at -80 °C until use.

Total RNA was treated with DNase (10 UI at 37 °C for 10 min; MBI Fermentas, Glen Burnie, MD, USA), and a total amount of 1 µg RNA was used for cDNA synthesis, employing iScript cDNA Synthesis kit (Bio-Rad).

Real-time PCR

PCR were performed with SYBR Green method in an iQ5 iCycler thermal cycler (Bio-Rad). Triplicate PCR were carried out for each sample analyzed. The reactions were set on a 96-well plate by mixing, for each sample, 1 µl diluted (1/20) cDNA, 5 µl 2× concentrated iQ SYBR Green Supermix (Bio-Rad), containing SYBR Green as a fluorescent intercalating agent, 0.3 µM forward primer and 0.3 µM reverse primer. The thermal profile for all the reactions was 3 min at 95 °C and then 45 cycles of 20 s at 95 °C, 20 s at 60 °C, and 20 s at 72 °C. Fluorescence monitoring occurred at the end of each cycle. Additional dissociation curve analysis was performed, and it showed one single peak in all cases.

β-actin and *gapdh* were used as housekeeping genes in each sample to standardize the results by eliminating variation in mRNA and cDNA quantity and quality. No amplification product was observed in negative control, and no primer-dimer formation was observed in the control templates. The data obtained were analyzed using the iQ5 optical system

Table 2 List of primers used for real-time PCR analyses.

Genes	Forward primer	Reverse primer
<i>lep</i>	AGCTCTCCGCTCAACCTGTA	CAGCGGGAATCTCTGGATAA
<i>kiss1</i>	ACAGACACTCGTCCCACAGATG	CAATCGTGTGAGCATGTCCTG
<i>kiss2</i>	ATTCTCTTCATGTCTGCAATGGTCA	TGCTTTCTCAGGTAAAGCATCATTG
<i>gnrh3</i>	TTAGCATGGAGTGGAAGGAAGGTTG	CTTTCAGAGGCAAACCTTCAGCAT
<i>tgfb1</i>	TCGCTTTGTCTCCAAGGACT	TGCAAGAGAGTTGCCATTTG
<i>bmp15</i>	AGGGTGACCGGATCACTATG	TGCTGCCAGACTTTTATAGCC
<i>gdf9</i>	CGACCACAACCACTCTCTCC	GGGACTGAGTGCTGGATGCC
<i>lhgr</i>	GGCGAAGGCTAGATGGCACAT	TCGCCATCTGGTTCATCAATA
<i>cbr11</i>	TCATGCTCTTCTGCCGATAC	CAAGGGACTTCTTGCTGACA
<i>paqr8</i>	CAACGAGCTGCTGAATGTGT	GGGCCAGTTCAGAGTGAGAC
β-actin	GGTACCCATCTCTGCTCCAA	GAGCGTGGCTACTCCTTCACC
<i>gapdh</i>	GTGTAGGCGTGGAAGTGTTG	GTGTAGGCGTGGAAGTGTTG

software version 2.0 (Bio-Rad). Modification of the gene expression is represented with respect to the control sampled at the same time of the treatment. In Table 2, the sequences of specific primers used were reported.

Follicles in vitro maturation assay

In vitro maturation assays were conducted as described previously (Clelland *et al.* 2006). The ovaries were teased into separate follicles using transfer pipettes (Samco Scientific Corp., San Fernando, CA, USA) without trypsinization. Therefore, stage IIIb follicles were separated according to diameters measuring (Selman *et al.* 1993, Wu *et al.* 2000) with an ocular micrometer under a dissecting microscope and sampled. Stage IIIb follicles derived from both the control and treated groups were incubated in 1 ml L15 medium at 25 °C in 24-well culture plates. For experimental purpose, incubations were carried out separately: stage IIIb follicles isolated from control groups ovary were incubated in L15 (CTRL) and L15+MIH (1 µg/ml) (MIH); stage IIIb follicles isolated from treated groups ovary were incubated in L15 (PROBIO) and L15+MIH (1 µg/ml) (PROBIO+MIH). Maturation was scored after 18 h of incubation. Follicles that underwent GVBD could be identified by their acquired translucency. Each experiment was conducted in four wells with ~20 follicles per well, and all the experiments were repeated three times.

Statistical analysis

Data presented are mean $\pm$ s.d. for the number of experiments. Student's *t*-test was used for comparison between the two experimental groups. $P < 0.05$ was considered significant.

In vitro maturation results were examined by one-way ANOVA followed by the Bonferroni's multiple comparison test, using a statistical software package Prism5 (GraphPad Software Inc., San Diego, CA, USA) with significance set at $P < 0.05$.

Declaration of interest

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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