Supplementary Information for

Transgenerational epigenetic sex determination: environment experienced by female zebrafish affects offspring sex ratio

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Tables S1

Gene	Sequencing primers	RT-qPCR primers
cyp19a1a	F: ATGTAGTTGTTGGGATATAAAAGTG	F: AAAGGGCTCATAACGGCACT
	R: CTTTAAATAAAAAAATCCACCACAACC	R: TGTGTGGTCGATGGTGTCTG
	S: TGTTGGGATATAAAAGTGT	
	F: TTTGTTTTGTTTTGAATGGGGATT	F: AAGTCCTGCAATGTCCAGCA
dmrt1	R: TCTCCCACCAACTCTAAAATACACAAATT	R: GGAGGAGTATGGAGTGGGCT
	S: TGTTTTGAATGGGGATTA	
	F: AGGAGAGAGGGGGAAAGATAG	F: CATCGCTAAGCCAAGTAGCC
foxl2a	R: AATTTTTTCCCTCTTCAAATATAATTTCAC	R: AGGGCCCAATTATTCACTCC
	S: GAGGGGAAAGATAGG	
actb1		F: ATGAAGTGCGACGTGGACAT
		R: CTGCATACGGTCAGCAATGC
eeflala		F: ACCCTTCCCGTGGATCA
		R: GTGGGTTTCAGACGCACA
rpl13a		F: GGGAAAGGAAAAGACCCACA
		R: CGCTCGGCCTTCAGTT
sequencing, sequences in bold are biotinylated in 5')		

Table S1: Primers used for bisulfite sequencing and qPCR analyses (F forward; R reverse; S

Figure S1: DNA methylation levels (%) of *cyp19a1a*, *dmrt1* and *foxl2a* in gonads of female (F) and male (M) zebrafish from the different experimental conditions: controls (C), exposed to cadmium (Cd) and exposed to high temperature (HT).



<u>Figure S2</u>: Box plots of the DNA methylation levels of *cyp19a1a* in gonads of females (F) and males (M) zebrafish from the different experimental conditions: controls (C), exposed to cadmium (Cd) and exposed to high temperature (HT). For each sex, means designed with different letters are significantly different (P<0.05, Tukey's HSD test or Dunn test with Bonferroni correction). Fore each condition, data from all generations were combined (n = 10 - 24).





<u>Figure S3:</u> Mean (\pm SE, n = 4) fish density determined at each generation (F0 to F3) at 170 dpf. Four aquaria were used per condition (C, control, blue; Cd, Cadmium, red). Means designed

<u>Figure S4:</u> DNA methylation level (%) of the gonadal *cyp19a1a* promotor. The mean methylation level determined in the first two CpG sites (CpG1-2) was highly correlated with the mean methylation level determined using all the six CpG analyzed (CpG1-6).



<u>Figure S5:</u> Pyrograms obtained from gonads of male or female zebrafish for cyp19a1a (A), dmrt1 (B) or foxl2a (C). For foxl2a, some polymorphisms were observed both in females and males from position 10 to 14.



