### Estrogen related receptor is required for the testicular development and for the normal sperm

### axoneme/mitochondrial derivatives in Drosophila males

Snigdha Misra<sup>1,2</sup>, Anuj Kumar Pandey<sup>1</sup>, Snigdha Gupta<sup>1,2</sup>, Ajay Kumar<sup>1,2</sup>, Priyanka Khanna<sup>1</sup>, Jai shankar<sup>3</sup>, Kristipati Ravi Ram<sup>1,2§</sup>

<sup>1</sup>Embryotoxicology Laboratory, Environmental Toxicology Group, CSIR-Indian Institute of Toxicology Research (CSIR-IITR), MG Marg, Lucknow, Uttar Pradesh, 226001, India. <sup>2</sup> Academy of Scientific and Innovative Research (AcSIR), CSIR-IITR campus, Lucknow, Uttar Pradesh, 226001, India. <sup>3</sup>Electron Microscopy Facility, CSIR-Indian Institute of Toxicology Research (CSIR-IITR), MG Marg, Lucknow, Uttar Pradesh, 226001, India. <sup>4</sup>Uttar Pradesh, 226001, India.



**Figure S1.** Confirmation and determination of the extent of knockdown of ERR through quantitative real time PCR based analysis of ERR transcript levels. The levels of ERR transcripts were reduced to 3.5 folds in miRNA based knockdown driven by testes-GAL4 (\*p<0.05, represented as ERR-miRNA), while 7.5 folds reduction was observed in knockdown males produced from UAS-ERR-TRiP (\*\*p<0.01, represented as ERR-TRiP), when compared to their controls. The  $\Delta$ ct values were determined through normalization against Act-5c, which was used as an internal control for the quality of the template.



**Figure. S2.** Quantitative real time PCR based analysis of ERR transcript levels in males with systemic knockdown of Sox100B. The levels of ERR transcripts in male pupae where Sox100B dsRNAi was driven through Act5c-GAL4/ Tb were comparable to control. The  $\Delta$ ct values were determined through normalization against Act-5c, which was used as an internal control for the quality of the template.

# **Supplementary methods**

Stock No.	Expression Pattern	
BL12608	Late spermatogonia, early	
	spermatocytes, cyst cells, pigment	
	cells, muscle sheath	
BL13134	Early spermatocytes, cyst cells, muscle	
	sheath	
BL12772	Transient in early spermatocytes, cyst	
	cells, pigment cells, muscle sheath	
BL6983	Adult male accessory gland, testes	
	sheath, cyst cells, larval salivary gland,	
	and adult female columnar follicle	
	cells	
BL6987	Adult ovary, amnioserosa, adult male	
	accessory glands and seminal vesicles.	
BL6989	Accessory glands, cyst cells and germ	
	cells	
BL1947	Male accessory glands	
Bam-Gal4	Germ cell/ Spermatogonia	
Tj-Gal4	Early cyst cells	
Nos-Gal4	Early germline cells	
Act5c-GAL4	Systemic expression	

Table S1. Expression patterns of the GAL4s used

## Conditions applied for Real time PCR

1 cycle of pre incubation (50°C for 2 min and 95°C for 10 min) followed by 40 amplification cycles (95°C for 15 seconds, and 60°C for 1 min) and melting curve detection (95°C for 5 sec, 60°C for 1 min). Experiments were preformed in duplicate for each sample for ten genes (please see Table S2 for primer sequences). The gene expression data were analyzed using the comparative  $2^{-\Delta\Delta}CT^1$  considering Act5c as the internal control. All results are given as Mean±Standard Error of the mean.

Table S2. Primers	used for R	Real time PCR
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Gene	Primer Sequences
Armadillo ( <i>arm</i> )	GATGAGGATCAGGTGGTAGTTTC
	CCATCTGAGGGCTGTTCATAAT
hopscotch ( <i>hop</i> )	GACATACCCTTTCTACCCTTTCC
	CCCTTGTAGACAGTGCCATAAT
always early (aly)	AATCTCCAGCGTCGCTATTC
	TCCTCGTCCTGCTTCTGATA
meiotic arrest (mia)	GCGTTCTCTATCCGCGATATT
	CCAGCCTCGTTTAGCAGATT
Bruce	CAGATGGCACCGTGGAAATA
	CCTATCCAAGTTCCGGCAATAA
fuzzy onions ( <i>fzo</i> )	GTCCTTCAATGTCTCTCCATACC
	CAATCCAGGCCGTAGATTAGTT
canonball (can)	ACACGTCGCAATCTCTTACTT
	CTCCTCCTTGTGCTTGTTCT

bag of marbles (bam)	CTGTTCATCGCCCAGAGATAC
	CGGGAAATAGGTCTCTGGATTG
benign gonial cell neoplasm	CTATGCCGTGGACACTAGAAAG
(bgcn)	GGAAGCTCTGTGCCGAAATA
Actin (Act-5c)	CGATTTGACCGACTACCTGATG
	GCACAGCTTCTCCTTGATGT
Sox100b	AGGGTCATGTCCAAACAGTATC
	GGGCTTCTTATCACTGTCCTTTA
ERR	CACAGCGCATGGAAAGAATATC
	GAACTCTGATCATCCAGCAGAA

## Immunoflorescence and confocal microscopy

The testes/ gonads from the male reproductive tract of control and knockdown males were dissected in physiological saline, and transferred separately to phosphate buffer saline (PBS), with 0.3% Triton-X (PBS with Triton-X, referred to as PBX). At the end of dissections, PBX was replaced with 4% paraformaldehyde prepared in 1x PBS, and the tissues were incubated for 1h. The tissues were then washed with 0.3% PBX thrice, for 15min each. The blocking solution comprising 3% bovine serum albumin (BSA) in 0.1% PBX was added, and tissues were incubated for 30min. The primary antibody was added, and tissues were incubated overnight at 4°C (For details of the name/ host/ dilutions of the primary antibodies used, please refer to Table S3). Subsequently, tissues were washed thrice, for 15min each with 0.3% PBX, and incubated at room temperature for 2h in secondary antibody (Alexa Fluor® 488 Rabbit Anti-Mouse; Alexa Fluor® 488 Goat Anti-Rabbit, Life technologies, USA), diluted in 0.1% PBX at 1:200 folds in dark. Tissues were washed thrice, for 15min each with 0.3% PBX prior to mounting them in vectashield (Vector labs, USA). Nuclei were stained with DAPI, which is pre-incorporated in the vectashield. The fluorescence was visualized under confocal microscope (TCS SPE, Leica, Germany), using in-built settings for the DAPI (excitation wavelength 340-380nm), and FITC (excitation wavelength 450-490nm). A minimum of five independent immunostaining batches, each having at least 5 replicates (pairs of testes) were analyzed in each case.

Antibody / Cat. No	Host	Dilution	References
bam-c DSHB-C1-252	Mouse	1:10	2
vasa	Rat	1:40	3
eya10H6-c DSHB-C1-385	Mouse	1:25	4
3A9 (323 or M10- 2)-c (α-spectrin) DSHB-C1-86	Mouse	1:25	5
Anti-Sox100B AAS77513C	Rabbit	1:25forImmunofluorescence1:1000 for western	M/s Antibody Verify, USA

Table S3. Primary antibodies used in this study

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- 4. Bonini, N. M., Leiserson, W. M. & Benzer, S. The eyes absent gene: genetic control of cell survival and differentiation in the developing Drosophila eye. *Cell* **72**, 379-395 (1993).
- 5. Dubreuil, R. R., Maddux, P. B., Grushko, T. A. & MacVicar, G. R. Segregation of two spectrin isoforms: polarized membrane-binding sites direct polarized membrane skeleton assembly. *Mol. Biol. Cell* **8**, 1933-1942 (1997).