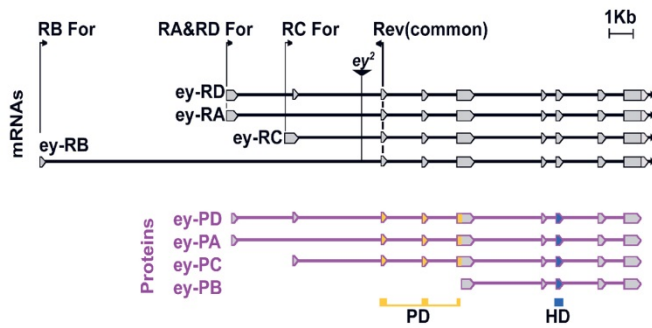
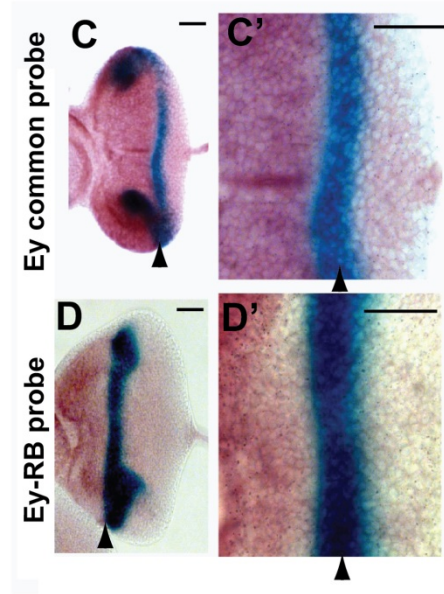
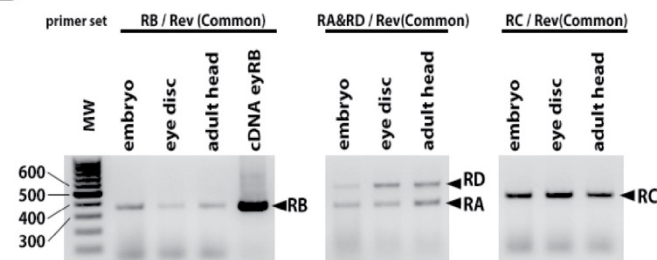


## SUPPLEMENTARY FILE

**A**

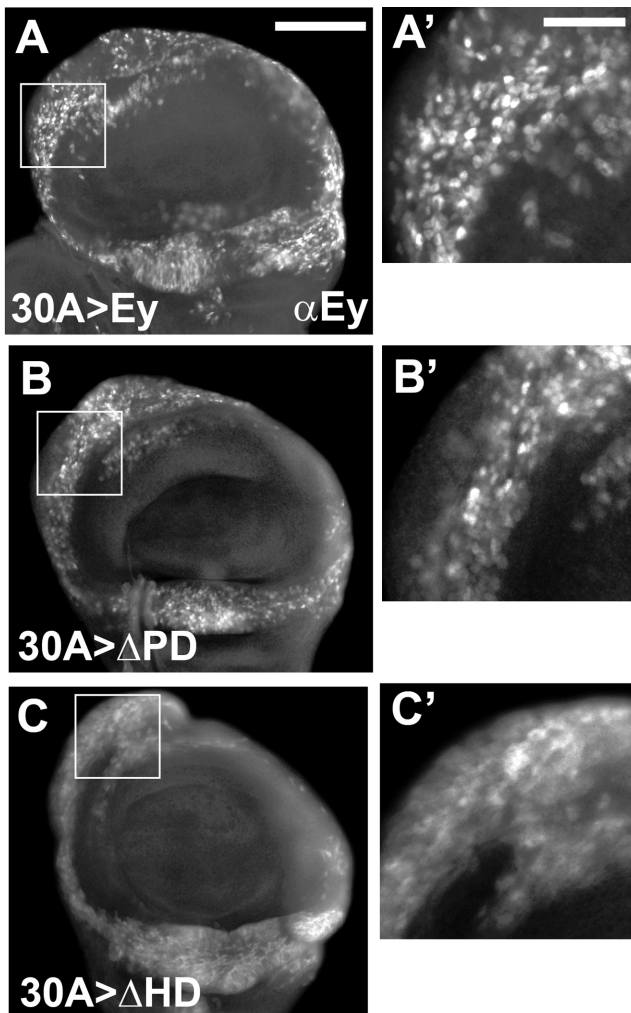


**B**



**Supplemental Figure1. Ey HD type isoform express in the anterior domain in the developing eye disc.** (A)Schematics of alternative spliced Ey. Exon (gray box) and intron (black line) structures are shown. Three forward primers used for RT-PCR are indicated with black forward oriented arrows. Common reverse primer position is indicated on reverse oriented arrow. Alternatively produced four Ey proteins are shown in below Ey-PA to Ey-PD in magenta color. Ey-PB protein contains only Homeodomain (highlighted with blue), other three alternatively produced Ey proteins contain both Paired Domain (highlighted with yellow) and Homeodomain. (B) RT-PCR analysis shows mRNA levels of ey alternative spliced products. Total RNA were isolated from O/N embryo collection, the 3<sup>rd</sup> instar eye discs, and adult heads were reversed transcription. All Flybase reported four Ey isoforms were detected in this analysis. (C and D) *in situ* hybridization with ey probe in the 3<sup>rd</sup> instar eye discs. Ey-RB (HD type isoform,

brown staining in D and D') expresses anterior to the MF (black arrows), this expression pattern is identical to the all other ey isoforms expression pattern (ey common, dark purple in C and C'). The morphogenetic furrow labelled with the expression of dpp-lacZ (blue). The scale bars are 50µm.



**Supplemental Figure 2. Expression levels of Ey deletion constructs.** (A) 30A>Ey wing discs stained with  $\alpha$ -Ey. (B and C) Expression of Ey deletion constructs  $\Delta$ PD and  $\Delta$ HD under 30A gal4 driver at identical condition to 30A>Ey. Images were taken at x20

magnification in (A-C). (A'-C') Photoshop enlarged images in boxed area in A-C are shown. The scale bars are 50µm in A-C, and 150µm in A'-C'.

## SUPPLEMENTARY MATERIALS AND METHODS

### RT-PCR

Total RNA was prepared from embryo using TRIZOL reagent followed by manufacturer's protocol. Reaction performed as described before (Tanaka-Matakatsu et al., 2009). The first exon of the Ey-RB mRNA used to generate in situ probe. The last 4 exon sequences were used for ey common in situ probe. Primer sequences are shown in the primer list.

### PRIMER LIST

Experiment	Primer	Sequence
Ey RT-PCR	Ey-RA&RD Forward	5'- TTCGCACGGCGTGCGTTTGG-3'
	Ey-RB Forward	5'- TCCAATCGATACTACAAAATACC-3'
	Ey-RC Forward	5'- GAATTCCAAGTACAAACTGAC-3'
	Ey-common Reverse	5'- CGAGAATTTTGCTCACACATCC-3'
Ey-RB in situ probe	Ey-RB unique reverse	5'-ATTTATATTACGAATGG-3' Used Ey-RB forward primer
EMSA	Ey2 (WT)Forward	5'- GATACAGTTTTCCAGCTCATTGCTTGT AATTGGGCACCAA-3'
	Ey2(WT) Reverse	5'-TTGGTGCCCAATTACAAGCAAATGAGCT GGAAACTGTATC-3'

construct	Primer name	Primer Sequence or origin of subcloning fragment
pUAST -Ey-PB	EyPB-XhoI Forward	5'- CTTCTT CGAGCAAATGCAGACAGCC-3'
	EyPB-XbaI Reverse	5'- CGCCTAGTCTAGACTAGACCCACGGTG-3'

Ey/pET28b		Ey coding sequence from pUAST-Ey was subcloned into pET28b vector
PD/pET28c	EyPD HindIII Forward	5'-CGGCCGAAGCTTTCAAATGTTTACATTGC-3'
	EyPD XhoI Reverse	5'-TCTAGGATCTCGAGCGTGCTTTGC-3'
HD/pET28b	EyHD HindIII Kozak Forward	5'-CGCCGCAAGCTTGCAAAAATGGAGGATGATC-3'
	EyHD XhoI Reverse	5'-AGTCATCTCGAGCTATGGTGTTCCTTCGC-3'
$\Delta$ PD/pET28b	$\Delta$ PD Forward	5'-GCGCAAAGGAGCAGCAAAGCACGGG-3'
	$\Delta$ PD Reverse	5'-CTGCTCCTTTTGCGCCAAAGGCCTTCC-3'
$\Delta$ HD/pET28b	$\Delta$ HD Forward	5'-AAGCTGCGAAACCAGCGAAGAACACC-3'
	$\Delta$ HD Reverse	5'-CGCTGGTTTCGCAGCTTCTTTCTTTTAG-3'
$\Delta$ PD $\Delta$ HD /pET28b		N-terminal $\Delta$ PD fragment was subcloned into $\Delta$ HD/pET28b plasmid
EyN /pGEX4T2	Ey5' BglII Forward	5' - CGCAGATCTATGTTTACATTGCAACC-3'
	EyN BglII Reverse	5' - CGCAGATCTGGAATTTGGTGTTCCTTC-3'
PD /pGEX4T2	PD $\beta$ -GST-BglII Forward	5' - CATGGCTAGATCTGGTCACAGTGGAGTAAATC-3'
	EyPD XhoI Reverse	5' - TCTAGGATCTCGAGCGTGCTTTGC-3'
N-PAI /pGEX4T2 #3		EcoRI fragment from N-term-PD/pGEX4T2
RED /pGEX4T2 #6		BamHI-XhoI fragment from PD/pGEX4T2
HD /pGEX4T2	EyHD-GST BglII Forward	5'-GCCAGAGATCTCAAATGGAGGATGATC-3'
	EyHD XhoI stop Reverse	5'-AGTCATCTCGAGCTATGGTGTTCCTTCGC-3'
HD h1-2 /pGEX4T2 #12		EcoRI digest from HD/pGEX4T2 (120bp fragment) and subcloned into pGEX4T2 at EcoRI site
HD h3 /pGEX4T2 #20		EcoRI/XhoI digest from HD/pGEX4T2 and subcloned into pGEX4T2