S60-6 Epigenetic regulation of winged eye capable of inducing eye to wing transformation in Drosophila

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In Drosophila, the developmental program of imaginal discs is determined. We previously identified winged eve (wge), the overexpression of which induced ectopic wings on the head. WGE localized on polyten chromosomes, and almost all Posterior sex comb (PSC) that one of Polycomb group (PcG) protein were coincident with some WGE binding sites. The analysis of genetic interaction between wge and Psc, Polycomb or Pleiohomeotic, revealed that wge antagonizes PcG genes. Surprisingly, wge also antagonizes trx (trithorax) that one of trithorax group (trxG) genes, which generally antagonize PcG genes. The phenomenon is also supported by effects of wge mutation on PcG / trxG response element (PRE / TRE) -mediated gene regulations to know whether wge maintains either silenced (PcG like) or active (trxG like) transcriptional states. wge is involved in maintenance of active transcriptional states on Fab-7, a genetically identified PRE / TRE of the bithorax-complex necessary to regulate transcription of the *Abdominal-B*, whereas, wge is involved in maintenance of silenced transcriptional states on PRE of vestigial, a target of Wge during eye to wing transformation. These results suggest that wge is the new type of epigenetic regulator different from PcG and trxG.