# **Drosophila** *Lyra* **Mutations Are Gain-of-Function Mutations of** *senseless*

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## ABSTRACT

The *Lyra* mutation was first described by Jerry Coyne in 1935. *Lyra* causes recessive pupal lethality and adult heterozygous *Lyra* mutants exhibit a dominant loss of the anterior and posterior wing margins. Unlike many mutations that cause loss of wing tissue (*e.g.*, *scalloped*, *Beadex*, *cut*, and *apterous-Xasta*), *Lyra* wing discs do not exhibit increased necrotic or apoptotic cell death, nor do they show altered BrdU incorporation. However, during wing disc eversion, loss of the anterior and posterior wing margins is apparent. We have previously shown that *senseless*, a gene that is necessary and sufficient for peripheral nervous system (PNS) development, is allelic to *Lyra.* Here we show by several genetic criteria that *Lyra* alleles are neomorphic alleles of *senseless* that cause ectopic expression of SENSELESS in the wing pouch. Similarly, overexpression of SENSELESS in the wing disc causes loss of wing margin tissue, thereby mimicking the Lyra phenotype. *Lyra* mutants display aberrant expression of DELTA, VESTIGIAL, WING-LESS, and CUT. As in *Lyra* mutants, overexpression of SENSELESS in some areas of the wing pouch also leads to loss of WINGLESS and CUT. In summary, our data indicate that overexpression of SENSELESS causes a severe reduction in NOTCH signaling that in turn may lead to decreased transcription of several key genes required for wing development, leading to a failure in cell proliferation and loss of wing margin tissue.

genesis have mutant phenotypes in which there are missing sectors of the wing margins. These include Tsakonas *et al.* 1999), which have been shown to play of nearby wing surface tissue. The presence of a duplicaimportant roles during fruitfly development. It is known tion of the chromosomal region carrying the wild-type that many of these so-called "wing scalloping" muta- $Lyra$  locus in a  $Lyra<sup>1</sup>$  background does not suppress the tions, including *cut* and *vestigial*, are caused by excessive Lyra phenotype (ABBOTT and SPREY 1990). Hence, the cell death in the prospective wing margins of late larvae dominant phenotype is not due to haploinsufficien cell death in the prospective wing margins of late larvae dominant phenotype is not due to haploinsufficiency.<br>
following a period of apparently normal development This suggested that Lyra is a gain-of-function mutation (FRISTROM 1968, 1969). In *Lyra* mutants, although there and is likely to be a neomorphic allele (MULLER 1932) is a significant reduction (10–20%) of the number of characterized by spatial and/or temporal misregulation is a significant reduction (10–20%) of the number of characterized by spatial and/or temporal misregulation cells in the adult wing, no evidence of apoptotic or  $\sigma$  expression of a gene product. In addition, the results cells in the adult wing, no evidence of apoptotic or of expression of a gene product. In addition, the results<br>necrotic cell death was found by transmission electron of clonal analysis with wild-type clones in  $Lyra/$  flies microscopy, acridine orange, or trypan blue staining in indicate a non-cell-autonomous function (ABBOTT<br>third instar and pupal discs (ABBOTT 1986). These and 1986), suggesting that the *Lyra* mutation may affect third instar and pupal discs (ABBOTT 1986). These and<br>other data have led to the suggestion that *Lyra* may<br>affect more fundamental parameters of cell growth and<br>specification. The *Lyra* mutant has therefore been of<br>inte

LIKE *Lyra*, many mutations that affect wing morpho-<br>
<u>J</u> genesis have mutant phenotypes in which there (COYNE 1935; ZHIMULEV and FELDMAN 1982; LINDSLEY<br>
e missing sectors of the wing margins. These include and ZIMM 1992). *vestigial*, *Notch*, *Delta*, *cut*, *apterous*, and others (Jack *et* a regular and predictable pattern of loss of the anterior *al.* 1991; Cohen *et al.* 1993; Kim *et al.* 1996; Artavanis- and posterior wing margins along with a small amount This suggested that *Lyra* is a gain-of-function mutation

does not account for margin loss in *Lyra* adult wings. Second, although *Lyra* mutants do not form anterior *Corresponding author:* Hugo J. Bellen, Department of Molecular and and posterior wing margins, there is a normal dorsal/ Human Genetics, Rm. T634, MS BCM235, Baylor College of Medicine, Houston, TX 77030. E-mail: hbellen@bcm.tmc.edu ventral compartment boundary as well as a zone of non-<sup>1</sup> These authors contributed equally to this work. proliferating cells (ZNC) along the "de facto" *Lyra* wing

margin. Third, a monoclonal antibody, which binds the *sens<sup>E2</sup>* red e/*TM3*, Sb e (H. Irick)<br>unidentified E1C antigen expressed in the largel wing *sens<sup>E53</sup>* red e/*TM3*, Sb e (H. Irick) unidentified E1C antigen expressed in the larval wing<br>margin precursor, enabled them to demonstrate that<br>an effect of *Lyra* on anterior and posterior wing margins<br> $\frac{\text{gens}}{\text{gens}}^{\text{SHS}} \text{red } e / TM3, \text{Sb } e \text{ (H. Irick)}$ <br>an effe is apparent early in the third larval instar. Fourth, mar- *vgBE-lacZ* and *vgQE-lacZ* (S. Carroll, Madison, WI) gin rescue experiments using clonal analysis showed<br>that wild-type Lyra is not required for bristle develop-<br>ment *per se*. Fifth, further analysis of shape and position<br> $TM6B$  (G. Mardon. Houston) with UAS-lacZ and UAS-sen of clones indicated that the wing margin, defined as a *C6)* (Noto *et al.* 2000) lines.<br>set of several rows of cells along either side of the dorsal/**Immunohistochemistry and antibody staining:** X-Gal stainset of several rows of cells along either side of the dorsal/<br>
ventral boundary, plays an important role in wing mor-<br>
phogenesis. This observation presaged the current para-<br>
digm that interactions among a number of gene ucts expressed in the margin region, often acting across S. Cohen), mouse anti-CUT (1:30, BODMER *et al.* 1987), rabbit the compartment boundary, serve to organize wing de-<br>anti-SCUTE (1:100, a gift from G. Panganiban), mo

*senseless* gene plays a key role in peripheral nervous Cy3, respectively). Biotinylated secondary antibodies were<br>system (PNS) development Its loss causes a severe loss from Vector Laboratories (Burlingame, CA) and were u system (PNS) development. Its loss causes a severe loss from Vector Laboratories (Burlingame, CA) and were used<br>of external peripheral sensory organs in embryos, and according to the manufacturer's instructions. Confocal i *Lyra* mutants, SENSELESS is ectopically expressed in *In situ* hybridization: *string* cDNA (a gift from B. Edgar) was the wing pouch and that this ectopic expression goes used as a template for digoxigenin-labeled RNA pr the wing pouch and that this ectopic expression goes used as a template for digoxigenin-<br>hand in bond with localized loss of DELTA MESTICIAL labeling kit; Roche, Indianapolis). hand in hand with localized loss of DELTA, VESTIGIAL, WINGLESS, and CUT expression. We therefore propose that the loss of wing margin in *Lyra* mutants is due <br>to ectopic expression of SENSELESS, which also causes RESULTS loss of expression of genes essential for wing margin *Lyra* **alleles are gain-of-function alleles of** *senseless***:** and wing development, including but not limited to SALZBERG *et al.* (1994, 1997) showed that the *senseless wingless*, *vestigial*, and *cut.* We suggest that it is this loss gene affects the development of the PNS and reportof normal pattern of gene expression in the developing ed three alleles: two ethyl methanesulfonate (EMS)-<br>wing margin that leads to failure of differentiation of induced alleles ( $\text{sens}^{M256}$  and  $\text{sens}^{M255}$ ) and on wing margin that leads to failure of differentiation of induced alleles ( $sens^{M256}$  and  $sens^{I225}$ ) and one induced by the sensory organ precursors in the wing discs and, later,  $P$ -element dysgenesis ( $sens^{I225/4}$ ). The *the sensory organ precursors in the wing discs and, later,* to the loss of wing margin tissue seen in *Lyra* wings. with these alleles was mapped by meiotic recombination However, the tissue loss because of ectopic expression to 3-40.5, where Lyra<sup>1</sup> maps (NoLo *et al.* 2000). As shown of SENSELESS in the wing margin is quite different in Table 1, *senseless* mutations *M256* and *I235* failed to from the excess PNS organs (bristles) that arise from complement the lethality of *Lyra<sup>1</sup>*, which is associated overexpression of SENSELESS in some areas of the fly with deletion 70A2-3;70A5-6 (LINDSLEY and ZIMM integument (Nolo *et al.* 2000). **deletion** 1992). The *Lyra<sup>1</sup>* deletion uncovers three essential com-

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- *yw; Df(3L)1228/4 P[lacZ, w<sup>+</sup>]/TM3, Ser Sb (SALZBERG <i>et al.* plement the *Lyra<sup>1</sup>* deficiency.
- 
- 
- 
- 5*lArB}A77.1M3*/*TM3, Sb ry*
- 

*PM6B* (G. Mardon, Houston) with *UAS-lacZ* and *UAS-sens (C5, C6)* (Noto *et al.* 2000) lines.

the compartment boundary, serve to organize wing de-<br>velopment (for review see BROOK *et al.* 1996).<br>Here we show that *Lyra* mutations correspond to neo-<br>morphic/gain-of-function mutations of *senseless*. The organism org

plementation groups: *l(3)70Aa*, *l(3)70Ab*, and *l(3)70Ac* (ZHIMULEV and FELDMAN 1982). Unfortunately, these MATERIALS AND METHODS mutants no longer exist. However, Holly Irick and Peter **Drosophila stocks:** The wild-type stock was Canton-S<br>(Bloomington Stock Center). The other stocks used in this<br>work are as follows:<br>Recause the  $Df(3L)BK10$  chromosome was marked with Because the *Df(3L)BK10* chromosome was marked with *Lyra<sup>l</sup>* we suspected that some of their lethals were in yw; P[lacZ, w<sup>+</sup>]64A sens<sup>M256</sup> th st cu sr e<sup>s</sup> ca/TM6, Hu P[w<sup>+</sup>, abdA-<br>lacZ] e Tb ca (SALZBERG et al. 1994)<br>w; P[lacZ, w<sup>+</sup>]64A sens<sup>2255</sup> th st cu sr e<sup>s</sup> ca/TM6, Hu P[w<sup>+</sup>, abdA-<br>failed to complement the Lyra<sup>1</sup> Df(3 some and isolated 8 lethal mutations that failed to com- *lacZ] eTb ca* (Salzberg *et al.* 1994)

1997)<br>
1997) As shown in Table 1, complementation tests for the<br>  $Ly^1/TM3$ , Sb (ABBOTT 1986)<br>  $Lyra^{\cos\theta}/TM3$ , Sb (P. Heizler, Strasbourg, France)<br>
1997 Ly / IND, 30 (ABBOTT 1960)<br>
Lyra<sup>8x67</sup>/TM3, Sb (P. Heizler, Strasbourg, France)<br>
P(hsneol l(3)neo19 (SPRADLING et al. 1999)<br>
alleles of *senseless* referred to as complementation group *l*(3)70Ad in FlyBase. This complementation group is pre*sensE1 red e*/*TM3, Sb e* (H. Irick, Bloomington, IN) sumably allelic to one of the lost *l(3)70A* complementa-

### **TABLE 1**

**Complementation tests**

	M256								1235 E1 E2 E53 E54 E58 E69 1228/4	$Ly^1$	$Ly^{Sx67}$	$delta^{130}$	P257
M256													
I235													
E1													
E2													
E53													
E54													
E58													
E69													
1228/4													
$\begin{array}{l} Ly^1\\ Ly^{8x67}\\ \end{array}$	$^{+}$	$^{+}$	$^{+}$	$^{+}$	$^{+}$	$^{+}$	$^{+}$	$^{+}$	$^{+}$	<b>NA</b>			
$delta^{130}$											$^+$		
P <sub>257</sub>	$^{+}$	$^{+}$	$^+$	$^+$	$^{+}$	$^{+}$	$^{+}$	$^+$		$^{+}$	$^{+}$		

 $M256$ , sens<sup> $M256$ </sup>, 1235, sens<sup> $225$ </sup>, E1, sens<sup>E1</sup>, E2, sens<sup>E2</sup>, E53, sens<sup>E53</sup>, E54, sens<sup>E54</sup>, E58, sens<sup>E58</sup>, E69, sens<sup>E69</sup>, 1228/4, *Df(3L)1228*/*4*; *delta130*, an imprecise excision #130 of *P{ry[*<sup>1</sup>*]* 5*lArB}A77.1M3*; *P257*, *P{hsneo}l(3)neo19*; NA, not applicable.

tion groups described by ZHIMULEV and FELDMAN function, regulatory mutations of *senseless*. First, the

loss-of-function allele of *senseless*. First, *Lyra<sup>1</sup>*/*sens* mutant embryos display either no loss of PNS neurons or a very showing that *Lyra<sup>SX67</sup>* is associated with *senseless*. Fourth, loss of wing margin phenotype in heterozygous flies phic/gain-of-function mutations of *senseless.* (*sens*/1), indicating that haploinsufficiency of *sens* does The complementation data, combined with data from not cause the Lyra phenotype, in agreement with the anti-SENSELESS-stained embryos, and the analysis of observations of LINDSLEY *et al.* (1972). Third, a second, the severity of the phenotypes in which there is loss of independently generated dominant allele of *Lyra*, PNS neurons in embryos support the allelic series that *LyraSX67*, interacts additively with *Lyra1* to produce a more is shown in Table 2. We propose to keep the name severe margin loss, but complements all *senseless* alleles *senseless*, which refers to the loss of PNS organs, and to (Table 1). Since *Lyra* mutations are dominant and their refer to *Lyra* alleles of *senseless* as neomorphic/gain-ofphenotype is not caused by haploinsufficiency of *sense-* function alleles that affect the wing margin. *less*, they are presumed to be either antimorphic (domi- **SENSELESS is ectopically expressed in the wing** nant negative) or neomorphic (gain of function) in **pouch of** *Lyra* **mutants:** We have previously shown that nature (Muller 1932). An antimorphic nature is most *senseless* is expressed in the sensory organ precursors unlikely since duplications of the chromosomal region (SOPs) of the embryonic and adult PNS (NoLo *et al.* do not ameliorate the phenotype associated with *Lyra<sup>1</sup>* 2000). In wild-type imaginal wing discs, SENSELESS is (Muller 1932; Abbott and Sprey 1990). Furthermore, expressed in the SOPs along the presumptive wing marthe *Lyra<sup>1</sup>/sens* mutants do not display obvious defects in the PNS, as would be expected if *Lyra* was a dominant- of SENSELESS is altered in wing discs of *Lyra<sup>1</sup>* and negative allele of *senseless*. We therefore conclude that *Lyra<sup>Sx67</sup>* we carried out *in situ* hybridization and immuno-*Lyra* mutations are neomorphic mutations. histochemical staining with antibodies raised against the

of the *Lyra* mutations, that is, that they are gain-of- in addition to the expression in SOPs, we observe a

(1982). The *sens* alleles were designated *E1*, *E2*, *E53*, cominant phenotype associated with  $L y r a^1$  could not *E54*, *E58*, *E64*, *E69*, and *E87.* (H. Irick and P. Cherbas, be recombined onto a *senseless* mutant chromosome, personal communication to FlyBase). All but *E64* and indicating that both mutations map at the same site and *E87* are still available. that the Lyra phenotype may be breakpoint dependent. The *Lyra<sup>1</sup>* deficiency *in trans* to other *senseless* alleles **11 Second**, molecular analyses show that the distal causes lethality but these mutant embryos do not display breakpoint of  $Df(3L)Ly<sup>1</sup>$  affects a genomic fragment that a severe loss of neurons as typically seen in homozygous contains the 3' end, including the 3' untranslated re*senseless* mutations (Nolo *et al.* 2000). Indeed, the follow- gion, of the *sens* gene (data not shown). Third, an X-raying observations suggest that the *Lyra*<sup>1</sup> mutation is not a induced revertant of *Lyra*<sup>SX67</sup>, *Lyra*<sup>SX67R12</sup>, is homozygous /*sens* mutant lethal and fails to complement all the *senseless* alleles, subtle loss, indicating that the deficiency associated with both *Lyra* mutations cause ectopic expression of SENSE-*Lyra*<sup>1</sup> does not result in the lack of the *senseless* gene LESS in wing imaginal discs (see below). These observaproduct. Second, none of the *senseless* alleles causes a tions strongly indicate that the *Lyra* alleles are neomor-

/*sens* mutants do not display obvious defects gin (Figure 1A). To determine if the expression pattern The following data support the neomorphic nature full-length SENSELESS protein. As shown in Figure 1B,

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**Characteristics of different** *senseless* **mutations**



ND, not determined; NA, not applicable;  $+++$ , wild-type levels;  $++$ , mildly reduced;  $+$ , strongly reduced;  $\pm$ , extremely reduced; \*, unable to establish, as the *Ly<sup>x67</sup>* carries at least one other early lethal mutation.

broad band of SENSELESS-positive cells in the anterior GAL4 rather specifically in the wing disc. Most and and posterior regions of the wing pouch of  $Lyra<sup>1</sup>/+$ mutant wing discs. As shown in Figure 1D, the ectopic sion of GAL4 are lethal in the presence of *UAS-senseless*. expression of SENSELESS surrounding the anterior and As shown in Figure 2, A and B, ectopic expression of posterior margin precursors in the wing pouch of SENSELESS in the wing disc using the *C1003-GAL4*  $Lyra<sup>W67</sup>/+$  is even more pronounced than in that of driver causes a phenotype that is similar to that observed  $Lyra<sup>1</sup>/+$  wing discs. The difference in expression levels is also observed with *in situ* hybridizations (data not affected whereas the rest of the wing is unaffected. As shown). These expression levels correlate positively with shown in Figure 2, C and D, expression of SENSELESS the loss of wing tissue in the anterior and posterior wing in a domain that corresponds to the wing margin using margin as the phenotype is more severe in  $Lyra<sup>3x67</sup>/+$  the *C96-GAL4* driver (Gustafson and Boullanne 1996) wings (Figure 1E) than in those of  $Lyra<sup>1</sup>/+$  (Figure 1C). Hence, these data indicate that the anterior and is not as severe as that induced by the more ubiquitous posterior margins of the presumptive wing are highly driver. These observations clearly show that ectopic exsensitive to ectopic SENSELESS expression and suggest pression of SENSELESS is sufficient to cause loss of wing that the wing margin loss in *Lyra* mutants may well be margin tissue. Conversely, they indicate that other areas triggered by ectopic SENSELESS. The of the wing disc do not respond to ectopic SENSELESS

genes under the control of GAL4 drivers that express over, ectopic expression of SENSELESS in wing discs,

possibly all GAL4 drivers that cause widespread expresin *Lyra* mutants in that the wing margins are severely also causes a loss of wing margin. In this case the loss **Ectopic expression of SENSELESS causes loss of wing** expression with tissue loss. Instead, as reported pre**margin:** To demonstrate that ectopic expression of viously (Nolo *et al.* 2000), we consistently observed that SENSELESS can mimic the Lyra phenotype, we con- ectopic SENSELESS causes scattered supernumerary structed flies that carried different *UAS-senseless* trans- bristles on nonmargin surfaces of adult wings. More-



FIGURE 1.—Overexpression of SENSELESS in *Lyra* mutants. Third instar wing discs were stained with anti-SENSELESS antibodies (A, B, and D). (A) Canton-S wing disc. Note that the expression is confined to many single cells that correspond to SOPs (Nolo *et al.*) 2000). (B)  $Lyra<sup>1</sup>/+$  wing disc. The arrows point to a

broad domain of SENSELESS expression that is never observed in wild-type discs. (C) Lyra<sup>1</sup>/+ wing. Wings of this genotype always show a much milder phenotype than the one shown in E. (D)  $\dot{L}y\alpha^{8\times67}/+$  wing disc. Note the ectopic expression of SENSELESS in a broad area of the wing pouch and beyond. (E)  $Lyra^{3x67}/+$  wing.



Figure 2.—Ectopic expression of SENSELESS affects the wing margin. Third instar wing discs (A and  $C$ ). Wings  $(B \text{ and } D)$ .  $(A)$ *C1003-GAL4*/*UAS-lacZ* wing disc stained with anti- $\beta$ -galactosidase. Note the scattered *lacZ* expression. (B) *C1003-GAL4*/*UAS-sens(C6)* wing. Although SENSELESS is expressed in many of the wing pouch cells, tissue loss is mostly observed at the wing margin. (C) *C96- GAL4*/*UAS-lacZ* wing disc stained with anti-β-galactosidase. *lacZ* expression is confined to the wing margin and adjacent cells. (D) *C96-GAL4*/*UAS-sens(C5)* wing displays some wing margin loss. The loss is more severe in the distal and posterior area of the wing.

extra bristles on the notum as well as parts of the wing wing discs in the anterior and posterior area of the wing surface. In leg discs, this driver caused a severe loss of pouch when compared to wild type (Figure 3G). We do all distal parts of the leg (data not shown). We conclude not know what underlies this altered pattern, but Figure that ectopic expression of SENSELESS causes very dif- 3A suggests that it may be due to loss of VESTIGIAL ferent phenotypes: loss of tissue in some areas of imagi- expression at the dorso-ventral boundary. Since loss-of-

development we tested four markers. The *vestigial* gene eration during pupal wing development. can be viewed as the wing selector gene as its lack of In addition to *vestigial*, *wingless* has also been shown

expression pattern of *lacZ* driven by the *vestigial* bound- LESS in second or early third instars results in the loss ary enhancer *vgBE* (WILLIAMS *et al.* 1994). As shown in of tissue from the wing margin (Couso *et al.* 1994; DIAZ-Figure 3, A and F, *lacZ* expression of the boundary BENJUMEA and COHEN 1995; NEUMANN and COHEN enhancer is almost entirely lost in the anterior and pos- 1996). The role of WINGLESS with respect to regulation terior portion of the wing pouch of *Lyra*<sup>1</sup> but is restored of VESTIGIAL expression at the dorso-ventral boundary in the revertant, which has the same pattern of expres- is still controversial (Go *et al.* 1998; KLEIN and MARTINEZ sion as the wild-type adult wing. The pattern is similar ARIAS 1999). However, it is fairly clear that NOTCH in *Lyra<sup>SX67</sup>* (data not shown), except that slightly more signaling is the primary inducer of *vgBE*. Hence, WINGprospective margin is missing, in agreement with the LESS expression in *Lyra* mutants may provide an indemore severe margin loss in the adult wing. *Lyra* has no pendent means to assess the effect of *Lyra* mutations on effect on the *vestigial* quadrant enhancer, *vg*QE (data wing development. As shown in Figure 3C, WINGLESS not shown), which controls later VESTIGIAL expression expression is severely reduced in the anterior and posteand growth of the nonmargin portion of the wing pouch rior domain of the wing pouch of *Lyra* mutants (com- VESTIGIAL is expressed throughout the wing blade but domain of the dorso-ventral boundary, where WINGnot in the prospective margin. As shown in Figure 3B, LESS expression is apparently normal (as is the *Lyra* immunocytochemical staining with the anti-VESTIGIAL wing margin), its expression is confined to a narrow

using a *dpp*-*GAL4* driver, resulted in large clusters of antibody shows a different pattern of expression in *Lyra* nal discs and extra sensory organs in others. function clones of *vestigial* (*vg*<sup>2</sup>/*vg*<sup>2</sup>) do not proliferate *Lyra* **mutations cause loss of DELTA, VESTIGIAL,** in the wing (Kim *et al.* 1996), the loss of wing margin **WINGLESS, and CUT expression:** To determine the tissue in *Lyra* mutants could be caused by a partial loss effects of *Lyra* mutations on the expression of key genes of VESTIGIAL expression at the anterior and posterior that have been shown to play important roles in wing wing boundary. This in turn may cause loss of cell prolif-

expression causes wing loss and its ectopic expression to play an essential role in wing development (Baker causes extra wing tissue (KIM *et al.* 1996). VESTIGIAL 1988). WINGLESS protein is secreted and is produced expression at the dorso-ventral boundary is essential to in a stripe of three to four cell rows stradling the dorsowing margin development. Furthermore, *vestigial* is a ventral boundary (Williams *et al.* 1993; Couso *et al.* marker for wing identity and has an important function 1994). The stripe of WINGLESS-expressing cells inin wing growth (KIM *et al.* 1996; NEUMANN and COHEN duces neighboring cells to differentiate into the bristles 1996; Klein and Martinez Arias 1999). that are present at the wing margin (Phillips and We tested the effect of both *Lyra* mutants on the WHITTLE 1993; Couso *et al.* 1994). Removing WING-(Williams *et al.* 1993, 1994; Kim *et al.* 1996). In this case pare with Figure 3H). With exception of the central



FIGURE 3.—Loss of vestigial, Wingless, Cut, and Delta in third instar wing discs of *Lyra* mutants. Third instar wing imaginal discs of *Lyra* mutants (A–E) and controls (F–J). (A and F) *Lyra*/+;*vgBE-lacZ*/+ (A) and a *Lyra*<sup>Sx67</sup> revertant (F) stained with X-gal. Note the loss of *lacZ* expression in the anterior and posterior wing margin in A. (B and G)  $Lyra<sup>8x67</sup>/+$  wing disc stained with anti-VESTIGIAL antibodies shows aberrant staining in mutants when compared to wild-type disc (G). (C and H)  $Lyra'/+$  and wild-type discs stained with anti-WINGLESS (C). (D and I)  $Lyra^{\omega s \sigma}/+$  (D) and wild-type disc (I) stained with anti-CUT. Note the loss of CUT staining in the anterior and posterior wing pouch (D). (E and J) Anti-DELTA staining of *Lyra<sup>Sx67</sup>*/+ (E) and control (J) discs.

domain in which levels of Wingless protein are reduced cause of high background levels. As shown in Figure severely. Since WINGLESS is an important secreted fac- 3E, there is an obvious reduction in the expression of tor for wing margin development, this reduction in ex- DELTA in the anterior and posterior wing pouch along pression in *Lyra* mutants may act in an additive fashion the presumptive wing margin (compare to Figure 3J). with the loss or severe reduction of VESTIGIAL expres- Hence, one of the key known activators of NOTCH sion. signaling at the dorso-ventral boundary is altered and

tissue in some *cut* and *Lyra* mutants (JACK and DELOTTO known markers that have previously been shown to be 1992) we also investigated CUT expression in *Lyra* mu- required for the development of the wing margin and tants. CUT is expressed in a row that is two to five cells the rows of bristles along the margin are not expressed wide at the dorso-ventral boundary (JACK *et al.* 1991; properly in *Lyra* mutants. In addition, the domains of BLOCHLINGER *et al.* 1993). This expression is largely expression that are affected in these mutants correoverlapping with that of WINGLESS and the *vestigial* spond to the domains that are affected in *Lyra* mutant boundary enhancer (WILLIAMS *et al.* 1994) but occurs discs and adult wings and are contained within the doin the mid-third instar, much later than either WING- mains in which SENSELESS is expressed ectopically. LESS or *vg*BE. Loss of CUT expression on both sides These data suggest that ectopic expression of SENSEof the wing boundary results in extensive notching of LESS in *Lyra* mutants may be able to downregulate the the margin (JACK *et al.* 1991; DORSETT 1993). *cut* has expression of several genes that play a pivotal role in been shown to be a direct target of *Notch*, but not of wing margin development, possibly by downregulating *wingless* (MICCHELLI *et al.* 1997). In addition, while the NOTCH signaling. initiation of WINGLESS expression is not dependent **Ectopic expression of SENSELESS affects WING**on *cut*, maintenance of WINGLESS expression is depen- **LESS and CUT expression:** To further investigate the dent on *cut* (MICCHELLI *et al.* 1997). As shown in Figure ability of SENSELESS to downregulate the expression of 3, D and I, CUT expression is essentially abolished in specific genes, we tested the effect of ectopic expression *Lyra* mutants in the anterior and posterior region of of SENSELESS on WINGLESS and CUT expression. the wing pouch. Ectopic expression of SENSELESS using the *C96-GAL4*

the regulation of the expression of *vg*BE, *wingless*, and is shown in Figure 4A. SENSELESS overexpression NOTCH signaling is affected. We stained *Lyra* wing discs ure 4B) and CUT protein levels (Figure 4C), although with anti-DELTA antibodies as anti-NOTCH antibody in both cases clusters of immunoreactive cells along immunohistochemical staining of wing discs failed be- the wing margin remain. Similarly, when using the *dpp-*

Given the similarities between the loss of wing margin reduced in its expression pattern. In summary, four

Since NOTCH signaling plays a prominent role in wing margin driver and staining with anti-SENSELESS *cut* (Go *et al.* 1998), we attempted to determine if causes a dramatic downregulation of WINGLESS (Fig-



Figure 4.—Ectopic expression of SENSELESS causes loss of WING-LESS and CUT expression. In all panels ectopic expression of SENSELESS is achieved using the *UAS-sens(C5)* transgene (Nolo *et al.* 2000). (A–C) The driver is *C96-GAL4*, causing ectopic expression in the wing margin area. (D–F) Ectopic expression is along the anterior-posterior wing boundary using the *dpp-GAL4* driver. (A–C) Ectopic expression of SENSE-LESS along the wing boundary (A) causes a severe reduction of WING-LESS immunoreactivity (compare Figure 4B with 3H) and CUT staining (compare Figure 4C with 3I). (D–F) Ectopic expression of SENSELESS along the anterior-posterior wing boundary (D) causes a small gap in WINGLESS (E) and CUT (F) expression. In addition, in other areas of the disc, CUT is ectopically expressed in cells where it is not normally expressed (F). This gap in WINGLESS and CUT expression causes a loss of the distal wing margin (data not shown).

the wing pouch that normally do not express CUT, as SCUTE and *string.* expected from previous observations (Nolo *et al.* 2000). *scute* is a proneural gene belonging to the *achaete*/ In summary, these data demonstrate that ectopic expres- *scute* complex and a basic Helix Loop Helix (bHLH) sion of SENSELESS in the wing margin is a potent re- transcription factor required for determination of SOPs pressor of expression of key players previously shown in the anterior wing margin (SKEATH and CARROLL to function in wing margin development. 1991). We observe a downregulation of SCUTE expres-

in *Lyra* mutants can be viewed as the sum of two compo- A and B). Indeed, in *Lyra<sup>x67</sup>* wings, there are few SOPs nents. The first component is an effect on margin deter- expressing SCUTE at the anterior wing margin (Figure mination in the developing wing disc. Indeed, our data 5B). This is in sharp contrast to ectopic expression of are in agreement with numerous observations showing *senseless* in other epithelial cells of the wing disc where that loss of NOTCH signaling causes loss of expression it causes induction of SCUTE expression (Nolo *et al.* in the wing margin of the patterning genes *wingless* and 2000). *vestigial* (Couso *et al.* 1995; Diaz-Benjumea and Cohen The failure to form SOPs in the wing discs of *Lyra* 1995; Kim *et al.* 1995, 1996; Rulifson and Blair 1995; mutants predicts that the set of two cell divisions re-De Celis *et al.* 1996; Doherty *et al.* 1996). The second quired for differentiation of margin bristles in the early component corresponds to an effect on cell prolifera- pupa will not take place. The reason for the loss of tion. Indeed, loss and gain of NOTCH signaling experi- the surrounding unspecialized margin cells in the adult ments have been shown to cause a severe decrease and wings of *Lyra* is not as obvious, but one hypothesis is increase in cell proliferation, respectively (Go *et al.* that these cells also fail to proliferate. To test this we 1998). Our data suggest that loss of DELTA causes a examined the mRNA expression pattern of *string. string* loss of NOTCH signal and a loss of cell proliferation in mRNA is normally expressed in the central cells of both

*GAL4* driver to ectopically express SENSELESS along *Lyra* wing development begins shortly after pupariation the anterior-posterior wing boundary (Figure 4D) we and continues during the first half of pupal developfind a precise disruption in the continuity of WINGLESS ment (ABBOTT and SPREY 1990). This is the time win-(Figure 4E) and CUT (Figure 4F) expression where the dow in normal development when differentiation of *dpp* stripe is normally expressed. This downregulation bristles and trichomes takes place as well. To further correlates with a loss of the distal tip of the wing (data examine how ectopic SENSELESS affects wing margin not shown). Note also that ectopic SENSELESS expres- specification and differentiation during *Lyra* wing develsion causes ectopic CUT expression in some cells of opment we have studied the expression pattern of

**Margin loss in** *Lyra* **mutants:** The loss of wing margin sion in the anterior pouch of the wing disc (Figure 5,

the wing margin. The reduction in cell proliferation in the anterior and posterior wing margin during the later



FIGURE 5.—Expression of SCUTE and *string* in *Lyra* wing<br>discs. Third instar wing discs of wild-type (A and C) and<br>*Lyra*<sup>8667</sup>/+ larvae (B and D). (A and B) Mutant discs show that *vestigial* can be viewed as a "wing se decreased levels of SCUTE expression in the anterior wing view that is supported by the observation that its ectopic margin (B). Hence, ectopic expression of SENSELESS can expression can rescue loss of WINGLESS (KLEIN and lead to an induction of SCUTE in some domains of the wing<br>disc (Noto *et al.* 2000) and loss of SCUTE in other areas. (C<br>and D) In situ hybridizations with *string*. Note the loss of *string*<br>disc causes also a failure of

even though margin cells are arrested at that time and developing wing margin. It has been proposed that the cell proliferation does not begin until early pupariation. main function of WINGLESS is to enforce gene expres-Our *in situ* experiments with *string* confirmed this ex- sion in the wing disc rather than to initiate it. Hence, pression pattern in wild-type discs. But in *Lyra* third the combined loss of *vestigial* expression at the boundary instar wing discs the mitosis-inducing phosphatase and the strong reduction in WINGLESS expression at STRING (Cdc25) is severely downregulated in the ante- the wing margin may affect cell proliferation and cell rior and posterior area of the prospective wing margin identity not only in the wing margin, but also in a few as indicated by *in situ* hybridization (Figure 5D). This cell rows adjacent to the anterior and posterior wing is consistent with an overall lack of proliferation in the margin. This model is in agreement with the observation anterior and posterior margin region. However, it is also that we find no alterations in the expression pattern of possible that the non-bristle-forming cells are present in the quadrant enhancer of *vestigial* in *Lyra* mutants (data the wing margin, but that they lose their capacity to not shown) and that *Lyra* wing discs exhibit a dramatic flatten and secrete margin elements (trichomes), which reduction in *string* expression in the cells along the serve as the visible hallmark of each cell. This could be dorso-ventral boundary (Figure 5, C and D). Since *st* caused by their lack of exposure to the sequence of has been shown to induce mitosis, and since *Lyra* mu-

dence that *Lyra* mutations are gain-of-function/neo- signaling. This decrease may explain the loss of WINGmorphic alleles of *senseless* that cause overexpression LESS, VESTIGIAL, and CUT expression, which have all of *senseless* in third instar imaginal discs. This ectopic previously been shown to depend on NOTCH signaling. expression of *senseless* causes a loss of anterior and poste-<br>
We propose that this defect in *Lyra* mutants underlies<br>
the effect on margin determination in the developing rior wing margin tissue. The data presented in this arti-<br>cle provide a molecular framework to understand this wing disc and the reduction in cell proliferation in early

cle provide a molecular framework to understand this wing disc and the reduction in cell proliferation in early<br>phenotype. pupae. Wingless is required for differentiation of bristles late we thank the Bloomington Stock Cen

proneural SOP determinants *acheate* and *scute.* In addition, CUT expression in third instar discs has been shown to be dependent on WINGLESS expression, while CUT is also required for the maintenance of WINGLESS expression (Neumann and Cohen 1996, 1997). Since CUT is essential for all wing margin bristles, both innervated and noninnervated (Jack *et al.* 1991), we propose that the combined reduction in WINGLESS and CUT expression in *Lyra* mutants may cause a secondary reduction in proneural gene expression in the wing margin, as revealed by SCUTE staining. This reduction in expression should lead to a loss of numerous bristles in the anterior and posterior wing margin. However, these observations do not provide a rationale for the loss of wing blade cells adjacent to the margin, which are also observed in *Lyra* mutants.

et al. 1996). Go et al. (1998) and KLEIN and MARTINEZ<br>in the anterior and posterior wing margin (D). ARIAS (1999) have recently proposed a model of wing development in which the *vg*BE is induced by NOTCH third instar larval stage (Johnston and Edgar 1998) signaling when and where WINGLESS is active at the dorso-ventral boundary (Figure 5, C and D). Since *string* proteins required for determination of the wing margin. tants exhibit no cell death and a loss of cells in pupal development, we propose a causal relationship between these observations. At the root of the Lyra phenotype may be the observation that the DELTA signal is im-The data presented in this article provide strong evi-<br>paired, which should lead to a decrease in NOTCH previously been shown to depend on NOTCH signaling.

BLAIR 1994). Indeed, high levels of WINGLESS are K. Norga for comments on the manuscript. R.N. is a research associate known to be required for the proper expression of the and H.J.B. is an investigator from the Howard Hughes Medical InstiHuda Zogbhi.

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