Mutation Notes



Drosophila melanogaster P{GAL4-Hsp70.PB} transposon insertion on 3rd chromosome creates mutations in *mth* and *Ptpmeg* genes.

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The third chromosome P element insertion P{w =GAL4-Hsp70.PB}89-2-1 (Bloomington stock number 1799) (Brand, Manoukian, and Perrimon, 1994), hereafter referred to as P{hs-GAL4}89, is being used as a heat-inducible GAL4 driver for the ubiquitous, conditional activation of UAS responders (Armstrong *et al.*, 2002; Chanut *et al.*, 2002; Kozlova and Thummel, 2002; Liu and Lehmann, 2008; Roman, He, and Davis, 2000; Seong, Ogashiwa, Matsuo, Fuyama, and Aigaki, 2001). To determine whether the P insertion in the P{hs-GAL4}89 line may potentially interfere with the expression of a gene or genes located at the genomic integration site, we amplified flanking genomic DNA by inverse PCR. Genomic DNA isolated from P{hs-GAL4}89 flies was digested with *Hpa*I, which cuts in the polylinker of the pCaSpeR3 transformation vector from which the P{hs-GAL4}89 element was derived (Brand *et al.*, 1994) and in flanking genomic DNA. Restriction fragments were ligated under dilute conditions to favor circularization of DNAs. The portion of the polylinker that remained attached to genomic DNA after *Hpa*I digestion was then cut by digestion with *Sal*I to generate linearized DNAs with P-element-derived ends. Linearized DNA was then amplified by PCR and sequenced using primers 5'-GGATCCCCGGGCGAG-3' and 5'-CCTGCAGCCCAAGCTT-3'.

PCR yielded a single product of about 0.8 kb. Sequencing of the PCR product indicated that the transposon resides in the Ptpmeg gene, located 543 bp upstream of the transcription start of transcripts RI and RJ, and within a large intron that is specific for transcripts RH and RK (Figure 1). This intron harbors three other genes, methuselah (mth) and methuselah-like 9 and 10 (mthl9, mthl10), that are transcribed in the opposite direction to Ptpmeg. The P{hs-GAL4}89 element is inserted after position +38 downstream of the putative transcription start site of the mth gene (Figure 1B). Consistent with this observation, microarray data indicate that mth RNA was 30-fold reduced in salivary glands of homozygous P{hs-GAL4}89 animals compared to control animals without the element (Liu and Lehmann, 2008). These findings are surprising, because mth is considered to be an essential gene. Homozygous mth null mutants die during the embryonic stage (Ja, Carvalho, Madrigal, Roberts, and Benzer, 2009). Despite the strong reduction, mth RNA was still detectable in homozygous animals (Liu and Lehmann, 2008), suggesting that basal expression of mth does not require sequences located immediately upstream of the transcription start site. It appears that a minimal amount of mth RNA sufficient for survival can be produced from an internal promoter. This interpretation is supported by the report of a positive transcriptional control element located +28 to +217 relative to the transcription start site of mth. A reporter gene that carries this element, but entirely lacks upstream sequences of *mth*, indeed exhibits basal transcriptional activity (H. Kim, Kim, Lee, Yang, and Han, 2006).

Finally, we wondered whether the GAL4 gene of the transposon is transcribed in the same direction as *mth* or *Ptpmeg*. The inverse PCR results had indicated that the P transposon is inserted in the same 5'-3' orientation as Ptpmeg. We determined the orientation of hs-GAL4 within the transposon by PCR with the help of primer 5'-CAGACACTTGGCGCACTTCGGT-3', which hybridizes within the GAL4 gene, and primers Casper1 (5'-GATCCCCGGGCGAGCTCGAAT-3') and Casper2 (5'-AACGCTACAAACGGTGGCGA-3'), which hybridize with vector sequences. Only Casper2, which was designed to give a PCR product if hs-GAL4 was oriented the same way as vector and Ptpmeg, but not Casper1, yielded a product. Thus, hs-GAL4 is transcribed in the same direction as Ptpmeg (Figure 1B).

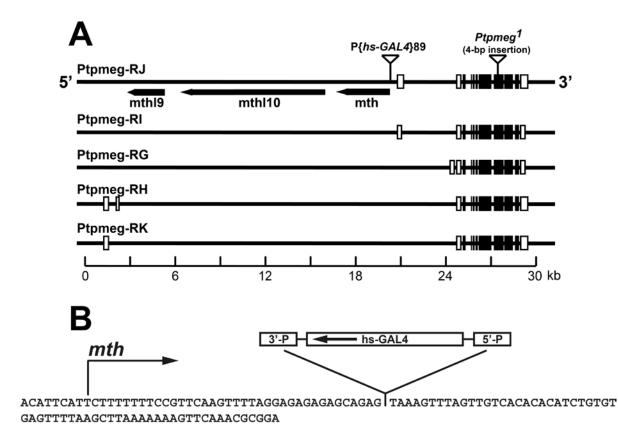


Figure 1. Genomic location and orientation of hs-GAL4 in the P{hs-GAL4}89 line. (A) Genomic organization at the P(hs-GAL4)89 integration site. Ptpmeg is transcribed from left to right, whereas the intronic genes, mth, mthl10, and mthl9, are transcribed from right to left. (B) P element and hs-GAL4 transgene are located in the indicated orientations within the non-coding 5'exon of *mth* close to the putative transcription start site predicted by Flybase.

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References: Armstrong, J.A., O. Papoulas, G. Daubresse, A.S. Sperling, J.T. Lis, M.P. Scott, and J.W. Tamkun 2002, EMBO J. 21: 5245-5254; Brand, A.H., A.S. Manoukian, and N. Perrimon 1994, Methods in Cell Biology 44: 635–654; Chanut, F., K. Woo, S. Pereira, T.J. Donohoe, S.-Y. Chang, T.R. Laverty, A.P. Jarman, and U. Heberlein 2002, Genetics 160: 623-635; Ja, W.W., G.B. Carvalho, M. Madrigal, R.W. Roberts, and S. Benzer 2009, Protein Science 18: 2203–2208; Kim, H., J. Kim, Y. Lee, J. Yang, and K. Han 2006, Molecules and Cells 21: 261-268; Kozlova, T., and C.S. Thummel 2002, Development 129: 1739-1750; Liu, Y., and M. Lehmann 2008, Fly 2: 92-98; Roman, G., J. He, and R.L. Davis 2000, Genetics 155: 1281–1295; Seong, K.H., T. Ogashiwa, T. Matsuo, Y. Fuyama, and T. Aigaki 2001, Biogerontology 2: 209– 217.