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**Transcriptional Regulation of the Cell Proliferation-Related Genes by 20-OH Ecdysone and Ursolic acid in *Drosophila***

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We previously found transcriptional inhibition of the proliferating cell nuclear antigen( PCNA ) and *D-raf*( *Drosophila* homolog of the human *c-raf-1* gene ) genes by ursolic acid in *Drosophila* Kc cells. In this study, we also confirmed the fact in vivo by using transgenic flies. To know the mechanism of repression of PCNA and *D-raf* promoter activities by ursolic acid, we examined the effect of 20-OH ecdysone similar to the structure of ursolic acid on expression of these genes. 20-OH Ecdysone also inhibit expression of the PCNA and *D-raf* genes. The inhibition effect of 20-OH ecdysone on expression of these genes was not shown at the tested concentration higher than 1  $\mu$ M in both *Drosophila* Kc cells and transgenic flies, while ursolic acid showed the dose dependent inhibition effect.

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**Tissue-Specific and Dietary Control of Alpha-Amylase Gene Expression in *Drosophila melanogaster***

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Amylase variants of *Drosophila melanogaster* were screened for spatial variation in adult and larva midgut  $\alpha$ -amylase(EC 3-2-1-1, 1,4- $\alpha$ -glucan glucanohydrolase: *Amy*). Enzyme activity was detected by DNSA method during several generations in each food component for dietary control. Every strain revealed a different activity in each restricted medium, indicating that each strains have different adaptation ability to carbon source. In this study, standard medium contained starch-like cornmeal gives less dietary stress to fly than two other carbon sources. The relationship of *Amy* genotype and midgut amylase-activity pattern(*Map*) analysis was shown expression of *MapP* indicated highly enzyme activity at posterior parts of region than that at anterior. In electrophoretic analysis, it revealed *Amy*<sup>1</sup> and *Amy*<sup>1-3</sup> was *MapA*<sup>123P<sup>00</sup>, *Amy*<sup>4-6</sup> was *MapA*<sup>123P<sup>12</sup>. This suggests that somehow *Amy* genes, or their products, are differentially recognized by products of the *Map* gene in addition to being differentially recognized in different parts of the midgut.</sup></sup>