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研究论文

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瓜实蝇嗅觉受体基因的克隆及表达谱分析

申建梅, 胡黎明, 宾淑英, 林进添

Cloning and expression profiling of an olfactory receptor gene in Bactrocera cucurbitae (Coquillett) (Diptera: Tephritidae)

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全文: PDF (0 KB) HTML (1 KB) 输出: BibTeX | EndNote (RIS) 背景资料

昆虫的嗅觉受体是一个高度变异的蛋白家族, 其中一类Or83b嗅觉受体在不同昆虫体内高度保守, 在昆虫的行为调控过程中起到十 分重要的作用。为进一步探讨Or83b受体的功能, 本研究利用RT-PCR和RACE方法克隆获得瓜实蝇Bactrocera cucurbitae (Coquillett) Or83b-like受体的全长cDNA序列, 命名为BcucOr83b-like(GenBank登录号: HM745934)。测序结果表明, BcucOr83b-like开放阅读框全长1 422 bp,编码473个氨基酸残基。氨基酸序列比对表明,此序列具有Or83b受体的典型特征, 序列中具有7个跨膜区和高度保守的C端区域。BcucOr83b-like与其他昆虫的Or83b具有较高的氨基酸序列一致性, 其中与桔小实 蝇Bactrocera dorsalis (Hendel) Or83b的序列一致性高达99.6%。对该基因在瓜实蝇成虫不同组织和发育时期表达量的荧光定 量PCR分析表明, BcucOr83b-like主要在瓜实蝇成虫触角中表达, 头部(去除触角)、 雌虫前足和翅中也有较高的表达; 瓜实 蝇在各个发育时期的表达水平不同, 在刚羽化雌成虫中的表达量最高。本研究为深入研究瓜实蝇Or83b受体的功能提供了理论依 据。

关键词:

Abstract:

Olfactory receptors in insects were found to be extremely diverse, but there is a kind of olfactory receptor, Or83b, which shares a high identity among different insects and plays an important role in the regulation of insect behavior. In order to research the function of Or83b, the gene encoding Or83b receptor from Bactrocera cucurbitae (Coquillett) was cloned and named as BcucOr83b-like (GenBank accession no. HM745934). The sequencing results showed that the open reading frame of BcucOr83b-like was 1 422 bp in length, encoding 473 amino acid residues. Homology analysis showed that the deduced amino acid sequence of BcucOr83b-like had the characteristics of Or83b, with seven putative transmembrane domains and conserved C domains. BcucOr83b-like had very high similarity of amino acid sequence with those from other insects and shared 99.6% identity with the Or83b sequence from Bactrocera dorsalis. Tissue expression pattern revealed that BcucOr83b-like transcript was observed clearly in the antenna of B. cucurbitae, and highly expressed in the head (with antennae removed), female foreleg and wings. The temporal expression pattern further indicated that BcucOr83b-like was also expressed in different developmental stages at different expression levels, with the highest level in female adult newly emerged. This study provides a basis for further researching the function of Or83b from B. cucurbitae.

Key words:

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