

ГЕНОМИКА.  
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ИДЕНТИФИКАЦИЯ ФРАГМЕНТА СТИГМАСПЕЦИФИЧНОЙ  
ЭКСПРЕССИИ В ПРОМОТОРЕ ГЕНА ХИТИНАЗЫ КЛАССА I СОИ<sup>1</sup>

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Уровень экспрессии гетерологичных генов в трансгенных растениях служит важным показателем эффективности работы генов. Тонкая настройка экспрессии трансгенов ограничена небольшим репертуаром известных на данный момент эффективных промоторов. Нами клонирован и охарактеризован тканеспецифичный фрагмент промотора гена хитиназы класса I сои (*GmChi1*). Промотор *GmChi1* (*GmChi1P*) клонировали из сои сорта Jungery. Последовательность промотора содержит ряд предполагаемых *cis*-действующих элементов, включая тканеспецифичные мотивы и мотивы, регулируемые в условиях стресса. По результатам гистохимического анализа самая высокая активность репортерного фермента  $\beta$ -глюкуронидазы (GUS), находящегося под контролем *GmChi1P*, обнаружена в корнях трансгенного растения, *Nicotiana tabacum* cv. NC89, на стадии формирования ростка с четырьмя листьями. Интересно, что обработка салициловой кислотой эффективно подавляла высокую активность GUS в корнях трансгенного табака. Делеционным анализом *GmChi1P* установлено, что последовательности, расположенные между позициями –719 и –382, содержат ключевые *cis*-элементы, ответственные за экспрессию репортерного гена *uidA* (кодирующего GUS) в листьях, корнях и местах поранения *Nicotiana tabacum*. Кроме того, согласно результатам флуориметрического анализа, активность укороченных промоторов: от *ChiP*(–1292) до *ChiP*(–719) – в корнях трансгенного табака значительно подавляется абсцизовой кислотой и полностью – салициловой. Обнаружено также, что промотор *ChiP*(–382) экспрессируется исключительно в рыльце цветков трансгенного растения. С использованием репортерного фермента GUS не обнаружено окрашивания в других органах цветка трансгенного *Nicotiana tabacum*, включая чашелистики, лепестки, пыльники, нити и завязи, а также ни в одной из вегетативных тканей. Полученные результаты свидетельствуют о том, что фрагмент промотора *ChiP*(–382) может быть использован в тканеспецифичной регуляции экспрессии генов и генной инженерии растений.

**Ключевые слова:** *GmChi1*, соя, промотор, трансгенные растения, *Nicotiana tabacum*, стигмаспецифичный промотор

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## Identification of Stigma-Specific Expression Fragment in the Promoter of the Soybean Chitinase Class I Gene

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The expression level of heterologous genes in transgenic plants serves as an important indicator of gene efficiency. The small number of currently known effective promoters, limits the possibilities in fine-tuning the expression of transgenes. We cloned and characterized a tissue-specific promoter fragment of the soybean chitinase class I gene (*GmChi1*). The *GmChi1* promoter (*GmChi1P*) was cloned from Jungery soybean. The promoter sequence contains a number of putative *cis*-acting elements, including tissue-specific and stress-regulated motifs. By histochemical analysis, the *GmChi1P*-controlled  $\beta$ -glucuronidase (GUS) reporter enzyme activity was shown to be highest in the roots of transgenic *Nicotiana tabacum* cv. NC89 at the four-leaf sprout formation stage. Interestingly, the high GUS activity in transgenic tobacco roots was effectively suppressed by salicylic acid (SA) treatment. Deletion analysis of *GmChi1P* revealed that the sequences located between positions –719 and –382 contain key *cis*-elements responsible for the reporter *uidA* gene expression (encoding GUS) in leaves, roots, and wounds of *Nicotiana tabacum*. In addition, fluorometric analysis showed that the activity of the shortened *ChiP*(–1292) to *ChiP*(–719) promoters in the roots of transgenic tobacco was significantly suppressed by abscisic acid and completely suppressed by SA. The *ChiP*(–382) promoter was also found to be expressed exclusively in the stigma of transgenic tobacco flowers. Using the GUS reporter enzyme, no staining was detected in other flower organs in transgenic *Nicotiana tabacum*, including sepals, petals, anthers, filaments, and ovaries, or in any vegetative tissues. The results indicate that the promoter fragment *ChiP*(–382) can be used in tissue-specific regulation of gene expression and plant genetic engineering.

**Keywords:** *GmChi1*, soybean, promoter, transgenic plant, *Nicotiana tabacum*, stigma-specific promoter