RESEARCH PAPER



Functional characterization of the three genes encoding 1-deoxy-p-xylulose 5-phosphate synthase in maize

Elizabeth Cordoba^{1,*}, Helena Porta^{1,*}, Analilia Arroyo^{1,†}, Carolina San Román¹, Luis Medina¹, Manuel Rodríguez-Concepción² and Patricia León^{1,‡}

¹ Departamento de Biología Molecular de Plantas, Instituto de Biotecnología, Universidad Nacional Autónoma de México, Av. Universidad 2001 Chamilpa. Apdo. Postal 510-3 Cuernavaca, Morelos 62210, Mexico

² Department of Molecular Genetics, Centre for Research in Agricultural Genomics (CRAG), CSIC-IRTA-UAB, 08034 Barcelona, Spain

* These authors contributed equally to this work.

[†] Present address: Centro de Investigación en Biotecnología Aplicada, Instituto Politécnico Nacional, Ex-Hacienda San Juan Molino Carrretera estatal Tecuexcomac Tepetitla Km. 1.5 Lardizabal, Tlaxcala, CP 90700, México.

[‡] To whom correspondence should be addressed. E-mail: patricia@ibt.unam.mx

Received 8 September 2010; Revised 9 November 2010; Accepted 15 November 2010

Plants produce many isoprenoids that are functionally

important in essential processes. Despite their diversity, all

isoprenoids are formed from two common precursors,

isopentenyl diphosphate (IPP) and its isomer dimethylallyl

diphosphate (DMAPP). Additionally isoprenoids are of

biotechnological, medical, and industrial importance (Lange

and Croteau, 1999; Römer et al., 2000, and references

within), prompting efforts to increase levels of these valu-

utilizes two independent pathways that have different

precursors. The acetate/mevalonate (MVA) pathway is

found in most eukaryotes. The second pathway known as

In plants, the biosynthesis of IPP and DMAPP

Abstract

Introduction

able compounds.

The 1-deoxy-D-xylulose 5-phosphate synthase (DXS) enzyme catalyses the first biosynthetic step of the 2-C-methyl-D-erythritol 4-phosphate (MEP) pathway. In plants the MEP pathway is involved in the synthesis of the common precursors to the plastidic isoprenoids, isopentenyl diphosphate and dimethylallyl diphosphate, in plastids. DXS is recognized as limiting this pathway and is a potential target for manipulation to increase various isoprenoids such as carotenoids. In *Zea mays* three *dxs* genes exist that encode plastid-targeted functional enzymes. Evidence is provided that these genes represent phylogenetically distinctive clades conserved among plants preceding monocot-dicot divergence. There is differential accumulation for each *dxs* gene transcript, during development and in response to external signals such as light. At the protein level, the analysis demonstrates that in *Z. mays*, DXS protein is feedback regulated in response to the inhibition of the pathway flow. The results support that the multilevel regulation of DXS activity is conserved in evolution.

Key words: 1-Deoxy-D-xylulose 5-phosphate synthase (DXS), isoprenoid biosynthesis, maize, 2-C-methyl-D-erythritol 4-phosphate (MEP) pathway, post-transcriptional regulation.

non 1. 1 9, 2 tol

the 2-*C*-methyl-D-erythritol 4-phosphate (MEP) pathway operates in the chloroplast of photosynthetic eukaryotes and also in most eubacteria and in apicomplexa parasites (Eisenreich *et al.*, 2004). Thus, the genes and enzymes of the MEP pathway are attractive targets to develop new antibacterial and antiparasitic drugs, and herbicides (Rodríguez-Concepción, 2004; Rohdich *et al.*, 2005). Although evidence supports a limited exchange of intermediates between these two pathways, it is clear that only one pathway is primarily responsible for the synthesis of particular isoprenoids. The MVA pathway provides the precursors for sesquiterpenes (C₁₅) and triterpenes (C₃₀) (Lichtenthaler, 1999; Eisenreich *et al.*, 2001). In contrast,

© The Author [2011]. Published by Oxford University Press [on behalf of the Society for Experimental Biology]. All rights reserved. For Permissions, please e-mail: journals.permissions@oup.com