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Redefining the hormonal control of tomato (*Solanum lycopersicum* cv. Micro-Tom) fruit development

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Thesis presented to obtain the degree of Master in
Science. Area: Plant Physiology and Biochemistry

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RESUMO

Redefinindo o controle hormonal do desenvolvimento do fruto do tomateiro (*Solanum lycopersicum* cv. Micro-Tom)

O controle hormonal do desenvolvimento do fruto do tomateiro (*Solanum lycopersicum* cv. Micro-Tom) já foi extensamente estudado, principalmente no amadurecimento. No entanto, ainda existem consideráveis lacunas em nosso conhecimento da função e distribuição espaço-temporal dos hormônios durante o desenvolvimento do fruto. Foram realizados ensaios histoquímicos com cinco linhas transgênicas de tomate cv. Micro-Tom que carregavam o gene repórter GUS fusionado a cinco diferentes promotores responsivos aos hormônios, auxina, citocinina (CKs), giberelina (GAs), ácido abscísico (ABA) e etileno, com o fim de redefinir a distribuição espaço-temporal deles ao longo do desenvolvimento do fruto, desde estágios iniciais pre-anteses até o estágio red ripe (RR). Os níveis de CKs foram altos na fase pre-anteses, inicialmente na placenta e posteriormente nos óvulos, indicando um papel importante deste hormônio no crescimento do ovário, pre-fertilização. Nos estágios iniciais, post-fertilização, o conteúdo de CKs foi alto nas sementes e depois nas camadas epidérmicas e subepidérmicas internas e externas do pericarpo. Alto conteúdo de auxina foi observado, durante todos o desenvolvimento do fruto, no pedicelo, sugerindo o transporte basípeto de auxina para a planta mãe. Os níveis de etileno aumentaram durante o crescimento do ovário antes da antese. Interessantemente, o nível de etileno não diminuiu imediatamente post-fertilização. Os conteúdos de GA e ABA foram baixos prévios a antese e nos estágios iniciais pre-fertilização, especificamente na placenta e pericarpo. Inversamente, o conteúdo desses hormônios aumentou na fase de expansão celular do crescimento do fruto. Os grãos de pólen que chegaram ao stigma, post deiscência das anteras, apresentaram altos níveis de ABA e auxina. Além disso, reportamos um antagonismo entre conteúdo de etileno e giberelina durante o desenvolvimento do fruto, onde o nível de etileno começou a decrescer durante a fase de expansão celular do crescimento do fruto, momento em que o nível de giberelina começou a aumentar na placenta e pericarpo.

Palavras-chaves: 1. Tomate 2. Desenvolvimento do fruto 3. Hormônios 4. Gene reporter GUS L.

ABSTRACT

Redefining the hormonal control of tomato (*Solanum lycopersicum* cv. Micro-Tom) fruit development

The hormonal control of tomato (*Solanum lycopersicum*) fruit development have been extensively studied, mainly during ripening. Notwithstanding, considerable gaps still exist in our understanding of the function and spatial-temporal distribution of hormones during fruit development. Here, we performed histochemical analysis of tomato *cv.* Micro-Tom transgenic lines harboring the gene reporter *GUS* fused to five different promoter responsive to the hormones, auxin, cytokinins (CKs), gibberellins (GAs), abscisic acid (ABA) and ethylene, in order to redefine their spatial-temporal distribution during tomato fruit development, from early pre-anthesis stages to red ripe (RR) stage. CKs levels were high during pre-anthesis, initially in placental tissues and subsequently in ovule, indicating a major function of this hormone during pre-fertilized ovary growth. At early post-fertilization stages, CKs contents were high in seeds, and afterward in the outer and inner epidermal and subepidermal layer of pericarp. High auxin content was observed, during all pre-anthesis and fruit development stages, in pedicel, suggesting a basipetal transport to the mother plant. Ethylene contents increased during pre-anthesis ovary growth. Interestingly, ethylene contents did not decrease, immediately, post-fertilization. The content of GAs and ABA was low at pre-anthesis and early post-fertilization stages, specific in placental tissues and pericarp. Conversely, these hormones accumulated mainly during cell expansion phase of fruit growth. Pollen grains that reached the stigma, post anther dehiscence, showed high content of auxin and ABA. In addition, we reported an antagonism between ethylene and GAs contents during tomato fruit development, where the level of ethylene started to decrease during the cell expansion phase of the fruit growth, moment when the level of GAs started to increase in placental tissues and pericarp.

Key-words: 1. Tomato 2. Fruit development 3. Hormones 4. Gene reporter *GUS* L.

1. INTRODUCTION

In angiosperms, embryonic development occurs concomitantly with the development of the ovary into a specialized organ, the fruit (Gillaspy et al., 1993). In tomato, a multicarpelar berry that is considered as a model system for fleshy climacteric fruit development (Azzi et al., 2015), fruit develops from the ovary, which consist in two or more carpels forming the locules, separated by a septum that fuses with each of the carpels, and ovules are attached to the central septum via placental tissue (Gasser and Robinson-Beer, 1993; Pattison et al., 2015).

Tomato fruit development is usually divided in four phases; fruit set, growth, maturation and ripening. The fruit set, defined as the transition of a quiescent ovary to a rapidly growing young fruit, depends on the successful completion of pollination and double fertilization, when one sperm fuses with the eggs cell to form the embryo and the other sperm fuses with the two haploid polar nuclei in the central cell to give rise to the endosperm (Raghavan, 2003). The fruit growth is divided in two stages: a very active period of cell division within the pericarp, mainly in the outer layer, which last about 7 to 10 days after successful fertilization, whereas after cell division phase during the following days until the fruit reach it final size, the fruit growth depend mainly on cell expansion. Fruit growth is followed by the fruit maturation, where it acquire the prerequisite competence to enter into the final development stage, ripening. Fleshy fruit such as tomato undergo a ripening process in which the biochemistry, physiology, and structure of the organ are developmentally altered to influence appearance, texture, flavor and aroma (Gillaspy et al., 1993; Giovannoni, 2004; de Jong et al., 2009a; McAtee et al., 2013; Kumar et al., 2014; Liu et al., 2015). It is worth noting that the four phases described above are well accepted by most author, although they do not take into account the initial ovary development before anthesis, where important events, such as the definition of the number of carpels, and thus the number of locules and the final size of the fruit, take place (Xu et al., 2015).

New evidences suggest that functions of plant hormones are not limited to a restricted phase in fruit development, and that there exists a complex crosstalk between them during the whole fruit development (Kumar et al., 2014; McAtee et al., 2013; Pattison et al., 2015). Increase of auxin contents upon pollination, mostly synthesized in the embryo and exported to pericarp, plays a major role in the regulation of fruit set and growth (Gillaspy et al., 1993; Serrani et al., 2008, 2007; Vriezen et al., 2008; de Jong et al., 2009a; Mariotti et al., 2011; Pattison and Catala, 2012; McAtee et al., 2013). This induction of fruit set and growth by increased auxin post-fertilization is, at least partially, due to an increase in gibberellins (GAs) biosynthesis (Serrani et al., 2008). Exogenous applications of either auxin or GAs induce parthenocarpic fruits that present, respectively, increased cell layers or greater cell expansion (Serrani et al., 2007), suggesting that auxin contributes to cell division in fruit, while gibberellin to cell expansion.

Application of synthetic cytokinins (CKs) induces parthenocarpic fruit through promotion of cell division (Ding et al., 2013). Additionally, the cytokinin transduction pathway is active in cell division phase of fruit development, but the first peak of cytokinin accumulation occurs during anthesis (Matsuo et al., 2012). These evidences suggest that cytokinin might be involved in the growth of ovaries until pollination and afterwards in the cell division period post-pollination.

Acid abscisic (ABA) and ethylene biosynthesis genes expression is attenuated post-pollination, which suggest an antagonist role of these two plant hormones in fruit set (Vriezen et al., 2008; Shinozaki et al., 2015). In climacteric fruit, such as tomato, ethylene production reaches its maximum at the onset of ripening, which indicates a main role of this hormone at this phase of fruit development (Liu et al., 2015). On the other hand, the facts that the expression of ethylene biosynthesis genes (*SLACS2*, *SLACS4* and *SLACOT*) is induced by exogenous ABA, and that

ABA maximum precedes ethylene release, reveals an ABA/ethylene interaction acting at the level of ethylene biosynthesis (Liu et al., 2015). These evidence imply that ABA may function as a trigger for the initiation of fruit ripening by inducing ethylene-mediated pathway and other ethylene-independent processes (Zhang et al., 2009; Mou et al., 2016). Moreover, Jones et al. (2002) and Trainotti et al. (2007) suggests a crosstalk between ethylene and auxin during onset of ripening.

Although the importance of hormones in fruit development is well known (Zhang et al., 2016), considerable gaps in our knowledge about hormonal networks and crosstalk between different hormones still exist (Kumar et al., 2014). Moreover, the spatial-temporal distribution of hormones and expression of genes related to hormonal metabolism and signaling during the entire fruit development, from pre-anthesis to red ripe stages, is poorly understood (Zhang et al., 2016). The aim of this work was to redefines the hormonal control of fruit development by using transgenic plant harboring the gene reporter GUS fused to five different promoter responsive to auxin, CKs, GAs, ABA and ethylene. Through histochemical analysis, we demonstrates the spatial-temporal distribution of the five classics plant hormones from 12 days before anthesis, referred to here as -12 days post anthesis (-12 DPA), to the red ripe (RR) stage in tomato *cv.* Micro-Tom fruit. According to our results, GAs and ABA have its main functions in the cell expansion phase of fruit development, reaching its maximum production at mature green stage and cell expansion phase, respectively, and presenting apparently no activity during fruit set and early fruit growth. Surprisingly, an increase in the ethylene contents was observed, during the ovary growth, at pre-anthesis stages without diminishing during early post-fertilization stages. Furthermore, CKs show a major role during ovary growth at pre-anthesis stages in placenta and ovules. We additionally, proposes an interplay between ethylene and GAs during the transition from cell division to cell expansion phase of fruit growth.

2. CONCLUSIONS

Taken together, the results presented here redefined the hormonal balance during tomato fruit development. In this working hypothesis, auxin and cytokinin are the main hormones in stimulating ovary/fruit development during pre-anthesis and early post-fertilization cell division phase, whereas GAs and ABA are responsible for fruit growth during cell expansion phase. On the other hand, ethylene may presents a role in cell division phase, suppressing GAs metabolism, giving that an antagonist temporal and spatial distribution of *EBS::GUS* and *GA2OX::GUS* was noticed, and that *Nr* fruits showed a higher *GA2OX::GUS* expression in placenta and pericarp compared to WT. The decrease of ethylene content during cell expansion phase may release ethylene repressed GAs metabolism, which in turns, promotes cell expansion. In seeds (ovules), the hormones balance, in general, parallels the pericarp, with the notable exception of ethylene, which seems not to accumulate in seeds, and the more persistent accumulation of auxin, probably reflecting their further role in germination.

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