Title: There are secrets along the pollen tube pathway, still in need to be discovered

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Abstract

Plant organogenesis is the charming process that embodies outstanding structures of refined control of gene expression. During this elegant process, a subtle communication occurs between neighboring cells, based on chemical signals, inducing cellular mechanisms of patterning and growth.

The path to be followed starts once the stigmatic cells recognize a compatible pollen grain, facing immediately self-incompatibility responses, and follows dealing with distinct players interacting on pollen tube growth and puzzling navigation along the transmitting tract. Pollen tube goes through a guidance process that starts at a preovular stage, with interacting players from the transmitting tissue. The ovular guidance stage considers the specific relation between the pollen tube and its ovule, divided in funicular and micropylar guidance steps, with plenty of receptors working in signalling cascades. Lately, just after the pollen tube passage beyond synergids, gametes fusion occurs and the aimed target, the developing seed, will start to mature.

The present review presents the existing knowledge of interactions and partners involved in the crucial biological process of pollen-pistil interaction giving rise to the new seed generation, the one aimed treasure.

The long lasting *abominable mystery*

When Charles Darwin found the puzzling phenomenon of angiosperms fossil record, an *abominable mystery* has been settled to be solved: how did the angiosperm dominance emerged and how flowers, their reproductive units, help them to thrive and evolve (Crepet, 2000)? And now, this long lasting mystery still needs to be solved. All the already gathered information about angiosperms life cycle, helped us to better understand this phenomenon, but every day, new players are being found, interacting during sexual plant reproduction process, to integrate the overall network of the male female crosstalk, that will lead to the generation of new seeds - the final aimed treasure.

This is, thus, a proposal to settle the overall map of this aimed treasure hunt, gather the known clues and trace the needed paths to reach this untangled mystery.

Remembering the basal, we need to retrace the scenario: angiosperms change from a sporophytic diploid generation to a gametophytic haploid generation. Inside the flowers, stamens and ovules generate two different types of spores, by meiotic divisions, the microspores and the megaspores, respectively. After mitotic divisions, these spores produce their gametophytes that will produce the gametes. The double fertilization event occurs when the two identical male gametes fuse with the two female gametes, oosphere and central cell, establishing a new diploid sporophytic generation and completing the plant's life cycle (Yadegari and Drews, 2004).

Arabidopsis thaliana is the important model plant, the well-known scenario to input all the needed quests, because it is small, with a good sequenced

genome and a rapid life cycle (Meinke, 1994). Inside A. thaliana carpel, ovules are enclosed within the ovary and in each ovule, there is a single megasporocyte that undergoes meiosis and produces four haploid megaspores, surviving only the functional megaspore (FM), after the others dyed by programmed cell death (PCD). After three successive mitotic divisions, the FM at the chalaza, forms a female polygonum gametophyte with seven cells and eight nuclei – the embryo sac (Yadegari and Drews, 2004; Palanivelu and Tsukamoto, 2012). At the micropylar pole near the opening of the two integuments, two synergids with their filiform apparatus, resulting from the complex invagination of the synergids cell wall, are present surrounding the egg cell. At the opposite pole, the chalazal pole, there are three antipodal cells, and the central zone is occupied by the central cell with two polar nuclei, that fuse before the arrival of the pollen tube (PT). Over 70% of the flowering plants share this type of female gametophyte (Maheshwari and Johri, 1950; Yadegari and Drews, 2004; Dresselhaus, 2006).

Inside the anther the male gametophytes are formed, also known as pollen grains. The mature pollen grain has, at the beginning, two haploid cells: a vegetative and a generative cell. The generative cell undergoes mitosis originating the two male gametes. In most plants, this mitosis occurs during the pollen tube growth, but, in Arabidopsis, it still occurs within the anther. The vegetative cell forms the PT that carries and releases the male gametes inside the female gametophyte (McCormick, 1993; Lord and Russell, 2002; McCormick, 2004).

Settling the treasure hunt map – double fertilization

The aimed treasure – the new seed generation – is only granted in flowering plants due to the double fertilization process, the unique key biological process that conceded the angiosperms success. The path to be followed starts once the stigmatic cells recognize a compatible pollen grain, it becomes properly hydrated, forming the PT, that will carry the two sperm cells into the ovule embryo sac (Figure 1). The rapid and controlled navigation of the PT is granted by a series of communications and consequent signalling cascades leading to various interactions between PT, sporophytic and gametophytic female tissues. The PT grows through the extracellular matrix of cells of the female reproductive tissue: stigma, style and transmitting tissue (TT) (Yadegari and Drews, 2004), until it gets the signals that direct it into an ovule. Once close to the ovule, the PT emerges from the septum, and the double fertilization moment starts. The PT that has grown along the funiculus, finally reaches the embryo sac entrance, the micropyle, and through the filiform apparatus of one of the two existing synergids, stops growing, ruptures and releases its two sperm cells. It is time for the process of simultaneous fertilization to occur, where one sperm cell fuses with the egg cell, generating the embryo while the other fuses with the central cell to form the endosperm, the basal tissue for the embryo nourishment (Palanivelu and Tsukamoto, 2012; Beale and Johnson, 2013; Dresselhaus and Franklin-Tong, 2013; Pereira et al., 2014). After PT entrance, the invaded synergid dies by PCD (Higashiyama et al., 2001; Dresselhaus, 2006; Dresselhaus e Franklin-Tong, 2013), and the persisting synergid will fuse with the endosperm after successful fertilization of both female gametes.

This plant organogenesis charming process will embody new structures of refined control of gene expression, based on a subtle communication occurring between neighbouring cells and on chemical signals that will induce cellular mechanisms of patterning and growth, tightly constrained in a complex regulatory network. Since the adhesion of the pollen grains to the stigma, every single step that will lead to the generation of the new seed, is being deeply studied, to map each player role in each tissue, in order to untangle this complex regulatory network of interacting genes.

Step by step: the clues

First interaction pollen-stigma

When a pollen grain lands at the stigma cells (Figure 2), it has to immediately face self-incompatibility (SI) responses, a pre-fertilization barrier avoiding the negative effects associated with inbreeding depression (Losdat *et al.*, 2014). Selective inhibition starts to prevent self-fertilization and also to grant the gene pool and genetic variation of plants (Kitashiba and Nasrallah, 2014). This process is controlled by the multiallelic *S*-locus in many plant species (Takayama and Isogai, 2005).

The complex exocytic subunit Exo70A1 in *A. thaliana* stigma, accepts a compatible pollen grain, acting on the polarized secretion of stigmatic cells to provide vesicles containing aquaporins, improving the water permeability, allowing the hydration of pollen and providing cell wall modifying enzymes to grant the stigma PT growth (Samuel *et al.*, 2009). Phosphatidylinositol-4-phosphate (PI4P) is crucial in this initial step, because mutants with low levels of PI4P on the stigma had slower hydration rates and maternal fertility defects due to failed pollinations (Chapman and Goring, 2010).

The VACUOLAR SORTING 41 protein (AtVPS41) is also a controlling factor, as the PTs of *atvps41* plants do not enter the female TT due to a compromised endocytic pathway (Hao *et al.*, 2016). The quiet studied signalling molecule Ca^{2+} can be an important intracellular signalling factor for the PT formation and its polarized growth (Iwano *et al.*, 2004; Pereira *et al.*, 2016a).

The 'minor' cell wall components and mysterious Arabinogalactan protein (AGP) family, always involved in distinct stages of the reproductive process, may also be essential regulatory components of PT growth (Lamport *et al.*, 2018). Apple blossom stigma receptivity depends on AGPs secretion and their disappearance is linked with PT growth (Losada and Herrero, 2012). The molecular properties and periplasmic location of the classical AGPs (Lamport *et al.*, 2006), make them the primary source of cytosolic Ca²⁺ oscillations, pectic plasticizers, working as Ca²⁺ signposts to the ovule (Lamport *et al.*, 2018). AGPs being present in the stigma cells and in growing PTs have roles in pollen–stigma interactions and pollen grain competence to initiate PT growth. AGP6, AGP11, AGP23 and AGP40, pollen specific AGPs, are included in the PT endosome machinery (Costa *et al.*, 2013; Pereira *et al.*, 2016a).

The gene, *O-FUCOSYLTRANSFERASE1* (*AtOFT1*) cannot be forgotten, as it plays a key role in PT penetration through the stigma-style interface and, according to genetic and cell biological evidences, a Golgi-localised fucosyltransferase system may be needed for PT growth through pistil tissues. Although yet unknown, the AtOFT1 substrates may include cell surface receptors or structural proteins secreted into the extracellular matrix. A nearly sterile *oft1* phenotype opens the path to find how putative proteins OFTs are utilized along pollen-pistil interactions (Smith *et al.*, 2018). Fasciclin-like AGPs (FLAs) are also recruited during this process as FLA3 is expressed in the pollen grains and PTs and it is involved in the microspores development affecting cellulose deposition (Li *et al.*, 2010).

The puzzling PT navigation along the TT

The rapid and controlled growth of PT is provided by an accurate communication network and signalling cascades during pollen-pistil interactions (Figure 3-4). The PT guidance process starts at preovular guidance, where any compatible PT travels in the extracellular matrix of TT or channel, from stigma to the ovary, later turning to ovular guidance, in a specific relation between one PT and its ovule (Higashiyama and Takeuchi, 2015; Kanaoka and Higashiyama, 2015; Mizuta and Higashiyama, 2018).

Mechanical, chemotropic, geometric and physiological orientation mechanisms may explain the precise growth of PTs towards the ovary (Heslop-Harrison, 1987; Lush, 1999; Chebli and Geitmann, 2007). There is a strict balance between cell turgor and cell wall resistance to keep the shape of PT, enabling the tip growth to expand and preventing the loss of the cell integrity (Winship et al., 2011). The accurate cell wall assembly are fundamental for PT growth. The polysaccharide pectin, an essential component of the primary cell wall, interferes with structure and flexibility during cell growth. Pectin methylesterases (PMEs) enzymes control spatially and temporally the deesterification of pectin, essential for PT integrity. PTs become unstable when pectin. The Vanguard1 (VGD1)-encoded loosing de-esterified pectin methylesterase mutant helps to demonstrate the importance of pectin in cell wall for PT integrity, as it presents *in vitro* bursting and *in vivo* delayed growth (Jiang et al., 2005; Bosch and Hepler, 2005). In addition, the Cl⁻ channel, MSL8 in Arabidopsis was shown to cause precocious PT burst before pollen arrival at the synergids (Li and Yang, 2018). Enzymatic production of reactive oxygen species (ROS) also affects PT growth. The characterization of the NADPH

oxidases RbohH and RbohJ (Respiratory burst oxidase homolog H and J) in Arabidopsis showed that the PTs tip growth was severely impaired in *rbohHrbohJ* double mutant. Both RbohH and RbohJ, whith Ca²⁺ binding EFhand motifs, showed Ca²⁺-induced ROS-producing activity at the PT tip of the plasma membrane. The PT slows down to couple the rate of exocytosis and cell expansion, possibly through the Ca²⁺ influx and K⁺ efflux (Lassig *et al.*, 2014; Kaya *et al.*, 2014). The affected amplitude and frequency of ion oscillation are detrimental to the precise coupling of tip growth and cell wall dynamics. The PT integrity depends on the cell wall composition and is controlled by its modifying enzymes and ROS, except for extensin (Jiang *et al.*, 2005; Woriedh *et al.*, 2013).

In *Torenia fournieri*, the PT has a unidirectional orientation (Higashiyama and Hamamura, 2008) maybe without having the need of the FG for the PT growth from stigma to style (Sogo and Tobe, 2005). With preovular guidance in mind, the chemocyanin of the stigma can be found to induce the PT chemotropism in *Lilium longiflorum*, with stigma cysteine-rich adhesins (SCAs) binding the PT, allowing the chemocyanin physical access to the plasma membrane, controlling and directing the PT growth (Park *et al.*, 2000; Kim *et al.*, 2003). The PT pectins may interact with the *A. thaliana* LIPID TRANSFER protein 5 (LTP5), facilitating its guidance (Chae *et al.*, 2009). In tomato, the CYSTEINE-RICH STIGMA SPECIFIC PROTEIN 1 (STIG1) interacts with a KINASE 2-type pollen receptor (LePRK2) and with phosphatidylinositol 3-phosphate (PI3P), as a signal peptide to promote PT growth (Tang *et al.*, 2004); In Arabidopsis, the PT growth is promoted by the pistil brassinosteroids (Vogler *et al.*, 2014), with the γ -aminobutyric acid (GABA) modulating the Ca²⁺ channels in the plasma membranes, regulating the PT signalling pathways (Palanivelu *et al.*, 2003; Yu

et al., 2014). During preovular guidance (Figure 3), the PT growth is, therefore, granted by several mechanisms (Mizuta and Higashiyama, 2018).

The TT connecting the stigma, style and ovules, supports the growth of PTs through an extracellular matrix (ECM) extremely rich in glycoproteins, polysaccharides, glycolipids and AGPs (Pereira *et al.*, 2015). The Arabidopsis TT is solid and composed of cylindrical cells (Lennon *et al.*, 1998; Erbar, 2003) that facilitates PT development and nutrition (Faure *et al.*, 2002; Crawford and Yanofsky, 2008). According to hormonal studies, the establishment of TT may be controlled by both auxin and cytokinin, which are responsible for the tissue polarity (Deb *et al.*, 2018) and thus, may play a key role along the complex interactions that control PT growth.

In *Nicotiana tabacum*, the AGP highly glycosylated TRANSMITTING TISSUE SPECIFIC (TTS) proteins are promoting PT growth and guidance to the ovules (Cheung *et al.*, 1995; Wu *et al.*, 2001). Transcription factors (TFs) like NO TRANSMITTING TRACT (NTT), HECATE 1 (HEC1), HEC2, HEC3, HALF FILLED (HAF), SPATULA (SPT) and water soluble chlorophyll proteins (AtWSCP) are important for TT development, because they induce the tissue PCD, after PT passage, which is also crucial for its own growth (Alvarez and Smyth, 2002; Crawford and Yanofsky, 2011; Boex-Fontvieille *et al.*, 2015; Mizuta and Higashiyama, 2018). The *ntt* mutants show abnormal development of the TT, with defects and slower growth of the PT, and a reduced fertilization rate (Crawford *et al.*, 2007). In addition, the *ntt* mutants have lower acidic polysaccharides, possibly related with AGPs contents, as AGPs are acidic glycoproteins, belonging to the TT.

PT may be guided by distinct cues, including K^+ , Cl^- , Ca^{2+} (Hepler *et al.*, 2006), glycoproteins (Sommer-Knudsen *et al.*, 1998), ROS (Foreman *et al.*, 2003),

nitric oxide (Prado *et al.*, 2016), peptides (Qu *et al.*, 2015) besides other complex signalling networks not yet elucidated (Leydon *et al.*, 2015).

In Arabidopsis, the Shaker family K^+ channel, SPIK, affects PT development. Loss of SPIK leads to precocious PT burst and consequent defected growth, suggesting a conserved role of K^+ homeostasis in PT growth and burst (Mouline *et al.* 2002).

AGPs are other guidance cues due to the evidences of their presence along the pathway from the stigma to the egg cell (Pereira *et al.*, 2014; Pereira *et al.*, 2016a). If pollen tubes grown *in vitro* acidify their growth medium (Feijo *et al.*, 1995), it may indicate that *in vivo*, they can dissociate AGP-Ca²⁺ of the TT, enabling PT to create a Ca²⁺ path to the ovule. The fertilization of multiple ovules can also be granted by the cooperation of multiple PTs (Heslop-Harrison *et al.*, 1985) together with the H⁺ efflux increase and the Ca²⁺ release from the locally abundant AGP-Ca²⁺ (Lamport *et al.*, 2018).

It is possible to find AGP1, AGP4 (JAGGER) and AGP19 in Arabidopsis TT, with high levels of expression along this tissue (Yang *et al.*, 2007; Pereira *et al.*, 2014; Pereira *et al.*, 2016a, Pereira *et al.*, 2016b). AGP19 has high expression in the style, ovary walls, TT and siliques, and the *agp19* mutants have low flower production and smaller siliques (Yang *et al.*, 2007). AGP12 and AGP15 also located along the PT pathway, can contribute nutritionally to the PT growth, from the top of the stigma to the base of the pistil, facilitating its movement, guiding to its targets, or even making it competent for being received by the embryo sac (Pereira *et al.*, 2014).

An AGP sidechain fragment designated AMOR, is the PT activation molecule for response to the synergids LURE guidance peptides in Torenia (Okuda *et al.*, 2009). AMOR was identified as a methyl-Oglucuronosyl-b-D-galactose

(Mizukami *et al.*, 2016), possibly a key component of the AGP Ca²⁺-binding motif, a Ca²⁺ carrier and good company for the AGP and Ca²⁺ couple (Lamport *et al.*, 2018). AMOR has a differential localization in AG sugar chains specific structures, indicating that AG biosynthesis enzymes are controlling AGPs functions and their sugar chains (Su and Higashiyama, 2018).

However, it is still not entirely clear how all TT components interact with PT and besides the importance of these interactions, the different components present in the TT, point to a contact-mediated ability provided by the stigma and style, for the PT to become aware of the signals that direct it to the ovule (Palanivelu and Preuss, 2006). The TT is thus, a key tissue for the quick growth of the PT throughout the pistil, but also to grant it the ability to move accurately to the ovule (Figure 4).

Ovular guidance: funicular and micropylar guidance

The PT has to emerge, exiting from the TT through a very narrow space between the cells of the septum (Palanivelu and Tsukamoto, 2012), facing all the quests and tissues that trigger its emergence (Higashiyama and Takeuchi, 2015), during a crucial transition phase from preovular to ovular guidance (Mizuta and Higashiyama, 2018). When PT emerges from the septum, it faces a funicular guidance phase from the placenta, along the funiculus and afterwards a micropylar guidance, where it navigates from the funiculus to the micropyle (Shimizu and Okada, 2000; Mizuta and Higashiyama, 2018). This pathway is dependent on signals from sporophytic tissues and from the female gametophyte (Pereira *et al.*, 2016a) (Figure 5).

In Arabidopsis, two potassium (K^+) transporters CATION HYDROGEN EXCHANGER 21 (CHX21) and CHX23 present in PTs, are able to block their growing along the funiculus. The *chx21* and *chx23* mutants show normal PT growth through the TT, but are unable to grow along the funiculus and into the

ovule, without perceiving the signals that come from the ovules (Lu *et al.*, 2011). PT competence for signalling is also dependent on the MITOGEN-ACTIVATED PROTEIN KINASE (MPK3) and MPK6 (Guan *et al.*, 2014), with defective PTs in the *mpk3mpk6* mutant along the funicular guidance phase. PTs can only pass through the TT with the controlling help of the PHYTOSULFOKINE (PSK), as its mutants show very low fertilization rate and loss of funicular orientation (Stührwohldt *et al.*, 2015). The knowledge path on the mechanisms that regulate this stage is now clearer, but sparse, and multiplying quests are trying to find new cues stage, especially under female control.

AAn ABNORMAL POLLEN TUBE GUIDANCE 1 (APTG1), a mannosyltransferase (Dai *et al.*, 2014), acts in the glycosylphosphatidylinositol (GPI) synthesis of the COBRA-LIKE 10 protein (COBL10) (Li *et al.*, 2013). The COBL10 protein, located at the PT end, plays a crucial role in its orientation and growth. Mutations in COBL10 generate gametophytic male sterility due to low PT growth and difficult directional sensing in the female TT. The apical pectin cap deposition and cellulose microfibrils were broken in *cobl10* PTs. (Li *et al.*, 2013).

After a funicular guided navigation period, the PT steps into the micropylar guidance stage. When reaching the micropylar region, PTs permeate through the opening between the ovules integuments until one of the synergid cells, an active and typically secretory cell (Huang and Russell, 1992). The *wuschel-7* (*wus-7*) mutant has a short inner integument (Lora *et al.*, 2018) and 35% of its ovules have defected embryo sacs. In 65 % of wus-7 ovules, PTs do not emerge in the micropyle in 65%, suggesting a partial effect of the inner integument in PT guidance. It is granted that the functional FG influences the PT attraction to the ovule (Ray *et al.*, 1997; Shimizu and Okada, 2000), thus being the structure that produces the ovular attracting molecules (Takeuchi and Higashiyama, 2016). Based on *Torenia fournieri* evidences, synergids are the female cells

responsible for the ovular PT attraction. *Torenia* has a bare embryo sac, which is projected to the outside of the ovule through the micropyle when mature, thus allowing an easy access to the embryo sac cells. It was shown that, *in vitro* PTs were attracted directly to the bare embryo sacs and no longer leaving the micropylar region, due to the embryo sac production of attraction signals (Higashiyama *et al.*, 2001). Later, a laser ablation performed on different gametophytic cells found that the synergids were providing the PT guidance by diffusible chemical signals (Higashiyama *et al.*, 1998; Higashiyama *et al.*, 2001; Higashiyama and Hamamura, 2008; Horade *et al.*, 2013).

An equivalent scenario happens in Arabidopsis with the MYB98 TF expressed in the synergids. The *myb98* mutant has an abnormal filiform apparatus, which blocks the secretion of attractive molecules, thus the PT cannot find the micropyle (Kasahara *et al.*, 2005; Punwani *et al.*, 2008). It is necessary to consider that the interaction between the egg and central cells is also important for the ovular guidance, since a laser disruption of the egg cell or the polar nucleus of the central cell can severely inhibit the PT attraction (Susaki *et al.*, 2015). Synergids are, therefore, together with the egg and central cell, regulating the PT attraction during the last stage of its journey, (Dresselhaus, 2006; Pereira *et al.*, 2016); Mizuta and Higashiyama, 2018).

So there are attractive molecules produced by the synergids, identified by several studies in different plant species. In *T. fournieri*, cysteine-rich peptides (CRPs) from the defensin-like proteins subgroup, TfLURE1 and TfLURE2 were identified as PTs attractive molecules inducing micropylar guidance (Okuda *et al.*, 2009), capable of attracting PTs also *in vitro* (Higashiyama, 2010). LURES were also identified in *A. thaliana* (AtLURES), *A. lyrata* (AlLURES) (Takeuchi and Higashiyama, 2012) and *T. concolor* (TcCRP1) (Kanaoka *et al.*, 2011). These small *A. thaliana* proteins, only expressed in the synergids, are secreted

through the filiform apparatus (Takeuchi and Higashiyama, 2012). The mutants magatama3 (Shimizu and Okada, 2000), myb98 (Kasahara et al., 2005) and central cell quidance (Chen et al., 2007) were found to be defective in micropylar guidance due to no AtLURE1 expression, giving to the AtLURE1 the control for the PT attraction (Takeuchi and Higashiyama, 2012). The novel interaction partner of CCG, the CCG-binding protein CBP1 seems to co-regulate the expression level of a subset of CRPs, including LURES, by recruiting the Mediator complex and RNA Pol II with CCG in the central cell (Li et al., 2015). Diverse genes expressed in the central cell and synergid, are downregulated in ccq ovules, such as MYB98, responsible for LUREs transcription regulation. Mutation of MAA3 affects PT attraction in A. thaliana. MAA3 encodes a Sen1plike RNA helicase and is expected to control target mRNA molecules, whose products are needed to guide PTs (Shimizu et al., 2008). Moreover, recombinant AtLURE1 peptides expressed in *Torenia* are capable of attracting Arabidopsis PTs, when in vitro Torenia ovules are placed close to the tip of A. thaliana PTs (Takeuchi and Higashiyama, 2012). These peptides are thus, enough to overcome reproductive barriers in ovule orientation, even among distant species (Mizuta and Higashiyama, 2018).

The unfertilized *A. thaliana* ovules send diffusible attractive molecules (Palanivelu and Preuss, 2006; Takeuchi and Higashiyama, 2012), but when the ovules are invaded by PTs, these molecules are no longer produced, so this may be enough to prevent the attraction of more PTs. But it may also be considered that PTs can rapidly release diffusible repellents, without no concrete evidence yet (Palanivelu and Preuss, 2006). The specific *Zea mays EGG APPARATUS1* gene (*ZmEA1*) of maize encodes a small protein present not only in the synergids, but also in the egg cell, filiform apparatus and micropylar nucellus, and is necessary for micropylar orientation (Márton *et al.*, 2005;

Márton *et al.*, 2012). No homologous candidate was found in *A. thaliana*, but there is no doubt about the importance of embryo sac cells for a successful PT guidance.

GAMETE-EXPRESSED3 (GEX3) encodes a membrane protein and is expressed, both in male and female tissues. Micropylar PT guidance defects were caused by aberrant expression of *GEX3* in the egg cell (Kanaoka and Higashiyama, 2015).

In turn, AGPs are also in ovule and synergid tissues as attractive and/or signalling molecules for the PT guidance, considering their immunolocalization evidences in ovules and in the synergids filiform apparatus of different species (Coimbra and Salema, 1997; Coimbra and Duarte, 2003; Coimbra *et al.*, 2007; Lopes *et al.*, 2016). AGP23 is intensively present upon fertilization, inside the embryo sac, where the PT bursts (Pereira *et al.*, 2016a).

The JAGGER protein, AGP4, is essential to prevent the polytubey occurrence. The *jagger* persistent synergid does not undergo PCD and more than one PT is attracted to the ovule, in a polytubey effect. JAGGER controls the persistent synergid death and ends the multiple PTs attraction (Pereira *et al.*, 2016b).

Arabidopsis has LURE1 receptors such as leucine-rich repeat (LRR) kinase type receptors (RLKs): MALE DISCOVERER 1 (MDIS1), MDIS1-INTERACTING LIKE KINASE RECEPTOR (MIK1, MIK2) (Wang *et al.* 2016) and POLLEN RECEPTOR-LIKE KINASE 6 (PRK6) (Takeuchi and Higashiyama, 2016). The recombinant AtLURE1 may induce heterodimerization of MDIS, MIK1 and MIK2 (Wang *et al.*, 2016). These three receptors, at the PT plasma membrane showed their knockout mutants with micropylar guidance defects. The other PT-specific receptors found by Takeuchi and Higashiyama (2016), the PRK (PRK6) receptors, at the PT tip, enable it to detect AtLURE1, controlling its elongation

and attraction. The detection of AtLURE1 by PRK6 is achieved in cooperation with other PRK family receptors like PRK1, PRK3 and PRK8, but PRK6 is the key receptor. PRK6 also interacts with pollen-expressed ROPGEFs (Rho of plant guanine nucleotide-exchange factors), which are important for PT growth through activation of the signalling switch Rho GTPase ROP1 (Kaothien *et al.*, 2005; Zhang and McCormick, 2007; Takeuchi and Higashiyama, 2017).

PT guidance is also affected by nitric oxide (NO) and D-serine. NO has diverse effects on plant development as a growth regulator, and acts as a negative chemotropic cue during *in vitro* PT growth and re-orientation, which was mediated by a cGMP signalling pathway (Prado *et al.*, 2004). Mutations in *A. thaliana NITRIC OXIDE SYNTHASE1 (AtNOS1*) generates low fertility because NO production is affected (Guo *et al.*, 2003), indicating some possible fuctions of NO during PT guidance. Further experiments with *semi-in vivo* isolated ovules and PTs confirmed that NO is necessary for ovular PT guidance in *Lilium longiflorum*, depending this signalling pathway on Ca²⁺ signalling (Prado *et al.*, 2008). Still it is unclear how the PT surface perceives NO.

The Ca²⁺ signalling network can be puzzling enough due to the existence of multiple Ca²⁺ channels, distinct regulation levels and plenty of Ca²⁺ PTs sensors. The Cyclic nucleotide-gated ion channel 18 (CNGC18) appears to be, among other Ca²⁺ channels expressed in pollen, an essential required Ca²⁺ channel in PT growth and guidance (Frietsch *et al.*, 2007). The two point-mutated *cngc18* mutants obtained showed that the cyclic nucleotide (cNMP) activation of CNGC18 is likely involved in ovular guidance, but not PT growth (Gao *et al.*, 2016). Thus, Ca²⁺ influx may regulate the PT growth and guidance in a double manner: basal influx for PT growth and the cNMP-activated influx for PT reorientation (Li *et al.*, 2018).

LOST IN POLLEN TUBE GUIDANCE 1 (LIP1) and LIP2 are also involved in the AtLURE1-dependent orientation mechanism (Liu *et al.*, 2013; Pereira *et al.*, 2016a; Higashiyama, 2018). LIPs do not bind directly to the attractive molecules, including the AtLURE1 peptides, due to the lack of an extracellular region, but they may form complexes with direct receptors of the LUREs peptides, acting in the signal transduction of the PT cytoplasm (Takeuchi and Higashiyama, 2016).

It is also important to map distinct proteins from the endoplasmic reticulum (ER), since the folding of these proteins may be important for the PT competence, enabling it to respond to attraction or orientation signals (Braakman and Herbert, 2013). POLLEN DEFECTIVE IN GUIDANCE 1 (POD1) is a ER protein that interacts with the chaperone CALRETICULIN 3 (CRT3) controlling the PT membrane proteins folding and regulating their signalling (Li *et al.*, 2011). Several molecules are acting in an intrinsic signalling pathway, during the micropylar guidance stage, however, their composition and entanglement is sparse (Pereira *et al.*, 2016a; Mizuta and Higashiyama, 2018) and are in need of new quests.

The aimed treasure is beyond synergids

Finally, the PT reaches the micropyle of its target ovule, entering the embryo sac through one of the synergids, stops its growth and bursts, releasing the two male gametes (Leshem *et al.*, 2013), that will fuse with the female ones, while the multiple PTs blocking system acts (Figure 6). The receptive synergid undergoes PCD shortly before the entry of the PT or even upon its entry (Faure *et al.*, 2002). In a complex signalling system and communication between the PT and the female tissues, the aimed treasure will be finally reached by generating a new seed, granting the reproductive success of flowering plants (Pereira *et al.*, 2016a).

Synergids gather the utmost point of PT attractive molecules, able to receive and rupture it. The first PT receptor found, from the synergid filiform apparatus, was a serine/threonine-type FERONIA/SIRENE (FER/SRN). *fer/srn* mutation leads to the continuous growth of the PT within the embryo sac, with no release of the male gametes (Huck *et al.*, 2003; Escobar-Restrepo *et al.*, 2007; Rotman *et al.*, 2008).

FER is a receptor like kinase (RLK), from the *Catharanthus roseus* CrRLK1L-1 subfamily, with an extracellular portion similar to a malectin domain, so its binding molecule can be a carbohydrate or a cell wall glycoprotein (Lindner *et al.*, 2012; Pereira *et al.*, 2016a). FER is thus, an essential factor to stop the PT growth, its reception, correct rupture and gametes release and may also be the link between extracellular events and intracellular orientation (Stegmann *et al.*, 2017).

LORELEI (LRE) is a GPI anchor protein predominantly expressed in the synergids and the *lre* mutant has a similar phenotype to the *fer/srn* mutation (Capron *et al.*, 2008). LRE and FER work together in the PT synergid reception with LRE acting as a co-receptor with FER (Hafidh *et al.*, 2016; Liu *et al.*, 2016). LRE and LLG1 (LORELEI-like GPI anchored protein 1) interact with FER in the ER lumen, acting as chaperones to conduct FER to the filiform apparatus and regulate PT rupture (Li *et al.*, 2015).

NORTIA (*NTA*) also has a *fer*-like phenotype (Kessler *et al.*, 2010). NTA acts in a FER dependent way, in the same signalling cascade. Its C-terminal calmodulin domain suggests ability for the Ca²⁺ oscillations perception in the synergids (Iwano *et al.*, 2012). Both *Ire* and *nta* mutants have a similar *fer* mutant phenotype, but not fully penetrant as this last one. They play important roles in PT-synergid interaction, but maybe acting redundantly with another yet

unknown factor, still need to be known in order to understand this signalling mechanism (Pereira *et al.*, 2016a).

In *A. thaliana*, in parallel to the FER signaling pathway, vacuolar acidification in synergid cells is impaired in low expression levels of AP1G, the γ -subunit of the tetrameric ADAPTOR PROTEIN1 (AP1). Thus, upon PT arrival, there is a delay in synergid cell degeneration. AP1 regulates protein sorting at the trans-Golgi network/early endosome. In synergids cells with functional loss of V-ATPases, the *ap1g* mutant has a defect in PT reception, indicating that acidification of the vacuole represents an important mechanism for its degeneration (Wang *et al.*, 2017).

Evidences were found on others female gametophyte CrRLK1L receptors in the PT perception (Galindo-Trigo *et al.*, 2018). HERCULES RECEPTOR KINASE1 (HERK1) and ANJEA (ANJ) are strongly localised at the filiform apparatus of the synergid cells, with a double mutant PT overgrowth. A yeast two hybrid assay showed that HERK1 and ANJ extracellular membrane domains interacted with LRE, leading the assumption that HERK1-LRE and ANJ-LRE may belong to a filiform apparatus receptor complex, responsible for mediating PT reception.

TURAN (TUN), encoding a UDP-glycosyltransferase protein, may also be involved in the PT reception, with a mutation revealing again a PT overgrowth phenotype (Lindner *et al.*, 2015). The TF VERDANDI (VDD) belongs to the Reproductive Meristem (REM) family, and *vdd* revealed defects in the cellular identity of the antipodals and synergids, with no PT explosion, and thus the continuation of PT growth after reaching the synergids (Matias-Hernandez *et al.*, 2010). Recently, VDD and VALKYRIE (VAL), also a REM family member, were shown to act as a complex regulating the receptive synergid death, and PT death, by bursting (Mendes *et al.*, 2016). A small group of chimeric AGPs, the EARLY NODULIN-LIKE (ENODLs), EN11-15, are essential for the PT reception in Arabidopsis, with high expression levels found in the funiculus and in the ovules. EN14 and EN15, are present in the synergids filiform apparatus. A five-fold mutant revealed a similar phenotype to the *fer* mutant, with, at least EN15 interacting with FER, and thus, maybe being involved in the signalling pathway activation, essential for the coordination of the male-female communication needed in the double fertilization process (Hou *et al.*, 2016).

There are also male players involved in the synergids PT reception and rupture, notoriously interacting in a coordinated way with the female players. The first identified set of pollen-specific MYB transcription factors MYB97, MYB101 and MYB120 act controlling the PT reception and regulating its gene expression, including genes encoding the secretion of small peptides (Liang *et al.*, 2013).

The Ca²⁺ pump ACA9 expressed in the PT was also shown to regulate PT growth and rupture in the synergid (Schiott *et al.* 2004).

ANXUR 1 (ANX1) and ANX2 are receptor like kinases from the PT and FER homologues that induce PT burst (Boisson-Dernier *et al.*, 2009). When the receptive synergid goes into PCD, responses of Ca²⁺ are activated (Boisson-Dernier *et al.*, 2013, Ngo *et al.*, 2014). The overexpression of ANX1/ANX2 receptors, at the plasma membrane of the PT tip, lead to the blockage of its growth by excessive activation of exocytosis and accumulation of secreted material by the cell wall. The *anx1anx2* double mutant stops the PT development, by premature tip rupture (Boisson-Dernier *et al.*, 2013). It is accepted that ANX blocks the PT rupture and premature gametes release, so it continues to grow in the female tissues, granting the cell wall integrity, until it reaches the FG (Boisson-Dernier *et al.* 2009; Miyazaki *et al.*, 2009; Pereira *et al.*, 2016a). The results from Muro *et al.* (2018) have shown that the PT tip

localization of ANX receptors is strictly mediated by two ANTH domaincontaining proteins via the clathrin-mediated endocytosis pathway. *phosphatidylinositol binding clathrin assembly protein 5a* (*picalm5a*) *picalm5b* double mutant exhibited an altered distribution of ANX receptors in the PT and an identical phenotype to *anx1anx2*: a premature burst of PTs during their growth.

It has to be an extremely coordinated mechanism of signalling between the female and male cells during these interactions. Once in the FG, the FER-dependent signalling cascade located in the synergid is activated to mediate the reception and fertilization, while the ANX-dependent signalling present in the PT is blocked, enabling the PT rupture and gametes release (Boisson-Dernier *et al.*, 2009; Miyazaki *et al.*, 2009). This is an interesting example, where duplicate sibling genes of an ancestral gene work on cells of the opposite sex during male-female interactions (Higashiyama, 2018).

RAPID ALKALINIZATION FACTORs (RALFs) are secreted peptides, working as extracellular signals and translated into the cell. These peptides are spread along the plant kingdom and Arabidopsis has 37 members (Murphy and De Smet, 2014), regulating various physiological processes (Stegmann *et al.*, 2017).

RALF peptides expression were searched in the PTs by Mecchia *et al.* (2017), and RALF4 and RALF19, the closest homologues, were chosen. The *ralf4ralf19* double mutant showed a similar phenotype to *anx1anx2*: no PT growth maintenance. So RALF4/19 are crucial for the PT growth, possibly acting upstream of ANX1/2.

Searching for more CrRLK1Ls involved in PT reception, Ge *et al.* (2017) found BUPS1 and BUPS2, new receptors predominantly expressed in PT, also with

bups1bups2 mutant having a similar phenotype to *anx1anx2*. Settling the map, it is possible to find RALF4/19 interacting with BUPS1/2 and these two peptides interacting with ANX1/2. Besides, BUPS1/2 and ANX1/2 interact with each other. Considering the same phenotype showed by *anx1anx2* and *bups1bups2* double mutants of premature PT rupture, it may be suggested a model in which ANX1/2 and BUPS/2 form a heterodimer. Quests are needed to find whether RALF4/19 binding induces heterodimerization of ANX and BUPS. RALF34 expressed only in the female tissues, is the closest homolog of RALF4/19 and induces PT rupture at a 2 nM concentration. RALF34 competes with RALF4/19 *in vitro*, in the BUPS1 and ANX interaction; so RALF34 can control PT rupture through female tissues, still to understand how this happens, and further studies will be needed (Liang and Zhou, 2018).

The autocrine signalling mediated by RALF4/19 and ANX-BUPS is critical to maintain a long distance PT growth and the tip cell wall integrity. Quests remaining to answer are how external autocrine and paracrine signals for the PT tip change during the long journey and how PTs respond to them (Higashiyama, 2018).

The improved understanding of the PT guidance and fertilization process includes a limited number of molecular players. Considering the FER and ANXUR1/2 description and their role in Ca²⁺ mediated fertilization, members of the CrRLK1L family had stepped into the regulation of polarized growth. ERULUS (ERU), also a CrRLK1L protein, has been described as a core root hair (RH) regulator, involved in the establishment of a functional apical [Ca²⁺]_{cyt} gradient (Bai *et al.*, 2014).

In vivo eru pollen are less competitive than WT pollen in terms of plant fertilization efficiency (Schoenaers *et al.*, 2017) and also with aberrant ovule

targeting of PTs, more mutant PTs growing around the funiculus with multiple turns before targeting the micropyle. This is a similar phenotype to the *lip1lip2* double mutant, which is defective in two PT plasma membrane localized RLKs (Liu *et al.*, 2013). LIP1 and LIP2 are crucial players in the receptor complex regulating PT guidance in response to the micropyle-secreted AtLURE1 signalling peptide. One could, thus, suggest that ERU could function as LURE receptor, and as such could regulate PT targeting in a complex with LIP1 and LIP2 (Schoenaers *et al.*, 2017).

After all the PT-synergid interactions, PT ruptures and releases its content, occurring the male-female gamete fusion (Figure 7). For the successful gametes fusion, three crucial players are interacting: GAMETE EXPRESSED 2 (GEX2), GENERATIVE CELL SPECIFIC 1/HAPLESS 2 (GSC1/HAP2) and EGG CELL1 (EC1) (Mori *et al.*, 2006; Besser *et al.*, 2006; Sprunck *et al.*, 2012). GEX2 encodes a membrane-expressed protein of the male gametes, containing extracellular immunoglobulin-like domains and required for gametes fusion (Mori *et al.*, 2014). The male gametes with the *gcs1hap2* mutation when delivered to the ovules cannot initiate fertilization, and the released gametes remain in the fusion site with the female gametes, leading eventually to the attraction of multiple PTs (Mori *et al.*, 2006; von Besser *et al.*, 2006). EC1 is a cysteine-rich protein, a possible activating molecule of the male gametes, to permit gametes fusion and *ec1* quintuple mutants show that successful male-female gamete interactions prevent multiple-sperm cell delivery (Sprunck *et al.*, 2012).

Recently, RBOHH and RBOHJ were found to act downstream of ANXs, controlling ROS production during PT growth (Boisson-Dernier *et al.*, 2013). Besides, the tube burst phenotype of both *anx1anx2* and *rbohhrbohJ* can be suppressed by a point mutation (MRIR^{240C}) in the receptor-like cytoplasmic

kinase MARIS (MRI) (Boisson-Dernier *et al.* 2015), while RALF4 and 19 were evidenced to function upstream of MARIS (Mecchia *et al.*, 2017). Thus, RALF-BUPS/ANX complex may be regulating downstream RBOHH, RBOHJ and MRI, to keep PT integrity during its growth. Moreover, RALF4 and 19 bind cell wall extensin LRX8, LRX9 and LRX11 expressed in PT, having a vital role in cell wall materials deposition (Mecchia *et al.*, 2017).

Ca²⁺ plays, once more, a preponderant role in regulating the release of male gametes and fertilization (Iwano *et al.*, 2012). Several molecules initiate their signalling mechanism and interaction after Ca²⁺ oscillations events. Specific signatures of this secondary messenger are essential to control the PT and the receptive synergid PCD, for male-female gametes fusion and double fertilization success (Denninger *et al.*, 2014; Ngo *et al.*, 2014).

Conclusion and outlook

The aimed treasure can finally emerge, after a stunning PT pathway, full of traces and signals of a complex genetic regulatory network, that step by step, establishes a precise and refined crosstalk, since the landing of the pollen on stigma to the generation of the new born seed. This proposal tried to settle the overall map of this aimed treasure hunt, gathering the known clues and, in a clearer and simpler step by step set of frameworks, state the acquired knowledge to a further establishment of links and filling of gaps.

For sure, a broader framework of signalling pathways should include the interaction between the hormonal signalling control of tissues domains positioning with the complex genetic regulatory network interacting along the double fertilization process, if we consider the reciprocal dependency of the PT on the TT.

Besides all the involved players, one could also highlight fundamental keys of this overall PT journey that will surely make it able to reach the aimed treasure, as it is the quiet studied signalling molecule Ca²⁺, fully present during all the stages of the double fertilization process, and their humble supporters, the AGPs family, responsible for vital interactions in each step.

Nevertheless, new key players are now being discovered to complete the overall perspective on the network of the male female crosstalk during this sexual plant reproduction process.

Conflict of Interest Statement

The authors declare that this review was written in the absence of any potential conflict of interest.

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Figure 1. Schematic representation of the treasure hunt scenario - an Arabidopis pistil. st, stigma; ov, ovules; pg, pollen grain; pt, pollen tube.





Figure 2. The players of pollen-stigma interactions and the clues for their roles.

Figure 3. The players involved in the preovular guidance stage and the clues for their roles.



Figure 4. The players of PT navigation along the TT and the clues for their roles.



Figure 5. The players involved in the ovular guidance stage, divided in funicular and micropylar guidance steps, and the clues for their roles.



Figure 6. The players involved during the PT- synergids interaction, and the clues for their roles.



Figure 7. The players involved during gametes fusion interaction, and the clues for their roles.