

Loss of LORELEI function in the pistil delays initiation but does not affect embryo development in *Arabidopsis thaliana*

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Double fertilization, uniquely observed in plants, requires successful sperm cell delivery by the pollen tube to the female gametophyte, followed by migration, recognition and fusion of the two sperm cells with two female gametic cells. The female gametophyte not only regulates these steps but also controls the subsequent initiation of seed development. Previously, we reported that loss of *LORELEI*, which encodes a putative glycosylphosphatidylinositol (GPI)-anchored protein, in the female reproductive tissues causes a delay in initiation of seed development. From these studies, however, it was unclear if embryos derived from fertilization of *lre-5* gametophytes continued to lag behind wild-type during seed development. Additionally, it was not determined if the delay in initiation of seed development had any lingering effects during seed germination. Finally, it was not known if loss of *LORELEI* function affects seedling development given that *LORELEI* is expressed in eight-day-old seedlings. Here, we showed that despite a delay in initiation, *lre-5/lre-5* embryos recover, becoming equivalent to the developing wild-type embryos beginning at 72 hours after pollination. Additionally, *lre-5/lre-5* seed germination, and seedling and root development are indistinguishable from wild-type indicating that loss of *LORELEI* is tolerated, at least under standard growth conditions, in vegetative tissues.

Double fertilization is unique to flowering plants. Upon female gametophyte reception of a pollen tube, the egg and

central cells of the female gametophyte fuse with the two released sperm cells to form zygote and endosperm, respectively and initiate seed development.¹ The female gametophyte controls seed development by (1) repressing premature central cell or egg cell proliferation until double fertilization is completed,¹⁻³ (2) supplying factors that mediate early stages of embryo and endosperm development^{1,4,5} and (3) regulating imprinting of genes required for seed development.^{1,6}

The molecular mechanisms underlying female gametophyte control of early seed development are poorly understood. Although much progress has been made in identifying genes and mechanisms by which the female gametophyte represses premature central cell or egg cell proliferation until double fertilization is completed and regulates imprinting of genes required for seed development,^{1,6} only a handful of female gametophyte-expressed genes that affect early embryo^{7,8} and endosperm⁹ development after fertilization have been characterized. This is particularly important given that a large number of female gametophyte-expressed genes likely regulate early seed development.⁵

We recently reported on a mutant screen for plants with reduced fertility and identification of a mutant that contained a large number of undeveloped ovules and very few viable seeds.¹⁰ TAIL-PCR revealed that this mutant is a new allele of *LORELEI* (*LRE*) [*At4g26466*].^{10,11} Four *lre* alleles have been reported;¹¹ so, this mutant was designated *lre-5*.¹⁰ The *Arabidopsis* *LORELEI* protein contains 165 amino acids and possesses sequence features indicative of a glycosylphosphatidylinositol

Key words: *LORELEI*, glycosylphosphatidylinositol (GPI)-anchored protein, embryogenesis, *DD45*, seed germination, primary and lateral root growth, seedling development

Submitted: 09/10/10

Accepted: 09/11/10

Previously published online:

www.landesbioscience.com/journals/psb/article/13598

DOI: 10.4161/psb.5.11.13598

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Addendum to: Tsukamoto T, Qin Y, Huang Y, Dunatunga D, Palanivelu R. A role for *LORELEI*, a putative glycosylphosphatidylinositol-anchored protein, in *Arabidopsis thaliana* double fertilization and early seed development. *Plant J* 2010; 62:571-88; PMID: 20163554; DOI: 10.1111/j.1365-313X.2010.04177.

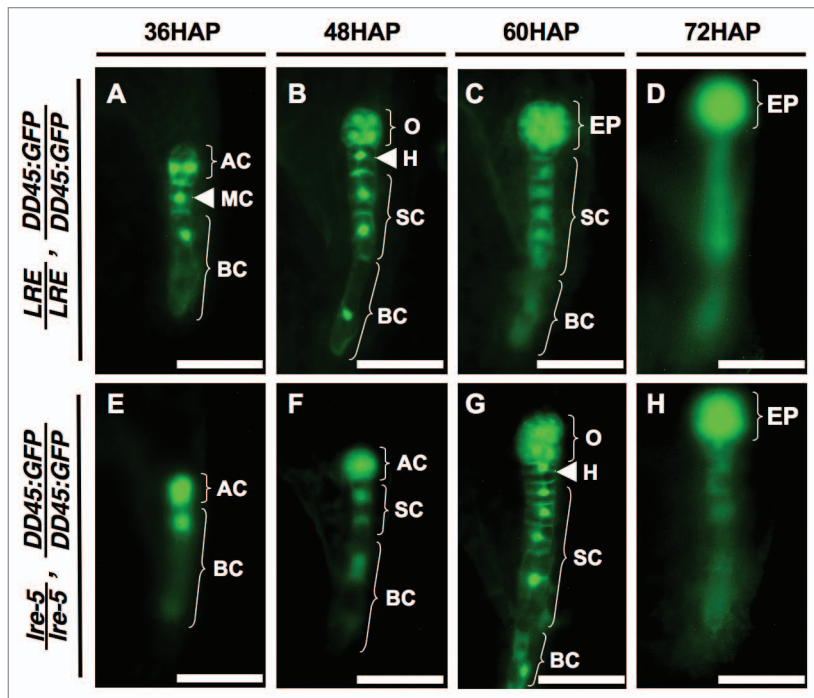


Figure 1. Analysis of embryo development in self-pollinated pistils. Fluorescent images of *DD45:GFP* expression in embryos in developing seeds in manually self-pollinated wild-type (A–D) or *lre-5/lre-5* pistils (E–H) carrying the *DD45:GFP/DD45:GFP* transgene and observed at the indicated HAP. In (D and H), the *DD45:GFP* expression in nuclei is no longer clearly visible due to thickening of the seed coat by 72 HAP. AC, MC and BC, apical, middle and basal cells, respectively, in a proembryo; O, octant stage of apical cell division; H, hypophysis; SC, suspensor cells; EP, embryo proper. Scale bars, 50 μ m.

(GPI)-anchor containing cell surface protein. GPI-anchors serve as an alternative to transmembrane domains for anchoring proteins in cell membranes and GPI-anchored proteins participate in many functions including cell-cell signaling.¹²

Loss of LORELEI Function in the Female Reproductive Tissues Results in a Delay of Seed Development Initiation

Most *lre-5* female gametophytes in *lre-5/lre-5* pistils do not allow pollen tube reception; consequently, these ovules remain unfertilized.¹⁰ The *lre-5* mutation is partially-penetrant and female gametophyte-specific; consequently, a majority of the *lre-5* female gametophytes (~73%) do not undergo normal pollen tube reception. In those that do, we demonstrated that initiation of seed development is delayed and that this delay originates after *lre-5/lre-5* ovules complete double fertilization.¹⁰ We identified this delay by pollinating *GRP23:GUS* pollen on *lre-5/lre-5* pistils and monitoring

GRP23:GUS expression in *lre-5/lre-5* ovules after fertilization. *GRP23:GUS* is expressed not only in pollen tubes but also in developing seeds from the zygote and endosperm nuclear proliferation stages.¹³ We reasoned that in *lre-5/lre-5* pistils, the ~27% of ovules that induce pollen tube reception¹⁰ would allow monitoring of post-fertilization *GRP23:GUS* expression and seed development. By 16 hours after pollination (HAP), 27.1% of all *lre-5/lre-5* ovules that induced pollen tube reception did not show *GRP23:GUS* expression in proliferating endosperm nuclei like that in wild type.¹⁰ Even by 24 HAP, 25.4% of all ovules continued to show GUS staining that is only indicative of pollen tube reception completion and lacked expression elsewhere in the embryo sac, indicating a delay in the initiation of early seed development. Importantly, the delay in initiation of seed development was not rescued by a wild-type *LORELEI* delivered through pollen (note that the *LORELEI* locus is wild-type in *GRP23:GUS* pollen). Together, these results suggested that

loss of *LORELEI* function in the female reproductive tissues causes the delay in initiation of seed development.

From these studies, however, it was unclear if embryos derived from fertilization of *lre-5* gametophytes continued to lag behind wild-type during seed development. Additionally, it has not been determined if the delay in initiation of seed development had any lingering effects during seed germination. Finally, it is not known if loss of *LORELEI* function affects seedling development given that *LORELEI* is expressed in eight-day-old seedlings.¹⁰ In this report, we used an embryo-expressed marker (*DD45:GFP*)¹⁴ to monitor *lre-5/lre-5* embryos during seed development. Additionally, we analyzed seed germination, seedling growth and root development in *lre-5/lre-5* plants to examine the effects of loss of *LORELEI* on these aspects of plant development.

Despite a Delay in Initiation *lre-5/lre-5* Embryos Recover and Complete Embryo Development

To examine the role of *LORELEI* in embryo development, we crossed *lre-5/lre-5* pollen onto pistils of *DD45:GFP/lre-5* plants¹⁴ and from the F2 progeny identified plants that are homozygous for both *lre-5* and the *DD45:GFP* transgene. We examined *DD45:GFP* expression from 36 HAP onwards in developing seeds within manually self-pollinated *lre-5/lre-5*, *DD45:GFP/DD45:GFP* siliques. Ovules closest to the stigma receive pollen tubes approximately six hours before those near the pedicle;¹⁵ therefore, only 15 rows of ovules from the stigma end were analyzed. In every wild-type seed, by 36 HAP, *DD45:GFP* expression was observed in the four-celled proembryo containing two apical cells, a middle and a basal cell (Fig. 1A). By 48 HAP, *DD45:GFP* expression was detected in the apical cells (octant-stage of cell division), hypophysis, suspensor and basal cells (Fig. 1B). Subsequently, by 60 HAP and 72 HAP, the embryo proper increases in size concomitant with an increase in suspensor length (Fig. 1C and D).

Unlike in wild-type, there was a delay in *lre-5/lre-5* embryo development at the earlier time points; by 36 HAP, the

DD45:GFP expression was detected in the apical and basal cells of an asymmetrically-divided zygote (Fig. 1E). By 48 HAP, *DD45:GFP* expression was detected in the apical and basal cells and in the two suspensor cells formed by the middle cell undergoing a transverse division (Fig. 1F). The lag in embryo development is also noticeable by 60 HAP as the apical cells in the mutant are only at the octant stage (Fig. 1G). However, by 72 HAP, the *lre-5/lre-5* embryos begin to reach similar stage of development as wild-type (compare size of embryo proper in Fig. 1D and H). Those *lre-5/lre-5* embryos that have not caught up with wild-type embryos by 72 HAP, similar to those reported in Figure 4 of our previous report,¹⁰ recover at a later time point in seed development (not shown). These results suggest that *lre-5/lre-5* embryos recover and overcome the delay in initiation of embryo development beginning from 72 HAP. The delay in embryo development phenotype reported here will also make it easy to characterize this phenotype in future experiments, as monitoring *DD45:GFP* expression in self-pollinated *lre-5/lre-5* pistils negates the need to cross the reporter gene-carrying pollen (such as *GRP23:GUS*) each time onto *lre-5/lre-5* pistils to detect this phenotype.¹⁰

The *lre-5/lre-5* Plants Do Not Exhibit any Detectable Defects during Seedling and Root Development

Previously we reported that in vegetative tissues, *LORELEI* is primarily expressed in eight-day-old seedlings.¹⁰ Yet, whether complete loss of *LORELEI* expression (*lre-5* is a null allele¹⁰) resulted in any defects during seedling and root development was not investigated. Additionally, it was not examined if the delay in seed development initiation caused by the loss of *LORELEI* function in pistils had any lingering effects during seed germination, seedling and root development. We determined the germination of stratified, wild-type and *lre-5/lre-5* seeds, eight days after plating on 1x MS growth medium; the rate of seed germination was nearly identical in both wild-type and *lre-5/lre-5* (Fig. 2A–C). The cotyledon leaves in *lre-5/lre-5* seedlings did not show any obvious defects and were

virtually indistinguishable from wild-type (not shown). The near uniformity and rate of *lre-5/lre-5* seed germination that is on par with wild-type is also consistent with the observation that nearly all *lre-5/lre-5* embryos recover and become equivalent to

wild-type embryos during embryo development (Fig. 1). We also examined root growth by plating stratified seeds on 1/2x MS medium and incubating the plates vertically in a growth chamber. The primary root growth in *lre-5/lre-5* seedlings, both

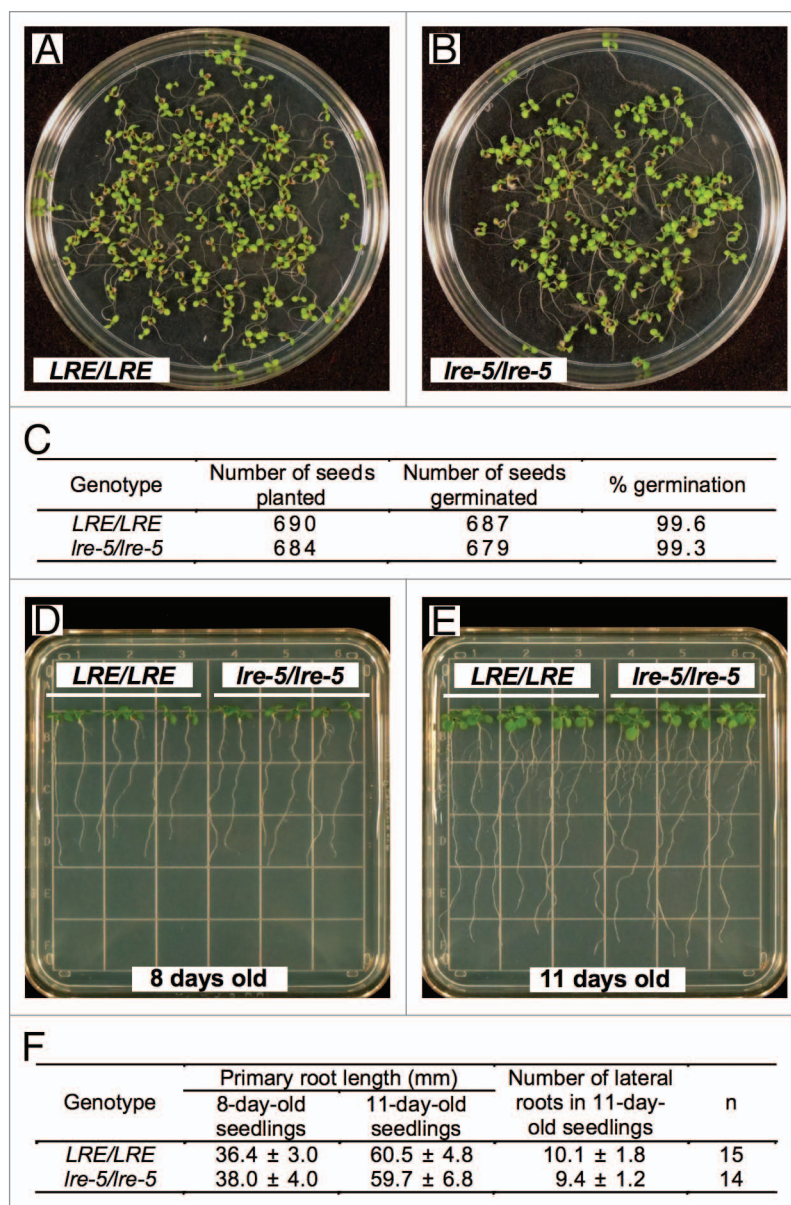


Figure 2. Analysis of germination and root development in seedlings. Photographs of germinated wild-type (A) and *lre-5/lre-5* (B) seedlings. Images in (A and B) were acquired eight days after plating the seeds on 1x MS medium (supplemented with 2% sucrose) and growing under continuous light at 21°C. The seeds used in this experiment were considered stratified as they were stored, since harvest, in 4°C with desiccant. (C) A table summarizing germination of wild-type and *lre-5/lre-5* seedlings. Photographs of root development, eight days (D) or 11 days (E) after plating six wild-type and six *lre-5/lre-5* seeds on the left and right side of the plate containing 1/2x MS medium (supplemented with 2% sucrose and 2.5 mM MES), respectively and growing under continuous light at 21°C. The seeds used in this experiment were considered stratified as they were stored, since harvest, in 4°C with desiccant. The area of each grid in the square plate is 13 mm². (F) A table summarizing primary root length and number of lateral roots in wild-type and *lre-5/lre-5* plants; n, total number of seedlings analyzed in this experiment.

8- and 11-days after plating, was indistinguishable from wild type (Fig. 2D–F). The number of lateral roots in 11-day-old *lre-5/lre-5* seedlings was also similar to wild type (Fig. 2F). These results indicate that LORELEI is not essential for seedling and root development in 11-day-old seedlings, at least in the plant growth conditions used in this experiment.

Acknowledgements

We thank Dr. Gary Drews for *DD45:GFP* seeds; Dr. Ramin Yadegari for providing access to the fluorescent microscope (Fig. 1); Shea Monihan and Dr. Karen Schumaker for assistance with root growth experiments (Fig. 2). We thank Dr. Jennifer Mach for critical reading of the manuscript. This work was supported by a NSF grant to R.P. (IOS-0723421).

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