The DP-E2F-like Gene *DEL1* **Controls the Endocycle in** *Arabidopsis thaliana*

Kobe Vlieghe,¹ Véronique Boudolf,¹ Gerrit T.S. Beemster,¹ Sara Maes,¹ Zoltan Magyar,^{1,2,3} Ana Atanassova,¹
Janice de Almeida Engler,^{1,4} Ruth De Groodt,¹ Dirk Inzé,^{1,*} and Lieven De Veylder¹ ¹Department of Plant Systems Biology

is widespread in nature. Well-known examples are fruit rable in number and size of abaxial epidermal cells (see fly polytene chromosomes and cereal endosperm. Al- Table S1 available with the Supplemental Data online). though endocycles are thought to be driven by the Similarly, no effects were seen on the first leaf pair. same regulators as those that control the G1-S transi- Obviously, *DEL1* **function did not affect cell division. In tion of the mitotic cell cycle, the molecular mecha- contrast, DNA ploidy levels varied significantly between nisms that differentiate mitotically dividing cells from wild-type and** *del1-1* **plants. Wild-type cotyledons disendoreduplicating ones are largely unknown. A novel played a typical pattern with C values ranging from 2C identified and is designated E2F7 in mammals [1, 2] endoreduplication. Cotyledons of** *del1-1* **plants had a and DP-E2F-like (DEL) in** *Arabidopsis thaliana* **[3–5]. significantly increased number of cells with 16C DNA We demonstrate that loss of** *DEL1* **function resulted content, as well as cells with a 32C DNA content; these** in increased ploidy levels, whereas ectopic expression

of DEL1 reduced endoreduplication. Ploidy changes

of E2F target genes encoding proteins necessary for

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and mammals [6–8], the physiological role of E2F7 or DEL proteins remained unclear. E2F and DP proteins only contain a single DNA binding domain and need to dimerize for high-affinity, sequence-specific DNA binding. By contrast, DEL proteins hold two domains highly **Department of Plant Systems Biology homologous to the DNA binding domain of E2Fs, which Flanders Interuniversity Institute for Biotechnology allows them to bind the canonical E2F binding site as Ghent University monomers [4, 5]. To study the function of DEL1 (also Technologiepark 927 known as EL3 [4] and E2Fe [5]) during plant develop-Gent B-9052 ment, we analyzed a T-DNA insertion mutation in the Belgium** *DEL1* **gene. The T-DNA of** *del1-1* **had been inserted 2Biological Research Center between the two DNA binding domains of DEL1 (Figure P.O. Box 521 1A). Because E2F7 and DELs need to have both DNA Szeged H-6701 binding domains to bind the E2F recognition sequence Hungary [1, 2, 4, 5],** *del1-1* **is most probably a null allele. The absence of full-length** *DEL1* **mRNA in the** *del1-1* **mutant was confirmed by reverse-transcriptase PCR analysis Summary (Figure 1B). The** *del1-1* **mutant did not differ morphologically from wild-type plants. For example, 12-day-old Endoreduplication or DNA replication without mitosis cotyledons of wild-type and** *del1-1* **plants were compaclass of atypical E2F-like proteins has recently been to 16C, where the peaks higher than 2C resulted from**

were detected exclusively in mitotically dividing cells,
we conclude that DEL1 is an important novel inhibitor
of the endocycle and preserves the mitotic state of
proliferating cells by suppressing transcription of
genes **levels were selected (Figure 1C). Homozygous** *DEL1OE* **Results and Discussion**
lines were phenotypically similar to those of wild-type Whereas the E2F-DP transcription factors are well-

known regulators of the G1-S transition, both in plants
 $DEL^{1/2E}$ line 2) and 19% (DEL1^{OE} line 4) reduced com**pared to that of wild-type plants as a result of a decrease** *Correspondence: dirk.inze@psb.ugent.be

³Present address: School of Biological Sciences, Royal Holloway, and the size (Table S1). Similarly, mature first leaves (22

University of London, Egham TW20 QEX, United Kingdom. **⁴ mine the cellular basis of this reduced size, we studied Present address: Institut National de la Recherche Agronomique,**

Unité Interactions Plantes-Microorganismes et Santé Végétale, B.P. 2078, F-06606 Antibes Cedex, France. analysis [10]. From day 4 until day 22 after sowing, leaves

binding domains are in gray. The triangle corresponds to the T-DNA

(B) Two-step RT-PCR analysis performed with equal amounts of *DEL1OE* **plants (Figures 2A and 2B; Tables S2 and S3). total RNA prepared from 8-day-old wild***-***type and** *del1-1* **seedlings The number of cells with a 4C DNA content was also**

(C) RNA gel blot analysis of wild*-***type (Col-0) and independent vestigated the effects of** *DEL1* **overexpression on the** *DEL1OE* **lines. Equal loading of the gel was confirmed by methylene endocycle in leaves (Figure 2D). Ploidy levels of the first**

(D) Change in transcript levels of replication genes (DNA polymerase
 α , DNA ligase I, and Replication factor c), DNA replication licensing

factors (MCM3) and E2F b in del1-1 (black) and DEL1^{0E}

line A (grav) All mo line 4 (gray). All measurements were performed on cotyledons of **10-day-old seedlings. Expression levels are compared with those exit from mitosis at day 10, the 2C population decreased, found in wild-type (WT) cotyledons of the same age. Data represent whereas the number of cells with 4C DNA content in** $average \pm SE(n = 3)$.

under the same conditions were harvested; leaf blade 16C DNA content could be measured. area and average cell area of abaxial epidermal cells In the *DEL1***OE lines, the ploidy distribution during the were determined. The total number of epidermal cells mitotic phase of leaf development, 8 days after sowing, was estimated as the ratio of leaf blade area and average resembled that found in wild-type plants. In contrast,**

cell area. Average cell division rates were calculated from the increase in cell number.

In the wild-type leaves, three developmental phases could be observed. First, until approximately 7 days after sowing, the cell number increased exponentially at maximal rates while cells retained a relatively constant cell size of approximately 80 μ m² (Figures 3B and 3C), **implying that expansion and division rates were balanced. In the second phase, between days 7 and 10, the division rate decreased (Figure 3D), testifying to the exit from the mitotic cell cycle. Simultaneously, average cell size started to increase (Figure 3C), indicating that the rate of cell expansion exceeded that of cell division. Between days 10 and 20, cell expansion continued in the absence of division, causing a 15-fold increase in cell size (Figure 3C). After day 20, leaves were mature and did not grow anymore (Figure 3A). When the development of the** *DEL1OE* **and wild-type leaves was compared, cell size (Figure 3C) and cell number (Figure 3D) were very similar during the mitotic phase of leaf development. Cell division rates were comparable in wildtype plants and the** *DEL1OE* **line 2 but were slightly lower in line 4. These data indicate that** *DEL1* **inhibits the mitotic cycle when it is expressed at high levels. In both** *DEL1OE* **lines, cell division rates dropped at the same pace between days 7 and 10, but they did so somewhat more slowly than in wild-type plants, suggesting that DEL1 might control the timing of the mitosis-to-endocycle transition. In accordance with the later exit from mitosis, the average cell size decreased below that of** the wild-type at day 9 (200 \pm 16 μ m² for wild-type, 112 ± 2 μ m² for DEL1^{0E} line 2, and 125 \pm 17 μ m² for *DEL1OE* **line 4) and remained smaller during the later stages of leaf development (Figure 3C). At maturity, cell area was approximately 25% smaller in the leaves of both** *DEL1OE* **lines. Taken together, these data show that the observed smaller leaf size in the** *DEL1OE* **transgenic lines of leaf development (Figure 3C). At maturity, cell**
 lines. Taken together, these data show that
 Plants lines predominantly originates from an inhibition of cell
 Plants Plants Plants predominantly or

(A) DEL1 gene structure. Black and gray boxes represent exons,
and lines indicate introns. Regions that encode to the two DNA by a reduced level of endoreduplication (Figures 2A–2C).
binding domains are in gray. The triang insert in the GABI-Kat 287D04 line (*del1-1*).
(B) Two-step RT-PCR analysis performed with equal amounts of DEL1^{0E} plants (Figures 2A and 2B: Tables S2 and S3). Finality and specifically all plugged and the DELT-country sequence

flanking the T-DNA insertion site. The actin 2 (ACT2) gene was used

as a loading control.

(C) RNA gel blot analysis of wild-type (Col-0) and independen blue staining of the membrane (bottom panel).
(D) Change in transcript levels of replication genes (DNA polymerase

development In wild-type leaves 70% of the cells had **SE (n 3). creased and became the most abundant population from day 12 onward. Simultaneously, cells with 8C DNA content could be detected. Later during leaf developof transgenic and wild-type plants grown side by side ment, a small, but reproducible, population of cells with**

Figure 2. Control of the Endoreduplication Level by DEL1 in *Arabidopsis* **(A–C) DNA ploidy distribution measured by flow cytometry of untransformed control plants (left panels),** *del1-1* **(center panels), and** *DEL1OE* **line 4 (right panels) 12 days after sowing. (A) Cotyledons. (B) Hypocotyls. (C) Leaves. (D) Ploidy level distributions during development of the first leaf in wild-type and** *DEL1OE* **lines. Leaves were harvested at the indicated time points. Data represent average** - **SD (n 2).**

(Figure 2D), whereas the 2C population of wild-type cells with a 16C DNA content together with two addileaves decreased at day 10. This reproducible increase tional endocycles, whereas the number of 2C cells was strongly reduced (Table 1). Expression of *DEL1OE* **probably reflects the accumulation of G1 cells that have in the** exited the G2 phase of the mitotic cycle, whereas part $E2Fa-DPa^{OE}$ background resulted in a reduced endore**of the nuclei of the wild-type leaves had already entered duplication, illustrating that DEL1 inhibits the endorethe first endocycle. Later during leaf development, the duplication phenotype induced by E2Fa-DPa. To ad-**2C DNA population of *DEL1*^{oE} plants decreased more dress the question of whether DEL1 could also suppress slowly than that of wild-type leaves. In the strongest the ectopic cell division phenotype caused by E2Fa-*DEL1* **DPa, we counted the number of abaxial epidermal pave-** *OE* **line, 2C cells remained the prevailing population for 15 days after sowing (Figure 2D). Moreover, almost ment cells of 6-day-old cotyledons (Table 2). Pavement cells of** *E2Fa-DPaOE* **no cells had a DNA content higher than 4C, even at the transgenic plants were significantly latest time point analyzed. Also in the** *DEL1OE* **line 2, a smaller than those of control plants. Despite their** clear effect on the endocycle was observed, as illus-

smaller size, cotyledons of E2Fa-DPa^{oE} plants had aptrated by the strong decrease in 8C cells $(12.4 \pm 1.1\%)$ compared to that of wild-type leaves (28.1 \pm 2.8%). The results obtained from the leaf kinematic growth analysis, cell size and an increased cell number that did not differ in combination with those of the ploidy measurements, from those of E2Fa-DPa^{OE} lines. These results show that **strongly indicate that DEL1 operates as a specific re- DEL1 does not suppress the ectopic cell divisions pressor of the endocycle. Only when the** *DEL1* **gene is caused by overexpression of both** *E2Fa* **and** *DPa* **and expressed highly above the endogenous level, the mi- again illustrate that DEL1 specifically inhibits the endototic cell cycle is inhibited as well. cycle.**

an increase of 2C cells was seen in the *DEL1OE* **lines** *E2Fa-DPaOE* **plants displayed an increased number of 1.1%) proximately 5-fold more epidermal cells. Cotyledons of** DEL1-E2Fa-DPa^{0E} triple transgenic lines had a reduced

DEL proteins have been postulated to act as negative DEL1 could repress the endocycle by preventing eiregulators of the E2F pathway [1-5]. Overexpression of ther mitotically dividing cells from endoreduplicating or **regulators of the E2F pathway [1–5]. Overexpression of ther mitotically dividing cells from endoreduplicating or** both *E2Fa* and *DPa* (*E2Fa-DPa^{OE}*) genes in *Arabidopsis* post-mitotic cells from initiating the endocycle. To dis**resulted in a dual phenotype: Ectopic cell division oc- tinguish between these possibilities, we examined the curred in certain cells, whereas others underwent exces- spatial expression of** *DEL1* **by in situ hybridization of sive endoreduplication [11]. To test whether DEL1 was sections from a 5-day-old etiolated hypocotyl (Figure able to suppress these phenotypes, we crossed an S2), which is a well-characterized model system for DNA** $E2Fa-DPa^{OE}$ line with either a $DEL1^{OE}$ (line 4) or an un-
endoreduplication in plants [12–14]. In contrast to *E2Fa*, **transformed control plant (Col-0). Offspring plants were no** *DEL1* **transcripts could be detected in cortex cells first analyzed by flow cytometry. As observed before, that undergo extensive endoreduplication but no mitotic**

confined to the vascular tissue that divides mitotically, E2F7 might play a crucial role in maintaining the normal as demonstrated by the expression of the mitosis-spe- euploid state of the cell.

cific, cyclin-dependent kinase gene *CDKB1***. The presence of** *DEL1* **transcripts in dividing cells and its absence in cells undergoing DNA endoreduplication supports a role for DEL1 as repressor of the endocycle in mitotically dividing cells.**

DEL1 resides in the nucleus and associates with the promoter of genes that contain the E2F consensus binding site [4, 5]. Consequently, DEL1 most probably represses the endocycle directly at the transcriptional level. Because *DEL1* **is specifically expressed in proliferating cells, DEL1 might preserve the mitotic state of dividing cells by repressing the transcription of genes required for the endocycle. Previously, the mammalian homolog of DEL1, E2F7, has been demonstrated to regulate the expression of only a subset of E2F-dependent genes [2]. By quantitative PCR, we tested whether the transcript levels of DNA replication genes containing an** *E2F-cis***-acting element in their promoter were altered in 10-day-old cotyledons of** *del1-1* **and** *DEL1OE* **plants. Of all six analyzed genes, which are regulated by** *E2Fa-DPa* **([11, 15]; unpublished data), four had increased transcript levels in the** *del1-1* **mutant, whereas their expression was reduced in the** *DEL1OE* **lines (Figure 1D). In contrast,** *MCM3* **and** *E2Fb* **transcript levels did not change in response to altered** *DEL1* **levels. These data suggest that DEL1, like its mammalian counterpart, regulates only a subset of the E2F target genes. Because the DEL1 protein specifically inhibits the endoreduplication program, genes with an altered expression level in the** *del1-1* **and** *DEL1OE* **lines presumably play a role in the endocycle, whereas genes with unaltered transcript levels might be specific for the mitotic cell cycle.**

In conclusion, we have identified DEL1 as a novel and specific repressor of the endocycle. Recently, we have demonstrated that the decision of cells to divide mitotically or to endoreduplicate is dictated by the activity of the mitosis-specific CDKB1;1 [16]. Therefore, it will be interesting to test whether DEL1 inhibits the endocycle Figure 3. Kinematic Analysis of Leaf Growth of the First Leaf Pair

of Wild-Type (Col-0) and *DEL1^{0E}* Plants

(A) Leaf blade area.

(B) Epidermal cell number on the abaxial side of the leaf.

(B) Epidermal cell number on **(C) Epidermal cell size on the abaxial side of the leaf. has been conserved evolutionarily. It has been noted (D) Average cell division rates of the epidermal cells on the abaxial before that E2F7 is to be found at a chromosomal loca**side of the leaf. Error bars give standard errors (n = 5).
 Symbols in (A), (B), and (C) are as in (D). the symbols of the property of the property of the symbols in (A), (B), and (C) are as in (D). property posis fo **nosis for pancreas cancer patients, marking** *E2F7* **as a possible tumor suppressor gene [2]. Because aneucell divisions. The** *DEL1* **hybridization signal was entirely ploidy predisposes cells to oncogenic transformation,**

Data represent average \pm SD (n=5 to 8).

ap 0.01 (comparison with Col-0).

^b p 0.01 (comparison with Col-0 *E2Fa-DPa***OE).**

p-values were derived from two-tailed Student's t-tests.

Line	Cotyledon size (mm ²)	Abaxial Pavement Cells	
		Size (μm^2)	Estimated Number
Col-0	2.4 ± 0.2	1954 ± 149	1307 ± 103
$DEL1^{OE} \times$ Col-0	1.8 ± 0.1^a	$1210 \pm 55^{\circ}$	1478 ± 73
Col-0 \times E2Fa-DPa ^{OE}	1.6 ± 0.1	265 ± 32	6820 ± 855
DEL1 ^{OE} \times E2Fa-DPa ^{OE}	1.9 ± 0.2	291 ± 58	7228 ± 833

Table 2. Size and Number of Abaxial Pavement Cells in Cotyledons of *E2Fa-DPaOE* **and** *DEL1-E2Fa-DPaOE* **Lines**

All measurements were performed on cotyledons harvested 6 days after sowing. The indicated values are means \pm SE (n = 9 to 15). **ap 0.01 (comparison with Col-0).**

p-values were derived from two-tailed Student's t-tests.

Supplemental Data including Experimental Procedures, two figures, inhibitors of Arabidopsis. Plant Cell *13***, 1653–1667. and four tables are available with this article online at http://www. 11. De Veylder, L., Beeckman, T., Beemster, G.T.S., de Almeida current-biology.com/cgi/content/full/15/1/59/DC1/. Engler, J., Ormenese, S., Maes, S., Naudts, M., Van Der**

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Supplemental Data Inzet and Complemental Data Inzet and Complemental Analysis of cyclin-dependent kinase

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	- **overexpressing plants reveals changes in the expression levels**
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