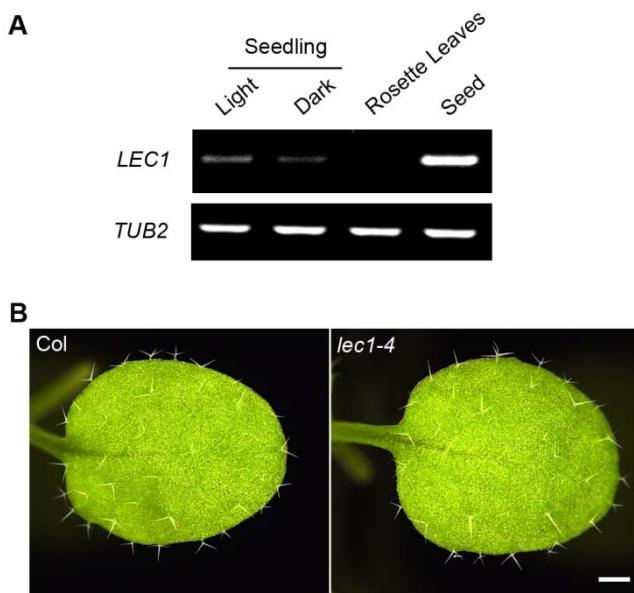


## *Supplementary Material*

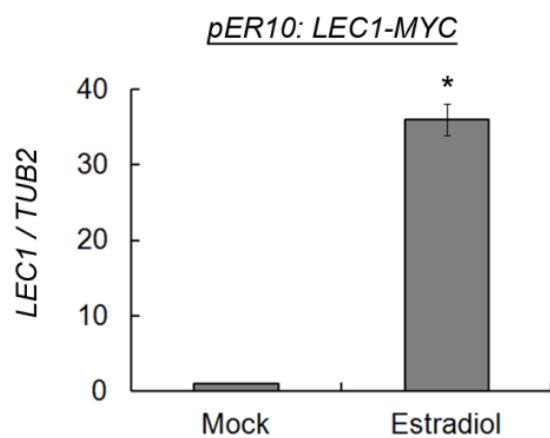
### ***Arabidopsis* LEAFY COTYLEDON1 controls cell fate determination during post-embryonic development**

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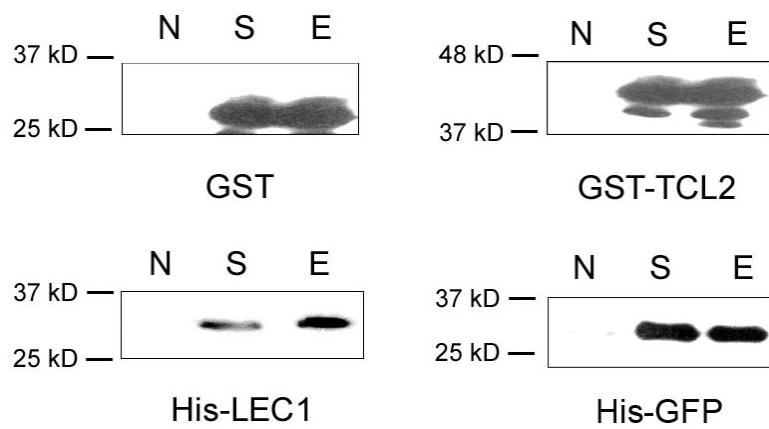
\* Correspondence: Corresponding Author: [houxl@scib.ac.cn](mailto:houxl@scib.ac.cn)



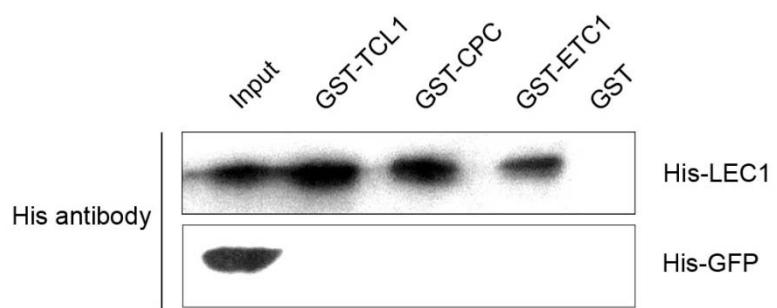
**Supplemental Figure S1. *LEC1* is not involved in trichome formation on rosette leaves.** (A) *LEC1* mRNA accumulated in developing seeds and 5-day-old light- and dark-grown seedlings but not in rosette leaves. *TUB2* was used as an internal control. (B) Trichome phenotype of the first pair of rosette leaves in *Col* wild type and *lec1-4* mutant showed no obvious distinction. Bar = 1 mm.



**Supplemental Figure S2. The chemical inducer estradiol significantly induces ectopic *LEC1* expression.** 5-day-old *pER10:LEC1-MYC* transgenic seedlings grown in the light with 10  $\mu$ M estradiol or mock treatment were used for gene expression analysis. *TUB2* was used as an internal control. Asterisk indicates significant difference between estradiol and mock ( $p < 0.05$ , by Student's *t*-test).



**Supplemental Figure S3. Western blot detection of GST- and His-tagged proteins.**  
N, no IPTG; S, soluble GST- or His-tagged fusion proteins supernatant; E, elution component.



**Supplemental Figure S4. Pull-down assay showing direct interaction between His-LEC1 and GST-TCL1, GST-CPC or GST-ETC1 fusion proteins *in vitro*.** His-LEC1 or His-GFP protein was incubated with immobilized GST or GST-fusion proteins, and immunoprecipitated fractions were detected by anti-His antibody.

**Supplemental Table S1. List of primers used in this Study.**

**Primers for quantitative real-time PCR**

Gene name	Primers (5'-3')
<i>LEC1</i>	F: AACCTATGGAGGAATGG R: TAATGGAAGACGAAGAGCC
<i>GL1</i>	F: TGGAACCGCATCGTCAGA R: ATTGCCGAGGAGCTTGTG
<i>GL2</i>	F: CAGCGAACCTATGTGGC R: CATGAAACTCTGGCAAGT
<i>GL3</i>	F: AATGAGGTGGTTATTGAGC R: ATTCCTGGTGTGCTATT
<i>EGL3</i>	F: TGAGGAGGAAAGAGCATC R: TATCCGTTAGACCAGCAT
<i>TTG1</i>	F: ATTACAACAACCGCATCG R: AAGGAGGACTGAACATTAGC
<i>TRY</i>	F: GGTGATAGGTGGGATTG R: TGAGGCTTGGTATGTTG
<i>CPC</i>	F: TGTCCGAAGAGGTGAGT R: CGTGTTCATAAGCCAATA
<i>ETC1</i>	F: AACCATTTGCCTCTTC R: TAAATCCCACCTTCACC
<i>ETC2</i>	F: TTGTCGGTAATAGGTGGG R: ACGTCGTCGTTGTGAGA
<i>TCL1</i>	F: CACTGTCATAAACAAACCAA R: CACGTCCCACCAACTCTTC
<i>TCL2</i>	F: AACCGTCTCGTCACCTT R: CTCGTCCTACCAACTCTTCC
<i>TUB2</i>	F: ATCCGTGAAGAGTACCCAGAT R: AAGAACCATGCACTCATCAGC

**Primers for constructs in yeast two-hybrid assays**

Gene name	Primers (5'-3')
<i>BD-LEC1</i>	F: AATT <u>CATATGATGGAACGTGGAGCTCC</u> T R: CG <u>CGGATCCTCACTTATACTGACCATA</u>
<i>AD-TCL2</i>	F: CCG <u>GAATTCATGGATAACACCAACCGT</u> R: CCG <u>CTCGAGTTAAGGAGGAGAAATAGA</u>

**Primers for constructs in pull-down assays**

Gene name	Primers (5'-3')
<i>GST-TCL1</i>	F: CCG <u>GAATTCATGGATAACACAAACCGT</u> R: CCG <u>CTCGAGTCATTGTGGGAGAAATA</u>
<i>GST-TCL2</i>	F: CCG <u>GAATTCATGGATAACACCAACCGT</u> R: CCG <u>CTCGAGTTAAGGAGGAGAAATAGA</u>

<i>GST-CPC</i>	F: CCGGAATTCATGTTCGTTCAGACAAG R: CCG <u>CTCGAGTC</u> ATTCCTAAAAAGTC
<i>GST-ETC1</i>	F: CCGGAATT <u>CATGAATA</u> ACGCAGCGTAAG R: CCG <u>CTCGAGTC</u> AACGTAATTGAGATCT
<i>His-LEC1</i>	F: CGCG <u>GATCC</u> ATGGAACGTGGAGCTCCCTT R: ACG <u>CGTCGAC</u> CTTATACTGACCATAATGGT
<i>His-GFP</i>	F: CG <u>CGGATCC</u> CATGAGTAAAGGAGAAGAAC R: ACG <u>CGTCGAC</u> TTTGTATAGTTCTACCAT

**Supplemental Table S2 Frequency of seedlings with (w/) or without (w/o) ectopic trichomes on cotyledons.**

		Seedlings w/o trichomes	Seedlings w/ trichomes	Frequency of seedlings w/o trichomes(%)	Frequency of seedlings w/ trichomes(%)
Col	Experiment1	15	0	100.00	0.00
	Experiment2	15	0	100.00	0.00
	Experiment3	15	0	100.00	0.00
	Mean	--	--	100.00	0.00
	SD	--	--	±0.00	±0.00
<i>lec1-4</i>	Experiment1	8	46	14.81	85.19
	Experiment2	10	42	19.23	80.77
	Experiment3	5	51	8.93	91.07
	Mean	--	--	14.32	85.68
	SD	--	--	±5.16	±5.17

**Supplemental Table S3 Frequency of different types of trichomes on cotyledons.**

<i>lec1-4</i>	Trichome branch point	Total		Frequency (%)			
		number of trichomes		Trichome branch point			
				0	1	2	
Experiment1	48	32	0	80	60.00	40.00	0.00
Experiment 2	24	36	1	61	39.34	59.02	1.64
Experiment 3	13	58	1	72	18.06	80.56	1.39
Experiment 4	30	40	0	70	42.86	57.14	0.00
Experiment 5	29	55	1	85	34.12	64.71	1.18
Experiment 6	34	42	1	77	44.16	54.55	1.30
Experiment 7	40	16	0	56	71.43	28.57	0.00
Mean	--	--	--	--	44.28	54.93	0.79
SD	--	--	--	--	±17.33	±16.79	±0.75