

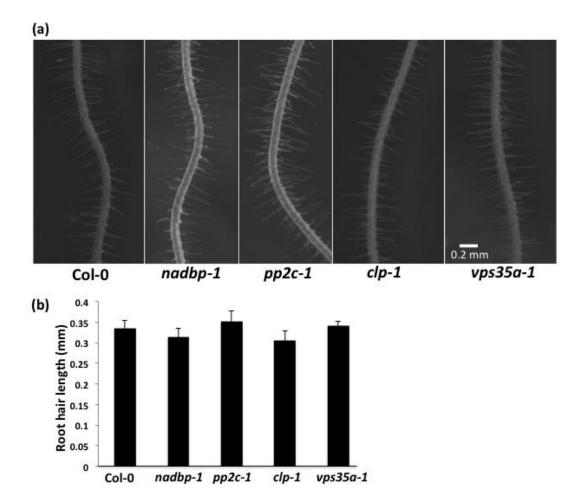
## New Phytologist Supporting Information

## Article title: **RSL4 promotes root hair elongation by transcriptionally regulating the expression of genes required for cell growth**

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The following Supporting Information is available for this article:



**Fig. S1** (a) Images of 6-d-old *Arabidopsis thaliana* wild type Col-0, *nadbp-1, pp2c-1, clp-1* and *vps35a-1* roots grown on MS plates with 0.5% Phytagel. (b) Root hair length measurements of 6-d-old wild type (Col-0), *nadbp-1, pp2c-1, clp-1* and *vps35a-1* plants (n = 15). Error bars represent SEM.



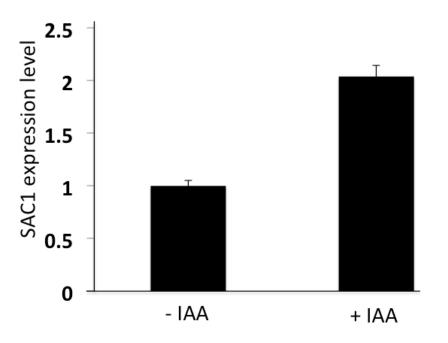
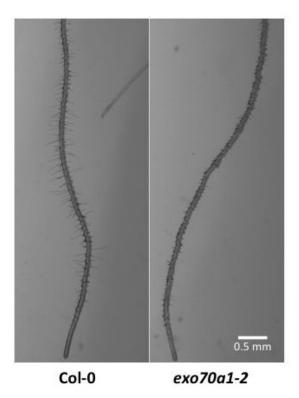


Fig. S2 SAC1 mRNA level in absence and presence of exogenously applied 100 nM IAA (n =

3). Error bars represent SEM.



**Fig. S3** Images of 6-d-old *Arabidopsis thaliana* wild type (Col-0) and *exo70a1-2* mutants with short root hairs grown on MS plates with 0.5% Phytagel.



## Table S1 List of primers used in this study.

## Primers used to confirm the GR: RSL4 fusion construct

Primers within the 5'UTR and promoter of RSL4

ATTB4: GGGGACAACTTTGTATAGAAAAGTTGATACGCGTTGGGCTTAAATG ANTI578: CCTAAAACCTAAACGCGGTG

Primers within the CDS+3'UTR of RSL4

**RDL2 KpnI:** CGGGGTACCATGGACGTTTTTGTTGATGGT **RDL2 BamHI**: CTTTTGTCTCGGCTTATGTGAGGATCCGCG

Primers within the GR sequence:

**GR forward:** CAAAGGGATTCAGCAAGCCAC **GR reverse**: CCTTACCTACTGCTTCCAGACATTT

**Table S2** 1022 genes in Arabidopsis thaliana that were expressed more than twice as high inDEX treated GR:RSL4 rsl2 rsl4 plants than in mock treated plants with a P value < 0.0001.</td>(See separate Excel file.)

 Table S3 259 genes in Arabidopsis thaliana that were expressed more than twice as high in

 DEX+CHX treated GR:RSL4 rsl2 rsl4 than in mock treated plants with a P value < 0.001. (See separate Excel file.)</td>

**Table S4** 130 genes in Arabidopsis thaliana whose expression was increased by more than

 two-fold in both DEX and DEX+CHX treated GR:RSL4 rsl2 rsl4 plants compared to mock

 treatment. (See separate Excel file.)



Table S5 List of mutants used in this study with details of T-DNA insertions

Mutant alleles	Gene name	Gene code	Insertion code
gbf2-2	GBF2	AT4G01120	SALK_206654
exo70a1-1, exo70a1-	EXO70A1		SALK_014826, SALK_135462
2		AT5G03540	
nadbp-1	NADBP	AT1G01800	SALK_009967
pp2c-1	PP2C	AT3G17090	SALK_203806
clp-1	CLP	AT1G26940	SALK_054035
cpk11-2	CDPK2/CPK11	AT1G35670	SALK_054495
prx7-1	PRX7	AT1G30870	GK_476A09-019131
vps35a-1	VPS35B/ LAZ4	AT1G75850	SALK_014345
sac1-2	SAC1	AT1G22620	SALK_070875