

***AtOPR3* specifically inhibits primary root growth in Arabidopsis under  
phosphate deficiency**

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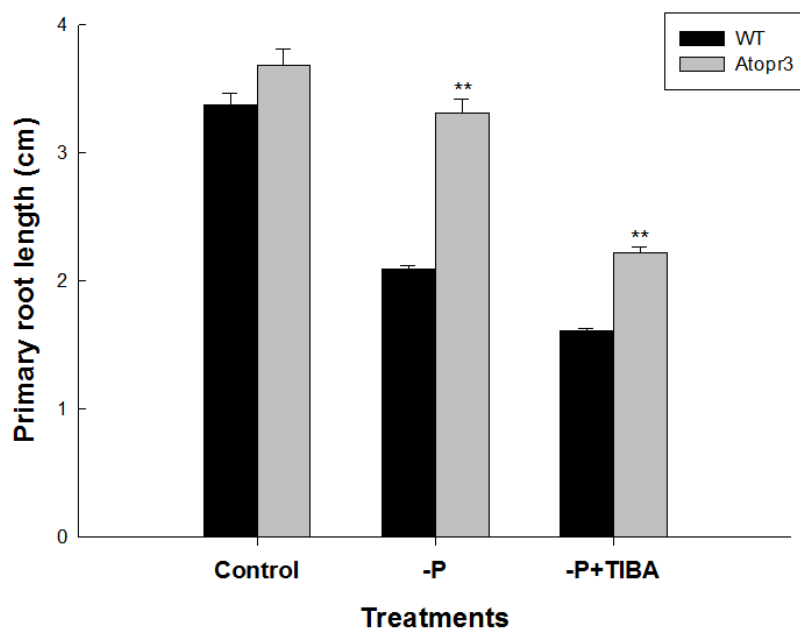
## Figure S1

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### Figure S1. The 5'-upstream region of *AtOPR3*.

The putative P1BS element, P1BS-like element, and the start codon were highlighted in red, green, and yellow, respectively.

**Figure S2**



**Figure S2. Auxin polar transport and primary root growth in *Atopr3* and WT plants under P deficiency.**

Seedlings were grown as described in Fig. 2. TIBA (5 $\mu$ M) was applied to inhibit auxin polar transport. Results were presented as means (n=30) with error bars (standard deviation). Asterisks indicated significant differences as determined by a t-test analysis (\*\*, P<0.01).

**Table S1. The relative length of the primary root in wild type *Arabidopsis* (*Ws*) and *opr3* mutant plants (*Atopr3*) without or with various concentrations of chemicals under P deficiency. The root length without the chemical treatment was set as 100%.**

Concentration ( $\mu\text{M}$ )	GA		PAC		Ancymidol		TIBA	
	WT	<i>Atopr3</i>	WT	<i>Atopr3</i>	WT	<i>Atopr3</i>	WT	<i>Atopr3</i>
<b>0</b>	100%	100%	100%	100%	100%	100%	100%	100%
<b>0.5</b>	-	-	-	-	-	-	-	-
<b>1</b>	-	-	-	-	-	-	92.5%	97.2%
<b>1.5</b>	-	-	-	-	-	-	-	-
<b>2</b>	-	-	-	-	-	-	-	-
<b>2.5</b>	114.3%	108.0%	70.9%	62.3%	78.9%	76.0%	-	-
<b>3</b>	-	-	-	-	-	-	83.1%	83.8%
<b>5</b>	116.2%	110.3%	77.0%	64.1%	74.2%	59.5%	77.0%	67.0%
<b>7.5</b>	135.7%	104.2%	76.5%	62.0%	66.7%	54.2%	-	-
<b>10</b>	100.7%	94.4%	-	-	-	-	-	-

**Table S2. Primers used to analyze gene expression levels via RT-qPCR in our studies.**

Gene	Primer sequence
<i>AtOPR3</i>	5'—3' AAGGTGTAGTTTCAGCCATAGGA
	5'—3' GATAGTGGGTCAGAATCAGTTGC
<i>AtACS2</i>	5'—3' AGATCGTCGAGAAAGCATCTG
	5'—3' GAAGAGGTGAGTGTGGTGAC
<i>AtCTR1</i>	5'—3' GCCACCCAGTGCTAATATGCC
	5'—3' CAAAGGAACCTGCTCCAATCT
<i>At20OX1</i>	5'—3' GATCCATCCTCCACTTTAGA
	5'—3' GTGTATTCATGAGCGTCTGA
<i>At20OX2</i>	5'—3' ACCGAGACTATTTCCGAGGATT
	5'—3' TGTTTGGCATGGAGGATAATG
<i>At20OX3</i>	5'—3' CCTATCTGCATATGGACTC
	5'—3' AACCTTCCCGAAATCTTC
<i>At3OX1</i>	5'—3' TCCGAAGGTTTCACCATCACT
	5'—3' TCGCAGTAGTTGAGGTGATGTTG
<i>At3OX2</i>	5'—3' CCTCGCGACTTCTCGAC
	5'—3' AATAATTTACAGTATTTGAGG
<i>At2OX1</i>	5'—3' CGGGAACTTTCAGAAACGC
	5'—3' ACATTCTTACCACCATTGG
<i>At2OX2</i>	5'—3' CCTAAAACCTCCGCCGTTTT
	5'—3' CCTTCATGTACTCCTCCACCGA
<i>AtLPR1</i>	5'—3' AGAAACTCCAAAATCAGGGACTACA
	5'—3' ATGGTCCAATATGTGACAATGGTAG
<i>ATLPR2</i>	5'—3' GAGGCATAAAGCAAGAGGAGC
	5'—3' GCAGGCAAGTCTCTGTGGAA
<i>ATPRD</i>	5'—3' CAAATGCGCTTCGTATCTCC
	5'—3' TAAAACAGCGTCTGCGTCTG
<i>ATPDR2</i>	5'—3' CTTTTCACTACGTGGTCCGTGG
	5'—3' TGTCTCCCGAGGATGCTGAAT
<i>ATPLD1</i>	5'—3' TGGATGGCAACCGCAAAGACAA
	5'—3' ATCGTTGTGTGTCCAGCTTCT
<i>ATPLD2</i>	5'—3' TTTGAGGACGGTCCAATTGCCA
	5'—3' ACAACACCGATCTCAGAGTCTCGT
<i>AtTUB4</i>	5'—3' AGAGGTTGACGAGCAGATGA
	5'—3' CCTCTTCTTCTCCTCGTAC