

# **Copper excess reduces nitrate uptake by *Arabidopsis* roots with specific effects on gene expression**

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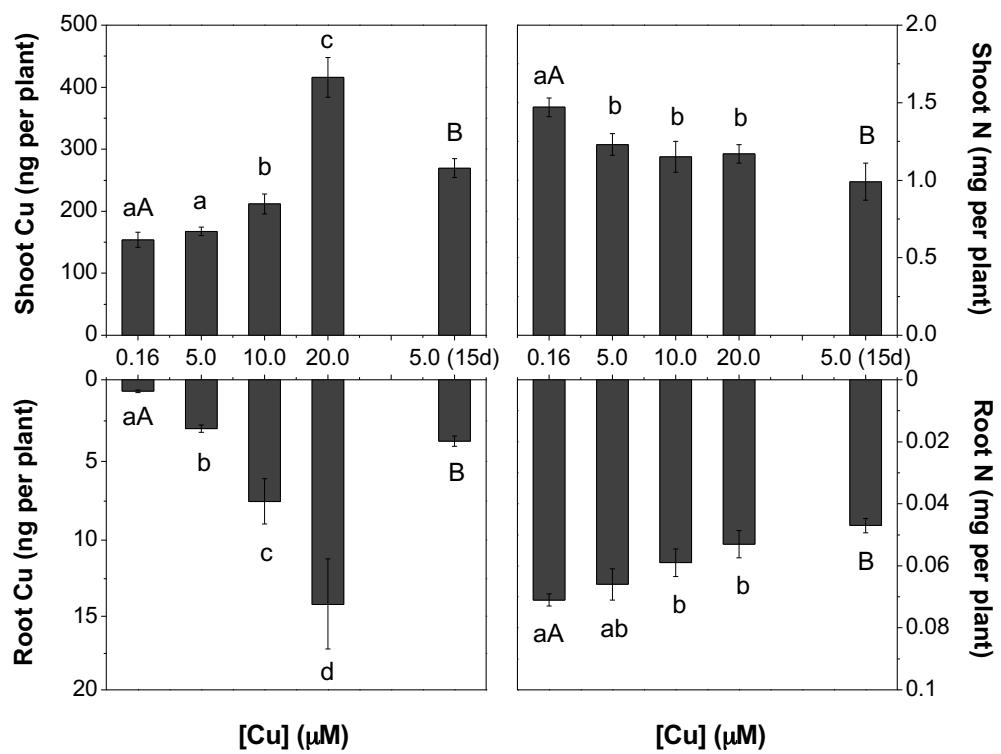
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## **SUPPLEMENTARY FILE**

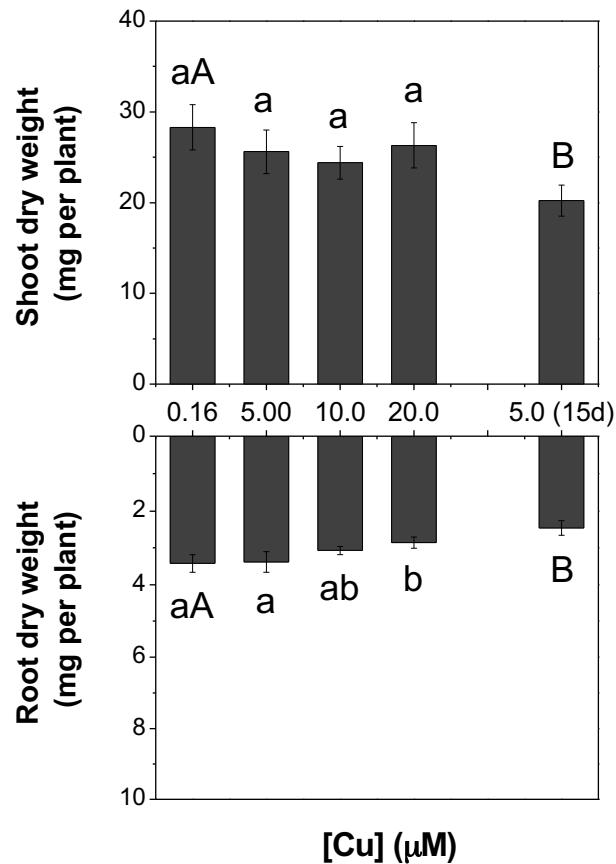
**Supporting Information Table S1.** Genes and primers analysed in roots of *Arabidopsis thaliana* grown under varying copper (Cu) concentrations.

Gene	Locus identification <sup>1</sup>	Forward primer (5' to 3')	Reverse Primer (5' to 3')
<b>Nitrate transporters</b>			
<i>NRT1.1</i> ( <i>NPF6.3</i> or <i>CHL1</i> )	AT1G12110	GGCCGTACTTGTGCCTTGA	CCCATTGGAATACTCGGCTCA
<i>NRT1.2</i>	AT1G69850	AGCTCAGGGAACACACCTTG	ACGCATTGCCCAAAGAGGT
<i>NRT1.5</i>	AT1G32450	CGAGCATTGACCTGGTGGTA	CGTTCCCTCTTCACTCTCGGT
<i>NRT2.1</i>	AT1G08090	CATCAAGGAAGCCTCCGGTT	TGTTGGGTGTGTTCTCAGGC
<i>NRT2.2</i>	AT1G08100	CGACGCTACGGAGCACTATT	GTTGCGTTCCCTTGTGGAC
<i>NRT2.4</i>	AT5G60770	CACGGAACAAGGGCTGACA	TCATCTCCGTGGAAGGCAAA
<i>NAR2.1</i> ( <i>NRT3.1</i> )	AT5G50200	CGCCACCGCGTCCCTAGATA	CTTGGCCTTCCTCTCATCG
<b>Plasma membrane H<sup>+</sup>-ATPase</b>			
<i>AHA1</i>	AT2G18960	TAGCTAGGCTTAGGGAGCTTCA	CAGTGTAGTGATGTCCTGCTGT
<i>AHA2</i>	AT4G30190	GAGCTGAGATCGCTAGGCTT	CTACACAGTGTAGTGACTGGAG
<i>AHA5</i>	AT2G24520	AACCCTCCACGGACTTCAAC	CTCTCTCAGCCTGACAACCTC
<b>Nitrate reductase</b>			
<i>NRI</i> ( <i>NIA1</i> )	AT1G77760	TGGTACGTCGTTGAAATCGC	TCCCTAAGCACAGCTTCAGTT
<b>bZIP transcription factors</b>			
<i>TGA1</i>	AT5G65210	GACAAGCGGCTCGAGGATT	GTTCACGATGTCGAGTTGCC
<i>TGA4</i>	AT5G10030	GGAGTTGCTGGTGAGTGG	TCCTTAGCCGCAAGACAAGA
<b>Lipoxygenase</b>			
<i>LOX1</i>	AT1G55020	GATGAGAGGAACGACGACGAG	CGCCTTCACTGCTCGGAAAC
<b>Housekeeping genes</b>			
<i>SAND</i>	AT2G28390	GGTGGATGTTGGAGAGCATT	CCAAACAAGAAGTGGATCCC
<i>YLS8</i>	AT5G08290	CTTCTTCCACATCTGCACTCC	CAGTTCACGGGATATTCAGC

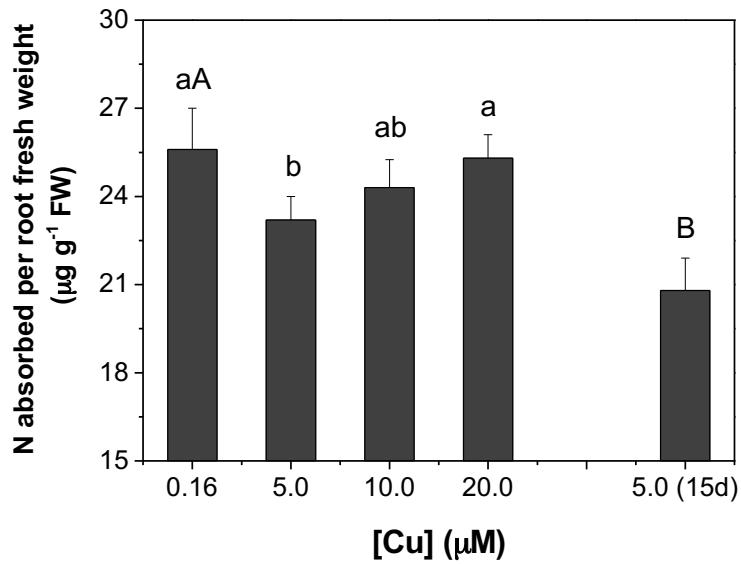
<sup>1</sup> NCBI data base - (<http://www.ncbi.nlm.nih.gov>)



**Fig. S1. Elevated copper in the media (Cu) enhanced Cu accumulation but reduced the accumulation of nitrogen (N) in shoot and root of *Arabidopsis thaliana*.** Plants were supplied with different Cu levels in the nutrient solution for 72 h or 15 days (15d). Mean (+/- SEM) are shown ( $n = 3$ ). For the Cu treatments for 72 h different lowercase letters indicate mean values are significantly different among the [Cu] (0.16, 5.0, 10.0 and 20.0  $\mu$ M) by Tukey's test ( $p < 0.05$ ). For the Cu treatments for 15 days different uppercase letters indicate mean values are significantly different between the [Cu] (0.16 and 5.0  $\mu$ M) by Tukey's test ( $p < 0.05$ ).



**Fig. S2. Elevated copper (Cu) negatively impacts dry weight of shoots and roots of *Arabidopsis thaliana*.** Plants were grown under hydroponic conditions for up to 72 h or 15 days (15d). Mean (+/- SEM) are shown (n = 3). For the Cu treatments for 72 h different lowercase letters indicate mean values are significantly different among the [Cu] (0.16, 5.0, 10.0 and 20.0  $\mu\text{M}$ ) by Tukey's test ( $p < 0.05$ ). For the Cu treatments for 15 days different uppercase letters indicate mean values are significantly different between the [Cu] (0.16 and 5.0  $\mu\text{M}$ ) by Tukey's test ( $p < 0.05$ ). White line in the image represent 2 cm.



**Fig. S3. Elevated copper in the media (Cu) disturbed N absorption per root fresh weight of *Arabidopsis thaliana*.** Plants were supplied with different Cu levels in the nutrient solution for 72 h or 15 days (15d). Mean (+/- SEM) are shown (n = 3). For the Cu treatments for 72 h different lowercase letters indicate mean values are significantly different among the [Cu] (0.16, 5.0, 10.0 and 20.0  $\mu\text{M}$ ) by Tukey's test ( $p < 0.05$ ). For the Cu treatments for 15 days different uppercase letters indicate mean values are significantly different between the [Cu] (0.16 and 5.0  $\mu\text{M}$ ) by Tukey's test ( $p < 0.05$ ).