

Fig. S1 Schematic representation of the experimental design

EXPERIMENTAL WORKFLOW

Identification of early cytokinin response proteins

Identification of early cytokinin response phosphoproteins

Contribution of individual cytokinin receptors to phosphoproteome regulation

Effect of calcium signaling inhibitors on phosphoproteome response to cytokinins

Identification of early cytokinin response proteins

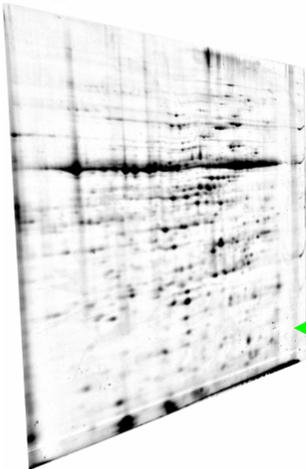
Arabidopsis thaliana (ecotype Columbia) seedlings were cultivated for 7 days on MS medium supplemented with ($5 \cdot 10^{-4}$ % v/v) DMSO at standard light intensity ($\sim 90 \mu\text{mol} \cdot \text{m}^{-2} \cdot \text{s}^{-1}$).



Seedlings were transferred on liquid MS medium supplemented with DMSO (mock; $5 \cdot 10^{-4}$ % v/v) or $5 \mu\text{M}$ cytokinin (BA, iP, TDZ or *t*-Z) dissolved in DMSO and incubated for 15 min.



Seedlings were rapidly harvested, frozen and ground in liquid nitrogen



Total *Arabidopsis* protein was extracted by standard acetone/TCA extraction and resolved by large (18 cm IPG strips, $500 \mu\text{g}$ of protein) 2-DE.

Identification of early cytokinin response proteins

2-DE

3 technical replicates for each biological replicate

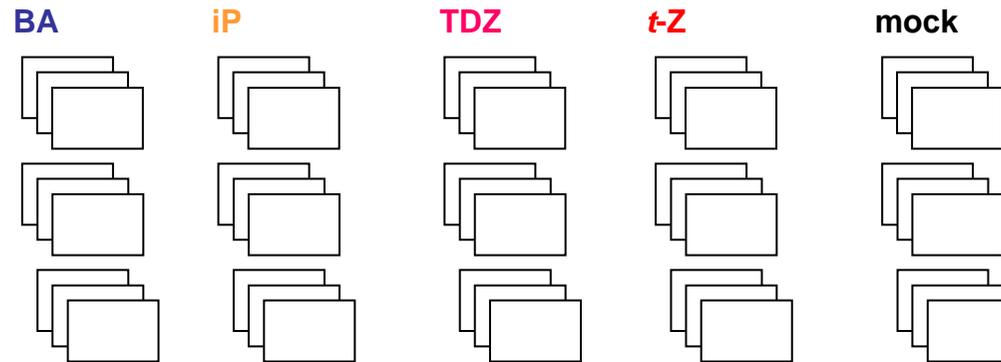


image analysis

45 2D gels

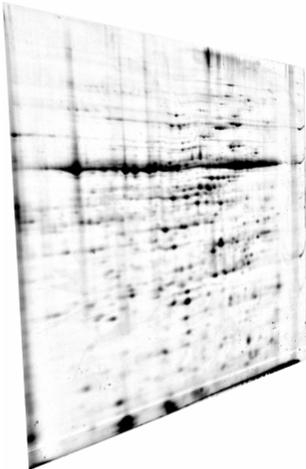
protein digest

53 protein spots reproducibly significant in at least 2 biological replicates

MALDI-TOF/TOF analysis

MASCOT search

67 identified proteins



Identification of early cytokinin response phosphoproteins

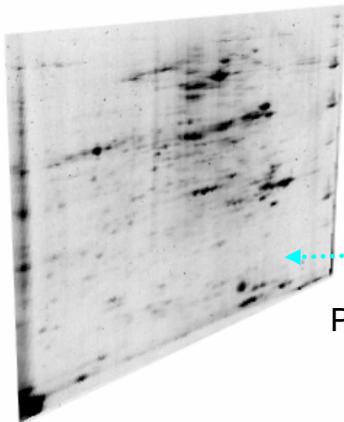
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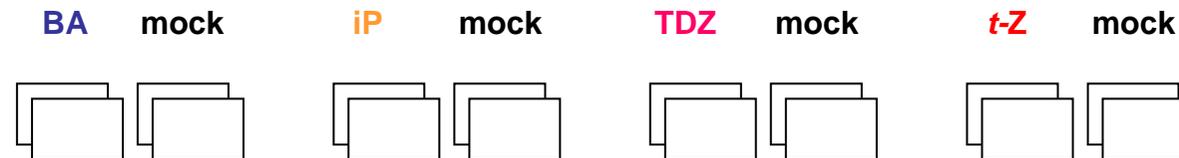
For phosphoproteome analysis, an isolation procedure was established based on PhosphoProtein Purification Kit (Qiagen). Collected phosphoprotein fractions were then resolved by 2-DE (7 cm IPG strip, $150 \mu\text{g}$ of protein per strip).

Identification of early cytokinin response phosphoproteins

2-DE

2 technical replicates for each biological replicate

Pilot experiments



Main experiments

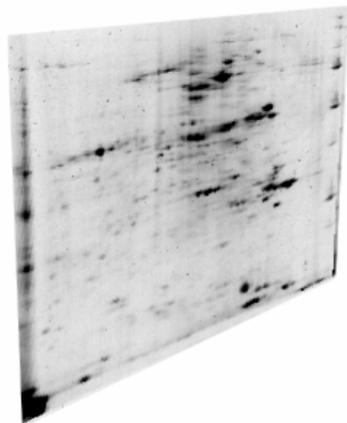
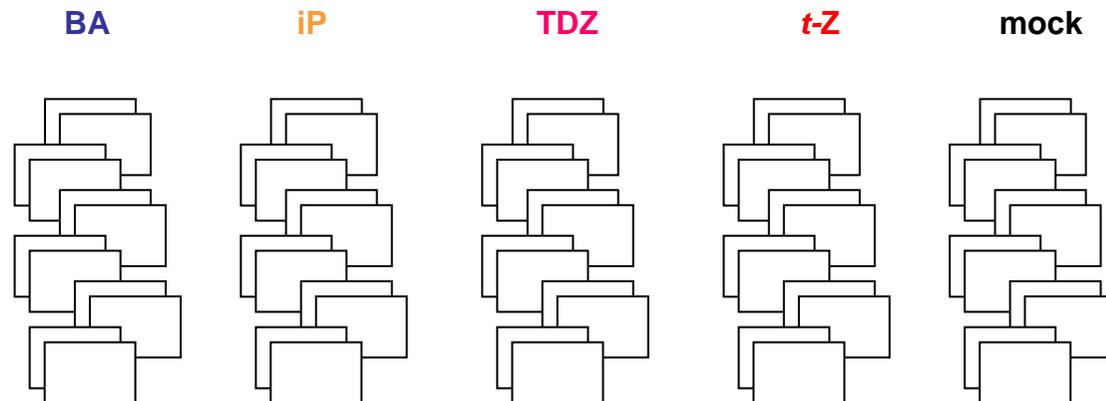


image analysis

76 2D gels

protein digest

31 protein spots reproducibly significant in at least 3 biological replicates

MALDI-TOF/TOF analysis

MASCOT search

29 identified proteins

Contribution of individual cytokinin receptors to phosphoproteome regulation

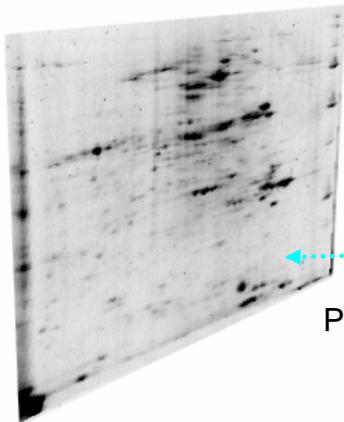
Arabidopsis thaliana **ahk2ahk3**, **ahk2cre1** and **ahk3cre1** double mutants seedlings were cultivated for 7 days on MS medium supplemented with (5.10⁻⁴ % v/v) DMSO at standard light intensity (~90 μmol.m⁻².s⁻¹).



Seedlings were transferred on liquid MS medium supplemented with DMSO (mock; 5.10⁻⁴ % v/v) or 5 μM *t*-Z dissolved in DMSO and incubated for 15 min.



Seedlings were rapidly harvested, frozen and ground in liquid nitrogen



For phosphoproteome analysis, an isolation procedure was established based on PhosphoProtein Purification Kit (Qiagen). Collected phosphoprotein fraction was then resolved by 2-DE (7 cm IPG strip, 150 μg of protein per strip).

Contribution of individual cytokinin receptors to phosphoproteome regulation

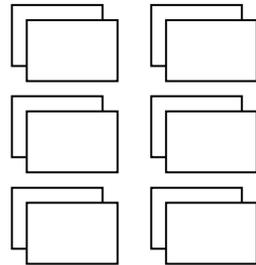
2-DE

2 technical replicates for each biological replicate

ahk2ahk3

t-Z

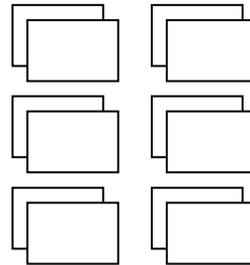
mock



ahk2cre1

t-Z

mock



ahk3cre1

t-Z

mock

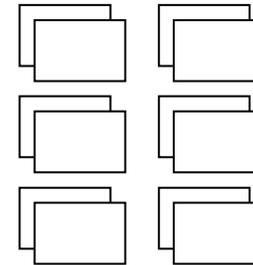
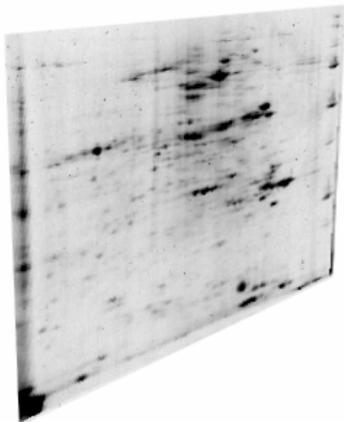


image analysis

36 2D gels

Comparison with early cytokinin response phosphoproteins

significant differences for 28 out of 31 proteins

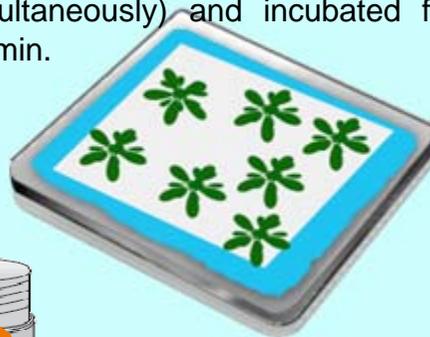


Effect of calcium signaling inhibitors on phosphoproteome response to cytokinins

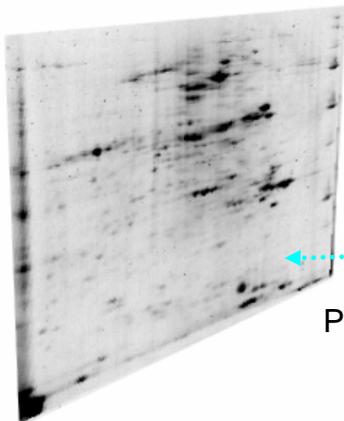
Arabidopsis thaliana (ecotype Columbia) seedlings were cultivated for 7 days on MS medium supplemented with (5.10⁻⁴ % v/v) DMSO at standard light intensity (~90 μmol.m⁻².s⁻¹).



Seedlings were transferred on liquid MS medium supplemented with inhibitors **INH**: 60 μM La³⁺, 30 μM D600 and DMSO (5.10⁻⁴ % v/v), **t-Z**: 5 μM t-Z dissolved in DMSO and **t-Z+INH** (inhibitors and cytokinins simultaneously) and incubated for 15 min.



Seedlings were rapidly harvested, frozen and ground in liquid nitrogen



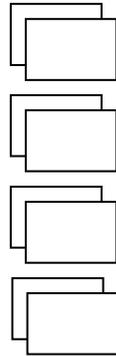
For phosphoproteome analysis, an isolation procedure was established based on PhosphoProtein Purification Kit (Qiagen). Collected phosphoprotein fraction was then resolved by 2-DE (7 cm IPG strip, 150 μg of protein per strip).

Effect of calcium signaling inhibitors on phosphoproteome response to cytokinins

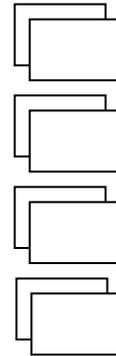
2-DE

2 technical replicates for each biological replicate

t-Z



INH+*t-Z*



INH

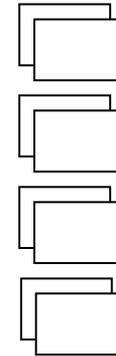


image analysis

24 2D gels

Comparison with early cytokinin response phosphoproteins

significant differences for 5 out of 31 proteins

