

Interactions of polyamines and phytohormones in plant response to abiotic stress

Oddziaływania poliamin i fitohormonów w odpowiedzi roślin na stres abiotyczny

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Table 1. Some representative evidence for interaction between PAs and plant hormones during plant growth and development. 6-BA, 6-benzylaminopurine; IAA, indole acetic acid; iP, isopentenyl adenine; NaSA, sodium salicylate.

Tabela 1. Niektóre reprezentatywne dowody interakcji między PA i hormonami roślinnymi podczas wzrostu i rozwoju roślin. 6-BA, 6-benzylaminopuryna; IAA, kwas indolooctowy; iP, izopentenyladenina; NaSA, salicylan sodu.

PAs-plant hormone crosstalk	Plant species	Exogenous treatment, gene mutation or over-expression	Effect (or outcome)	References
ABA	Maize	Exogenous ABA	Increased PAO activity, enhanced <i>PAO</i> expression	[50]
	Tomato	Exogenous ABA	Induced expression of <i>SIPAO2-4</i> genes, reduced expression <i>SIPAO1</i> gene and <i>SIPAO6-7</i> genes	[42]
	Cucumber	Exogenous ABA	Up-regulated expression of <i>CsSAMS1</i> and <i>CsSAMS2</i> in leaves, increased expression of <i>CsSAMS1</i> and inhibited expression of <i>CsSAMS2</i> in roots	[51]
	Rice	Exogenous ABA	Enhanced expression of <i>OsPAO1</i>	[52]
	<i>Arabidopsis</i>	Exogenous ABA	Up-regulated expression of <i>ADC2</i> , <i>ARGAH1</i> , <i>ARGAH2</i> , <i>AIH</i> , <i>CPA</i> , <i>SPDS2</i> , <i>SPMS</i> , <i>SAMDC1-2</i> , <i>PAO1-4</i> , <i>CuAOγ1/CuAO1</i> , <i>CuAOδ</i>	[43]
	<i>Arabidopsis</i>	Exogenous Spd and exogenous Spm	Increased ABA content, up-regulated <i>NCED3</i> expression	[53]
Auxin	Tomato	Exogenous IAA	Stimulated expression of <i>SIPAO2</i> , <i>SIPAO4</i> and <i>SIPAO7</i> genes, reduced expression of <i>SIPAO6</i>	[42]
	<i>Arabidopsis</i>	Exogenous IAA	Up-regulated expression of <i>ADC2</i> , <i>ACL5</i> , <i>SAMDC4/BUD2</i> , <i>PAO1</i> , <i>PAO2</i> , <i>PAO5</i> , reduced expression of <i>CuAOα2</i> and <i>CuAOα3/CuAO2</i>	[43]
	<i>Arabidopsis</i>	<i>acl5-1</i>	Up-regulation of genes related to auxin biosynthesis (<i>YUCCA2</i>), methylation (<i>IAMT1</i>), transport (<i>PIN-FORMED1</i> and <i>PIN6</i>) and signal transduction (<i>MP/ARF5</i>)	[47]
BRs	Tomato	Exogenous EBL	Modulated PAs content in plant organs	[54]
CKs	Cucumber	Exogenous KN	Increased Put content, decreased Spd content, reduced SAMDC activity and stimulated PAO activity	[55]
	Tomato	Exogenous 6-BA	Stimulated expression of <i>SIPAO2-5</i> genes and reduced expression of <i>SIPAO6-7</i> genes	[42]
	Rice	Exogenous iP	Up-regulated expression of <i>OsPAO1</i> , <i>OsPAO3</i> , <i>OsPAO5</i> and <i>OsPAO7</i>	[52]
	<i>Arabidopsis</i>	<i>bud2</i>	Enhanced CKs production, hypersensitivity to exogenous CKs	[56]

ETH	Tomato	Exogenous ethephon	Stimulated expression of <i>SIPAO1-2</i> genes and <i>SIPAO4</i> gene, decreased expression of <i>SIPAO5-7</i> genes	[42]
	<i>Arabidopsis</i>	Exogenous ACC	Enhanced expression of <i>ADC2</i> and reduced expression of <i>CuAOα2</i>	[43]
	Peach	Exogenous Spd	Decreased ETH content, enhanced expression of <i>ETR1</i> and <i>ERS1</i> in fruits	[46]
	Olive (ARB cv.)	Exogenous Put	Negatively regulated expression of <i>OeACO2</i> and <i>OeCTR1</i> , and positively regulated <i>OeERS1</i> expression in fruits	[57]
	Olive (ARB cv.)	Exogenous Spd	Up-regulated expression of <i>OeERS1</i> , down-regulated expression of <i>OeCTR1</i> and <i>OeEIL2</i> in fruits	[57]
	Grapes	Exogenous guazatine	Put accumulation, stimulated expression of <i>EIN3</i> and <i>EBF2</i> genes	[58]
	tomato	<i>Nr</i>	Enzymes of Put production responded to exogenous SA in a light- and ET-dependent manner	[44]
GAs	Pea	Exogenous GA3	Increased <i>ADC</i> expression, decreased <i>ODC</i> expression	[59]
	Grape	Exogenous GA3	Enhanced free Put content	[60]
	Tomato	Exogenous GA3	Up-regulated expression of <i>SIPAO2-3</i> genes and <i>SIPAO5</i> gene	[42]
	Rice	Exogenous GA	Up-regulated expression of <i>OsPAO5</i> and <i>OsPAO7</i>	[52]
	<i>Arabidopsis</i>	<i>ADC2</i>	Lower GA1, GA4 and GA9 content, reduced expression of genes related to GA biosynthesis (<i>AtGA20ox1</i> , <i>AtGA3ox1</i> and <i>AtGA3ox3</i>)	[45]
	Tomato	<i>ySAMdc</i>	Enhanced expression of genes encoding GA 2-oxidase and GA 20-oxidase and during fruit ripening	[61]
	<i>Arabidopsis</i>	Exogenous Put	Up-regulation <i>GA3ox1</i> expression	[53]
JAs	<i>Arabidopsis</i>	Exogenous Spm	Reduced GA1 content, down-regulation <i>GA3ox1</i> expression	[53]
	Tomato	Exogenous MeJA	Enhanced expression of <i>SIPAO1-2</i> genes and decreased expression of <i>SIPAO4</i> and <i>SIPAO6-7</i> genes	[42]
	Cucumber	Exogenous MeJA	Modulated expression of <i>CsSAMS1</i> and <i>CsSAMS2</i> genes in roots and leaves during treatment	[51]
	Rice	Exogenous JA	Up-regulated expression of <i>OsPAO1-3</i> and <i>OsPAO6-7</i>	[52]
	<i>Arabidopsis</i>	Exogenous MeJA	Enhanced expression of <i>ARGAH1</i> , <i>ARGAH2</i> , <i>PAO3</i> and <i>CuAOα3/CuAO2</i>	[43]
	<i>Arabidopsis</i>	Exogenous Spm	Up-regulation of JA biosynthesis genes (<i>LOX1-4</i> , <i>AOS</i> , <i>AOC1</i> , <i>AOC2</i> , <i>OPR3</i> , <i>OPCL1</i> , <i>CYP94B3</i>) and JA signaling genes (<i>JAZ1</i> , <i>JAZ5</i> , <i>JAZ6</i> , <i>JAZ7</i> and <i>JAZ10</i>)	[62]
	<i>Arabidopsis</i>	Exogenous tSpm	Up-regulation of JA biosynthesis genes (<i>LOX1-4</i> , <i>AOS</i> , <i>AOC1</i> , <i>AOC2</i> , <i>OPR3</i> , <i>OPCL1</i> , <i>CYP94B3</i>), JA signaling genes (<i>JAZ1</i> , <i>JAZ5</i> , <i>JAZ6</i> , <i>JAZ7</i> , <i>JAZ10</i> and <i>MYC4</i>) and JA response marker genes (<i>VSP2</i> and <i>PDF1.2</i>)	[62]
SA	<i>Arabidopsis</i>	Exogenous Spd	Lower JA content, reduced <i>AOS</i> expression	[53]
	<i>Arabidopsis</i>	Exogenous Spm	Lower JA content, enhanced <i>AOS</i> expression	[53]
	Tomato	Exogenous SA	Stimulated expression of <i>SIPAO2-4</i> genes and reduced expression of <i>SIPAO6-7</i> genes	[42]
	Cucumber	Exogenous SA	Enhanced expression of <i>CsSAMS1</i> and <i>CsSAMS2</i> in leaves	[51]
	<i>Arabidopsis</i>	Exogenous SA	SA modulated PAs content and PA metabolism gene expression	[63]

<i>Arabidopsis</i>	<i>mpk6-2</i>	Decreased Put content	[63]
<i>Arabidopsis</i>	Exogenous NaSA	Up-regulated expression of <i>ADC2</i> , <i>AIH</i> , <i>SPMS</i> and <i>PAO1</i> , down-regulated expression of <i>CuAOα2</i> , <i>CuAOα3/CuAO2</i> and <i>CuAOγ2</i>	[43]
<i>Arabidopsis</i>	Exogenous PAs	Modulation of PAs biosynthesis and catabolism gene expression in <i>eds5</i> and <i>sid2</i> mutants	[53]

Table 2. A summary of evidence for interaction between PAs and plant hormones in response to abiotic stresses.

Tabela 2. Podsumowanie dowodów na interakcje między PA i hormonami roślinnymi w odpowiedzi na stresy abiotyczne.

Abiotic stress	PAs-plant hormone crosstalk	Plant species	Exogenous treatment, gene mutation or over-expression	Effect (or outcome)	References
Drought, osmotic stress	ABA ↔ PAs	<i>Vicia faba</i>	Exogenous PAs	Induced stomatal closure	[71]
	ABA ↔ PAs	<i>Vicia faba</i>	Exogenous ABA	Higher CuAO and NADPH oxidase activity leading to enhanced H ₂ O ₂ production	[71]
	ABA ↔ PAs	<i>Arabidopsis</i>	Exogenous ABA	Higher expression of <i>AtPAO2</i> in guard cells	[72]
	ABA ↔ PAs	<i>Arabidopsis</i>	Exogenous PAs	Induced stomatal closure, enhanced NO and ROS levels in guard cells due to NADPH oxidase and AO activity	[73]
	ABA ↔ PAs	<i>Arabidopsis</i>	<i>aba2</i> , <i>abi1</i>	No observed dehydration-inducible expression of <i>ADC2</i> , <i>SPDS1</i> and <i>SPMS</i>	[74]
	ABA ↔ PAs	<i>Arabidopsis</i>	-	Presence of ABREs in promoter regions of <i>ADC2</i> , <i>SPDS1</i> and <i>SPMS</i> genes	[75]
	ABA ↔ PAs	<i>Arabidopsis</i>	-	Enhanced expression of ABA-inducible and drought-responsive genes (<i>RD29A</i> and <i>RD22</i>) and PAs biosynthesis genes (<i>ADC1</i> , <i>ADC2</i> , <i>SPDS1</i> , <i>SPDS2</i> , <i>SPMS</i> and <i>SAMDC1</i>)	[66]
	ABA ↔ PAs	Rice	-	Presence of putative cis-acting elements, such as ABRE, LTRE, MYB and W-box, in the promoter region of <i>SAMDC</i> gene	[76]
	ABA ↔ PAs	Rice	<i>OsHSFA3</i>	Higher ABA level, increased PAs content, up-regulation of <i>ADC1</i> , <i>ADC2</i> , <i>SPDS1</i> and <i>SPMS</i> expression	[77]
	ABA ↔ PAs	<i>Lotus tenuis</i>	<i>ADC</i>	Increased Put content, expression of <i>NCED</i> gene and ABA accumulation	[78]
	ABA ↔ PAs	Wheat	Exogenous Put	Enhanced expression of <i>NCED</i>	[79]
		Wheat (drought-susceptible cv.)	Exogenous Spm	Further escalation in ABA level	[80]
	ABA ↔ PAs	Wheat (drought-resistant cv.)	Exogenous Spd	Reduced ABA level	[80]

	ABA ↔ PAs	Wheat	Exogenous ABA	Enhanced expression of <i>ADC</i> , decreased expression of <i>SPDS</i> and <i>PAO</i>	[79]
	ETH ↔ PAs	Wheat	Exogenous Spd and exogenous Spm	Reduction of ETH evolution rate, increased CKs and ABA level	[81]
	ETH ↔ PAs	Wheat	Exogenous Put	Increased ETH evolution rate, excessive ABA accumulation	[81]
	JAs ↔ PAs	Rice	Exogenous MeJa	Negatively regulation of <i>OsAdc</i> , <i>OsSamdc</i> and <i>OsSpds</i> genes	[82]
	JAs ↔ PAs	Maize	Exogenous Spd	Decreased MeJA content	[37]
	CKs ↔ PAs	Maize	Exogenous Spd	Increased ZR content	[37]
	GAs ↔ PAs	Wheat	Exogenous Spd and exogenous Spm	Higher GA (GA1+GA4) content during seed germination	[81]
	GAs ↔ PAs	Bentgrass	Exogenous Spm	Increased GA1 content	[83]
	GAs ↔ PAs	Bentgrass	Exogenous Spd	Increased GA4 and GA20 content	[83]
Salt stress	ABA ↔ PAs	<i>Arabidopsis</i>	<i>SAMDC1</i>	Improved expression of ABA-induced genes (<i>NCED</i> and <i>RD29A</i>)	[19]
	ABA ↔ PAs	<i>Arabidopsis</i>	<i>spms</i> , <i>acl5/spms</i>	Decreased <i>NCED3</i> expression	[19]
	ABA ↔ PAs	<i>Arabidopsis</i>	<i>atpao5-3</i>	Higher ABA content, up-regulated expression of <i>NCED</i> and <i>RD29B</i>	[62]
	BRs ↔ PAs	Tomato	Exogenous EBL	Increased PAs level in the middle leaf and fruits, higher Spm level in the stem higher (Spd+Spm)/Put ratio in the fruits	[54]
	BRs ↔ PAs	Lettuce	Exogenous DI-31	Enhanced PAs accumulation in shoots, reduced Spd and Spm content in roots	[84]
	BRs ↔ PAs	Canola	Exogenous EBL	Increased accumulation of Put in cotyledons and reduced in hypocotyls and radicles, reduced H ₂ O ₂ content, decreased DAO activity and increased PAO activity	[12]
	ETH ↔ PAs	Rice	Exogenous Put	Enhanced ETH production	[65]
	ETH ↔ PAs	Tobacco	<i>AS-ZmPAO</i>	Up-regulation of <i>ACCSyn</i> and <i>ACCOx</i> gene expression	[4]
	ETH ↔ PAs	Tomato	<i>Nr</i>	Lower Spm content, enhanced DAO, and PAO activity	[44]
	JAs ↔ PAs	Cucumber	Exogenous Spd	Decreased JA content	[85]
	JAs ↔ PAs	<i>Arabidopsis</i>	<i>atpao5-3</i>	Increased JA accumulation	[62]
	CKs ↔ PAs	<i>Vigna sinensis</i>	Exogenous KN	Increased PAs content	[86]
	GAs ↔ PAs	Cucumber	Exogenous Spd	Enhanced <i>GT-3b</i> expression, increased GA3 level, increased activity of GA3-oxidase and GA20-oxidase, upregulated expression of genes related to GA biosynthesis	[31]
Heavy metal stress	BRs ↔ PAs	Radish	Exogenous EBL	Increased Put content, decreased Spd content	[54]
	BRs ↔ PAs	Mustard	Exogenous EBL	Increased PAs level in leaves	[87]
	BRs ↔ PAs	Mustard	Exogenous EBL and SA	Increased Spd level in roots	[88]
	ETH ↔ PAs	Wheat	Exogenous Put	Improved root elongation, decreased ACS activity and ETH level	[6]

	ETH ↔ PAs	Cucumber	Exogenous 2-OHMT	Higher PAs content, reduction of PAO activity, increased CS-ERS expression	[7]
	ABA ↔ PAs	Tomato	Exogenous Put	Increased ABA level and <i>LeNCED1</i> expression, role of ABA in Put-induced tolerance	[33]
	ABA ↔ PAs	<i>Arabidopsis</i>	<i>adc</i>	Reduced ABA accumulation, lower <i>NCED</i> expression, down-regulation of ABA-regulated genes	[10]
	ABA ↔ PAs	<i>Arabidopsis</i>	<i>adc</i>	Reduced ABA level, decreased ABA-dependent gene induction, decreased freezing tolerance	[89]
Low-temperature stress	JAs ↔ PAs	Zucchini	Exogenous MeJA	Higher Put content, lower Spd and Spm content	[90]
	JAs ↔ PAs	Rice	Exogenous MeJA	Shoots: increased Put and Spm content, reduced Spd content, increased ADC activity, reduced ODC and SAMDC activity; roots: increased Put and reduced Spd content, increased ADC and SAMDC activity, reduced ODC activity	[30]
	CKs ↔ PAs	<i>Indica-japonica</i> hybrid rice	Exogenous Spd	Higher ZR content	[91]
	GAs ↔ PAs	<i>Indica-japonica</i> hybrid rice	Exogenous Spd	Higher GA3 content	[91]