

## Inducing salt tolerance in castor bean through seed priming

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**Supplementary Table 1.** Summary of the results of the ANOVA for the main effects of salinity on seed germination (G), dead seeds (DS), non-germinated seeds (NGS), normal seedlings (NP), germination speed index (GSI), average time of germination (ATG), root DM (RDM), hypocotyl DM (HDM), root/hypocotyl DM ratio (RDM/HDM), root length (RL), hypocotyl length (HL), root/hypocotyl length ratio (RL/HL), sodium (Na<sup>+</sup>) and potassium (K<sup>+</sup>) content and K<sup>+</sup>/Na<sup>+</sup> content ratio in the roots and hypocotyl of castor bean seeds subjected to different values of  $\Psi$ s obtained with NaCl salt

Source of variation	df	Mean Square					
		G	DS	NGS	NP	GSI	ATG
Salinity	4	4012.7**	3*	3805.7**	4343.3**	90.77**	11.31**
Error	15	49.73	0.73	49.93	62.53	0.90	1.36
CV (%)		13.16	171.27	15.40	15.75	13.42	13.53
Source of variation	df	Mean Square					
		RDM	HDM	RDM/HDM	RL	HL	RL/HL
Salinity	3	65.30**	392.99**	1.11**	625.21**	949.29**	417.02**
Error	11	1.78	3.15	0.11	8.31	5.09	2.26
CV (%)		21.63	12.16	51.26	42.80	43.59	72.4
Source of variation	df	Root			Hypocotyl		
		Na <sup>+</sup>	K <sup>+</sup>	K <sup>+</sup> /Na <sup>+</sup>	Na <sup>+</sup>	K <sup>+</sup>	K <sup>+</sup> /Na <sup>+</sup>
Salinity	3	632054**	73447**	13.7**	174683**	3920.94**	65.28**
Error	11	1394.11	9129.78	0.018	1512.97	597.60	0.89
CV (%)		6.06	21.97	7.4	15.45	13.50	33.52

\* p < 0.05; \*\* p < 0.01; <sup>ns</sup> p > 0.05

**Supplementary Table 2.** Summary of the results of the ANOVA for the main effects of salinity and time on fresh mass of castor bean seeds subjected to different values of  $\Psi$ s obtained using NaCl salt

Source of variation	df	Mean Square	
		Wet mass	Wet mass
Salinity (S)	3	0.148448**	0.099109**
Time (T)	24	0.139922**	0.134580**
S x T	72	0.006417**	0.170922**
S / 0h	3	0 <sup>ns</sup>	0.275835**
S / 1h	3	0.001304 <sup>ns</sup>	0.294480**
S / 3h	3	0.001693 <sup>ns</sup>	0.342410**
S / 6h	3	0.002958 <sup>ns</sup>	0.386110**
S / 12h	3	0.001269 <sup>ns</sup>	0.560261**
S / 24h	3	0.005056 <sup>ns</sup>	0.770868**
S / 36h	3	0.004944 <sup>ns</sup>	0.933924**
S / 48h	3	0.004792 <sup>ns</sup>	1.130545**
S / 60h	3	0.003568 <sup>ns</sup>	1.446836**
S / 72h	3	0.005436 <sup>ns</sup>	0.587753**
S / 84h	3	0.006392 <sup>ns</sup>	0.371084**
S / 96h	3	0.016550 <sup>ns</sup>	0.349136**
S / 108h	3	0.031344 <sup>ns</sup>	0.017887
S / 120h	3	0.048145**	2.65
Error	900		
CV (%)			

\* p < 0.05; \*\* p < 0.01; <sup>ns</sup> p > 0.05

**Supplementary Table 3.** Summary of the results of the ANOVA for electrical conductivity (EC), normal seedlings (NP), hypocotyl length (HL), root length (RL), root/hypocotyl length ratio (RL/HL), hypocotyl DM (HDM), root DM (RDM) and root/hypocotyl DM ratio (RDM/HDM) of castor bean seeds not primed and primed with H<sub>2</sub>O, NaCl, CaCl<sub>2</sub>, KCl, NaNO<sub>2</sub>, NaNO<sub>3</sub>, KH<sub>2</sub>PO<sub>4</sub>, NaSiO<sub>3</sub>, PEG-6000 or H<sub>2</sub>O<sub>2</sub>

Source of variation	df	Mean Square			
		EC	NP	HL	RL
Priming	10	1055.42**	505.81*	49.11*	105.63**
Error	44	49.49	201.81	20.80	33.51
CV (%)		11.11	29.71	52.46	48.88

  

Source of variation	df	Mean Square			
		RL/HL	HDM	RDM	RDM/HDM
Priming	10	3.23**	1630.68**	143.66 <sup>ns</sup>	0.026**
Error	44	1.20	248.83	89.43	0.004
CV (%)		67.48	12.73	22.03	18.72

\* p < 0.05; \*\* p < 0.01; <sup>ns</sup> p > 0.05

**Supplementary Table 4.** Summary of the results of the ANOVA for the main effects of osmopriming on seed germination (G), dead seeds (DS), non-germinated seeds (NGS), normal seedlings (NP), germination speed index (GSI), average time of germination (ATG), root DM (RDM), hypocotyl DM (HDM), root/hypocotyl DM ratio (RDM/HDM), root length (RL), hypocotyl length (HL), root/hypocotyl length ratio (RL/HL), sodium (Na<sup>+</sup>) and potassium (K<sup>+</sup>) content and the K<sup>+</sup>/Na<sup>+</sup> content ratio in roots and hypocotyl of castor bean seeds subjected an Ψs of -0.6 MPa obtained with NaCl salt

Source of variation	df	Mean Square					
		G	DS	NGS	NP	GSI	ATG
Priming	4	1083.15**	31.4**	1160**	781.85**	73.38**	19.07**
Error	35	52.54	4.68	54.94	71.68	1.22	0.31
CV (%)		13.65	47.06	17.52	18.53	14.26	7

  

Source of variation	df	Mean Square					
		RDM	HDM	RDM/HDM	RL	HL	RL/HL
Priming	4	108.62**	525.78**	0.00024**	148.46**	31.04**	54.66**
Error	35	4.41	10.95	0.000011	5.58	0.76	8.02
CV (%)		11.46	12.44	10.49	39.32	45.93	69.77

  

Source of variation	df	Mean Square					
		Root			Hypocotyl		
		Na <sup>+</sup>	K <sup>+</sup>	K <sup>+</sup> /Na <sup>+</sup>	Na <sup>+</sup>	K <sup>+</sup>	K <sup>+</sup> /Na <sup>+</sup>
Priming	4	13723 <sup>ns</sup>	49175**	0.05**	6880*	5432**	0.0091 <sup>ns</sup>
Error	35	17400	6375	0.002	1856	986	0.0034
CV (%)		11.22	12.12	8.05	10.62	11.79	8.89

\* p < 0.05; \*\* p < 0.01; <sup>ns</sup> p > 0.05