Supplementary material

The phosphatidylcholine-hydrolyzing phospholipase C NPC4 plays a role in response of Arabidopsis roots to salt stress

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Locus	UPL	Name	Oligo Sequence
At1g07230	UPL39	NPC1 L	ctcgccaagattgtccaga
		NPC1 R	gagttgagaggccagttgga
At2g26870	UPL76	NPC2 L	tctccgagttcgctgttttc
		NPC2 R	tgacgtccctgagtgtacaaa
At3g03520	UPL93	NPC3 L	atcgaacctggaacggtgt
		NPC3 R	agtgttgccggaattgatg
At3g03530	UPL46	NPC4 L	tgcacaattaagcgaatttcaa
		NPC4 R	ttcttcgtttttgtagtctcctttt
At3g03540	UPL122	NPC5 L	attacgagaaactcaccaagacaa
		NPC5 R	cattttgaccggatttgaca
At3g48610	UPL92	NPC6 L	aatcccgatgggaatactgg
		NPC6 R	ggtccttttgcttcacttacga
At3g18780	UPL30	Actin2 L	ccgctctttctttccaagc
		Actin2 R	ccggtaccattgtcacacac
At4g05320	UPL10	UBQ10 L	tcaccggaaagaccatcact
		UBQ10 R	cggtgggataccctctttg
	1		

At4g26080ABI1 LccatccatcattcctgatccABI1 RccatccacacgcttcttcaAt5g57050ABI2 LABI2 RgagatcgatgaatcagagtgaAT5g27150NHX1 LNHX1 RgcaccaagcaaggtacttagAt5g07920DGK1 LDGK1 RaaccagtattccttccgcaAt5g52310RD29A/LTI78/COR78 LRD29A/LTI78/COR78 RgtgctctgtttggctcctcAt1g14660NHX8 LNHX8 RcaatccaacccatagccaggAt3g11410PP2CA LPP2CA RaagacggatgtaacggaagttgccaggtAt5g66400RAB18 LCaatccaacccatagccaggtAt5g66400RAB18 LCaatccaacccatagccaggtCaatgagaagttgccaggtAt5g66400RAB18 LCaatgagaagttgccaggtCaatgagaagttgccaggtCaatgagaagttgccaggtCaatgagagagagtgccaggtCaatgagagagtgccaggtg	9
ABI1 RccatctcacacgcttcttcaAt5g57050ABI2 LgagatcgatgaatcagagtgaABI2 RcattagtgactcgaccatcaagAT5g27150NHX1 LtcttgtattcggagagggtgNHX1 RgcaccaagcaaggtacttagAt5g07920DGK1 LDGK1 RaaccagtattccttccgcaDGK1 RaaccagtattccttccgcaAt5g52310RD29A/LTI78/COR78 LtcaccggaactttctcatccRD29A/LTI78/COR78 RAt1g14660NHX8 LAt3g11410PP2CA LPP2CA RaagacggatgtaacggaactcgggAt5g66400RAB18 LtcacggaagttgccaggtRAB18 Rtatccaatcacccttette	9
At5g57050AB/2 LgagatcgatgaatcagagtgaAB/2 RcattagtgactcgaccatcaagAT5g27150NHX1 LtcttgtattcggagagggggNHX1 RgcaccaagcaaggtacttagAt5g07920DGK1 LDGK1 RagccagtttatgaacaaagtcoDGK1 RaaccagtattccttccgcaAt5g52310RD29A/LTI78/COR78 LtcaccggaactttctcatccRD29A/LTI78/COR78 RgtgctctgttttggctcctcNHX8 RAt1g14660NHX8 LNHX8 RcaatccaacccatagccagAt3g11410PP2CA LtctgaggcggagtgtaacggaagtgtaacgcaaccAt5g66400RAB18 LtcaaggagagttgccaggtAt5g66400RAB18 Ltatccaatccccttcttc	g
AB/2 RcattagtgactcgaccatcaagAT5g27150NHX1 LtcttgtattcggagagggtgNHX1 RgcaccaagcaaggtacttagAt5g07920DGK1 LagccagtttatgaacaaagtcoDGK1 RaaccagtattccttccgcaAt5g52310RD29A/LTI78/COR78 LtcaccggaactttctcatccRD29A/LTI78/COR78 RgtgctctgttttggctcctcAt1g14660NHX8 LatcttacaaaggaaactcgtggNHX8 RcaatccaacccatagccagAt3g11410PP2CA LtctgaggccgagatacggPP2CA RaagacggatgtaacgcaaccAt5g66400RAB18 Ltctacaaggaagttgccaggt	:
AT5g27150NHX1 LtcttgtattcggagaggtgNHX1 RgcaccaagcaaggtacttagAt5g07920DGK1 LDGK1 RagccagtttatgaacaaagtcoDGK1 RaaccagtattccttccgcaAt5g52310RD29A/LTI78/COR78 LRD29A/LTI78/COR78 RgtgctctgttttggctcctcAt1g14660NHX8 LNHX8 RcaatccaacgagatacggNHX8 RcaatccaacccatagccagAt3g11410PP2CA LPP2CA RaagacggatgtaacgcaaccAt5g66400RAB18 LBAB18 Rtatccaataccccttcttc	:
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At5g52310RD29A/LTI78/COR78 LtcaccggaactttctcatccRD29A/LTI78/COR78 RgtgctctgttttggctcctcAt1g14660NHX8 LatcttacaaaggaaactcgtggNHX8 RcaatccaacccatagccagAt3g11410PP2CA LtctgaggccgagatacggPP2CA RaagacggatgtaacgcaaccAt5g66400RAB18 Ltctaaggagaagttgccaggt	
RD29A/LTI78/COR78 RgtgctctgttttggctcctcAt1g14660NHX8 LatcttacaaaggaaactcgtggNHX8 RcaatccaacccatagccagAt3g11410PP2CA LtctgaggccgagatacggPP2CA RaagacggatgtaacgcaaccAt5g66400RAB18 Ltcaaggagaagttgccaggt	
At1g14660 NHX8 L atcttacaaaggaaactcgtgg NHX8 R caatccaacccatagccag At3g11410 PP2CA L tctgaggccgagatacgg PP2CA R aagacggatgtaacgcaacc At5g66400 RAB18 L tctaaggagaagttgccaggt	l
NHX8 RcaatccaacccatagccagAt3g11410PP2CA LtctgaggccgagatacggPP2CA RaagacggatgtaacgcaaccAt5g66400RAB18 LtcaaggagaagttgccaggtRAB18 Rtgtccatcatccccttettc	
At3g11410 PP2CA L tctgaggccgagatacgg PP2CA R aagacggatgtaacgcaacc At5g66400 RAB18 L tcaaggagaagttgccaggt RAB18 R tdtccatcatccccttettc	
PP2CA RaagacggatgtaacgcaaccAt5g66400RAB18 LtcaaggagaagttgccaggtRAB18 Rtdtccatcatccccttettc	
At5g66400 RAB18 L tcaaggagaagttgccaggt	
RAR18 R tatccatcatcatcatcttc	
At2g47770TSPO Ltttgctctgtttggtttggg	
TSPO R cattaacagcggctacaaagg	
At2g01980 SOS1 L agcgaccgattcgtcttct	
SOS1 R ttagctccatattcgagagatcc	
At5g35410 CIPK24/SOS2 L gtgcagtttttgatggaattga	
CIPK24/SOS2 R gccttttaacaaaatcctgtcg	
At5g24270 CBL4/SOS3 L tatttgatgtgaagcgaaatgg	
CBL4/SOS3 R cgatgaatccagtttgtcgt	
At2g03760 SOT12 L cccagatttcgatttctcca	
SOT12 R tcccaaaatgccataaggac	
At1g50030 TOR L tcgaagggcaaagtacgac	



Figure S1. Histochemical analysis of P_{NPC3} :GUS expression under salt stress in Arabidopsis plants. A) Effect of salt on P_{NPC3} :GUS expression pattern in the main root of ten-day-old seedlings. The plants were grown on agar and transferred in liquid nutrient solution with or without 100 mM NaCl for 24 h. B) Effect of salt on P_{NPC3} :GUS expression pattern in roots of five-week-old Arabidopsis plants. The plants were grown hydroponically in modified half strength Hoagland's solution and exposed to 100 mM NaCl for 4 h.



Figure S2. The position of T-DNA insertions in two Arabidopsis *npc4* mutant lines. The SALK line 046713 was marked as *npc4-1* and the GABI-KAT line 571E10 was marked as *npc4-2*.



Figure S3. Relative level of gene expression of ion transporter genes *NHX1*, *NHX8* (A), phospholipid signalling related genes *TOR* and *DGK1* (*B*) and of *RD29/LT178/COR78* gene (C) in WT, *npc4-1* and *npc4-2* under 100 mM NaCl treatment. The transcript levels of genes were measured by quantitative real-time PCR in roots of non-treated plants and in plants treated with 100 mM NaCl for 3, 6, 9 and 12 h. *Actin2* and *UBQ10* were used as an internal

control. The expression of the genes in non-treated controls in particular time was set to 1. Data represents means + SE, n = 2 discrete samples from one biological experiment. This experiment was repeated twice with similar results.