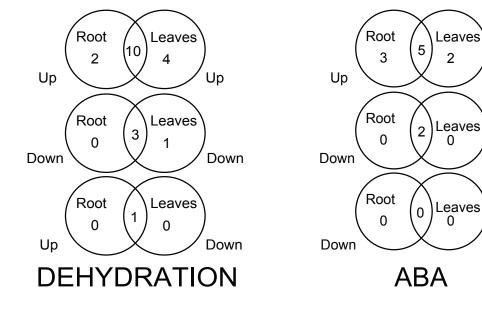
A	#	Gene	Dry		ABA		ABA
			Leaves	Roots	Leaves	Roots	responsiveness
	1	CaNAC02	Down	Down	Down	Down	Dependent
	2	CaNTL1/CaNAC04	Down	Down	Down	Down	Dependent
	3	CaNAC05	Up				Independent
	4	CaNAC06	Up	Up	Up	Up	Dependent
	5	CaNAC16	Up	Up			Independent
	6	CaNTL2/CaNAC19	Up	Up			Independent
	7	CaNAC21	Up			Up	Dependent
	8	CaNAC24	Down	Up	Up	Up	Dependent
	9	CaNAC27	Up	Up			Independent
	10	CaNTL3/CaNAC31					
	11	CaNTL4/CaNAC33					
	12	CaNAC39					
	13	CaNAC40	Up	Up		Up	Dependent
	14	CaNTL5/CaNAC41	Up				Independent
	15	CaNAC43	Up	Up		Up	Dependent
	16	CaNTL6/CaNAC44		Up			Independent
	17	CaNAC46	Down	Down	Up		Dependent
	18	CaNAC47	Up	Up			Independent
	19	CaNAC50	Up	Up	Up	Up	Dependent
	20	CaNAC52	Up	Up	Up	Up	Dependent
	21	CaNTL7/CaNAC57	Up		Up		Dependent
	22	CaNAC67	Up	Up	Up	Up	Dependent
	23	CaNTL8/CaNAC71					

Β



Up

Down

Up

Figure S3. Expression of 23 selected *CaNAC* genes in chickpea roots and leaves under dehydration and ABA treatments. (A) Summary of the results of the expression data. (B) Venn diagram analysis of dehydration- and ABA-responsive CaNAC genes in roots and leaves of chickpea plants. The ABAand/or dehydration-responsive genes were defined as those whose expression is altered by at least 2-fold (P < 0.05) at 2 and/or 5 h of dehydration and/or ABA treatment.