A

#	Gene	Dry		ABA		ABA responsiveness
		Leaves	Roots	Leaves	Roots	
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*					

B

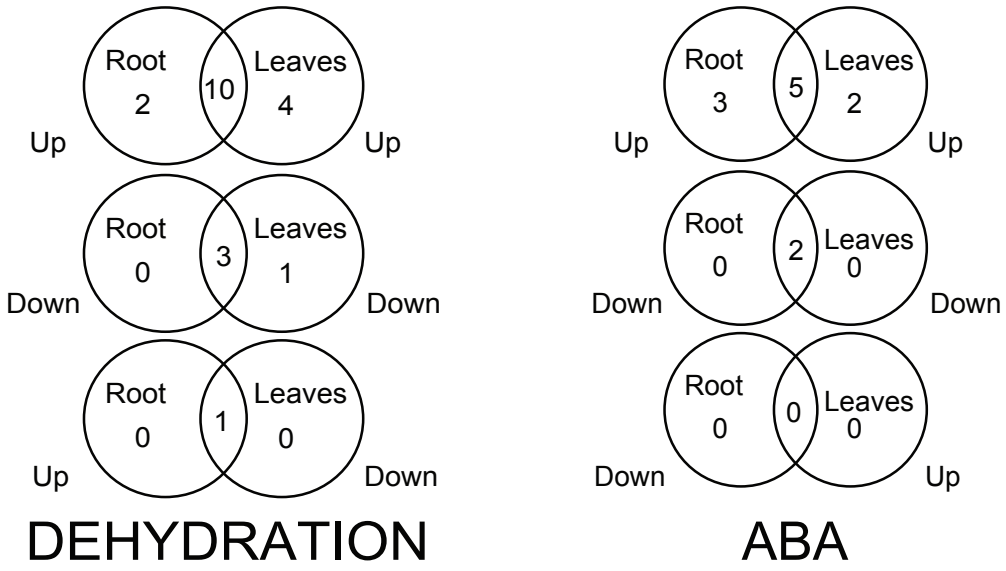

Figure S3. Expression of 23 selected *CaNAC* genes in chickpea roots and leaves under dehydration and ABA treatments. (A) Summary of the results of the expression data. (B) Venn diagram analysis of dehydration- and ABA-responsive CaNAC genes in roots and leaves of chickpea plants. The ABA- and/or dehydration-responsive 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.