

Table S4. Tissue distribution profile of chickpea ARF genes according to the number of expressed sequence tags (ESTs) present in NCBI's EST Database.

Gene Name	Gene ID	EST tissue	Accession	Genotype	Reference
<i>CaARF1</i>	LOC101492112	root	HO067475	RIL from the cross ICC4958xICC1882	[1]
<i>CaARF1</i>	LOC101492112	leaf	FE670495	XJ-209	[2]
<i>CaARF2</i>	LOC101513952	leaf	FE670500	XJ-209	[2]
<i>CaARF3</i>	LOC101501408	root	GR403988	ICCV2	[3]
<i>CaARF7</i>	LOC101504978	leaf	ES560336	Pusa-362	[4]
<i>CaARF12</i>	LOC101496441	leaf	ES560336	Pusa-362	[4]
<i>CaARF14</i>	LOC101498188	shoot	HO066200	ICC4958	[1]
<i>CaARF14</i>	LOC101498188	embryo	JK709419	ICCV2	[5]
<i>CaARF15</i>	LOC101500671	leaf	FE671045	XJ-209	[2]
<i>CaARF15</i>	LOC101500671	root	GR403988	ICCV2	[3]
<i>CaARF20</i>	LOC101489666	root	HO064721	ICC4958	[1]
<i>CaARF23</i>	LOC101505502	root	HO064459	ICC4958	[1]
<i>CaARF23</i>	LOC101505502	root	HO062185	ICC4958	[1]
<i>CaARF23</i>	LOC101505502	root	GR399001	ICC4958	[3]

[1] Deokar, A.A., Kondawar, V., Jain, P.K. et al. Comparative analysis of expressed sequence tags (ESTs) between drought-tolerant and -susceptible genotypes of chickpea under terminal drought stress. *BMC Plant Biology* (2011) 11:70

[2] Gao, W.R., Wang, X-S., Liu, Q.Y. et al. Comparative analysis of ESTs in response to drought stress in chickpea (*C. arietinum* L.). *Biochem Biophysical Res Comm* (2008) 376:578-583

[3] Varshney RK, Hiremath PJ, Lekha P, et al. A comprehensive resource of drought- and salinity-responsive ESTs for gene discovery and marker development in chickpea (*Cicer arietinum* L.). *BMC Genomics* (2009) 10:523

[4] Jaiswal, P., Cheruku, J.R., Kumar, K. et al. Differential transcript accumulation in chickpea during early phases of compatible interaction with a necrotrophic fungus *Ascochyta rabiei*. *Mol Biol Rep* (2012) 39: 4635.

[5] Gupta S, Garg V, Bhatia S (2015) A New Set of ESTs from Chickpea (*Cicer arietinum* L.) Embryo Reveals Two Novel F-Box Genes, *CarF-box_PP2* and *CarF-box_LysM*, with Potential Roles in Seed Development. *PLoS ONE* 10(3): e0121100.