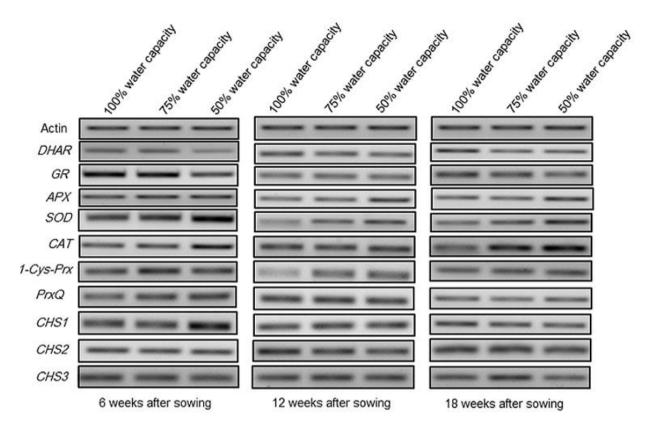
## **Supplementary Materials**



**Figure S1.** Semi-quantitative RT-PCR of antioxidant enzymes genes *DAHR*, *GR*, *APX*, *SOD*, and *CAT*, and expression of *1-Cys-Prx* and *PrxQ*, additionally, expression of chalcone synthase genes (*CHS1*, *CHS2*, and *CHS3*) from cDNA of milk thistle (*Silybum marianum* L.) leaves treatments with 100% water capacity, 75% water capacity and 50% water capacity at different time points. Samples standardized to actin. RT-PCR product were separated on agarose gels (1.5%) and visualized by ethidium bromide stain. *DHAR* dehydroascorbate reductase; *SOD* superoxide dismutase; *CAT* catalase; *APX* ascorbate peroxidase; *GR* glutathione reductase; *1-Cys-Prx* 1-cysteine peroxiredoxin; *PrxQ* peroxiredoxin Q.

**Figure S2.** Schematic presentation of flavonolignan biosynthetic pathway. CHS, chalcone synthase; CHI, chalcone isomerase; F3H, flavanone 3-hydroxylase; F3'H, flavonol 3'-hydroxylase). The end product of the pathway is silybin.

**Table S1.** Primers sequences for semi quantitative and quantitative RT-PCR of the drought-related genes in milk thistle (*Silybum marianum* L.).

Gene name	Reference Sequence	5' - 3' primer sequence	TA
Actin	AB181991	CTCTGACAATTTCCCGCTCA	58 ℃
		ACACGCTTCCTCATGCTATCC	
CHS1	JN182805.1	GTGTATACCAAGCCGATTATCCA	58 ℃
		GTGTGAATACATACACATGCGCT	
CHS2	JN182806.1	GTGCCTGCGTGTTGTTCATC	58 ℃
		CTGTGAAGGACCACCGTCTC	
CHS3	JN182807.1	GGGAGGAAGGGCTCACATTT	58 ℃
		ATGCCCAATGGAAAAACCGC	
DHAR	AY074784	TCAAGAACGTGACAAAGGTGG	58 ℃
		TAACGGTGGTGATGGCAAAT	
SOD	MG893090.1	TTCGCCATGCTGGTGATCTT	58 ℃
		CATGGACAACTACGGCCCTT	
CAT	GU984379	GGCTGCTTGAAGTTGTTCTCCT	58 ℃
		CTGCTAGTACCTCCTGATCCGTT	
APX	KU747079.1	TGGCCTGCTCTTCCTCTAGT	58 ℃
		CATGCCACGCTAATCGAAGC	
GR	KX828561.1	CAACGCGCTTTGGTAACTCC	58 ℃
		GGGCCCTAATGAAGTGGAGG	
1-Cys-Prx	AY304482	AATGGACCAAGGACATCGAG	58 ℃
		GACGGGTACAGGAAGCTCAG	
PrxQ	AY789643	ACTTCACGCTCAAGGACCAG	58 ℃
		CCGCCTTCTTGTACTTCTCG	