

Supplementary data**"Type A" Response Regulators are involved in the plant-microbe interaction****Fariba Rafiei, Zohreh Torabi, Esmaeil Ebrahimie**

Supplementary Table 1. Selected microarray data from GSE33636 to find co-over expressed genes in *Medicago truncatula-Sinorhizobium meliloti* interaction

name	Treatment	Sample name	Sample Treatment
GSM831907	Nod-LCOs 6h roots biological replicate 1	GSM831910	Nod-LCOs 24h roots biological replicate 1
GSM831908	Nod-LCOs 6h roots biological replicate 2	GSM831911	Nod-LCOs 24h roots biological replicate 2
GSM831909	Nod-LCOs 6h roots biological replicate 3	GSM831912	Nod-LCOs 24h roots biological replicate 3
GSM831913	Nod-LCO control 6h roots biological replicate 1	GSM831916	Nod-LCO control 24h roots biological replicate 1
GSM831914	Nod-LCO control 6h roots biological replicate 2	GSM831917	Nod-LCO control 24h roots biological replicate 2
GSM831915	Nod-LCO control 6h roots biological replicate 3	GSM831918	Nod-LCO control 24h roots biological replicate 3

Microarray data were normalized in Expression console, and were analyzed in FlexyArray. To obtain co-over expressed probsets in the experiment, firstly, Bayesian t-test analysis were run to find the probsets which over-expressed in Nod-LCOs 6h compare to Nod-LCO control (comparison 1). The same analysis was run on Nod-LCOs 24h compared to Nod-LCO control 24h (comparison 2). Over expressed data were imported to Excel, and co-over expressed probsets between two comparisons were selected to further analysis in Plexdb.

Supplementary Table 2. List of co-over expressed genes encoded DNA binding proteins in *Medicago truncatula*/bacterium interaction.

Up regulated DNA binding genes in <i>R. solanacearum</i> - <i>M. truncatula</i> interaction		Up regulated DNA binding genes in <i>S. meliloti</i> - <i>M. truncatula</i> interaction	
Prob set	TAIR annotation	Prob set	TAIR annotation
Mtr.32159.1.S1_at; Mtr.51829.1.S1_at; Mtr.51829.1.S1_s_at	response regulator 5	Mtr.32159.1.S1_at; Mtr.51829.1.S1_S-at	response regulator 5
Mtr.5343.1.S1_s_at	response regulator 7	Mtr.5343.1.S1_at	response regulator 7
Mtr.174.1.S1_at	response regulator 8	Mtr.174.1.S1_at	response regulator 8
Mtr.12619.1.S1_at	response regulator 9	Mtr.17273.1.S1_at; Mtr.33808.1.S1_at	response regulator 9
Mtr.5335.1.S1_at	response regulator 15	Msa.3058.1.S1_at	Ethylene-responsive transcription factor 9
Mtr.19570.1.S1_at	ethylene response factor 1	Mtr.19570.1.S1_at	Ethylene-responsive transcription factor 1B
Mtr.10822.1.S1_at	splicing factor 33	Mtr.35480.1.S1_at	Ethylene-responsive transcription factor RAP2-11
Mtr.15332.1.S1_at	MYB domain protein 14	Mtr.22332.1.S1_at	Transcription factor MYB1R1
Mtr.1885.1.S1_at	basic helix-loop-helix (bHLH) DNA-binding superfamily protein (bHLH18)	Mtr.39233.1.S1_at	Auxin response factor 16
Mtr.16587.1.S1_at	basic helix-loop-helix (bHLH) DNA-binding superfamily protein (bHLH25)	Mtr.45719.1.S1_at ; Mtr.8357.1.S1_at	Transcription factor bHLH30
Mtr.37926.1.S1_a; Mtr.45719.1.S1_at	basic helix-loop-helix (bHLH) DNA-binding superfamily protein	Mtr.32254.1.S1_at	Zinc finger protein 6

Up regulated DNA binding genes in <i>R. solanacearum</i> / <i>M. truncatula</i> interaction		Up regulated DNA binding genes in <i>S.meliloti</i> / <i>M. truncatula</i> interaction	
Prob set	TAIR annotation	Prob set	TAIR annotation
Mtr.37926.1.S1_at; Mtr.45719.1.S1_at	basic helix-loop-helix (bHLH) DNA-binding superfamily protein	Mtr.32254.1.S1_at	Zinc finger protein 6
Mtr.41077.1.S1_at	zinc finger protein 5	Mtr.38349.1.S1_at	NAC domain-containing protein 42
Mtr.32254.1.S1_at	zinc finger protein 6	Mtr.43750.1.S1_at	Nuclear transcription factor Y subunit A-10
Mtr.38581.1.S1_at	zinc finger (AN1-like) family protein	Mtr.4282.1.S1_at	Nuclear transcription factor Y subunit B-3
Mtr.18592.1.S1_at	C2H2-type zinc finger family protein		
Mtr.44379.1.S1_at; Mtr.52180.1.S1_at	C2H2 and C2HC zinc fingers superfamily protein		
Mtr.1814.1.S1_at	RING-H2 zinc finger protein RHA4a		
Mtr.35677.1.S1_at; Mtr.44939.1.S1_at	RING-H2 finger A2A		
Mtr.48864.1.S1_at	Dof-type zinc finger DNA-binding family protein		
Mtr.12511.1.S1_at	heat shock transcription factor B3		
Mtr.39929.1.S1_at; Mtr.44462.1.S1_at	17.6 kDa class II heat shock protein		
Mtr.44719.1.S1_at	heat shock transcription factor A2		
Mtr.11053.1.S1_at; Mtr.29902.1.S1_at;	WRKY DNA-binding protein 33		
Mtr.32279.1.S1_at; Mtr.52252.1.S1_at			

Up regulated DNA binding genes in *R. solanacearum*/*M. truncatula* interaction

Prob set	TAIR annotation
Mtr.48109.1.S1_at	WRKY DNA-binding protein 40
Mtr.32312.1.S1_at	WRKY DNA-binding protein 50
Mtr.24228.1.S1_at	NmrA-like negative transcriptional regulator family protein
Mtr.11007.1.S1_at; Mtr.1814.1.S1_at; Mtr.42240.1.S1_at;	RING/U-box superfamily protein
Mtr.43917.1.S1_at; Mtr.8934.1.S1_at	
Mtr.35248.1.S1_at	Integrase-type DNA-binding superfamily protein
Mtr.40776.1.S1_at	SC35-like splicing factor 33

Supplementary Table 3. List of up regulated Expansin genes in *M. truncatula*-bacterium interaction.

Up regulated genes encoded Expansins in <i>R.</i>		Up regulated genes encoded Expansins in <i>S.</i>	
<i>solanacearum-M. truncatula</i> interaction		<i>meliloti-M. truncatula</i> symbiosis	
Prob set	TAIR annotation	Prob set	TAIR annotation
Mtr.19007.1.S1_at;	expansin A10	Mtr.41561.1.S1_at	expansin A4
Mtr.8533.1.S1_s_at			
Msa.2686.1.S1_at;	expansin A15		
Mtr.8532.1.S1_at;		Mtr.31597.1.S1_at	expansin A7
Mtr.8532.1.S1_s_at			
		Mtr.8533.1.S1_s_at	expansin A10
		Msa.2686.1.S1_at;	expansin A15
		Mtr.8532.1.S1_at;	
		Mtr.8532.1.S1_s_at	
		Mtr.37590.1.S1_s_at	expansin A8
		Mtr.31598.1.S1_at	expansin A18
		Mtr.33549.1.S1_at	expansin B2