

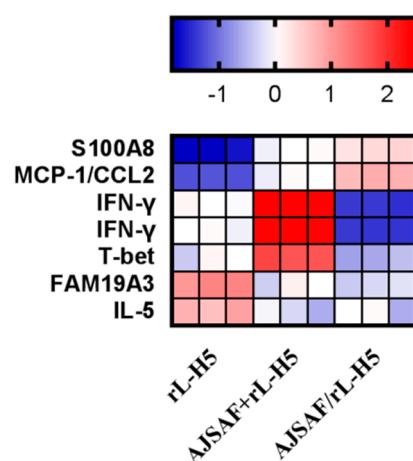
*Supplementary Materials*

# Mechanisms of mixed Th1/Th2 responses in mice induced by *Albizia julibrissin* saponin active fraction by *in silico* analysis

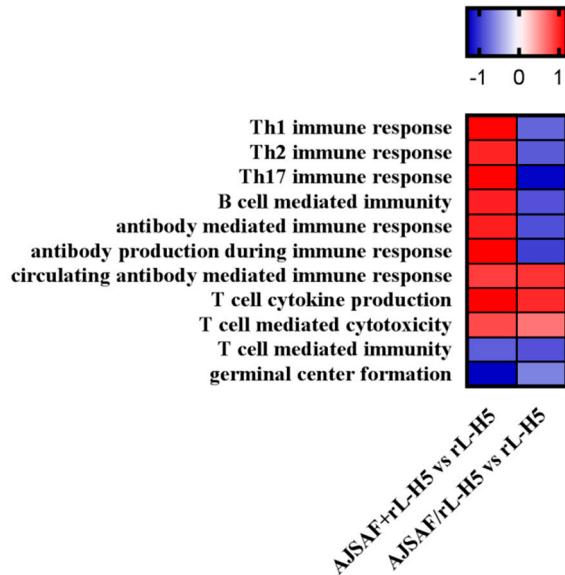
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**Table S1.** Primers used for qRT-PCR.

Gene	EntrezGeneID	Accession	Primer sequence	Size/bp
GAPDH	14433	NM_008084.3	5'-AGCCTCGTCCCGTAGACAA-3' 5'-AATCTCCACTTGCCACTGC-3'	104
S100A8	20201	NM_013650	5'-GGAGTTCCTGCGATGGTGA-3' 5'-GGCCAGAAGCTCTGCTACTC-3'	78
MCP-1/CCL2	20296	NM_011333	5'-GTCTGTGCTGACCCAAGAACG-3' 5'-TGGTTCCGATCCAGGTTTTA-3'	62
IFN- $\gamma$	15978	NM_008337	5'-TCTTGAAGACAATCAGGCCATCA-3' 5'-GAATCAGCAGCCACTCCTTTCC-3'	233
T-bet	57765	NM_019507	5'-ATTGGTTGGAGAGGAAGCGG-3' 5'-GCACCAGGTTCTGACTGTA-3'	129
FAM19A3	329731	NM_183224	5'-AGTAGCGAGAACGACTCCCA-3' 5'-TCCGGCTTAGAGCACCTAGA-3'	132
IL-5	16191	NM_010558	5'-GCTGGCCTCAAACCTGGTAATGTA-3' 5'-GGCAATGGTGCATGTCTGTAACCTC-3'	100



**Figure S1.** Heatmap of verified genes in microarray analysis.



**Figure S2.** Heatmap of gene sets on “adaptive immune response (GO: 0002250)” using gene set enrichment analysis (GSEA).



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