

Table S7. Enriched DEGs related to PRR, IRF, and ISG

	gene	protein	Expression change fold (log2FC)			
			3 h	12 h	24 h	36 h
	dhx58	Probable ATP-dependent RNA helicase DHX58 (LGP2)			2.37	4.42
PRR	ifih1	Interferon-induced helicase C domain-containing protein 1 (MDA5)			1.06	2.05
	nlr5	NOD-like receptor C5			3.28	4.61
	ptx3a	Pentraxin-related protein PTX3			2.32	2.72
	irf1b	Interferon regulatory factor 1		1.60	2.29	2.46
	irf3	Interferon regulatory factor 3		1.22	2.03	2.89
IRF	irf7	Interferon regulatory factor 7				1.57
	irf9	Interferon regulatory factor 9				1.11
	Irf10	Interferon regulatory factor 10				2.01
	LOC122871629	E3 ubiquitin/ISG15 ligase TRIM25			4.64	
	spock2	Testican-2 (IFN induced)			4.02	
	LOC122868453	Interferon-induced protein 44			3.7	
	LOC122878108	Interferon-induced protein 44			3.00	5.26
	LOC122878107	Interferon-induced protein 44				4.76
	rsad2	Radical S-adenosyl methionine domain-containing protein 2 (IFN induced) (Viperin)			3.32	4.07
ISG	LOC122862285	Interferon-induced GTP-binding protein Mx			3.27	
	LOC122878212	interferon-induced protein with tetratricopeptide repeats 1-like			2.48	5.19
	LOC122868174	Interferon-induced protein with tetratricopeptide repeats 5-like				1.35
	ifi35	Interferon-induced 35 kDa protein homolog				2.30
	LOC122869735	Erythroid membrane-associated protein E3 ubiquitin-protein ligase TRIM21			4.27	6.87