

geneID	Gene.name	logFC	FDR	Gene.description
23687956	NA	3.592568	3.507680e−11	NA
110676293	LYSC11	2.912345	1.348776e−10	C-Type Lysozyme (Lys-A). [Source:VB Community Annotation]
5563663	CLIPB35	2.894428	2.497743e−10	Clip-Domain Serine Protease family B. [Source:VB Community Annotation]
5564288	CTLMA14	4.068112	9.294342e−09	C-Type Lectin (CTL) – mannose binding. [Source:VB Community Annotation]
5565977	CLIPB46	5.052688	1.500756e−08	Clip-Domain Serine Protease family B. [Source:VB Community Annotation]
5564201	CLIPB15	3.090297	3.158017e−06	Clip-Domain Serine Protease family B. [Source:VB Community Annotation]
5574112		3.420640	5.838037e−05	GTP cyclohydrolase i [Source:VB Community Annotation]
5575350		6.377948	6.413055e−05	serine protease [Source:VB Community Annotation]
110680407	NA	3.546184	7.073212e−05	NA
5574170		4.205679	7.375455e−05	serine protease [Source:VB Community Annotation]
5566832	SRPN3	7.019244	1.660931e−04	Serine Protease Inhibitor (serpin) likely cleavage at T/I. [Source:VB Community Annotation]
5566117	NA	7.533905	4.658121e−04	NA
5578692		2.289610	5.054028e−04	Clip-domain serine protease [Source:UniProtKB/TrEMBL;Acc:Q1HQL3]
5570330	NA	2.469965	6.114231e−04	NA
5572333		6.199043	6.334413e−04	clip-domain serine protease, putative [Source:VB Community Annotation]
5563725		4.673286	1.503534e−03	serine protease inhibitor, serpin [Source:VB Community Annotation]
5572968	RpS2	2.348363	2.738978e−03	40S ribosomal protein S2 [Source:UniProtKB/TrEMBL;Acc:Q1HRV1]
5570814	NA	5.454934	2.882210e−03	NA
110678604		3.234094	3.746354e−03	
5569658	CLIPD1	4.807636	6.394790e−03	Clip-Domain Serine Protease family D [Source:VB Community Annotation]
5574952		4.804509	6.579870e−03	metalloproteinase, putative [Source:VB Community Annotation]
5576150		3.549113	7.787932e−03	lipase 1 precursor [Source:VB Community Annotation]
110676173	NA	5.887140	7.787932e−03	NA
5572428		5.111612	7.787932e−03	macroglobulin/complement [Source:VB Community Annotation]
5563616	CLIPB34	3.394840	7.904755e−03	Clip-Domain Serine Protease family B. [Source:VB Community Annotation]
5569420	GNBPA1	6.984448	8.711446e−03	Gram-Negative Binding Protein (GNBP) or Beta-1 3-Glucan Binding Protein (BGBP). [Source:VB Community Annotation]
5574109	NA	4.330156	1.031100e−02	NA
5572603	NA	7.104501	1.053915e−02	NA
5579417	Tf1	6.461632	1.120992e−02	transferrin [Source:VB Community Annotation]
110676739	NA	5.478952	2.948212e−02	NA
5579377	NA	6.551186	3.356480e−02	NA
5571998	PGRPS1	4.875659	3.356480e−02	Peptidoglycan Recognition Protein (Short) [Source:VB Community Annotation]
5577757	NA	4.170598	3.535674e−02	NA
5575056	CTLMA13	4.309369	4.106625e−02	C-Type Lectin (CTL) – mannose binding. [Source:VB Community Annotation]
5565454		2.370246	4.121954e−02	
5577410	NA	2.695940	4.275048e−02	NA
5568791		3.603029	4.586390e−02	