

geneID	Gene.name	logFC	FDR	Gene.description
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]
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]
5578692		2.289610	5.054028e−04	Clip−domain serine protease [Source:UniProtKB/TrEMBL;Acc:Q1HQL3]
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]
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]
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]
5579417	Tf1	6.461632	1.120992e−02	transferrin [Source:VB Community Annotation]
5571998	PGRPS1	4.875659	3.356480e−02	Peptidoglycan Recognition Protein (Short) [Source:VB Community Annotation]
5575056	CTLMA13	4.309369	4.106625e−02	C−Type Lectin (CTL) − mannose binding. [Source:VB Community Annotation]