

geneID	Gene.name	logFC	FDR	Gene.description
5563663	CLIPB35	3.056835	1.124804e−12	Clip–Domain Serine Protease family B. [Source:VB Community Annotation]
5564201	CLIPB15	4.138850	5.369934e−11	Clip–Domain Serine Protease family B. [Source:VB Community Annotation]
23687956	NA	3.405927	5.808029e−11	NA
5564288	CTLMA14	4.008050	4.204730e−09	C–Type Lectin (CTL) – mannose binding. [Source:VB Community Annotation]
5578692		2.584706	3.666582e−07	Clip–domain serine protease [Source:UniProtKB/TrEMBL;Acc:Q1HQI3]
5570115		7.103515	5.128507e−06	trypsin–eta, putative [Source:VB Community Annotation]
5563616	CLIPB34	5.524026	5.206469e−06	Clip–Domain Serine Protease family B. [Source:VB Community Annotation]
5574170		5.399507	1.024645e−05	serine protease [Source:VB Community Annotation]
5575350		6.447690	3.502794e−05	serine protease [Source:VB Community Annotation]
5565977	CLIPB46	3.613238	4.703270e−05	Clip–Domain Serine Protease family B. [Source:VB Community Annotation]
5573138	NA	3.633692	6.897477e−05	NA
5572333		6.148834	6.897477e−05	clip–domain serine protease, putative [Source:VB Community Annotation]
5568624		3.362300	8.124267e−05	
5569417		2.761228	1.370572e−04	
110674010	NA	2.737801	3.415007e−04	NA
5575395		3.028820	7.532299e−04	prohibitin, putative [Source:VB Community Annotation]
5568791		4.918363	9.053252e−04	
5574109	NA	7.234961	1.592778e−03	NA
5567561	CLIPB42	3.234920	2.091499e−03	Clip–Domain Serine Protease family B. [Source:VB Community Annotation]
5575054		4.590308	2.091499e−03	serine protease, putative [Source:VB Community Annotation]
5575056	CTLMA13	6.063663	2.318155e−03	C–Type Lectin (CTL) – mannose binding. [Source:VB Community Annotation]
5563566	CLIPB1	2.942619	2.366135e−03	Clip–Domain Serine Protease family B. [Source:VB Community Annotation]
5578083		2.710441	4.111619e−03	F–spondin [Source:VB Community Annotation]
5580206	NA	4.688291	4.111619e−03	NA
5576866	NA	2.978338	4.780044e−03	NA
5576674		3.552481	5.215740e−03	ATP–binding cassette sub–family A member 3, putative [Source:VB Community Annotation]
110676129	NA	6.889119	5.499543e−03	NA
5578648		3.625562	6.456572e−03	
110680407	NA	2.951290	6.456572e−03	NA
5569658	CLIPD1	4.901962	6.456572e−03	Clip–Domain Serine Protease family D [Source:VB Community Annotation]
5570931	CLIPB22	4.088559	7.345263e−03	Clip–Domain Serine Protease family B. [Source:VB Community Annotation]
5578380		2.238029	7.469922e−03	bm–40 precursor [Source:VB Community Annotation]
5567077	CLIFE8	4.873646	7.905326e−03	Clip–Domain Serine Protease family E. Protease homologue. [Source:VB Community Annotation]
110676221	NA	7.573899	8.013958e−03	NA
5571998	PGRPS1	3.944134	1.065119e−02	Peptidoglycan Recognition Protein (Short) [Source:VB Community Annotation]
5570984	NA	7.023570	1.065119e−02	NA
5565795		3.036948	1.182498e−02	
5566117	NA	5.201916	1.195026e−02	NA
23687765	NA	6.809386	1.307059e−02	NA
5573598	CLIPD6	3.269684	1.437808e−02	Clip–Domain Serine Protease family D [Source:VB Community Annotation]
5577757	NA	4.789861	1.744637e−02	NA
5579417	Tf1	4.804349	1.744637e−02	transferrin [Source:VB Community Annotation]
5569420	GNBPA1	6.633641	1.868872e−02	Gram–Negative Binding Protein (GNBP) or Beta–1 3–Glucan Binding Protein (BGBP). [Source:VB Community Annotation]
5570814	NA	5.049566	2.199206e−02	NA
5567079		4.958922	2.627590e−02	
5572428		3.995275	3.557840e−02	macroglobulin/complement [Source:VB Community Annotation]
5565454		2.341725	3.725316e−02	
5575325		3.517778	3.993694e−02	
5564141		3.213174	4.922646e−02	Niemann–Pick Type C–2, putative [Source:VB Community Annotation]
5579873		3.020483	4.935718e−02	