

Table S1. Transcripts that vary significantly in BMMs exposed to Pam3CSK4 or *L. pneumophila* treated with IFN- γ vs. IFN- γ + 2DG.

Location	PAM + IFN- γ FPKM	L.p. + IFN- γ FPKM	PAM + IFN- γ + 2DG FPKM	L.p. + IFN- γ + 2DG FPKM	log ₂ fold change PAM #DIV/0!	log ₂ fold change L.p.	Gene
chr3:121723536-121735052	8.64	40.80	0.00	0.00	5.34	2.93	F3
chr10:24914852-24927760	330.60	207.90	2.82	2.82	1.73	6.87	Arg1
chr10:99263230-99267489	51.60	57.56	1.67	1.67	2.19	4.95	Dusp6
chr11:9117979-9136170	378.41	400.15	22.16	22.16	16.15	4.09	Upp1
chr8:65618039-66473349	65.13	33.38	4.18	4.18	4.00	3.96	March1
chr12:119158487-119238276	19.50	12.96	1.64	1.64	2.44	3.57	Itgb8
chr13:60895350-60897447	98.59	132.14	9.22	9.22	13.67	3.42	Ctla2b
chr2:128698860-128803160	22.88	25.86	2.16	2.16	3.90	3.40	Mertk
chr13:117220572-117274415	81.92	73.16	7.79	7.79	12.61	3.39	Emb
chr5:43868826-43912389	208.64	107.66	20.31	20.31	14.14	3.36	Cd38
chr14:60634728-60764556	26.37	27.59	2.65	2.65	5.08	3.32	Spata13
chr9:124102182-124109140	15.99	25.65	1.63	1.63	4.28	3.29	Ccr2
chr16:10782304-10785536	115.02	113.26	12.45	12.45	18.68	3.21	Socs1
chr7:43526933-43533175	30.95	22.50	3.38	3.38	1.66	3.20	Cd33
chr12:32173396-32208649	25.18	16.27	3.00	3.00	2.38	3.07	Pik3cg
chr4:132732965-132757171	147.14	106.91	17.72	17.72	14.48	3.05	Smpd13b
chr5:123863569-123865516	79.28	95.76	9.69	9.69	20.41	3.03	Niacr1
chr10:56377299-56390419	76.79	35.79	9.43	9.43	5.28	3.03	Gja1
chr13:43785106-43803132	37.04	157.83	4.70	4.70	98.43	2.98	Cd83
chr6:29272487-29276390	127.10	143.20	16.23	16.23	38.78	2.97	Hilpda
chr19:40659646-40742515	39.22	47.47	5.09	5.09	11.21	2.94	Entpd1
chr10:10335702-10472314	55.22	30.34	7.18	7.18	4.46	2.94	Adgb
chr11:62248983-62266580	123.70	155.86	16.22	16.22	29.61	2.93	Adora2b
chr5:43818892-43843468	71.44	57.94	9.40	9.40	9.24	2.93	Bst1
chr9:7272513-7283333	352.87	137.78	47.51	47.51	22.29	2.89	Mmp13
chr11:117965741-117969830	262.49	306.84	35.50	35.50	94.93	2.89	Socs3
chr6:123227866-123247024	151.03	133.87	21.24	21.24	26.78	2.83	Clec4n
chr7:3485746-3502752	159.87	96.29	23.43	23.43	23.92	2.77	Tarm1
chr13:97241104-97253040	17.85	23.35	2.74	2.74	5.14	2.71	Enc1
chr11:83116844-83122659	109.82	148.71	17.36	17.36	26.50	2.66	Sfn1
chr17:17887823-17893952	569.21	405.70	90.54	90.54	49.52	2.65	Fpr2
chr17:17875768-17883951	233.86	190.30	38.79	38.79	25.60	2.59	Fpr1
chr12:31958478-32061279	31.62	24.11	5.41	5.41	3.49	2.55	Prkar2b
chr12:85473900-85477478	109.40	181.78	19.23	19.23	57.55	2.51	Fos
chr2:117279992-117342877	28.14	21.39	5.01	5.01	5.37	2.49	Rasgrp1
chr18:60393061-60443899	38.13	72.99	6.80	6.80	12.16	2.49	EMBL-EBI AK149718.1
chr15:74979722-75048837	203.51	318.72	37.53	37.53	87.21	2.44	Ly6a
chr9:118606708-118901003	18.55	6.20	3.48	3.48	2.30	2.42	Itga9
chr2:164948238-164955846	40.93	31.23	7.80	7.80	7.27	2.39	Mmp9
chr16:75858793-75909273	96.32	139.27	18.98	18.98	32.44	2.34	Samsn1
chr7:38183216-38197565	26.84	48.18	5.34	5.34	5.68	2.33	RIKEN cDNA 1600014C10
chr4:132781761-132796364	112.02	119.12	22.44	22.44	31.54	2.32	Themis2
chr19:21778341-21858450	48.39	42.18	9.98	9.98	6.90	2.28	Cemp2
chr9:41012959-41157668	29.20	18.98	6.47	6.47	6.32	2.17	Ubash3b
chr5:137787801-137858049	197.74	153.77	44.00	44.00	22.16	2.17	Zcwpw1
chr12:80107759-80113013	16.27	22.12	3.87	3.87	2.84	2.07	Zfp361i
chr8:92855349-92919279	75.81	97.27	18.55	18.55	17.46	2.03	Lpcat2
chr11:48985328-48992246	26.32	69.92	6.72	6.72	13.40	1.97	Tgtp1
chr11:44454570-44470548	34.39	40.38	9.35	9.35	8.33	1.88	Ublcp1
chr7:144415253-144538029	9.19	16.69	2.59	2.59	2.62	1.83	Pde3b
chr3:27317076-27339665	5.40	16.81	1.74	1.74	2.59	1.64	Tnfrsf10
chr17:44096293-44105808	8.18	14.92	2.71	2.71	2.14	1.60	Enpp4
chr8:45395664-45410539	15.12	33.76	5.17	5.17	6.57	1.55	Tlr3
chr11:83175185-83190251	71.33	191.83	26.41	26.41	24.21	1.43	Sfn4
chr19:21391306-21472661	11.06	32.62	6.57	6.57	6.32	0.75	Gda
chr6:7675170-7693182	73.21	24.00	165.25	165.25	132.37	-1.17	Asns
chr4:59805649-59904832	5.90	2.86	17.84	17.84	14.42	-1.60	Snx30
chrX:36328408-36366856	9.53	4.48	30.18	30.18	22.38	-1.66	Lonrf3
chr3:108424774-108445259	20.26	12.28	70.09	70.09	65.73	-1.79	Sars
chr3:100468062-100489192	3.45	3.65	14.71	14.71	26.76	-2.09	Fam46c
chr10:127514938-127522444	5.23	4.16	22.87	22.87	29.79	-2.13	Shmt2
chr16:22857844-22879634	39.83	50.36	195.50	195.50	234.30	-2.30	Dnajb11
chr12:116405401-116463531	1.97	2.87	9.82	9.82	14.79	-2.32	Ncapg2
chr7:132557474-132576398	24.59	26.95	127.96	127.96	123.99	-2.38	Oat
chr11:114922780-114934386	9.25	13.23	53.38	53.38	47.90	-2.53	-1.86 Cd300lb
chr2:121413901-121438686	24.63	23.63	149.03	149.03	124.87	-2.60	Pdia3
chr12:91805907-91849157	26.09	27.07	159.39	159.39	150.35	-2.61	Sel1l
chr7:45522738-45526268	8.36	35.02	51.65	51.65	77.48	-2.63	-1.15 Ppp1r15a
chr14:118937931-118981702	12.60	14.45	82.25	82.25	73.67	-2.71	Dnajc3
chr13:95627176-95891922	3.81	7.23	27.27	27.27	22.09	-2.84	Iggap2
chr2:152337424-152344060	13.31	5.55	106.51	106.51	83.01	-3.00	Trib3
chr18:65800577-65817657	16.65	48.02	140.92	140.92	196.46	-3.08	-2.03 Sec11c
chr13:110395043-110400843	21.74	43.21	195.08	195.08	288.27	-3.17	Plk2
chr12:17266594-17324730	61.82	47.19	606.19	606.19	664.54	-3.29	-3.82 Pdia6
chr14:103814624-103844524	2.83	2.98	27.82	27.82	24.59	-3.30	-3.04 Ednrb
chr9:106887414-106891938	19.43	19.31	201.31	201.31	236.22	-3.37	-3.61 Manf
chr6:47796140-47813512	31.57	35.74	330.67	330.67	344.11	-3.39	-3.27 Pdia4
chr9:44379489-44392576	24.95	23.65	269.93	269.93	281.33	-3.44	-3.57 Hyou1
chr16:92391952-92466146	5.64	13.03	63.96	63.96	77.68	-3.50	-2.58 Rcan1
chr2:84936608-84958509	3.33	2.78	41.74	41.74	39.64	-3.65	-3.84 Slc43a3
chr12:83987860-83993875	2.60	1.65	33.74	33.74	23.37	-3.70	-3.83 Acot2
chr12:44205896-44210068	17.95	25.83	259.96	259.96	223.40	-3.86	-3.11 Dnajb9
chr10:127290792-127311786	14.13	14.59	217.63	217.63	284.18	-3.94	-4.28 Ddit3
chr15:88819645-88826718	10.05	18.32	184.55	184.55	257.10	-4.20	-3.81 Creld2
chr16:17130137-17132383	15.35	11.96	331.95	331.95	311.42	-4.43	-4.70 Sdf211
chr8:94386499-94395358	6.33	3.36	225.67	225.67	175.88	-5.16	-5.71 Herpud1