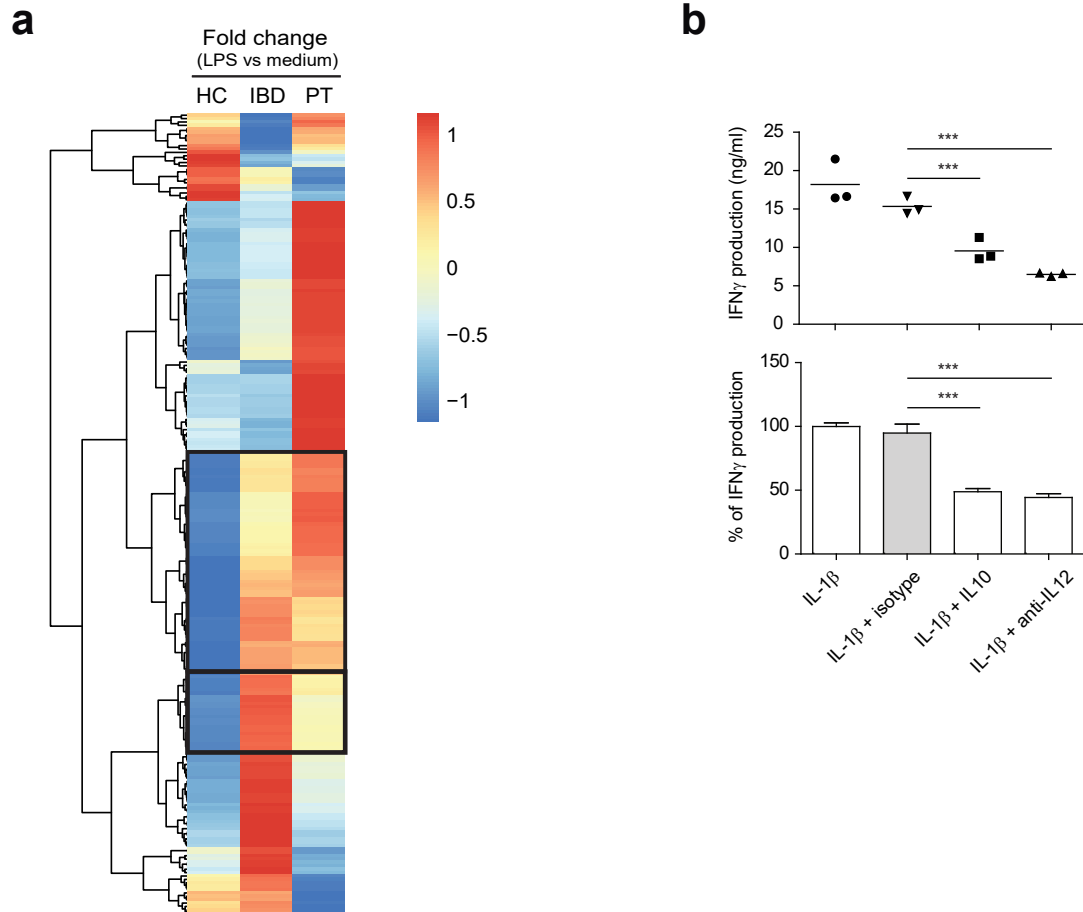


Supplementary Figure S1. IL-10 cannot inhibit IFN γ secretion by CD4 $^+$ T cells in a direct manner.

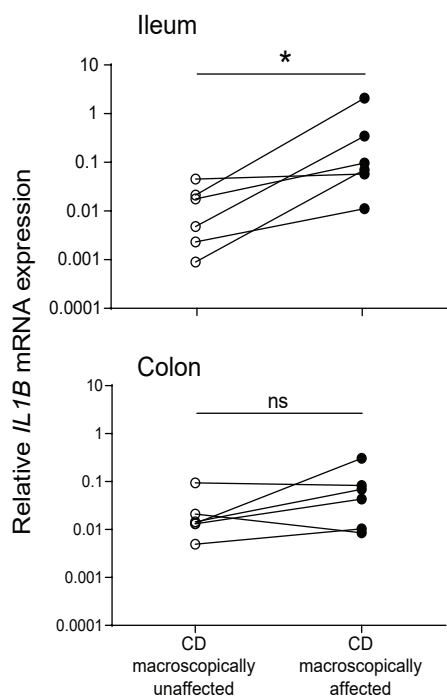
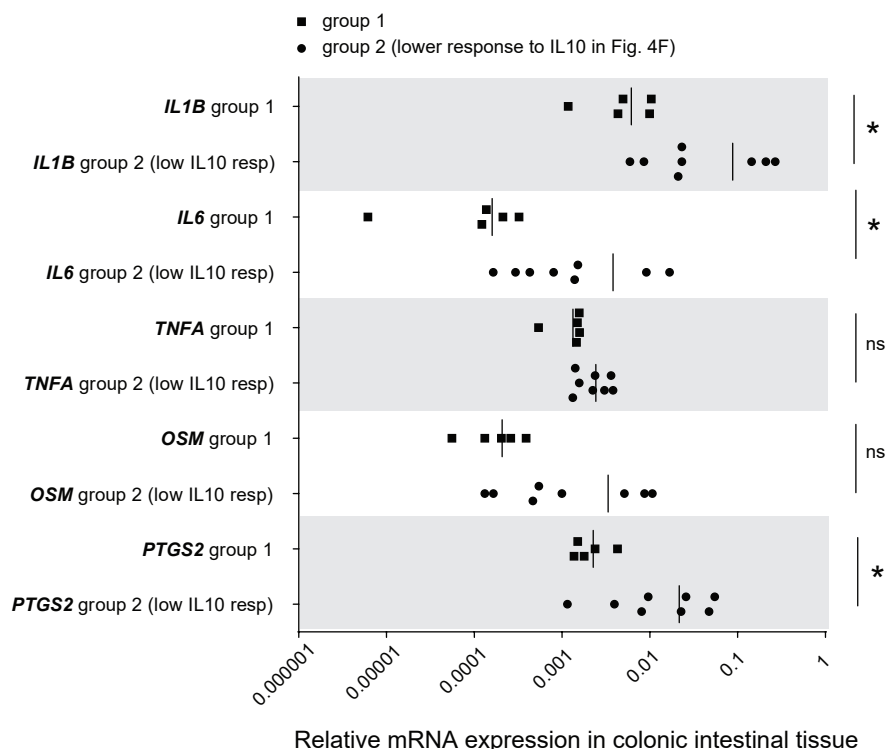
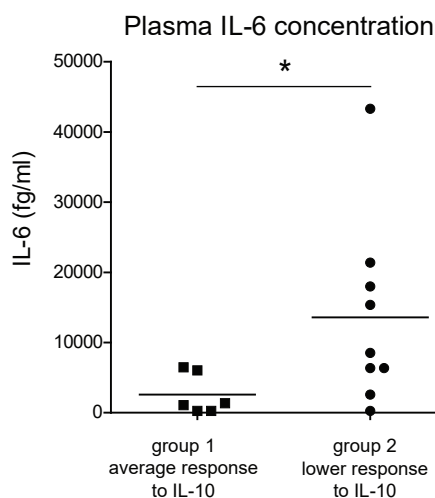
(a, b) CD4 $^+$ T cells from adult healthy controls (n=4) were stimulated with (a) anti-CD3/CD28 beads (bead-to-cell ratio 1:4) or (b) plate-bound anti-CD3 antibody (0.1, 1, 10 μ g/ml) in the presence or absence of IL-10. After 24 h or 48 h, supernatants were assayed for IFN γ using an ELISA. (c) Monocyte-derived DCs from an *IL10RA*-deficient patient or healthy controls were preincubated with LPS or LPS/IL-10 for 24 h. Allogeneic MLRs were performed by co-culturing moDCs and CD4 $^+$ T cells with or without IL-10. After 72 h, supernatants were assayed for IFN γ using an ELISA. (d) Representative dotplots of data shown in Fig.1e. CD14 $^+$ monocytes and T cells were analyzed by flow cytometry for expression of IL-10RA and IL-10RB chains (gated on singlets).

Results are mean \pm SEM (a, b) or mean \pm SD (c) of a representative of at least 2 independent experiments. ***P<0.001 using unpaired Student's t test.



Supplementary Figure S2. (a) Gene expression profile of dendritic cells from pediatric IBD patients partially overlaps with *IL10RA*-deficient dendritic cells. Heat map of differentially expressed genes showing color-coded fold change between LPS stimulated and unstimulated moDCs from healthy controls (n=3), *IL10RA*-deficient patient, and pediatric IBD patients (n=3).

(b) IL-1 β stimulates IFN γ release by human CD4 $^+$ T cells indirectly through induction of IL-12 production (data related to Figure 3 of the manuscript). Allogeneic MLRs were performed using moDCs and CD4 $^+$ T cells from healthy individuals. SEB was added in the presence of IL-1 β , with or without IL-10, neutralizing IL-12 antibodies, or isotype control. After 72 h, supernatants were assayed for IFN γ using an ELISA. Top graph depicts 1 representative experiment of n=3. Bottom depicts combined data of n=3 experiments where IFN γ production is expressed as percentage relative to the IL-1 β condition (100%) in each experiment.

a**b****c****d**

group 1: average IL-10 responder	age at diagnosis	diagnosis	Paris classification (Location);(behavior)	exposed to anti-TNF/biological 1=yes/0=no	exposed to other medication	resection
1-female	14	CD	L1+L4a; B1 + p	1; IFX	aza; 6-TG	0
2-female	16	CD	L2; B1	0	aza	0
3-female	18	CD	L3; B1	0	aza	0
4-male	15	CD	L1; B1	0	aza; mesa	0
5-female	13	UC	E4; S0	0	mesa	0
6-male	13	UC	E1; S0	0	mesa; aza	0
group 2: low IL-10 responder	age at diagnosis	diagnosis	Paris classification (Location);(behavior)	exposed to anti-TNF/biological 1=yes/0=no	exposed to other medication	resection
1-male	8	CD	L3+L4a; B1 + p	1; IFX	aza	0
2-male	12	CD	L3+L4a; B1 + p	1; IFX; ADA	aza	0
3-male	10	UC->CD	L2; B1	1; IFX;	aza	0
4-female	13	CD	L1+L4a; B1 + p	0	budesonide	0
5-male	9	UC	E4; S1	1; IFX; ADA	aza;mesa	0
6-female	12	UC	E2; S0	1; IFX; VEDO	mesa; aza	0
7-male	13	IBDU	E4;S0	1; IFX		1
8-female	9	UC	E4; S0	0	mesa	0
9-female	13	UC	E4;S1	0	mesa; aza	0

UC, ulcerative colitis; CD, Crohn's disease; IFX, infliximab; ADA, adalimumab; VEDO, vedolizumab; aza, azathioprine; mesa, mesalazine; 6-TG, 6-thioguanine.

Paris Classification: *CD-Disease location*: (L1) involvement of the terminal ileum only, with limited or no cecal disease; (L2) colonic involvement only; (L3) involvement of both the terminal ileum and colon; (L4) isolated upper GI disease, defined as macroscopic and/or radiologic abnormalities proximal to the terminal ileum. Isolated upper GI disease (L4) is further separated into esophagogastrointestinal disease (L4a), jejunal/proximal ileal disease (L4b), or both L4a and L4b. Upper GI disease (i.e. L4a, L4b, L4ab) can also coexist with L1, L2, or L3.

CD-Disease behavior: is registered as the presence of strictures, intraabdominal fistulas, and/or intraabdominal abscesses and categorized disease according to the Paris classification: (B1) non-stricturing, non-penetrating disease; (B2) stricturing disease; (B3) penetrating disease (excluding isolated perianal or rectovaginal fistulas); (p) perianal disease. *UC-Disease location*: (E1) proctitis only; (E2) left-sided colitis, inflammation distal to the splenic flexure; (E3) extensive colitis, inflammation proximal to the splenic flexure but distal to the hepatic flexure; and (E4) pancolitis, defined as inflammation extending proximal to the hepatic flexure.

UC-Disease behavior: (S1) has experienced acute severe colitis; (S0) has not experienced acute severe colitis.

Supplementary Figure S3. Enhanced *IL1B* expression and suboptimal IL-10 function in pediatric IBD patients.

(a) *IL1B* mRNA expression was measured in biopsies from macroscopically affected intestine (closed symbol) and biopsies from unaffected regions (open symbol) derived from treatment-naive CD patients. Paired samples per individual patient are shown. (b) mRNA expression was measured in biopsies from group 1 and group 2 (lower response to IL-10) patients analyzed in Fig. 4f. (c) IL-6 concentrations in plasma of group 1 and group 2 (lower response to IL-10) patients analyzed in Fig. 4f. (d) Patient disease characteristics of group 1 and group 2 (lower response to IL-10) patients analyzed in Fig. 4f.

Supplemental references referred to in Methods:

51. Trapnell C, Pachter L, Salzberg SL. TopHat: discovering splice junctions with RNA-Seq. *Bioinformatics (Oxford, England)* 2009; **25**(9): 1105-1111.
52. Trapnell C, Williams BA, Pertea G, Mortazavi A, Kwan G, van Baren MJ *et al.* Transcript assembly and quantification by RNA-Seq reveals unannotated transcripts and isoform switching during cell differentiation. *Nature biotechnology* 2010; **28**(5): 511-515.

PT.LPS vs.PT.unstim

tracking_id	Healthy Controls						Inflammatory bowel disease patients						IL10RA-deficient patient		logFC (PT LPS/PT unstim)	logCPM (counts per million)	PValue	FDR
	HC1 unstim	HC1 LPS	HC2 unstim	HC2 LPS	HC3 unstim	HC3 LPS	IBD1 unstim	IBD1 LPS	IBD2 unstim	IBD2 LPS	IBD3 unstim	IBD3 LPS	PT unstim	PT LPS				
CMFK2	81.982	170.55	3.91011	231.745	2.91992	220.241	5.64159	241.614	5.79375	249.077	2.43557	241.427	1.73704	292.617	7.33933061	6.4568128	1.00474E-09	1.69315E-05
IL1B	1.93472	139.946	2.52815	551.64	1.08036	905.961	0.583326	535.264	0.208481	521.021	2.08817	1188.28	0.795252	1245.66	10.5986027	10.54995098	1.7774E-09	1.69315E-05
IL8	1.47831	174.614	9.29087	923.455	3.77777	1400.42	2.09274	552.801	2.30063	627.965	5.2537	561.494	7.59637	2218.94	10.87455864	10.87455864	3.83314E-09	2.2433E-05
IL2RA	2.02511	21.3285	2.00383	85.7318	0.602627	37.1544	0.497805	93.6372	0.310094	44.0758	0.305028	31.5818	0.0647704	68.1972	9.94161615	8.95259E-09	3.31152E-05	3.31152E-05
GCHI	1.06736	101.155	0.624818	107.441	1.58384	324.266	0.72337	254.959	0.521104	140.148	1.97782	250.509	1.4361	341.411	7.892843632	8.822916238	9.40048E-09	3.5819E-05
GBP5	4.31311	156.852	5.34422	241.239	4.12013	181.925	0.70541	510.597	3.24626	309.74	3.1358	326.974	3.46742	283.333	6.346923793	9.263179212	1.36594E-08	4.3373E-05
HERC5	2.459	132.591	1.46792	206.02	2.07946	255.035	3.15648	310.101	2.30222	194.984	0.883408	187.578	1.22088	204.096	7.612929321	8.809834134	1.69063E-08	4.6014E-05
IDO1	8.78675	417.533	2.61067	683.778	4.84584	727.883	2.79992	1337.41	2.56386	508.59	3.13196	973.858	2.5964	1360.92	9.028447746	6.26725E-05	6.26725E-05	6.26725E-05
CXCL10	0.933688	1128.07	1.24918	2666.28	0.721847	1942.87	1.29245	3334.76	0.682744	1454.95	3.54997	3776.3	1.25077	2719.47	11.07100718	10.59718287	8.16024E-08	0.00172743
CFB	1.68546	74.9856	1.86127	103.668	1.77782	140.199	2.41863	280.854	5.34691	344.682	3.40021	213.657	1.93193	128.173	6.046278238	7.466696491	1.03526E-07	0.00197237
DDX58	7.41079	716.037	6.76623	474.405	5.8043	500.557	8.17065	641.447	7.42127	401.027	2.89581	314.894	9.75315	575.472	5.880252492	10.54424841	1.14962E-07	0.00199115
MCOLN2	11.9283	72.8657	15.0701	175.784	14.3921	248.826	6.83454	231.336	5.56241	156.747	8.59947	154.488	8.9296	222.339	4.635592378	8.525316579	1.49962E-07	0.000234464
IFIT3	19.8136	1236.52	8.43754	1199.41	7.4065	1326.21	16.9743	1781.87	34.7479	2116.53	7.75625	1599.57	11.8134	1628.35	7.104927402	11.02122959	1.60594E-07	0.000234464
IL1B	15.9829	1367.71	36.3603	2575.98	21.8116	2868.16	9.11188	2301.12	15.6688	3082.94	5.45178	9642.2	14.3804	7633.21	9.050455838	12.44138693	1.72251E-07	0.000234464
IFI4L4	2.0609	75.9017	0.76707	153.714	0.429261	186.201	2.45213	188.886	1.60398	136.524	0.169206	83.474	0.499117	183.549	8.51352221	9.192044285	2.40323E-07	0.000273167
PDE4B	1.14035	14.0707	1.7322	40.1181	1.101	34.6159	0.790558	53.7205	1.94624	36.3354	1.30986	63.2904	2.1449	31.111	5.928073437	4.25187E-07	0.000273167	0.000273167
DDX80	3.11665	47.072	2.6048	73.7511	3.4851	72.6714	4.28868	113.708	1.61875	57.2071	1.47104	50.803	1.75422	68.7645	5.288729873	7.857818075	2.497E-07	0.000273167
EREG	0.675609	18.2128	1.00504	28.7771	0.714368	46.1869	1.00395	35.4144	0.541499	23.3	1.03693	0.860157	63.7557	7.452386106	6.334469218	7.452386106	2.58083E-07	0.000273167
DAS3	8.55228	167.954	5.69021	236.709	6.98077	185.543	9.98077	268.584	9.03897	231.594	11.4558	261.721	4.27714	197.523	5.526311549	9.504694648	2.8931E-07	0.000277744
PARP1	45.3236	411.363	39.4066	568.469	52.2405	720.549	42.6087	792.853	30.3352	563.173	22.8825	508.789	33.5102	764.927	4.510291045	11.71325463	3.14386E-07	0.000277744
RARL1	0.115158	66.0161	1.52818	151.97	0.331247	153.312	0.331247	79.3266	1.44533	208.596	0.136684	88.9808	0.572664	88.8519	7.186852164	4.102105099	3.199E-07	0.000277744
IL15RA	1.60472	88.575	1.86616	97.2408	0.901437	69.3594	0.643528	134.133	0.732031	140.576	3.30913	286.664	0.472716	188.298	8.63885914	7.287789434	3.21099E-07	0.000277744
CCL20	0.179329	41.4614	0.30599	160.445	1.65235	458.782	0.131773	164.391	0.127808	147.735	0.848409	343.977	0.99507	537.586	7.594977585	7.594977585	3.39571E-07	0.000277744
ELOVL7	1.23782	12.7537	0.752417	17.4175	0.5289	34.5079	0.834508	31.7419	0.895871	21.9939	1.6631	33.3113	0.941681	40.7774	5.433057659	6.410556577	3.98977E-07	0.000277744
HERC6	2.66324	46.5265	2.15795	78.029	2.92419	83.7858	3.34351	96.156	2.82091	83.3675	1.84529	57.7233	2.85088	90.3456	4.984264449	7.575341715	1.49962E-07	0.000277744
IFIT2	11.4341	815.788	3.82584	782.753	1.79597	917.48	9.60049	1141.25	8.26707	926.108	2.00618	817.981	3.49775	946.19	8.076047012	10.77545332	4.85752E-07	0.00036159
GBP4	1.94152	85.7592	1.13638	127.678	0.984902	86.7133	0.9856	197.12	0.43737	91.4466	1.06919	139.694	0.73918	95.9952	7.014293668	8.331384042	4.77042E-07	0.00036159
USP18	1.78187	112.536	0.973316	132.708	0.652027	129.67	1.42496	166.996	1.81715	193.234	0.239739	175.675	1.22324	184.359	7.225879736	7.577363015	5.3036E-07	0.000349173
CXCL1	0.0391699	18.4554	0.201502	141.925	0.540937	480.91	0.0288148	111.117	0.421178	286.745	0.75662	160.879	0.108196	10.3689502	9.113118413	5.31494E-07	0.000349173	0.000349173
RSAD2	25.4073	1071.31	13.9299	1587.34	10.9554	1874.77	11.6158	1663.95	21.1752	1812.66	4.54674	1099.71	14.9955	2115.13	7.137862706	11.948303024	5.96638E-07	0.000378905
OAS2	27.3949	548.81	16.0245	582.204	13.0076	372.482	3.4615	792.5	26.5642	638.148	13.3137	619.063	12.3442	10.61828	5.769809987	10.12591553	0.000530678	0.000530678
CYP7B1	0.256674	2.99174	0.0888434	14.7991	0.16588	12.8135	0.0887486	27.2972	0	5.31118	0.080078	15.4336	0	12.3627	10.96450912	3.926158073	1.16877E-06	0.000695854
MX2	7.21066	242.66	4.29574	265.482	2.69066	181.664	12.3587	376.606	9.72176	345.109	7.38359	376.93	2.42336	249.489	6.8812728	6.805897324	1.78763E-06	0.001026926
XAF1	17.9324	164.33	16.9228	235.479	16.8344	367.302	26.9311	312.958	31.2917	289.965	7.91662	208.329	16.109	375.288	4.570266136	9.52729265	1.83264E-06	0.001026926
CBP1	10.7061	358.933	10.117	483.139	16.5487	505.93	108.413	1030.34	1082.41	7.93361	382.391	13.287	780.278	959.551	5.392125723	5.392125723	2.31029E-06	0.001257589
IFIH1	25.649	422.74	27.96	585.984	37.6604	899.928	28.466	757.118	21.0448	543.537	22.849	559.676	57.6679	1258.23	4.445444726	11.29129818	2.44493E-06	0.00128426
SOD2	38.6141	1617.84	72.51	3248.93	45.2183	2230.54	58.1319	3267.71	54.8371	2754.39	211.108	3040.18	83.3227	2675.37	5.031152438	10.76030345	2.52159E-06	0.00128426
VEP1	0.0153544	0.612639	0.018718	1.31588	0.0383576	0.379083	0.0297081	1.08711	0.113695	0.708994	0.0340226	0.339854	0	0.955757	9.73986723	2.700577102	5.19511E-06	0.00128426
IFI4	13.0212	137.865	6.58444	316.729	9.80888	506.729	13.0768	326.332	7.50058	260.632	2.72034	232.781	10.5973	484.859	5.402456617	8.617752339	2.94347E-06	0.00137921
PARP9	16.0622	132.255	12.6896	173.792	17.5716	256.579	16.3398	225.615	13.8411	171.904	4.88264	148.65	17.7061	257.887	7.700693789	9.043786242	3.37001E-06	0.00137921
PLSCR1	2.3564	22.9885	1.00265	50.6654	1.86428	84.345	2.74222	70.5523	0.971906	53.2043	1.13802	55.3229	1.94415	59.1421	4.920844871	6.104059686	5.44995E-06	0.002532497
PTGS2	0.331051	33.4102	1.08206	128.75	0.604018	208.147	0.504792	118.037	0.12307	113.97	1.22155	368.772	1.96779	84.1261	8.740412441	10.99045567	6.69438E-06	0.002532497
NFKB1	41.8609	238.411	50.1332	335.146	40.3206	312.434	27.893	432.577	31.5264	326.146	50.7973	428.626	48.3476	528.765	5.449157654	10.2941414	6.4272E-06	0.002724517
DENN5A	12.1335	101.699	11.7905	103.958	13.3049	122.392	13.2442	187.724	7.96193	114.339	23.291	236.157	12.548	166.686	3.22851059	8.902046971	6.47429E-06	0.002724517
MX1	19.5717	743.35	13.0624	763.214	5.10095	548.274	32.7324	1025.72	27.8244	981.816	7.40385	797.968	10.4121	748.068	6.177705021	10.07942692	6.56471E-06	0.002724517
TRIM22	10.1125	70.3008	11.3631	118.247														

PT.LPS.vs.PT.unstim

UBE2L6	33.6059	275.257	30.3464	246.187	42.4375	246.226	35.2496	287.899	39.5067	419.165	42.1091	256.275	31.7841	305.846	3.26637065	7.649680867	3.01353E-05	0.007176722
NEURL3	0.511471	25.7904	0.683452	45.2833	0.495176	47.0774	0.0755936	42.3528	0.662499	118.287	0.299574	32.0688	0.430611	53.9957	6.93925364	5.491174567	3.18513E-05	0.007315292
NUS1	6.56626	348.984	54.7166	410.268	33.2516	265.326	35.8635	473.079	28.409	384.939	31.7423	316.323	28.5677	282.179	3.45749398	5.042163929	3.1962E-05	0.007315292
MARPK8	4.02680	17.0705	3.93225	25.0346	4.32118	46.7369	3.65224	38.7767	2.06738	1.5298	4.41152	67.9522	3.84857	56.2018	3.862097595	6.524296786	3.21085E-05	0.007315292
SLAMP1	1.02386	19.3963	2.31051	69.1317	2.80523	98.5511	0.463904	55.1629	1.4693	163.999	1.44312	41.1681	1.3562	87.3773	6.000617959	5.484327198	3.2253E-05	0.007315292
TRIM25	12.0094	97.9525	12.8833	110.504	15.4053	97.3677	13.32	118.444	15.0018	140.163	14.5559	135.549	12.7706	128.517	3.328649047	8.777350399	3.292E-05	0.007315292
LOC731424	2.95904	68.3344	2.20001	60.3014	7.48642	156.281	0.784315	80.4009	3.50507	210.44	4.91415	146.099	4.43961	147.354	5.04957699	8.083447373	3.39502E-05	0.00749211
TRDR7	14.9253	83.4482	13.9014	92.2454	18.1716	116.928	12.5785	117.014	8.15544	81.2553	6.7151	83.2499	12.4414	115.935	3.217825444	8.001756563	3.4232E-05	0.00749211
TNFAIP6	1.08152	179.489	2.32117	372.351	3.11728	1233.23	2.53052	1537.4	0.499604	406.533	14.1646	1458.35	6.22805	1612.59	8.013423905	10.11668205	3.46056E-05	0.00749211
IRF7	2.10088	36.1961	1.02149	27.827	0.804086	29.673	2.29487	30.8213	4.374	70.4026	2.55384	58.1796	1.42064	36.4738	4.673385739	5.160850089	3.6865E-05	0.007891601
KCNQ3	0.035528	0.044094	0.0494879	0.273907	0.0640964	0.231918	0.058157	0.24519	0.0850986	0.190267	0.0746679	0.247949	0.0279003	1.52006	5.701173427	3.186226215	3.87128E-05	0.008195078
STAT1	42.4661	323.136	27.5325	318.2	51.5636	345.03	41.1623	427.067	30.1108	308.469	27.5222	278.017	33.203	230.232	3.188084149	9.430661182	3.9311E-05	0.008230265
ADORA2A	0.158648	4.17911	0.508231	14.9245	0.0219748	5.20777	0.14111	20.0346	0.0515838	18.6572	0.278361	28.9604	0	11.7504	10.99941102	3.961072216	4.05205E-05	0.008391275
PNPT1	4.46643	27.6765	4.18386	53.0907	6.12252	88.8312	4.04137	59.6606	2.80279	49.3568	2.20526	38.8031	0.05144	68.1896	3.751938212	7.489109257	4.80646E-05	0.008465616
CD80	26.9651	145.991	30.1584	291.573	21.5036	235.288	22.9096	439.533	12.1852	208.459	5.04439	193.152	23.4174	333.586	8.30517439	8.998290865	4.88729E-05	0.00905592
IL1A	2.96888	39.1826	4.58958	144.276	5.84667	402.657	1.42723	150.072	0.745423	97.4968	3.8574	532.745	5.12021	806.106	7.29585077	10.27573197	5.15283E-05	0.010245092
BRIP1	1.42903	15.9447	1.30052	23.8855	1.42372	24.3759	1.46715	34.5567	0.662865	11.1044	1.03957	16.0436	0.931623	20.542	4.457869448	6.58214188	5.16234E-05	0.010245092
UDSP5	9.28854	119.725	10.8891	151.122	14.4735	165.424	5.85635	141.933	15.0817	211.163	8.43202	139.042	12.5674	21.242	4.068984948	8.189142259	5.63597E-05	0.01106274
RNF213	44.7205	387.826	34.0103	330.728	37.8932	241.644	32.8556	472.566	27.8598	307.497	40.8476	420.501	28.8614	336.076	3.53441564	12.00531629	5.69047E-05	0.01106274
ARHGAP18	68.2143	14.2775	94.8505	13.3843	74.0161	14.6409	76.0028	9.88732	43.9471	9.40395	36.8049	7.67561	62.75	7.1454	-3.159611547	7.014841492	6.89339E-05	0.011341498
CD40	29.3855	367.548	64.1272	626.417	42.8198	495.326	25.2317	579.997	82.5727	91.4748	39.0428	110.375	50.2701	548.283	8.85450739	8.85450739	6.1708E-05	0.011756603
CHS1	8.8959	12.6886	0.348627	9.44366	2.1879	23.6374	0.625984	10.645	0.552391	16.5179	1.08521	10.3114	0.0786133	6.25023	6.267521377	4.309588684	6.34656E-05	0.011917433
TKB	4.62811	29.0836	3.18042	41.8759	2.5337	35.9395	2.66132	43.8668	2.91627	75.7198	8.73651	94.5299	2.01985	62.8245	4.951093773	5.514391433	6.51786E-05	0.012177442
PLAT	0.224779	11.6373	0.251377	24.7834	0.0549633	14.9511	0.176553	28.6672	0.0143407	13.5674	0.0618383	41.9174	0.206718	21.74	6.729803023	5.174600418	6.80284E-05	0.012583272
TAF1	14.9048	137.148	18.1956	153.453	9.2878	75.1192	13.3391	159.59	13.2708	166.981	20.3599	153.238	9.60905	107.637	6.943529298	7.497834539	6.93325E-05	0.01270118
MGC39372	1.44184	37.6672	1.23727	27.6202	0.497735	18.9448	0.21241	11.1443	0	7.88426	0.211456	17.5282	0	9.93545	9.589720395	2.550235906	7.09045E-05	0.01286545
GK	4.92263	16.4789	4.66646	24.6911	6.64996	30.9976	3.94943	26.5572	3.29964	21.4918	4.29751	26.643	5.34084	36.6627	9.589720395	6.754735968	7.36452E-05	0.013236691
APOBEC3G	4.56503	53.0791	4.81541	55.9795	3.60468	39.124	4.94481	59.9152	1.62027	30.0772	7.64037	74.9045	2.21247	35.6407	3.999703019	5.111616241	7.50219E-05	0.013249671
IFITM3	8.0589	374.422	11.2987	348.787	17.4088	119.425	4.57908	239.536	7.36817	513.147	1.57722	279.6	1.35951	238.642	7.427579484	5.976285546	7.50184E-05	0.013249671
APOL6	5.85916	37.3337	4.43382	60.5983	8.53713	66.588	4.89823	71.4309	4.0331	68.5802	2.30085	50.3072	6.78873	70.9913	3.383788895	8.758299709	7.71156E-05	0.013478966
CNNA1	13.7065	85.0074	8.58858	106.044	16.2623	162.491	6.99801	133.647	11.4832	125.916	9.58503	156.191	12.7119	215.029	7.71623949	9.35071E-05	0.01440728	0.01440728
EIF2AK2	13.7662	92.982	9.10247	151.058	8.45919	137.008	14.353	160.848	9.94938	126.357	5.03602	84.3737	6.56487	103.425	3.968301835	7.01452578	8.4134E-05	0.01440728
TNIP3	0.254806	20.467	0.394598	47.9728	1.05658	458.085	0.414408	191.708	1.28529	443.675	1.60604	21.9151	1.52087	163.922	7.64471458	7.681821244	8.92183E-05	0.015176675
C15orf26	0.0823786	0.61866	0.0705991	1.72328	0.0379672	3.85001	0	0.911532	0	1.95871	0.0643515	1.38326	0.035351	5.14168	6.825196266	1.995942154	9.16519E-05	0.015340091
SHB	2.37448	0.662938	2.41047	0.429512	3.0356	0.442268	1.70315	0.214146	4.13193	0.651049	3.47071	0.261796	3.09521	0.166152	-4.202599735	3.412507964	9.17893E-05	0.015340091
OASL	1.38621	128.692	0.862304	152.263	0.276225	27.4361	1.6261	174.558	1.96105	122.228	1.97879	214.817	0.233231	69.7624	8.147470984	6.175785602	9.61038E-05	0.015830968
OCI3L3	0.134185	68.4197	1.07915	52.9535	0.737636	14.7124	0.244757	39.377	1.20787	1.20787	0.632973	6.17067	0	146.86	5.62855512	9.69875E-05	0.015830968	
RARRR51	0.943394	3.32457	0.287886	3.81696	0.574915	11.4093	1.69946	28.8943	2.18125	17.0318	0.60907	17.8808	1.30516	41.6313	4.982096408	5.030351994	9.72194E-05	0.015830968
FAM94	2.25915	25.9586	2.72021	61.0615	2.69519	32.3046	2.15359	61.1298	1.79636	27.4021	2.93073	36.827	4.37108	48.765	3.47838479	7.066136994	9.88893E-05	0.01596443
SERPINB1	15.6586	66.6081	14.2957	97.4586	16.2577	98.078	19.8991	189.515	10.8554	99.7388	16.8764	126.885	14.7918	117.531	2.98843344	7.512847936	0.000108085	0.01619804
GBP1P1	0.0366852	0.84243	0	10.7001	0.0506512	5.66125	0.108067	27.9453	0.394587	7.62094	0	25.7378	0	14.0231	10.106145648	3.02248187	0.000119223	0.01893398
MT2A	24.9978	848.452	12.2389	766.504	28.7932	504.455	28.7045	870.776	50.4758	1359.28	107.307	1561.81	21.8994	749.641	5.06953921	6.573613575	0.000122525	0.019252354
KIAA0040	8.84903	75.9637	13.8365	93.6265	12.2846	68.8909	9.60917	105.186	16.9728	169.449	6.58926	111.271	8.37221	97.4732	3.548250339	8.0139995	0.000129741	0.020260639
IRAK2	0.721779	15.1172	1.90827	27.9395	1.17511	27.805	0.69729	34.2878	1.56709	33.8908	3.83024	43.0975	2.34451	41.6387	3.145796958	6.331722217	0.00014181	0.021968524
SAMD1L	25.6451	139.116	22.7668	242.949	21.2339	262.477	25.7703	332.118	13.1541	179.512	5.84297	167.208	18.7612	232.929	3.631677835	9.932861393	0.000146811	0.022376958
TNFRSF9	0.52234	3.82366	0.55507	12.0927	0.429368	4.94852	0.44103	13.3844	0.511976	10.0865	0.504679	5.00797	0.598435	11.9894	4.316889002	5.352112224	0.000146811	0.022376958
IRG1	0.0395275	12.9945	0.0854106	30.5297	0.0820767	52.2846	0.085255	101.314	0.0855588	73.1715	0.370105	126.826	0.154015	51.7641	8.326450565	5.790520493	0.000148357	0.022425588
APOBEC3A	0.794366	74.4349	0.576361	198.417	0.48329	112.346	1.68233	291.374	1.16493	241.921	1.78959	436.442	0.903425	253.675	7.413862278	7.413862278	0.000152352	0.02283258
GBP2	17.8652	117.435	15.1571	142.972	34.5362	248.397	25.0681	385.664	29.5215	241.1	22.0148	335.056	28.0816	263.152	3.226432941	8.610177661	0.000175048	0.026044685
LYN	39.1683	148.937	36.3828	205.939	38.4329	187.281	34.6147	273.168	32.2424	186.914	29.2886	209.229	31.8911	188.402	5.559174694	8.923805286	0.000176807	0.026044685
WNT5A	1.28404	6.22763	4.08497	16.06														

PT.LPS.vs.PT.unstim

SERP1G1	0.870351	72.1422	0.443194	51.3972	0.945684	18.2931	1.36332	75.6466	1.03744	54.151	1.22991	62.6594	0.938336	36.2233	5.287456617	5.100107506	0.000364646	0.043150571	
CCl8	0	32.727	0.185945	67.8265	0.349087	221.135	0.452237	169.32	0.62167	211.176	0.0847473	87.9884	0.930942	281.829	8.222040117	7.379306296	0.000368069	0.043286796	
SRP2	0.71137	68.1181	1.3341	58.7087	1.2872	42.241	1.453	78.345	0.897764	51.7901	9.71097	83.8874	2.34611	67.034	4.82170961	5.21823481	0.00037618	0.044252391	
XRN1	11.4337	64.0701	12.4346	106.249	18.3658	158.116	11.5467	151.596	7.24118	92.5769	1.62617	77.801	14.6474	160.119	3.355122581	6.288823658	0.000382225	0.044403387	
FAM186B	0.0952605	2.03656	0.168127	2.63032	0.017916	1.3983	0.050468	3.24184	0	1.70202	0.127309	2.63946	0.0177235	1.82649	6.348234931	1.521798123	0.000399204	0.046034763	
CCl15	0.0418282	67.0611	0.517919	21.8843	1.13336	172.212	0.254279	70.2574	0	22.6197	0.838998	173.996	0.582372	126.487	7.25735356	5.969706564	0.000409527	0.04700181	
RIPK2	8.60837	44.9557	10.9012	72.8568	13.2755	107.791	9.49547	84.5621	8.08516	65.2709	8.61497	88.371	14.1919	141.757	3.181263254	7.703208039	0.000413956	0.047252659	
DCP1A	11.3353	55.3514	13.3994	62.6466	12.339	52.608	10.464	70.9659	10.6527	65.5068	10.8113	67.0269	13.5156	75.912	2.481215168	8.182546234	0.000420359	0.04769115	
TRIM31	0	4.52614	0.0177936	3.18573	0.0855159	4.57342	0.0152425	1.187	0	8.27186	0	4.89496	0	4.3001	9.178079882	2.13797899	0.000438928	0.049491953	
PPP3CC	8.2109	30.1914	9.62465	37.3933	11.5118	45.3846	8.56973	52.9804	7.43023	38.1268	6.6455	51.7428	10.1386	64.2088	2.66274631	6.459504253	0.000445629	0.049495881	
MIR155HG	0.356034	7.77128	0.612761	50.39	0.943204	43.5801	0.481745	37.3399	1.1518	15.8857	0.347768	18.1117	1.49585	52.74	5.129416184	5.27488004	0.000444247	0.049495881	
TNFAIP2	22.3351	159.296	19.2873	208.673	12.5802	115.739	17.5459	199.582	30.7065	219.733	33.91	231.207	11.302	102.931	3.184680447	7.990795614	0.000461248	0.051091243	
LINC00158	0	0.666794	0	2.82393	0	4.64052	0	3.276	0	0.866115	0	2.78206	0.0382948	9.23399	7.561052517	2.728894933	0.000464005	0.05109952	
TAP2	9.51658	28.6412	12.4384	46.1936	8.2052	33.847	7.79299	37.1392	5.82132	37.3235	4.35748	29.0614	5.88459	39.8056	2.766758711	7.11459515	0.000469651	0.05145799	
BCL2L14	1.2925	24.1179	0.67415	16.4703	0.80171	22.6735	0.214695	52.3425	0	4.42123	0.366139	31.8529	0.115757	14.2078	6.860786747	3.778022253	0.000474251	0.051631009	
IFIT5	3.26961	33.2656	3.41476	53.6107	5.0067	55.1432	5.79838	52.8469	4.09231	47.1213	1.3117	48.8017	6.67404	63.9286	7.244504676	7.244504676	0.000485447	0.05254963	
CCXCL3	0	3.39065	0.0339389	6.29228	0	18.8234	0.145731	6.55506	0	17.3762	1.16251	15.3262	0.356788	60.0351	7.338978631	5.020451615	0.00049013	0.053111719	
CGN	0.022876	1.85805	0.106022	2.0503	0.0317264	1.24916	0.0623311	2.61156	0	1.02625	0.0712684	2.39561	0	0.937159	8.405404461	1.363572698	0.000502202	0.053452225	
FGD5	6.76752	2.26243	10.0115	1.81346	9.15993	6.75145	0.773146	1.95959	13.2978	1.33715	19.9569	13.2978	1.33715	9.37637	1.16715	-3.005631666	5.074139998	0.000506789	0.05364078
WTAP	14.6009	52.3729	18.3532	104.305	18.2643	115.05	15.3337	146.474	18.2527	120.487	10.8385	84.9996	19.4142	121.112	2.19629253	6.982707723	0.000516194	0.054334382	
TNIP1	47.5245	269.351	43.803	233.678	31.6654	157.132	33.6932	353.591	58.9716	397.917	100.554	506.964	29.4726	266.554	3.185374819	8.589055782	0.000533933	0.055882329	
HHEX	4.08618	0.793419	5.52714	0.212633	5.18438	0.491388	7.57912	0.856822	4.12783	0.132842	5.04049	0.569449	5.07672	0.155046	-4.957801989	2.182909061	0.000542433	0.056472727	
IFI6	46.4375	767.347	19.0321	740.075	16.2727	834.712	99.1916	681.56	161.515	2265.56	30.8894	783.939	16.1501	602.966	5.222498757	7.786424519	0.000548386	0.056781766	
TRIM69	0.647026	1.81762	0.186036	3.49181	0.384322	1.76844	0.26203	3.16275	0.299762	3.44281	0.217358	2.27551	0.200407	4.33429	4.588941282	2.102655982	0.000554511	0.057105689	
TMOD2	0.885201	0.841522	0.596308	2.12627	0.349925	1.66457	0.26875	3.1702	0.237887	2.30751	0.112874	1.20821	0.206465	2.75142	3.896105754	3.896105754	0.000562017	0.057562402	
C10TNF1	0.0895004	0.797308	0.0129152	1.21005	0	0.740237	0.0442352	3.04403	0	2.4954	0	2.76388	0	1.86092	8.4279192789	1.437567448	0.000573	0.058375879	
TNFAIP3	11.3683	79.319	13.6709	110.076	9.35011	80.094	6.09508	139.038	11.7572	125.183	23.3361	178.878	15.3366	134.181	3.172178179	8.555977166	0.00059264	0.060058431	
PER3	12.2953	4.82119	11.5948	3.29249	13.8968	3.91764	11.9256	1.99337	9.17039	2.13493	10.3859	2.66494	14.0388	1.87296	-2.906708857	5.743361023	0.000599662	0.060450512	
ZNF618	3.45802	16.0853	3.49056	16.0712	3.41002	10.3424	3.19219	16.2232	1.80452	8.1618	3.89611	15.2654	1.00244	6.68054	2.732248573	5.261568737	0.00061446	0.061632169	
AINOCE3F	1.34447	11.6227	1.10502	8.16292	0.892045	3.74395	1.43531	8.10608	0.470446	5.64517	0.889908	13.8563	0.260892	7.00919	4.062799926	3.55611476	0.000623471	0.062190423	
LINC000515	1.4455	31.4924	2.09188	87.5534	5.01083	83.6698	0.787803	122.53	0	49.1019	1.53398	77.7314	7.24539	283.16	5.286756234	5.352705338	0.000636574	0.063166698	
DOCK4	11.4313	56.5607	16.6093	63.8752	23.3084	94.1833	16.4801	95.835	19.4951	66.5119	21.143	90.3645	20.6495	101.478	2.294588422	9.120645223	0.000654281	0.064409643	
BLZF1	6.58129	34.7642	8.90077	51.9568	10.6608	48.55	8.55473	51.3546	6.09051	40.9102	3.75681	37.8587	9.24787	66.2825	2.838046146	6.484846454	0.000674582	0.066248112	
CCl4	19.9455	623.22	66.1666	953.55	63.3498	1166.07	19.0072	1143.52	82.2801	4620.2	40.5543	2292.14	60.1564	2178.25	5.180877814	9.195418824	0.000678522	0.066270957	
ADAR	37.2267	149.217	40.109	158.812	43.0428	126.528	41.2732	186.192	40.9532	169.007	39.8268	179.898	33.3386	163.175	2.287838387	9.449099595	0.000681771	0.066270957	
AKT1S1	2.12903	20.3241	2.82399	28.7475	1.39247	17.8845	2.1988	19.0378	2.48737	20.5001	2.80239	19.189	1.06809	8.95363	3.058978588	3.65393366	0.000691275	0.068853632	
AKT5C	0.160325	2.83367	0.00671033	12.2117	0	3.45012	0.029414	11.7977	0.0083921	2.47673	0.0185119	0.0302653	7.32604	5.317521949	2.7221949	0.000709392	0.067451038	0.000709392	
TBC1D14	13.3712	7.03128	13.7454	4.2484	11.2074	3.05455	13.8421	2.55389	10.3179	3.6098	13.0672	2.78984	12.2665	1.97304	-2.66686203	6.213994397	0.000723313	0.068248999	
TOR1B	10.4111	68.304	14.1422	89.1997	11.2678	73.1313	12.6138	65.2237	10.054	49.1528	12.8272	102.744	15.2472	96.5758	2.661117914	7.330439268	0.000730641	0.068600852	
SLC25A28	8.33332	57.3025	6.64877	37.3558	5.08637	41.5367	7.12112	71.4165	13.6236	87.1534	9.63453	86.2305	8.11352	68.6306	5.86257904	0.000735952	0.000735952	0.06872952	
TNC	0.386516	14.404	0.320892	25.8499	0.554811	36.7574	0.11236	26.3419	1.67787	54.0624	0.290522	24.2471	0.999729	78.0311	6.281746571	8.538445471	0.000748396	0.070586382	
DNAAF1	0.0335946	1.81259	0.0436063	2.04433	0.0465201	13.1699	0.0746707	10.3974	0	12.9878	0.392968	30.9828	0.065398	17.65	7.94357599	4.471869452	0.000761464	0.07182337	
SOCS3	3.77529	38.7117	3.91146	58.8047	2.25613	38.321	1.92102	54.1974	6.40484	234.304	6.61435	105.665	3.48862	62.1087	4.150414228	6.541014361	0.000775076	0.072385981	
GJB2	2.09501	35.6979	3.32035	81.1491	2.82794	49.8821	1.7998	59.1304	5.572057	57.6069	8.89852	47.9203	2.47201	43.6986	4.1389664	5.786557461	0.000782547	0.072435433	
GCNT4	0.0578716	2.3146	0.240968	5.64794	0.497318	9.87191	0.111676	8.84606	0.175899	4.92141	0.0135301	5.62839	0.301826	13.056	5.413610293	4.59774134	0.000785952	0.072435433	
TNFSF15	0.217003	22.3233	0.370384	36.0163	0.267586	55.9863	0.27806	24.303	0.438237	29.3799	1.51234	96.3177	2.09656	163.392	6.298256126	9.22040762	0.000790514	0.072435433	
APOL1	6.1135	58.643	5.66571	43.4819	4.56135	27.0995	5.25031	55.6157	6.86283	65.7379	5.35575	41.2906	4.45635	33.2179	2.882386655	5.795736899	0.000791644	0.072435433	
NBN	12.3241	56.1189	14.181	84.5786	25.066	197.585	11.6128	129.075	7.81362	84.119	9.04313	100.028	24.1999	227.919	3.233609628	9.286629998	0.000819615	0.072435433	
MYZAP	1.89688	0.036546	3.17799	0.752862	1.60916	0.26339	1.88646	0.394265	3.35068	0	0.85966	0	2.11157	0	-8.42145425	1.379528669	0.000799067	0.072494422	
B4GALT5	12.139	51.5015	14.3336	85.2236	13.8678	73.7047	9.7961	93.4071	15.9037	138.034	10.1446	107.282	15.7447	103.298	8.23826577	8.23826577	0.000821282	0.0741567	
OPTN	21.1708	133.692	17.3052	134.621	24.5518	227.393	8.80478	209.077	39.0362	399.555									