#### **Online Supplementary Material:**

#### **Experimental Procedures**

**Cells:** Fresh adult whole bone marrow (BM) was purchased from All Cells, Inc. (Emeryville, CA). Unmatched human peripheral blood was purchased from the Stanford Blood Bank according to an IRB-approved protocol. All blood and bone marrow samples were collected in heparin sulfate anticoagulant, stored at room temperature for 4-6 hours, and then separated over Ficoll-Paque Plus (Amersham Biosciences) using Accuspin tubes (Sigma-Aldrich, St. Louis, MO) to remove erythrocytes, platelets, and granulocytes. Cells were used fresh prior to 34-parameter mass cytometry experiments, or frozen in FCS with 10% DMSO. Cells were rested at 37°C, 5% CO<sub>2</sub> for 1 hour in RPMI with 10% FCS (supplemented with 2mM EDTA in the case of frozen samples), 1X L-glutamine and 1X penicillin/streptomycin (Invitrogen).

**Antibodies:** A summary of all mass cytometry antibodies, reporter isotopes and concentrations used for analysis can be found in Methods Table 2. With the exception of CD3-QDot 655 (Invitrogen, Carlsbad, CA), primary conjugates of mass cytometry antibodies were prepared 100 µg at a time using the MaxPAR antibody conjugation kit (DVS Sciences, Toronto, Canada) according to the manufacturer's recommended protocol. Following labeling, antibodies were diluted in Candor PBS Antibody Stabilization solution (Candor Bioscience GmbH, Wangen, Germany) to between 0.1 and 0.3 mg/mL and stored long-term at 4°C. Each antibody clone and lot was titrated to optimal staining concentrations using cell lines and primary human samples.

For the comparison of mass cytometry and fluorescence flow cytometry data, commercially available, fluorescently labeled antibodies were used. The pre-methanol staining panel consisted of: (1) Pacific Blue CD8 (clone RPA-T8); (2) V500 CD4 (clone RPA-T4) (BD Biosciences, San Jose, CA); (3) Qdot 605 CD45RA (clone MEM-56) (Invitrogen); (4) Alexa Fluor 700 CD56 (NCAM) (BioLegend, San Diego). The post-methanol staining panel consisted of: (1) PE CD3 (clone UCHT1); (2) PerCP-Cy5.5 CD20 (clone H1(FB1)); (3) PE-Cy7 CD33 (clone 67.6); (4) Alexa Fluor 488 phospho-STAT3; (5) Alexa Fluor 647 phospho-STAT5 (all from BD Biosciences, San Jose, CA).

**Stimulation of peripheral blood and bone marrow with phospho-signaling analysis:** Stimulation and cellular staining protocols were based on procedures previously described by (1). Briefly, after resting cells 1 hour, cytokines were

added and cells were returned to the 37°C incubator for 15 minutes. A summary of stimuli and inhibitors can be found in Methods Table 1. Where indicated, chemical inhibitors were added to cells for the final 30 minutes of the rest, prior to stimulation. To halt signal transduction, cells were fixed with formaldehyde (PFA; Electron Microscopy Sciences, Hatfield, PA) added directly to growth media at a final concentration of 1.6% for 10 minutes at room temperature. Cells were then centrifuged at 500g for 5 minutes and washed once with staining media (PBS with 0.5% BSA, 0.02% sodium azide) to remove residual PFA, and blocked with Purified Human Fc Receptor Binding Inhibitor (eBioscience Inc., San Diego, CA) following manufacturer's instructions. Surface marker antibodies were added yielding 50 or 100 uL final reaction volumes and stained at room temperature for 30min. Following staining, cells were washed 2 more times with cell staining media then permeabilized with 4°C methanol for at 10 min at 4°C, then optionally stored at -80°C for later use. Cells were then washed twice in cell staining media to remove remaining methanol, and then stained with phospho-specific antibodies in 50 or 100 µL for 30 min at room temperature. For fluorescence analysis, cells were then washed once in cell staining media and analyzed using a BD LSRII (BD Biosciences, San Jose, CA). For mass cytometry analysis, cells were washed once in cell staining media, stained with 1 mL of 1:4000 <sup>191/193</sup> Ir DNA intercalator(2) (www.dvssciences.com; DVS Sciences, Richmond Hill, Ontario, Canada) diluted in PBS with 1.6% PFA for 20 mins at room temperature. Cells were then washed once with cell staining media and then finally with PBS alone. Care was taken to assure PBS and buffers preceding CyTOF<sup>™</sup> analysis were not contaminated with metals in the mass range above 200 AMU analysis range. In cases where cells were not treated with methanol, the iridium intercalator was used at 1:2500. Mass cytometry samples were diluted in dH<sub>2</sub>O to approximately  $10^6$  cells per mL and then analyzed on a CyTOF<sup>TM</sup> mass cytometer(3)(DVS Sciences, Toronto, Canada) equilibrated with dH<sub>2</sub>O.

**Mass cytometry viability (membrane permeability) staining**: Where indicated, cells were viability stained for mass cytometry analysis during the 15 minute stimulation period. Here, the viability dye was mixed 1:20 with stimulation media. Viability dye was made fresh by creating a solution of 30 mM RhCl<sub>3</sub> (Sigma-Aldrich, St. Louis, MO) and 60 mM DOTA-NHS ester (Macrocyclics, Dallas TX) in 50% v/v DMSO in L-Buffer (DVS Sciences, Toronto, Canada). Dasatinib-treated cells from the unmatched bone marrow were viability stained in a 3-fold higher concentration of viability dye (i.e. 4.5 mM final concentration) due to a technical error.

Mass cytometry measurement: A review of initial mass cytometry analysis

methods can be found by Ornatsky et al.(4). Here, cells were acquired on the CyTOF<sup>™</sup> mass cytometer (www.dvssciences.com; DVS Sciences, Richmond Hill, Ontario, Canada) at an event rate of approximately 500 cells per second. The instrument was run in high-resolution mode (Mass resolution ~700) with internally calibrated dual-count detection. Noise reduction and cell extraction parameters were: cell length 10-65, lower convolution threshold 10. A cell subtraction value was set to -100. After acquisition, the effect of the cell subtraction setting was negated by subtracting a value of 100 from every channel of each FCS file using the FlowCore package for R (5). Also using FlowCore, the cadmium 110, 111, 112, and 114 values were summed to create a single representative channel for the CD3-QDot 655 used in the mass cytometry analysis. For both BM samples, data collection was carried out over multiple days, during which a systematic decay in the signal intensity was observed over a scale of hours. To help minimize this, each day standard cleaning of the instrument was performed. To account for remaining variability between samples, the intensity of each parameter was modeled over time to account for systematic signal decay, and to allow normalization of samples from different days to a consistent baseline. Biological replicates of the unstimulated condition were collected at the beginning and end of each day, and used to calculate a linear regression that estimates the dependence of signal intensity on absolute event number. The change in signal intensity over time for each channel of each unstimulated sample was estimated using the median intensity in overlapping sliding windows of 5000-10,000 cells each, sliding in increments of 1,000 from the first to the last cell in the sample, yielding 395 data points from a typical FCS file with 400,000 events. Values between 0 and 2 were common when antigen expression is at or below the threshold of detection therefore, to avoid their predominance, values < 2 were excluded from the line fitting procedure. Lines with equations s=m\*n+b were fit to the unstimulated samples each day, where s is signal, n is the cell number ("event number") and m and b are fit parameters signifying the slope of the decay curve and the maximal signal intensity on that day ('y-intercept'), respectively. The individual events were multiplied by s N/(m\*n+b), where s N is a constant representing the signal at the last cell, thus normalizing each event to the signal level at the end of the last day.

**Data visualization:** All cell density plots shown were created in Cytobank(*6*) (www.cytobank.org, Cytobank, Inc., Menlo Park, CA). For mass cytometry data, all parameters except time and cell length were displayed with an arcsinh transformation and a scale augment of 5 ranging from -20 to 10000. Cell length

and time were displayed on linear scales. LSRII data was displayed using default Cytobank scales (arcsinh with a scaling argument of 150, ranging from -200 to 262,144). Justification for the use of these different transformations is provided in Fig. S2.

Phospho-signaling and mass cytometry data analysis: Normalized. background-subtracted FCS files were imported into Cytobank for single cell and population gating. Cytobank was also used to create the heatmaps comparing fluorescence and mass cytometry measurements using log2 ratio of the stimulated population phospho-STAT mean fluorescent intensity (MFI) versus the unstimulated control. For larger heatmaps, the median intensity values from the indicated cell populations (Fig. 3A) were exported from Cytobank and imported into Spotfire (TIBCO Software, Inc., Palo Alto, CA) to create of heatmaps, drug response X-Y plots, and perform hierarchical clustering. Clustering was performed using UPGMA with Euclidean distance measurements and normalization by means. For spanning tree visualization and comparison, singlet-gated FCS files were exported from Cytobank and analyzed using SPADE in Matlab v.R2010a (The Mathworks, Inc., Natick, MA) as described below.

For all of the mass cytometry analysis, signaling induction was calculated using the average of the median scaled arcsinh values of 5 untreated replicates, minus the median scaled arcsinh value of a stimulated condition. To minimize the noise in measurements where the data is close to zero (the limit of detection on the mass cytometry scale), the median intensities in were scaled (divided by 5) prior to archsinh calculations. An example calculation and comparison of arcsinh difference to fold change is shown below:

	For	examp	le:
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Sample	Number	ofNumber	ofMedian scaled arcsinh intensity of pSTAT5 in
_	cells	incells	inNode X (median(arcsinh(intensity/5)))
	downsam	oledupsample	d
	Node X	Node X	
Unstim 1	54	3132	1.4
Unstim 2	62	3286	1.2
Unstim 3	81	4050	1.3
Unstim 4	71	3692	1.6
Unstim 5	87	4437	1.5
TPO	54	3240	4.5
Unstim Ave	rade:		1.4
Signaling Induction of pSTAT5 by TPO:		STAT5 by TPO	4.5 - 1.4 = 3.1

Arcsinh vs. fold-change:

	<u> </u>		
Unstimulated	Stimulated	Fold Change	Arcsinh difference
Intensity	Intensity		(calculated above)
5	10	2	0.6
5	25	5	1.4
5	50	10	2.1
5	250	50	3.7
5	500	100	4.4

Statistical analysis of signaling responses in manually gated bone marrow was performed as follows: For each experimental condition, the arcsinh-transformed median intensity of each of 18 intracellular antibodies was compared to the distribution of the arcsinh-transformed median intensities of unstimulated replicates (n=5) in each of 24 manually gated cell populations (resulting in a total of 432 observations per condition). Independent one-sample *t*-tests were performed with 4 degrees of freedom and *p*-values were adjusted for 432 multiple comparisons (Bonferroni). All signaling responses with adjusted *p*-values below 0.05 are listed in Table S3.

**SPADE analysis and tree construction**. Given an FCS file as input, SPADE first performs a density-dependent down-sampling step based on n parameters specified by the user. This step heavily down-samples dense populations corresponding to abundant cell types while preserving rare cell types with relatively sparse distributions. The resulting dataset is density-normalized, but retains the multidimensional structure of the original FCS file. Using the density-normalized dataset, rare and abundant cell types are equally likely to form clusters in the subsequent clustering step. After down-sampling, an unsupervised agglomerative hierarchical clustering step consolidates down-sampled cells into clusters of phenotypically similar cells based on n parameters. SPADE then

connects the centroids of the cell clusters using a minimum spanning tree (MST) (7). The topology of this tree representation reveals the underlying relationship and similarity of cell nodes in n dimensional space. All stimulated BM and PBMC mass cytometry datasets were singlet-gated in Cytobank, exported as FCS files and imported into SPADE for analysis. SPADE analysis was performed with the default configuration (arcsinh cofactor = 5, normalization = OFF, down-sampling scaling factor = 5, sparsity exclusion percentile = 1, target number of cells = 20000, max number of cells = 50000, target number of clusters = 200). Fifteen (15) 'static' parameters that were not expected to change between experimental conditions were used for down-sampling and tree construction: DNA (Ir191), Cell length, and the 13 'core' surface antibody parameters. All conditions from a single experiment were processed simultaneously so the resulting tree structure would capture all cell surface subpopulations present in the entire dataset. To reduce the effect of counting error, the *n* dimensional coordinates of each node were then used to up-sample the data from the original FCS file, thereby populating each node with every cell from the original FCS file that shares the phenotype with the cells in the node. Here, for each cell in the original data file we computed the distance to every individual cell in the pooled down-sampled dataset from which the tree was constructed (L1 distance based on the 13 surface markers). Then, based on which one was its nearest neighbor (NN), the cell in the original file was assigned to the same cell cluster (tree node) as its downsampled NN. The same approach was used to overlay the results of the 'immunophenotype' staining panel on the tree constructed from the 29 conditions with the 'functional' panel. The set of 13 'core' surface markers, which were shared between the two staining panels, allowed assignment of the 'immunophenotype' cells to the tree. These cells were used to compute the median intensities of the complementary 18 surface antigens shown in Fig. 2 and S4B. For cellular signaling response calculations, the median arcsinh intensity of up-sampled cells within each node (i.e. cluster) for each original FCS file (i.e. each experimental condition) was used for signaling induction calculations and visualizations. See above example table for archsinh calculations.

**Principal component analysis (PCA) of phenotypic progression:** We identified two subsets of cells involved in B-cell development. One subset included HSC, MPP, Pro-B cells and early Pre-B cells. The other included late Pre-B cells, Immature B-cells, Naïve Mature B-cells and IL3RA+ Mature B-cells. For each subset, we ran PCA(8) over the cell surface markers, and examined the coefficients of the first principal component. For the first subset (Fig. 4), the first component explained 22.8% of the variation in the data; the three highest coefficients for this component were CD38 (-0.67), CD20 (0.5) and CD45RA

(0.38). For the second subset (Fig. 5D), the first component explained 23.9% of the variation in the data; the three highest coefficients for this component were CD19 (0.55), CD33 (-0.54) and CD34 (-0.51). For each cell, we defined the progression value as the projection of the cell markers values on to the first principal component. In each subset, we divided the cells to 17 overlapping windows according to their progression values. The windows were in equal distances across the progression trajectory, had width of 20%, and 75% overlap between consecutive windows. Notably, the plots remain smooth even if no overlapping windows are used. For each marker, the mean in each window was used to plot the progressions in Fig. 4 and 5.

**Public data distribution:** For replicate human bone marrow samples analyzed here, single-cell gated FCS files are publicly available on www.cytobank.org for download.

**Methods Table 1**. Molecules used for stimulation and inhibition of biochemical responses in human PBMCs and bone marrow. All cytokines are the human sequence.

Short		Final			
name	Stimulation/Inhibitor	Concentration	Supplier		
	Goat anti-human IgG (BCR		Southern Biotech, Birmingham		
	cross-linking)	10 µg/mL	AL		
	Goat anti-human IgM (BCR		Southern Biotech, Birmingham		
PCP	cross-linking)	10 µg/mL	AL		
DUK	Goat F(ab)2 anti-human Ig κ		Southern Biotech, Birmingham		
	(BCR cross-linking)	10 µg/mL	AL		
	Goat F(ab)2 anti-human Ig $\lambda$		Southern Biotech, Birmingham		
	(BCR cross-linking)	10 µg/mL	AL		
Flt3L	Flt-3 Ligand	50 nh/mL	PeproTech, Rocky Hill, NJ		
IL-2	IL-2	100 ng/mL	PeproTech, Rocky Hill, NJ		
IL-3	IL-3	20 ng/mL	PeproTech, Rocky Hill, NJ		
IL-6	IL-6	100 ng/mL	BD Biosciences, San Jose, CA		
IL-7	IL-7	20 ng/mL	BD Biosciences, San Jose, CA		
IL-10	IL-10	100 ng/mL	PeproTech, Rocky Hill, NJ		
LPS	LPS (Ultrapure)	1 µg/mL	Invitrogen, Carlsbad, CA		
G-CSF	G-CSF	20 ng/mL	PeproTech, Rocky Hill, NJ		
GM-CSF	GM-CSF	20 ng/mL	PeproTech, Rocky Hill, NJ		
IFNα	IFNα	5000 U/mL	PeproTech, Rocky Hill, NJ		
IFNα			PBL InterferonSource,		
	IFNα/AD (Cat. #11200-1)	5000 U/mL	Piscataway, NJ		
PMA/lono.			Sigma-Aldrich, St. Louis, MO,		
	PMA	50 nM	St. Louis, MO		
			Sigma-Aldrich, St. Louis, MO,		
	lonomycin	1 µg/mL	St. Louis, MO		
PVO4	Pervanadate (Sodium	125 µM in 0.53			
	orthovanadate)	$mM H_2O_2$	Calbiochem, San Diego, CA		

SCF	SCF	100 ng/mL	PeproTech, Rocky Hill, NJ
TNFα	ΤΝϜα	20 ng/mL	PeproTech, Rocky Hill, NJ
TPO	TPO	50 ng/mL	PeproTech, Rocky Hill, NJ
Dasatinib	Dasatinib	100 nM	LC Laboratories, Woburn MA
JAKi	JAK Inhibitor I	5 µM	Calbiochem, San Diego, CA
U0126	U0126	20 µM	Calbiochem, San Diego, CA

**Methods Table 2**. A summary of antibodies used for mass cytometry analysis. Also indicated is the staining panel the antibody was used in.

Antigen Target	Clone	Supplier	Elemental	Final	Immunophe	Functional
(Human)	Number		Isotope	Concentratio	notype	Panel 2
0540	11140	<b>B</b> : 1	0.1450	n (µg/mL)	Panel 1	
CD10	HI10a	Biolegend	Gd156	3	X	
CD117	104D2	Biolegend	YD1/1	1	X	×
CD11b	ICRF44	Biolegend	Nd144	3	X	Х
CD11C	3.9	Biolegend	Sm154	5	X	×
CD123	915	BD Biosciences	Eu151, Gd160	1	X	Х
CD13	L138	BD Biosciences	Er168	1	X	
CD14	M5E2	Biolegend	Gd160	2	X	
CD15	W6D3	Biolegend	Dy164	1	X	
CD16	3G8	Biolegend	H0165	2	X	
CD161	HP-3G10	Biolegend	Nd150	5	X	X
CD19	HIBI9	BD Biosciences	N0142	1.5	X	X
CD20	2H7	BD Biosciences	Sm147	3	X	Х
CD235a/b	HIR2	Biolegend	Pr141	2	X	
CD3	UCH11	Biolegend	Pr141	3	Х	X
CD3 Qdot 655	54.1	Invitrogen	Cd110, 111, 112, 114	1:500 dilution		X
CD33	WM53	Biolegend	Nd148, Gd158	1.5	Х	Х
CD34	8G12	BD Biosciences	Nd148	3	Х	Х
CD38	HIT2	Biolegend	Tb159, Er167	1	Х	Х
CD4	RPA-T4	Biolegend	Nd145	3	Х	Х
CD41	HIP8	Biolegend	Sm152	1	Х	
CD44	G44-26	BD Biosciences	Er166	1	Х	
CD45	HI30	Biolegend	In115	2	Х	Х
CD45RA	HI100	Biolegend	La139	3	Х	Х
CD47	B6H12	BD Biosciences	Yb172	2	Х	
CD56	B159	BD Biosciences	Er170	2	Х	
CD56	HCD56	Biolegend	Er170	2	Fig. 1 Compari	son
CD61	VI-PL2	BD Biosciences	TM169	0.25	X	
CD7	M-T701	BD Biosciences	Er167	2	Х	
CD8a	RPA-T8	Biolegend	Nd146	1.5	Х	Х
CD90	5E10	Biolegend	Er170, Yb176	5	Х	Х
CXCR4	12G5	Biolegend	Lu175	3	Х	
HLADR	L243	Biolegend	Yb174	2	Х	
IgM	G20-127	BD Biosciences	Eu153, Dy164	2	Х	
lkB alpha	L35A5	Cell Signaling Technology	Er166	2		Х
Ki67	B56	BD Biosciences	Sm152	2		Х
Btk/Itk (pY551/pY511)	24a/BTK	BD Biosciences	Gd158, Yb171	2		Х
Čreb (pS133)	87G3	Cell Signaling Technology	Yb176	1.5		Х
CrkL (pY207)	polyclonal	Cell Signaling Technology	Lu175	2		Х
Erk1/2	20A	BD Biosciences	Eu151	1		Х
U3 (nS20)		Piologond	Er169	0.5		v
MADKADK 2	2787		E1100 Eu153	1		X
(pT334)	2101	Technology	Luiss			^
M <sup>2</sup> · · · · /						1

NfkB (pS536)	K10- 895.12.50	BD Biosciences	Ho165	1	Х
p38 (pT180/pY182)	36/p38	BD Biosciences	TM169	2	Х
PLCg2 (pY759)	K86-689.37	BD Biosciences	Pr141	2	Х
S6 (pS235/pS236)	N7-548	BD Biosciences	Yb172	2	Х
Shp2 (pY580)	polyclonal	Cell Signaling Technology	Sm154	2	Х
SLP-76 (BLNK) (pY128)	J141- 668.36.58	BD Biosciences	Gd160, Dy164	1	Х
Src (pY418)	K98-37	BD Biosciences	Yb174	2	Х
STAT3 (pY705)	4/P-STAT3	BD Biosciences	Tb159	2	Х
STAT5 (pY694)	47	BD Biosciences	Nd150	2	Х
ZAP70/Syk (pY319/pY352)	17a	BD Biosciences	Gd156	1	X

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#### Supplemental Figure Legends:

# Figure S1 – Comparison of mass and fluorescence cytometry profiling of immune cell response patterns in two PBMC samples.

(A–B) Representative antibody surface staining results and cell population definitions ("gating") for fluorescence (A) and mass (B) cytometry analysis of fixed PBMCs from Donor A. Identical analysis of Donor B is shown in Fig. 1B and 1C of the main text. \*Pearson correlation between frequencies measured by fluorescence or mass cytometry, including both donors (r = 0.99, p < 0.000001, two-tailed *t*-test) as shown in Table S1 and (C).

(C) Linear regression of paired frequencies measured by fluorescence or mass cytometry. Data includes frequencies of 12 populations from each of 2 donors (r = 0.99, p < 0.000001, two-tailed *t*-test) (Table S1).

(D) Heatmap summary of induced STAT phosphorylation in immune populations from PBMC Donor A. Immune populations are defined in (A) and (B) (*column headers* refer to *blue polygons* in (A) and (B)). Responses to the indicated stimuli in each row were measured by fluorescence (*top*) and mass (*bottom*) cytometry. Color scale indicates the difference in log<sub>2</sub> mean intensity of the stimulated condition compared to the unstimulated control. Signaling responses of Donor B are shown in Fig. 1E of the main text. \*\*Pearson correlation between signaling induction measured by fluorescence or mass cytometry, including both donors (pSTAT3: r = 0.92; p < 0.000001, two-tailed *t*-test; pSTAT5: r = 0.89, p < 0.000001, two-tailed *t*-test) as shown in (E) and (F).

(E) Linear regression of paired pSTAT3 signaling induction values calculated from intensities measured by fluorescence or mass cytometry. Data includes signaling intensities of 45 unique combinations of populations and stimuli (9 populations x 5 stimuli) from each of 2 donors (r = 0.92; p < 0.000001, two-tailed *t*-test).

(F) Linear regression of paired pSTAT5 signaling induction values calculated from intensities measured by fluorescence or mass cytometry. Data includes signaling intensities of 45 unique combinations of populations and stimuli (9 populations x 5 stimuli) from each of 2 donors (r = 0.89; p < 0.000001, two-tailed *t*-test).

# Figure S2 – Justification for Arcsinh transformation of display scale in mass cytometry analysis.

CD4 and CD8 expression as measured by mass (top) and fluorescence (bottom) cytometry for T cells as gated in Fig. 1B and C of the main text. Shown is the same data on a linear scale as well as on arcsinh scales with a scaling

argument of 150, 5, or 1 in Cytobank software. The  $log_{10}$  scales displayed are compressed according to the arcsinh transformation. An arcsinh scaling factor of 150 works well for digital fluorescence data while and augment of 5 works well for mass cytometry data. Both of these are default on cytobank.org.

### Figure S3 – High dimensional single-cell surface analysis is difficult to display in a 2-dimensional space with dot plots.

A summary of possible 2-parameter dot plots for a 31-parameter cell surface immunophenotype of healthy human bone marrow obtained using mass cytometry.

### Figure S4 – Expression of immunophenotype surface markers overlaid onto the SPADE plots of healthy human bone marrow.

(A) The expression of the 13 core surface markers used to construct the SPADE MST in Fig. 2 of the main text.

(B) The expression of an additional 18 surface markers from the grafted 31 surface marker analysis of the same sample. These 18 surface markers were not used in the SPADE plot and their localized expression is based solely on the shared expression patterns of the 13 core surface markers.

### Figure S5 – Manual gating strategy for healthy human bone marrow using traditional 2D cytometry analysis methods.

The gating hierarchy is shown for 24 manually gated cell populations in used in the heatmaps of Fig. 3 and 5 of the main text. All gates were applied with Boolean "AND" logic in Cytobank software. Therefore, each cell population is defined by the combination of gates in its respective row.

### Figure S6 – Hierarchically clustered heatmap of intracellular functional marker dynamics induced by 13 biological and chemical stimuli.

Single-cell data from healthy human bone marrow was divided into 24 traditionally gated cell populations, and hierarchically clustered based on signaling induction and cell population. Each row in the heatmap represents the phosphorylation level of one phospho-protein in response to one stimulation condition.

### Figure S7 – Venn diagram of statistically significant signaling responses among manually gated bone marrow populations from two donors.

Independent one-sample *t*-tests (see Experimental Procedures) were performed to identify statistically significant signaling responses for each of 22 unique experimental conditions in bone marrow samples from each of 2 healthy

donors. The effect of each experimental condition was measured using 18 intracellular markers in 24 manually gated cell populations, for a total of 432 signaling responses per condition. After correction for multiple comparisons, an average of 554 significant signaling responses were observed for each donor, of which 248 were significant for both (45% concordance). All 1,108 significant signaling responses are listed in Table S3.

# Figure S8 – A comprehensive summary of dynamics of 18 intracellular functional markers in a healthy human bone marrow sample, overlaid on SPADE plots.

(A) Induced changes in 18 functional markers by 13 *ex vivo* stimuli were overlaid onto the SPADE MST using the 13 core surface markers as anchors.

(B) A subset of 6 conditions (Unstimulated, BCR, IL-7, Flt3L, PMA/Ionomycin., PVO<sub>4</sub>) are shown after 30 minutes pretreatment with 0.1  $\mu$ M dasatinib.

(C) A subset of 2 conditions (Unstimulated, GCSF) are shown after 30 minutes pretreatment with 5  $\mu$ M JAK inhibitor I. Signaling induction was calculated as the difference of arcsinh medians of the indicated *ex vivo* stimuli compared to the unstimulated control for each node.

# Figure S9 – Confirmatory analysis of a second healthy human bone marrow.

A comprehensive summary of 18 intracellular functional markers and 13 extracellular markers in a second healthy human bone marrow sample.

(A) Induced changes in 18 functional markers by 13 *ex vivo* stimuli were overlaid onto the SPADE MST using the 13 core surface markers as anchors. Signaling induction was calculated as the difference of arcsinh medians of the indicated *ex vivo* stimuli compared to the unstimulated control for each node.

(B) A subset of 6 conditions (Unstimulated, BCR, IL-7, Flt3L, PMA/Ionomycin, PVO<sub>4</sub>) are shown after 30 minutes pretreatment with 0.1  $\mu$ M dasatinib. Calculated as in (A).

(C) A subset of 2 conditions (Unstimulated, GCSF) are shown after 30 minutes pretreatment with 5  $\mu$ M JAK inhibitor I. Calculated as in (A).

(D) A subset of 2 conditions (Unstimulated, PMA/Ionomycin) are shown after 30 minutes pretreatment with 20  $\mu M$  U0126, an inhibitor of MEK. Calculated as in (A).

(E) The expression of the 13 core surface markers used to construct the SPADE MST in (A) through (E).

#### Figure S10 – Signaling dynamics in response to the chemotherapeutic

#### agent dasatinib.

(A) Heatmap of the effect of dasatinib on intracellular signaling induction by five biological or chemical stimuli in 24 traditionally gated cell populations (Fig. S5). Suppression index was calculated as the signaling induction by the stimulus in the presence of dasatinib minus the induction by the stimulus alone.

(B) A complete spectrum of responses to dasatinib was observed. Induced signaling by pervanadate is shown with (Y-axis) and without (X-axis) dasatinib pre-treatment. Each data-point indicates induced phosphorylation of 18 functional markers (*colors*) in each of the 24 manually gated cell types. Suppression index is overlaid as the background of the plot. Points that fall along the white transition were unaffected by the presence of dasatinib. Points that fall to the lower right (*orange*) or to the upper left (*purple*) were suppressed or potentiated by dasatinib, respectively.

### Figure S11 – Workflow diagram of optimization steps leading to final experimental design and published bone marrow dataset.

The final set of antibodies and stimuli used in the bone marrow experiments were determined through an iterative process of testing and titrating individual reagents, combining panels of reagents, then revising those panels based on biological experiments. The steps to optimize different components of the final experimental design are shown (*blue* – surface antibodies; *orange* – intracellular antibodies; *red* – stimulation conditions; *green* – viability indicator).

#### Supplemental Tables:

**Table S1** – A summary of gated cell population frequencies for comparison of fluorescence and mass cytometry in Fig. 1 of the main text.

**Table S2** – A summary of justifications for putative cell population assignments for the SPADE plot in Fig. 2 of the main text.

**Table S3** – Statistically significant signaling responses observed in bone marrow samples from either of two healthy donors.

#### Figure S1



### Figure S2



CD8

#### Figure S3





Erythrocyte

3.7

-3.7

































Promyelocyte CD38<sup>mid</sup> CD3- platelet CD38- CD3- platelet CD38<sup>mid</sup> CD3<sup>mid</sup> platelet

Erythroblast Erythrocyte

-3.9 3.9

`Immature B Mature B

IL-3Rα+ mature B

Plasma cell

















CD38<sup>mid</sup> CD3<sup>mid</sup> platelet

**Erythroblast** Erythrocyte

-3.1

3.1




























































Monocyte Pro-monocyte Pre-DC GMP-

**Plasmacytoid DC** 

Myelocyte -

Promyelocyte CD38<sup>mid</sup> CD3- platelet<sup>1</sup> CD38-CD3-platelet CD38<sup>mid</sup> CD3<sup>mid</sup> platelet

Naive CD8+T Mature CD8+T Naive CD4+T Mature CD4+T Immature B Mature B IL-3Rα+ mature B Plasma cell

**Erythroblast** Erythrocyte -3.5

3.5



























0.2

























## Figure S6





















































































































































































































































































































































































































































































































































































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## Figure S9A
































































































































































































































## Figure S9A















































## Figure S9A






































Figure S9A



















Figure S9A











































































































































## Figure S9A





















## Figure S9B



Figure S9B



## Figure S9B



## Figure S9B 141–pPLCgamma2 --- Dasatinib+PMAiono vs Ref Ratio

































































































































Figure S9B









Figure S9B






































Figure S9B























Figure S9B













# Figure S9C




















Figure S9C













Figure S9C







Figure S9C

























Figure S9C



Figure S9C





Figure S9C



Figure S9C



## Figure S9D





## Figure S9D


















Figure S9D














































































### Figure S10

-2

-2.5

-1

-0.5

0



Suppression of

PVO₄→ pSrcFK

3

3.5

4

4.5

0.5 1 1.5 2 2.5 Signaling Induction by Stimulus

## Figure S11



# **Table S1**: Frequency of cell populations gated forcomparison of fluorescence and mass cytometry signalinganalysis of healthy PBMC samples (n=2)

			Frequency of S	Single Cell Gate
		Gate #	Fluorescence	Mass Cytometry
	CD33 Monocytes	1	13.5	11.4
	CD33- Lymphocytes		80.3	79.9
	CD20+ B Cells	2	18.8	18.3
	CD33-CD20- Lymphocytes		80.5	76.2
	NKT Cells	7	0.8	4.2
PBMC A	CD56mid NK Cells	8	1.7	5.6
(Fig. S2)	CD56hi NK cells	9	0.1	0.6
	CD3+ T Cells		81.0	77.5
	CD8+CD45RA+ Tcells	5	14.4	17.3
	CD8+CD45RA- T Cells	6	11.4	15.4
	CD4+CD45RA+ Tcells	3	22.4	28.6
	CD4+CD45RA- T Cells	4	28.4	36.4
	CD33 Monocytes	1	38.8	31.3
	CD33- Lymphocytes		49.7	57.1
	CD20+ B Cells	2	18.5	18.4
	CD33-CD20- Lymphocytes		80.4	79.2
	NKT Cells	7	0.6	2.7
PBMC B	CD56mid NK Cells	8	14.0	19.9
(Fig. 1)	CD56hi NK cells	9	1.3	2.0
	CD3+ T Cells		61.1	59.3
	CD8+CD45RA+ Tcells	5	14.3	12.5
	CD8+CD45RA- T Cells	6	21.9	26.1
	CD4+CD45RA+ Tcells	3	18.4	17.7
	CD4+CD45RA- T Cells	4	37.4	39.8

Table S2: Evidence for manua	I annotations of node	groups on SPADE tree.
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Group name	Reference	Evidence for	Evidence against
HSC	1	<ul> <li>Lin- CD45lo CD34+ CD38+ CD45RA- CD90- Ki67+</li> </ul>	
MPP	1	<ul> <li>Lin- CD45lo CD34+ CD38+ CD45RA- CD90- Ki67+</li> </ul>	Some CD19+ contamination
Pro-B	2	CD34+, CD38++     Clusters near Pre-B L cells	
Pre-B I	2	CD45lo CD34+ CD38+ CD19+ CD20-	Upsampled tree shows CD10-
Pre-B II	2	<ul> <li>CD45mid CD34het CD38+ CD19+ CD123- CD10+</li> </ul>	Some CD34+ contamination
Immature B	2	<ul> <li>CD45hi CD34- CD38+ CD19+ CD20+ CD123mid CD10+ IgM+</li> </ul>	
Mature B	2	<ul> <li>CD45hi CD34- CD38- CD19+ CD20+ CD123- CD10- IgMhet</li> </ul>	
IL3Ra+ mature B	2	CD45hi CD34- CD38- CD19+ CD20+     CD123+ CD10- IaMhet	
Plasma cell	3	CD45lo CD19+ CD20- CD38+++	
NK	4	<ul> <li>Directly observed as CD45hi CD45RA+ CD38+ CD3- CD19-</li> <li>Upsampled tree shows CD56+ CD7+ CD161+ CD16+</li> </ul>	
NKT	4	<ul> <li>CD3+ CD4- CD8-</li> <li>Upsampled tree shows CD161+.</li> </ul>	
Memory CD4 T	9	<ul> <li>CD45hi CD3+ CD19- CD33- CD4+ CD8- CD45RA-</li> </ul>	
Naive CD4 T	9	<ul> <li>CD45hi CD3+ CD19- CD33- CD4+ CD8- CD45RA+</li> </ul>	
Memory CD8 T	9	CD45hi CD3+ CD19- CD33- CD8+ CD4- CD45RA-	
Naive CD8 T	9	<ul> <li>CD45hi CD3+ CD19- CD33- CD8+ CD4- CD45RA+</li> </ul>	
CMP	1	<ul> <li>Lin- CD34+ CD38+ CD45RA- CD123-</li> </ul>	
Monoblast	5	CD45mid CD4lo CD11b- CD34- CD38+ CD33+ CD123lo CD14mid CD15mid CD16- CD13lo/mid CD117lo/- HLADR++	Lacks CD34
Monocyte	2	CD33+ CD14+ CD11b+ HLADB+	
Pro-monocyte	5	CD45mid/+ CD4mid CD11b- CD34- CD38mid CD33+ CD123lo CD14lo CD15lo/- CD16lo/- CD3mid CD117het HLADR++	
Pre-DC	6	CD45mid/+ CD45RA+ CD4+ CD11b- CD34- CD38mid CD33lo CD123lo CD14- CD15mid CD16- CD13- CD117mid HLADR-	
GMP	1	<ul> <li>CD45mid CD11b- CD34+ CD38+ CD45RA+ CD33mid CD123+ CD14- CD15- CD16- CD13+ CD117++ HLADR-</li> <li>Flt3L and TPO responsive</li> </ul>	
Plasmacytoid DC	7	CD38mid CD33mid CD11blo CD123++	Upsampled tree shows HLADR-
Myelocyte	2	CD15++ CD13- CD14lo CD16lo CD123lo CD33mid CD11bmid HLADR-	
Promyelocyte	2	<ul> <li>CD38mid, CD45mid, CD11b-, CD33mid</li> <li>Upsampled tree shows CD117mid, CD15mid, CD16-, CD13lo.</li> </ul>	Pro-myelocytes should be CD13+ according to Reference 2

CD38mid CD3- platelet	8	<ul> <li>CD38mid CD3- DNAlo CD45lo</li> <li>Upsampled tree shows some nodes are CD61+</li> <li>Nearly all nodes are basally pBtk high and all respond to TPO on Ki67, pBtk, and pS6</li> </ul>	
CD38- CD3- platelet	8	<ul> <li>CD38- CD3- DNAlo CD45lo</li> <li>A few nodes are basally pBtk high, but all respond to TPO on Ki67, pSrc, and pS6</li> </ul>	Upsampled tree shows no nodes are CD61+
CD38mid CD3mid platelet	8	<ul> <li>CD38mid CD3mid DNAlo CD45lo</li> <li>Upsampled tree shows some nodes are CD61+</li> <li>All nodes respond to TPO by upregulating Ki67 and downregulating pBtk</li> </ul>	Upsampled tree shows no nodes are CD61+
Erythroblast	2, 5	<ul> <li>DNAmid CD45lo CD38mid</li> <li>Upsampled tree shows CD235het, CD117mid</li> <li>Pro-erythroblasts should be CD117+ CD235- according to Reference 2</li> <li>These nodes generally respond to SCF on pSyk/Zap70 more than any other CD45lo cell type.</li> </ul>	
Erythrocyte	2	<ul> <li>DNAlo CD45- CD38-</li> <li>Upsampled tree shows CD235++, CD117lo</li> <li>Erythrocytes should be CD117- CD235+ according to Reference 2</li> <li>These nodes don't respond much to SCF</li> </ul>	

#### **References for Supplemental Table 2:**

- 1. Majeti et al. Identification of a hierarchy of multipotent hematopoietic progenitors in human cord blood. Cell Stem Cell (2007) vol. 1 (6) pp. 635-45
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- Crosby and Poole. Interaction of Bruton's tyrosine kinase and protein kinase Ctheta in platelets. Cross-talk between tyrosine and serine/threonine kinases. J Biol Chem (2002) vol. 277 (12) pp. 9958-65
- 9. Seder and Ahmed. Similarities and differences in CD4+ and CD8+ effector and memory T cell generation. Nat Immunol (2003) vol. 4 (9) pp. 835-42

# **Table S3:** Significant perturbation-induced signaling responses in healthybone marrow (one-sample *t*-test, P < 0.05 after Bonferroni correction)

					Marrow 1		Marrow 2		
				Marrow 1	Bonferroni-	Marrow 2	Bonferroni-		Dataset(s) with
				unadjusted t-	corrected t-test	unadjusted t-	corrected t-test	Direction	significant shift
Inhibitor	Stimulus	Antibody	Population	test P value	P value	test P value	P value	of Shift	(Bonferroni P < 0.05)
	BCR	168-pH3	CD11b- Monocytes	1.692E-0	6 7.310E-04	n.s.	n.s.	Up	Marrow 1 Only
	BCR	169-pP38	CD11bmid Monocytes	5.699E-0	6 2.462E-03	n.s.	n.s.	Up	Marrow 1 Only
	BCR	151-pERK1/2	Plasma cell	6.336E-0	6 2.737E-03	n.s.	n.s.	Up	Marrow 1 Only
	BCR	168-pH3	CD11bmid Monocytes	1.075E-0	5 4.645E-03	n.s.	n.s.	Up	Marrow 1 Only
	BCR	141-pPLCgamma2	Immature B	1.132E-0	5 4.892E-03	n.s.	n.s.	Up	Marrow 1 Only
	BCR	176-pCREB	Mature CD38lo B	1.344E-0	5 5.805E-03	9.884E-06	4.270E-03	Up	Both
	BCR	168-pH3	CD11bhi Monocytes	1.849E-0	5 7.986E-03	n.s.	n.s.	Up	Marrow 1 Only
	BCR	168-pH3	Mature CD38lo B	2.900E-0	5 1.253E-02	4.368E-05	1.887E-02	Up	Both
	BCR	141-pPLCgamma2	Mature CD38mid B	3.136E-0	5 1.355E-02	1.072E-04	4.630E-02	Up	Both
	BCR	141-pPLCgamma2	Mature CD38lo B	6./38E-0	5 2.911E-02	1.815E-06	7.841E-04	Up	Both
	BCR	176-pCREB	Mature CD38mid B	6.822E-0	5 2.947E-02	2.053E-05	2.053E-05 8.868E-0	Up	Both
	BCR	164-pSLP-76	Mature CD38lo B	7.248E-0	5 3.131E-02	5.0/1E-06	2.191E-03	Up	Both
	BCR	164-pSLP-76	Immature B	7.616E-0	5 3.290E-02	n.s.	n.s.	Up	Marrow 1 Only
	BCR	159-pSTAT3	Immature B	1.078E-0	4 4.656E-02	n.s.	n.s.	Up	Marrow 1 Only
	BCR	156-pZAP70/Syk	Immature B	n.s.	n.s.	1.210E-05	5.228E-03	Up	Marrow 2 Only
	BCR	156-pZAP70/Syk	Mature CD38lo B	n.s.	n.s.	3.440E-05	1.486E-02	Up	Marrow 2 Only
	BCR	156-pZAP70/Syk	Pre-B I	n.s.	n.s.	5.413E-05	2.338E-02	Up	Marrow 2 Only
	BCR	151-pERK1/2	HSC	n.s.	n.s.	6.131E-05	2.649E-02	Down	Marrow 2 Only
	BCR	151-pERK1/2	Mature CD38lo B	n.s.	n.s.	7.461E-05	3.223E-02	Up	Marrow 2 Only
	Flt3L	168-pH3	CD11b- Monocytes	6.431E-0	6 2.778E-03	n.s.	n.s.	Up	Marrow 1 Only
	Flt3L	176-pCREB	Plasmacytoid DC	3.973E-0	5 1.717E-02	n.s.	n.s.	Up	Marrow 1 Only
	Flt3L	152-Ki67	Plasma cell	n.s.	n.s.	2.290E-06	9.892E-04	Up	Marrow 2 Only
	Flt3L	176-pCREB	MPP	n.s.	n.s.	6.636E-06	2.867E-03	Up	Marrow 2 Only
	Flt3L	176-pCREB	HSC	n.s.	n.s.	1.763E-05	7.618E-03	Up	Marrow 2 Only
	Flt3L	151-pERK1/2	HSC	n.s.	n.s.	1.841E-05	7.955E-03	Up	Marrow 2 Only
	Flt3L	176-pCREB	CMP	n.s.	n.s.	4.204E-05	1.816E-02	Up	Marrow 2 Only
	GCSF	159-pSTAT3	HSC	2.145E-0	8 9.265E-06	2.796E-05	1.208E-02	Up	Both
	GCSF	169-pP38	CD11bmid Monocytes	1.192E-0	6 5.147E-04	n.s.	n.s.	Up	Marrow 1 Only
	GCSF	159-pSTAT3	CD11bhi Monocytes	3.222E-0	6 1.392E-03	3.479E-07	1.503E-04	Up	Both
	GCSF	159-pSTAT3	CD11bmid Monocytes	4.077E-0	6 1.761E-03	1.700E-05	7.342E-03	Up	Both
	GCSF	159-pSTAT3	Immature B	5.152E-0	6 2.226E-03	2.962E-06	1.280E-03	Up	Both
	GCSF	159-pSTAT3	CD11b- Monocytes	9.124E-0	6 3.942E-03	3.785E-05	1.635E-02	Up	Both
	GCSF	169-pP38	CD11bhi Monocytes	1.274E-0	5 5.502E-03	n.s.	n.s.	Up	Marrow 1 Only
	GCSF	159-pSTAT3	MPP	2.169E-0	5 9.368E-03	5.384E-06	2.326E-03	Up	Both
	GCSF	141-pPLCgamma2	Platelet	2.741E-0	5 1.184E-02	n.s.	n.s.	Up	Marrow 1 Only
	GCSF	168-pH3	CD11b- Monocytes	3.388E-0	5 1.463E-02	n.s.	n.s.	Up	Marrow 1 Only
	GCSF	169-pP38	CD11b- Monocytes	4.201E-0	5 1.815E-02	n.s.	n.s.	Up	Marrow 1 Only
	GCSF	176-pCREB	HSC	4.383E-0	5 1.893E-02	n.s.	n.s.	Up	Marrow 1 Only
	GCSF	174-pSrcFK	CD11bhi Monocytes	8.398E-0	5 3.628E-02	n.s.	n.s.	Up	Marrow 1 Only
	GCSF	175-pCrkL	Platelet	9.149E-0	5 3.953E-02	n.s.	n.s.	Up	Marrow 1 Only
	GCSF	174-pSrcFK	CD11bmid Monocytes	9.229E-0	5 3.987E-02	n.s.	n.s.	Up	Narrow 1 Only
	GCSF	150-pSTATS	IVIPP	9.839E-0	5 4.250E-02	n.s.	n.s.	Up	Iviarrow 1 Only
	GCSF	159-pSTAT3	Iviyelocyte	n.s.	n.s.	8.998E-07	3.887E-04	Up	Marrow 2 Only
	GCSF	159-pSTAT3	Megakaryocyte	n.s.	n.s.	3.067E-06	1.325E-03	Up	Marrow 2 Only
	GCSF	150-pSTAT5	HSC	n.s.	n.s.	3.941E-05	1.703E-02	Up	Marrow 2 Only
	GCSF	168-pH3	Immature B	n.s.	n.s.	7.642E-05	3.301E-02	Up	Marrow 2 Only
	GMCSF	150-pSTAT5	CD11bmid Monocytes	3.400E-0	6 1.469E-03	1.437E-05	6.210E-03	Up	Both
	GMCSF	150-pSTAT5	CD11bhi Monocytes	5.564E-0	6 2.404E-03	2.182E-05	9.426E-03	Up	Both
	GIVICSE	150-pSIAI5	CD11D- Monocytes	6.452E-0	b 2./87E-03	2.764E-05	1.194E-02	Up	BOTH
	GMCSF	150-pSTAT5	Plasmacytoid DC	8.801E-0	6 3.802E-03	5.661E-05	2.445E-02	Up	Both
	GIVICSE	150-pSIAI5		1.038E-0	5 4.485E-03	n.s.	n.s.	Up	Iviarrow 1 Only
	GMCSF	1/6-pCREB	Plasmacytoid DC	1.226E-0	5 5.295E-03	in.s.	n.s.	Up	Marrow 1 Only
	GMCSF	168-pH3	CD11b- Monocytes	1.358E-0	5 5.867E-03	n.s.	n.s.	Up	Marrow 1 Only
	GMCSF	150-pSTAT5	HSC	1.360E-0	5 5.875E-03	6.819E-05	2.946E-02	Up	Both
	GMCSF	150-pSTAT5	MPP	2.341E-0	5 1.011E-02	n.s.	n.s.	Up	Marrow 1 Only
	GMCSF	168-pH3	Plasmacytoid DC	2.724E-0	5 1.177E-02	2.073E-05	8.957E-03	Up	Both
	GMCSF	169-pP38	CD11bmid Monocytes	5.265E-0	5 2.274E-02	n.s.	n.s.	Up	Marrow 1 Only
	GMCSF	169-pP38	CD11bhi Monocytes	5.365E-0	5 2.318E-02	in.s.	n.s.	Up	Narrow 1 Only
	GMCSF	1/1-pBtk/ltk	IVIATURE CD38lo B	6.431E-0	5 2.778E-02	in.s.	n.s.	Up	Marrow 1 Only
	GMCSF	168-pH3	CD11bhi Monocytes	6.527E-0	5 2.820E-02	n.s.	n.s.	Up	Marrow 1 Only

					Marrow 1		Marrow 2		
				Marrow 1	Bonferroni-	Marrow 2	Bonferroni-	Direction	Dataset(s) with
Inhibitor	Stimulus	Antibody	Population	test P value	P value	test P value	P value	of Shift	(Bonferroni P < 0.05)
	GMCSF	159-pSTAT3	HSC	6.673E-05	2.883E-02	n.s.	n.s.	Up	Marrow 1 Only
	GMCSF	171-pBtk/Itk	HSC	9.202E-05	3.975E-02	n.s.	n.s.	Up	Marrow 1 Only
	GMCSF	168-pH3	Immature B	n.s.	n.s.	8.760E-06	3.784E-03	Up	Marrow 2 Only
	GMCSF	168-pH3	Myelocyte	n.s.	n.s.	2.526E-05	1.091E-02	Up	Marrow 2 Only
	GMCSF	152-Ki67	Plasma cell	n.s.	n.s.	4.128E-05	1.783E-02	Down	Marrow 2 Only
		169-pP38	Megakaryocyte	n.s.	n.s.	5.396E-05	2.331E-02	Up	Marrow 2 Only
	IENIad	108-PH3		6 7715 09	2 0255 05	8.310E-US	3.592E-02	Up	Marrow 1 Only
	IFNad	159-p31A13	Immature B	1 048F-07	4 527F-05	5 171F-06	2 234F-03	Un	Both
	IFNad	159-pSTAT3	Plasma cell	2.060E-06	8.897F-04	n.s.	n.s.	Un	Marrow 1 Only
	IFNad	150-pSTAT5	Naïve CD4+ T	3.833E-06	1.656E-03	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	150-pSTAT5	MEP	5.990E-06	2.588E-03	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	150-pSTAT5	Naïve CD8+ T	7.130E-06	3.080E-03	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	159-pSTAT3	CD11bhi Monocytes	7.224E-06	3.121E-03	2.705E-06	1.169E-03	Up	Both
	IFNad	159-pSTAT3	Mature CD4+ T	8.840E-06	3.819E-03	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	150-pSTAT5	Pre-B I	1.151E-05	4.971E-03	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	159-pSTAT3	CD11bmid Monocytes	1.235E-05	5.334E-03	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	159-pSTAT3	Mature CD8+ T	1.585E-05	6.847E-03	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	159-pSTAT3	CD11b- Monocytes	1.710E-05	7.389E-03	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	159-pSTAT3	NK	1.784E-05	7.706E-03	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	159-pSTATS	Naive CD8+ I	1.884E-05	8.138E-03	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	150-pSTAT5	Matura CD4 LT	2.047E-05	8.843E-03	n.s.	n.s.	Up	Marrow 1 Only
	IFNau	159-p3TAT3	Naïve CD4+ T	2.134L-03	9.217L-03	7 314F-05	3 160F-02	Un	Both
	IFNad	159-p51A13	Plasmacytoid DC	2.135E-05	1.085E-02	n s	5.100L-02	Un	Marrow 1 Only
	IFNad	150-pSTAT5	Mature CD8+ T	3.568E-05	1.542E-02	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	150-pSTAT5	Mature CD38mid B	3.757E-05	1.623E-02	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	174-pSrcFK	CD11bmid Monocytes	3.888E-05	1.680E-02	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	174-pSrcFK	CD11bhi Monocytes	4.042E-05	1.746E-02	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	169-pP38	CD11bmid Monocytes	4.084E-05	1.764E-02	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	159-pSTAT3	Pre-B II	4.147E-05	1.791E-02	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	174-pSrcFK	CD11b- Monocytes	4.191E-05	1.810E-02	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	150-pSTAT5	HSC	4.262E-05	1.841E-02	9.153E-05	3.954E-02	Up	Both
	IFNad	159-pSTAT3	MPP	4.635E-05	2.002E-02	3.724E-05	1.609E-02	Up	Both
	IFNad	159-pSTAT3	CMP	5.230E-05	2.259E-02	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	159-pSTAT3	Mature CD38lo B	5.586E-05	2.413E-02	n.s.	n.s.	Up	Marrow 1 Only
	IFINAU	150-PSTATS		5.800E-05	2.508E-02	n.s.	n.s.	Up	Marrow 1 Only
	IFNau IFNad	150-p3TAT3	MEP	6 270E-05	2.363E-02 2 709E-02	n s	n s	Up	Marrow 1 Only
	IFNad	174-nSrcFK	Pre-B II	7 249F-05	3 132F-02	n s	n.s.	Un	Marrow 1 Only
	IFNad	166-lkBalpha	MEP	9.622E-05	4.157E-02	n.s.	n.s.	Up	Marrow 1 Only
	IFNad	159-pSTAT3	Myelocyte	n.s.	n.s.	5.755E-06	2.486E-03	Up	Marrow 2 Only
	IFNad	159-pSTAT3	Megakaryocyte	n.s.	n.s.	9.913E-06	4.282E-03	Up	Marrow 2 Only
	IFNad	152-Ki67	Plasma cell	n.s.	n.s.	8.560E-05	3.698E-02	Down	Marrow 2 Only
	IL3	150-pSTAT5	Plasmacytoid DC	5.519E-06	2.384E-03	5.155E-05	2.227E-02	Up	Both
	IL3	176-pCREB	Plasmacytoid DC	9.437E-06	4.077E-03	n.s.	n.s.	Up	Marrow 1 Only
	IL3	150-pSTAT5	MEP	1.203E-05	5.196E-03	n.s.	n.s.	Up	Marrow 1 Only
	IL3	169-pP38	CD11bmid Monocytes	1.409E-05	6.085E-03	n.s.	n.s.	Up	Marrow 1 Only
	IL3	1/4-pSrcFK	CD11bmid Monocytes	1.826E-05	7.886E-03	n.s.	n.s.	Up	Marrow 1 Only
	IL3 II 2	159-pSTAT3	Immature B	2.392E-05	1.033E-02	n.s.	n.s.	Up	Marrow 1 Only
	IL3 II 2	174-pSrCFK	CD11b- Monocytes	3.0/1E-05	1.327E-02	n.s.	n.s.	Up	Marrow 1 Only
	IL3 II 3	159-µ31A15 168-nH3	Plasmacytoid DC	3.201E-03	1.417E-02	6 516F-06	11.5. 2 815F-03	Up	Both
	113	174-nSrcFK	CD11bbi Monocytes	4 314F-05	1.705E-02	n s	2.813L-03	Un	Marrow 1 Only
	113	172-pS6	Plasmacytoid DC	6.594F-05	2.848F-02	n.s.	n.s.	Un	Marrow 1 Only
	IL3	174-pSrcFK	Plasmacytoid DC	7.904E-05	3.415E-02	n.s.	n.s.	Up	Marrow 1 Only
	IL3	159-pSTAT3	Plasmacytoid DC	8.049E-05	3.477E-02	n.s.	n.s.	Up	Marrow 1 Only
	IL3	150-pSTAT5	CD11bmid Monocytes	n.s.	n.s.	2.847E-05	1.230E-02	Up	Marrow 2 Only
	IL3	150-pSTAT5	CD11b- Monocytes	n.s.	n.s.	2.998E-05	1.295E-02	Up	Marrow 2 Only
	IL3	150-pSTAT5	HSC	n.s.	n.s.	8.769E-05	3.788E-02	Up	Marrow 2 Only
	IL7	150-pSTAT5	Naïve CD4+ T	2.308E-06	9.970E-04	2.796E-05	1.208E-02	Up	Both
	IL7	150-pSTAT5	Naïve CD8+ T	3.347E-06	1.446E-03	3.046E-05	1.316E-02	Up	Both
	IL7	150-pSTAT5	Mature CD4+ T	7.826E-06	3.381E-03	1.458E-05	6.297E-03	Up	Both
	IL7	150-pSTAT5	Mature CD8+ T	9.972E-06	4.308E-03	3.868E-05	1.671E-02	Up	Both
		169-0438	CD11bbi Monocytes	1.454E-07	6.281E-05	n.s.	n.s.	Up	Marrow 1 Only
	105	160-0020		3.1/1E-U/ 1 210E 00	1.3/UE-U4	n s	n.s.		Marrow 1 Only
		166-1kBalnha	CD11bmid Monocytes	1 333E-06	5.093E-04	n s	n s	Down	Marrow 1 Only
	5		Correction monocytes	1.3231-00	J.7 1 JL-04			20001	

					Marrow 1		Marrow 2		
				Marrow 1	Bonferroni-	Marrow 2	Bonferroni-	Direction	Dataset(s) with
Inhibitor	Stimulus	Antibody	Population	test P value	P value	test P value	P value	of Shift	(Bonferroni P < 0.05)
	LPS	169-pP38	CD11b- Monocytes	7.408E-06	3.200E-03	n.s.	n.s.	Up	Marrow 1 Only
	LPS	166-IkBalpha	CD11b- Monocytes	1.254E-05	5.418E-03	n.s.	n.s.	Down	Marrow 1 Only
	LPS	168-pH3	CD11b- Monocytes	1.652E-05	7.139E-03	n.s.	n.s.	Up	Marrow 1 Only
	LPS	166-IkBalpha	MEP	5.279E-05	2.281E-02	n.s.	n.s.	Down	Marrow 1 Only
	LPS	168-pH3	CD11bmid Monocytes	9.318E-05	4.026E-02	n.s.	n.s.	Up	Marrow 1 Only
		153-PMAPKAPK2	CD11bmid Monocytes	n.s.	n.s.	2.561E-05	1.106E-02	Up	Marrow 2 Only
		153-piviAPKAPK2	CD11b- Monocytes	n s	11.S. n s	4.450E-05	1.925E-02	Up	Marrow 2 Only
	PMAiono	176-pCRFB	Naïve CD4+ T	1.146F-07	4.952F-05	1.873E-04	4.092F-04	Un	Both
	PMAiono	151-pERK1/2	Plasma cell	1.497E-07	6.468E-05	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	172-pS6	Mature CD4+ T	5.610E-07	2.423E-04	5.481E-05	2.368E-02	Up	Both
	PMAiono	168-pH3	CD11b- Monocytes	6.731E-07	2.908E-04	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	172-pS6	Naïve CD8+ T	7.120E-07	3.076E-04	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	168-pH3	Naïve CD4+ T	1.399E-06	6.046E-04	1.132E-06	4.890E-04	Up	Both
	PMAiono	174-pSrcFK	CD11bhi Monocytes	1.935E-06	8.359E-04	2.727E-05	1.178E-02	Down	Both
	PMAiono	172-pS6	Naïve CD4+ T	1.953E-06	8.439E-04	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	168-pH3	Plasmacytoid DC	1.968E-06	8.502E-04	3.073E-07	1.328E-04	Up	Both
	PIVIAIONO	165-PNFKB	Naive CD8+ I	4.226E-06	1.826E-03	n.s.	n.s.	Up	Marrow 1 Only
	PIVIAIONO	160 pD29	CD11bmid Monocytes	4.299E-06	1.857E-03	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	103-pr 36	Pre-B I	4.400L-00	2 081E-03	n s	n s	Un	Marrow 1 Only
	PMAiono	165-nNFkB	Naïve CD4+ T	5 104F-06	2.081E-03	4 986F-05	2 154F-02	Un	Both
	PMAiono	168-pH3	Mature CD4+ T	7.999E-06	3.456E-03	2.127E-07	9.189E-05	Up	Both
	PMAiono	168-pH3	Mature CD38lo B	8.514E-06	3.678E-03	1.752E-06	7.569E-04	Up	Both
	PMAiono	172-pS6	Plasmacytoid DC	9.519E-06	4.112E-03	6.476E-05	2.798E-02	Up	Both
	PMAiono	172-pS6	Pre-B II	1.047E-05	4.523E-03	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	168-pH3	CD11bhi Monocytes	1.103E-05	4.764E-03	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	169-pP38	CD11bhi Monocytes	1.263E-05	5.458E-03	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	176-pCREB	Plasmacytoid DC	1.268E-05	5.477E-03	6.021E-05	2.601E-02	Up	Both
	PMAiono	168-pH3	Immature B	1.320E-05	5.701E-03	1.334E-07	5.761E-05	Up	Both
	PMAiono	174-pSrcFK	CD11bmid Monocytes	1.444E-05	6.240E-03	n.s.	n.s.	Down	Marrow 1 Only
	PIVIAIONO	176-PCREB	Pre-BI	1.527E-05	6.596E-03	1.11/E-06	4.828E-04	Up	Both
	PMAiono	100-PHS 150-pSTAT2	Immature B	1.744E-05	7.555E-03	1.005E-05	4.599E-05	Up	Marrow 1 Only
	PMAiono	151-nFRK1/2	CD11h- Monocytes	1.754E-05	7.373E-03	n s	n.s.	Un	Marrow 1 Only
	PMAiono	169-pP38	Mature CD8+ T	2.028E-05	8.760E-03	n.s.	n.s.	Un	Marrow 1 Only
	PMAiono	176-pCREB	Mature CD38lo B	2.131E-05	9.207E-03	4.630E-06	2.000E-03	Up	Both
	PMAiono	172-pS6	CD11b- Monocytes	2.385E-05	1.030E-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	176-pCREB	Naïve CD8+ T	2.534E-05	1.095E-02	1.998E-05	8.632E-03	Up	Both
	PMAiono	151-pERK1/2	HSC	2.761E-05	1.193E-02	4.767E-08	2.060E-05	Up	Both
	PMAiono	169-pP38	Mature CD4+ T	2.874E-05	1.242E-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	172-pS6	Immature B	3.057E-05	1.320E-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	168-pH3	HSC	3.146E-05	1.359E-02	6.630E-06	2.864E-03	Up	Both
	PIVIAIONO	172-ps6	GIVIP	3.195E-05	1.380E-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	151-PERK1/2	CD11bmid Monocytes	3.472E-05	1.500E-02	n.s.	n.s.	Up Up	Marrow 1 Only
	PMAiono	151-nFRK1/2	Immature B	3.494E-05	1.505E-02	6 763E-05	2 922F-02	Un	Both
	PMAiono	168-pH3	MPP	3.892E-05	1.681E-02	1.161E-05	5.017E-03	Up	Both
	PMAiono	169-pP38	HSC	4.274E-05	1.846E-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	172-pS6	Mature CD8+ T	4.475E-05	1.933E-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	172-pS6	HSC	4.974E-05	2.149E-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	151-pERK1/2	CD11bhi Monocytes	5.029E-05	2.172E-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	176-pCREB	Mature CD38mid B	5.254E-05	2.270E-02	4.807E-06	2.077E-03	Up	Both
	PMAiono	172-pS6	Mature CD38lo B	5.596E-05	2.418E-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	151-pERK1/2	MEP	8.178E-05	3.533E-02	2.752E-07	1.189E-04	Up	Both
	PMAiono	153-pMAPKAPK2	Naive CD4+ T	8.230E-05	3.555E-02	2.142E-05	9.253E-03	Up	Both
	PIVIAIONO	172-p56	MEP	8.475E-05	3.661E-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAione	151-pERK1/2		8.490E-05	3.008E-02	1.25/E-05	5.429E-03	Up	Both
	PMAiono	176-pCRFR	Pre-B II	9 205F-05	3.777E-02	6 702F-06	2 895F-02	Un	Both
	PMAiono	169-pP38	Naïve CD8+ T	9.776F-05	4.223F-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	153-pMAPKAPK2	Naïve CD8+ T	1.005E-04	4.340E-02	1.696E-05	7.328E-03	Up	Both
	PMAiono	166-IkBalpha	CD11bhi Monocytes	1.119E-04	4.836E-02	n.s.	n.s.	Up	Marrow 1 Only
	PMAiono	151-pERK1/2	Mature CD4+ T	n.s.	n.s.	1.035E-08	4.470E-06	Up	Marrow 2 Only
	PMAiono	151-pERK1/2	Mature CD38lo B	n.s.	n.s.	8.653E-08	3.738E-05	Up	Marrow 2 Only
	PMAiono	151-pERK1/2	Naïve CD8+ T	n.s.	n.s.	3.374E-07	1.457E-04	Up	Marrow 2 Only
	PMAiono	151-pERK1/2	Mature CD8+ T	n.s.	n.s.	8.333E-07	3.600E-04	Up	Marrow 2 Only
	PMAiono	176-pCREB	CMP	n.s.	n.s.	1.213E-06	5.241E-04	Up	Marrow 2 Only

				Marrow 1	Marrow 1 Bonferroni-	Marrow 2	Marrow 2 Bonferroni-		Dataset(s) with
Inhibitor	Stimulus	Antibody	Population	unadjusted t- test P value	corrected t-test P value	unadjusted t- test P value	corrected t-test P value	Direction of Shift	significant shift (Bonferroni P < 0.05)
	PMAiono	151-pERK1/2	MPP	n.s.	n.s.	2.087E-06	9.015E-04	Up	Marrow 2 Only
	PMAiono	176-pCREB	MPP	n.s.	n.s.	2.639E-06	1.140E-03	Up	Marrow 2 Only
	PMAiono	151-pERK1/2	СМР	n.s.	n.s.	2.969E-06	1.283E-03	Up	Marrow 2 Only
	PMAiono	151-pERK1/2	NK	n.s.	n.s.	2.996E-06	1.294E-03	Up	Marrow 2 Only
	PMAiono	151-pERK1/2	Mature CD38mid B	n.s.	n.s.	3.107E-06	1.342E-03	Up	Marrow 2 Only
	PMAiono	153-рМАРКАРК2	GMP	n.s.	n.s.	3.757E-06	1.623E-03	Up	Marrow 2 Only
	PMAiono	151-pERK1/2	Pre-B II	n.s.	n.s.	4.605E-06	1.989E-03	Up	Marrow 2 Only
	PMAiono	151-pERK1/2	GMP	n.s.	n.s.	4.846E-06	2.094E-03	Up	Marrow 2 Only
	PMAiono	153-pMAPKAPK2	Mature CD4+ I	n.s.	n.s.	5.543E-06	2.394E-03	Up	Marrow 2 Only
	PIVIAIONO	153-PIVIAPKAPKZ	Mature CD4+ T	n.s.	n.s.	6.162E-06	2.662E-03	Up	Marrow 2 Only
	PMAiono	170-PCKED 153-pMADKADK2	Immature B	n s	n s	0.494E-00 7 814E-06	2.800E-03	Up	Marrow 2 Only
	PMAiono	156-n7AP70/Svk	Immature B	n s	n s	7.814E-00 8 498E-06	3.570E-03	Un	Marrow 2 Only
	PMAiono	176-pCREB	NK	n.s.	n.s.	8.620E-06	3.724E-03	Up	Marrow 2 Only
	PMAiono	168-pH3	Myelocyte	n.s.	n.s.	9.188E-06	3.969E-03	Up	Marrow 2 Only
	PMAiono	156-pZAP70/Syk	Pre-B I	n.s.	n.s.	1.064E-05	4.597E-03	Up	Marrow 2 Only
	PMAiono	172-pS6	Plasma cell	n.s.	n.s.	1.180E-05	5.098E-03	Up	Marrow 2 Only
	PMAiono	153-рМАРКАРК2	Mature CD38lo B	n.s.	n.s.	1.253E-05	5.412E-03	Up	Marrow 2 Only
	PMAiono	153-рМАРКАРК2	Pre-B II	n.s.	n.s.	1.373E-05	5.932E-03	Up	Marrow 2 Only
	PMAiono	141-pPLCgamma2	MPP	n.s.	n.s.	1.373E-05	5.932E-03	Up	Marrow 2 Only
	PMAiono	168-pH3	Mature CD8+ T	n.s.	n.s.	1.471E-05	6.355E-03	Up	Marrow 2 Only
	PMAiono	141-pPLCgamma2	MEP	n.s.	n.s.	1.561E-05	6.743E-03	Up	Marrow 2 Only
	PMAiono	156-pZAP70/Syk	Naïve CD4+ T	n.s.	n.s.	1.815E-05	7.839E-03	Up	Marrow 2 Only
	PIVIAIONO	153-PMAPKAPK2	Pre-B I	n.s.	n.s.	1.836E-05	7.930E-03	Up	Marrow 2 Only
	PIVIAIONO	176-PCREB	GIVIP	n.s.	n.s.	1.994E-05	8.612E-03	Up	Marrow 2 Only
	PMAiono	103-μΝΡΚΔ 153-ηΜΔΡΚΔΡΚ2	MEP	n s	n s	2.244E-05	9.090E-03	Un	Marrow 2 Only
	PMAiono	176-pCRFB	MEP	n.s.	n.s.	2.944F-05	1.272F-02	Un	Marrow 2 Only
	PMAiono	151-pERK1/2	Pre-B I	n.s.	n.s.	3.005E-05	1.298E-02	Up	Marrow 2 Only
	PMAiono	165-pNFkB	Plasmacytoid DC	n.s.	n.s.	3.139E-05	1.356E-02	Up	Marrow 2 Only
	PMAiono	153-рМАРКАРК2	Plasmacytoid DC	n.s.	n.s.	3.376E-05	1.459E-02	Up	Marrow 2 Only
	PMAiono	168-pH3	MEP	n.s.	n.s.	3.730E-05	1.612E-02	Up	Marrow 2 Only
	PMAiono	153-рМАРКАРК2	NK	n.s.	n.s.	3.809E-05	1.646E-02	Up	Marrow 2 Only
	PMAiono	141-pPLCgamma2	Pre-B I	n.s.	n.s.	4.106E-05	1.774E-02	Up	Marrow 2 Only
	PMAiono	153-рМАРКАРК2	HSC	n.s.	n.s.	4.125E-05	1.782E-02	Up	Marrow 2 Only
	PMAiono	141-pPLCgamma2	Plasmacytoid DC	n.s.	n.s.	4.217E-05	1.822E-02	Up	Marrow 2 Only
	PMAiono	153-pMAPKAPK2	Mature CD38mid B	n.s.	n.s.	4.244E-05	1.833E-02	Up	Marrow 2 Only
	PMAiono	176-pCREB	Mature CD8+ I	n.s.	n.s.	4.245E-05	1.834E-02	Up	Marrow 2 Only
	PIVIAIONO	152-KI67		n.s.	n.s.	4.810E-05	2.078E-02	Up	Marrow 2 Only
	PIVIAIONO	168-рнз 168-рнз		n.s.	n.s.	5.591E-05	2.415E-02 3.150E-02	Up	Marrow 2 Only
	PMAiono	156-n74P70/Svk	Mature CD4+ T	n s	n s	7.292L-03 8 141E-05	3.130L-02	Un	Marrow 2 Only
	PMAiono	172-pS6	MPP	n.s.	n.s.	9.625E-05	4.158E-02	Up	Marrow 2 Only
	PMAiono	153-pMAPKAPK2	CD11bmid Monocytes	n.s.	n.s.	1.027E-04	4.436E-02	Up	Marrow 2 Only
	PMAiono	152-Ki67	CD11bhi Monocytes	n.s.	n.s.	1.027E-04	4.438E-02	Up	Marrow 2 Only
	PMAiono	176-pCREB	Immature B	n.s.	n.s.	1.101E-04	4.758E-02	Up	Marrow 2 Only
	PMAiono	168-pH3	GMP	n.s.	n.s.	1.110E-04	4.797E-02	Up	Marrow 2 Only
	PMAiono	156-pZAP70/Syk	MPP	n.s.	n.s.	1.141E-04	4.928E-02	Up	Marrow 2 Only
	PVO4	169-pP38	CD11bmid Monocytes	3.162E-09	1.366E-06	5.029E-05	2.172E-02	Up	Both
	PVO4	159-pSTAT3	Immature B	2.703E-08	1.168E-05	2.431E-08	1.050E-05	Up	Both
	PVO4	164-pSLP-76	Immature B	4.046E-08	1.748E-05	8.025E-13	3.467E-10	Up	Both
	PVO4	174-pSrcFK	CD11bmid Monocytes	5.887E-08	2.543E-05	9.801E-07	4.234E-04	Up	Both
	PVO4	151-PERK1/2	Plasma cell	6.167E-08	2.664E-05	n.s.	n.s.	Up	Narrow 1 Only
		169-pP38	Immature P	9.895E-08	4.275E-05	7.292E-05	3.150E-02	Up	Both
		141-pPLCgammaz		1.154E-07	4.900E-05	2.709E-09	1.170E-00	Up	Both
	PVO4	170-perceb	CD11h- Monocytes	1.214L-07	6 213E-05	6.895E-06	2 979F-03	Un	Both
	PVO4	174-pSrcFK	CD11bhi Monocytes	1.542E-07	6.661E-05	2.563E-07	1.107E-04	Up	Both
	PVO4	169-pP38	CD11b- Monocytes	1.543E-07	6.664E-05	9.719E-05	4.199E-02	Up	Both
	PVO4	141-pPLCgamma2	Mature CD4+ T	1.784E-07	7.707E-05	2.818E-10	1.217E-07	Up	Both
	PVO4	164-pSLP-76	Naïve CD4+ T	2.178E-07	9.408E-05	4.722E-10	2.040E-07	Up	Both
	PVO4	164-pSLP-76	Naïve CD8+ T	2.428E-07	1.049E-04	1.812E-10	7.827E-08	Up	Both
	PVO4	141-pPLCgamma2	Mature CD8+ T	2.441E-07	1.055E-04	1.071E-09	4.626E-07	Up	Both
	PVO4	164-pSLP-76	Mature CD4+ T	3.434E-07	1.483E-04	4.516E-11	1.951E-08	Up	Both
	PVO4	141-pPLCgamma2	Naïve CD8+ T	3.762E-07	1.625E-04	1.791E-07	7.738E-05	Up	Both
	PVO4	164-pSLP-76	Plasma cell	4.887E-07	2.111E-04	8.924E-05	3.855E-02	Up	Both
	PV04	164-pSLP-76	LU11bmid Monocytes	5.981E-07	2.584E-04	2.260E-08	9.762E-06	Up	Both
	۲۷04	104-µSLP-70	iviature CD8+ I	0.096E-07	2.893E-04	2.110E-10	9.112F-08	υp	вош

				Marrow 1	Marrow 1 Bonferroni-	arrow 1 Arrow 2			Dataset(s) with
				unadjusted t-	corrected t-test	unadjusted t-	corrected t-test	Direction	significant shift
Inhibitor	Stimulus	Antibody	Population	test P value	P value	test P value	P value	of Shift	(Bonferroni P < 0.05)
	PVO4	164-pSLP-76	CD11b- Monocytes	6.742E-07	2.912E-04	3.772E-08	1.629E-05	Up	Both
	PVO4	159-pSTAT3	CD11bmid Monocytes	6.884E-07	2.974E-04	1.467E-06	6.335E-04	Up	Both
	PVO4	174-pSrCFK	Pre-B II	7.402E-07	3.197E-04	5.139E-06	2.220E-03	Up	Both
	PV04	164-pSI P-76	Pre-B I	9.722F-07	4.200F-04	3.711F-05	1.603F-02	Un	Both
	PVO4	141-pPLCgamma2	CD11b- Monocytes	1.157E-06	4.998E-04	1.953E-08	8.436E-06	Up	Both
	PVO4	141-pPLCgamma2	Plasma cell	1.167E-06	5.043E-04	2.685E-07	1.160E-04	Up	Both
	PVO4	175-pCrkL	Immature B	1.305E-06	5.638E-04	6.213E-08	2.684E-05	Up	Both
	PVO4	169-pP38	Mature CD8+ T	1.466E-06	6.335E-04	7.564E-05	3.268E-02	Up	Both
	PVO4	141-pPLCgamma2	Platelet	1.518E-06	6.559E-04	4.331E-06	1.871E-03	Up	Both
	PVO4	141-pPLCgamma2	Pre-B I	1.593E-06	6.882E-04	1.707E-09	7.373E-07	Up	Both
	PVO4	141-pPLCgamma2	CD11bmid Monocytes	2.126E-06	9.184E-04	3.644E-08	1.574E-05	Up	Both
		150-PSTATS		2.224E-06	9.608E-04	5.946E-06	2.509E-03	Up	Both
	PV04	164-pSLP-76	Myelocyte	2.648F-06	1.135E-03	1.490E-07	1.512F-08	Un	Both
	PVO4	169-pP38	Mature CD4+ T	2.784E-06	1.203E-03	3.612E-05	1.561E-02	Up	Both
	PVO4	175-pCrkL	CD11bmid Monocytes	3.069E-06	1.326E-03	3.864E-06	1.669E-03	Up	Both
	PVO4	174-pSrcFK	Naïve CD8+ T	3.079E-06	1.330E-03	2.323E-05	1.004E-02	Up	Both
	PVO4	169-pP38	Pre-B I	3.506E-06	1.514E-03	2.962E-05	1.280E-02	Up	Both
	PVO4	154-pSHP2	CD11bmid Monocytes	3.724E-06	1.609E-03	8.827E-07	3.813E-04	Up	Both
	PVO4	169-pP38	MEP	3.781E-06	1.634E-03	2.851E-05	1.232E-02	Up	Both
	PVO4	172-pS6	Naïve CD8+ T	3.845E-06	1.661E-03	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	151-pERK1/2	CD11b- Monocytes	3.854E-06	1.665E-03	6.794E-05	2.935E-02	Up	Both Marrow 1 Only
		164 pSLD 76		3.905E-00	1.713E-03	11.5. 5 1925 07	2 2205 04	Up	Roth
	PV04	141-nPI (gamma?	СМР	4.370L-00	2 141F-03	3.183L-07	2.239L-04	Un	Both
	PVO4	174-pSrcFK	Mature CD4+ T	5.492E-06	2.373E-03	9.698E-06	4.190E-03	Up	Both
	PVO4	151-pERK1/2	Immature B	5.593E-06	2.416E-03	5.000E-07	2.160E-04	Up	Both
	PVO4	151-pERK1/2	CD11bmid Monocytes	5.701E-06	2.463E-03	4.519E-05	1.952E-02	Up	Both
	PVO4	154-pSHP2	Pre-B I	5.785E-06	2.499E-03	1.965E-06	8.487E-04	Up	Both
	PVO4	141-pPLCgamma2	Myelocyte	5.867E-06	2.535E-03	3.794E-11	1.639E-08	Up	Both
	PVO4	150-pSTAT5	Naïve CD8+ T	5.997E-06	2.591E-03	6.373E-05	2.753E-02	Up	Both
	PVO4	154-pSHP2	CD11b- Monocytes	6.063E-06	2.619E-03	6.402E-07	2.766E-04	Up	Both
	PVO4	1/4-pSrCFK	Immature B	6.079E-06	2.626E-03	1.4/1E-06	6.354E-04	Up	Both
		164-pSLP-76		6.38E-06	2.809E-03	4.259E-00	1.840E-03	Up	Both Marrow 1 Only
	PV04	172-pS6	Mature CD4+ T	6.923E-00	2.991F-03	n.s.	n.s.	Un	Marrow 1 Only
	PVO4	175-pCrkL	Pre-B I	7.929E-06	3.425E-03	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	159-pSTAT3	CD11b- Monocytes	7.989E-06	3.451E-03	5.035E-06	2.175E-03	Up	Both
	PVO4	169-pP38	СМР	8.134E-06	3.514E-03	1.469E-05	6.345E-03	Up	Both
	PVO4	176-pCREB	Pre-B I	8.312E-06	3.591E-03	7.987E-06	3.450E-03	Up	Both
	PVO4	174-pSrcFK	Naïve CD4+ T	8.596E-06	3.714E-03	9.634E-06	4.162E-03	Up	Both
	PVO4	156-pZAP70/Syk	Naïve CD8+ T	8.794E-06	3.799E-03	4.722E-06	2.040E-03	Up	Both
	PVO4	175-pCrkL	Naïve CD8+ T	9.261E-06	4.001E-03	1.443E-05	6.232E-03	Up	Both
	PVO4	150-pSTAT5	CD11D- Monocytes	1.108E-05	4.785E-03	9.093E-06	3.928E-03	Up	Both Marrow 1 Only
	PVO4	176-pCREB	HSC	1.457F-05	6.296F-03	1.252F-05	5.410F-03	Up	Both
	PVO4	175-pCrkL	CD11b- Monocytes	1.562E-05	6.749E-03	1.084E-05	4.685E-03	Up	Both
	PVO4	141-pPLCgamma2	MPP	1.757E-05	7.590E-03	3.354E-08	1.449E-05	Up	Both
	PVO4	169-pP38	MPP	1.773E-05	7.660E-03	5.082E-05	2.195E-02	Up	Both
	PVO4	141-pPLCgamma2	MEP	1.868E-05	8.068E-03	3.541E-09	1.530E-06	Up	Both
	PVO4	154-pSHP2	Naïve CD8+ T	1.934E-05	8.355E-03	4.097E-06	1.770E-03	Up	Both
	PVO4	176-pCREB	Naïve CD8+ T	1.996E-05	8.621E-03	3.574E-05	1.544E-02	Up	Both
	PVO4	169-pP38	Naïve CD4+ T	2.016E-05	8.708E-03	n.s.	n.s.	Up	Marrow 1 Only
	PV04	129-021413		2.021E-05	8./30E-03	1.112E-06	4.805E-04	Up	BUTH Marrow 1 Ophy
		170-PCKED	Mature CD38lo B	2.024E-05	8 918F-03	n.s.	n s	Down	Marrow 1 Only
	PVO4	169-pP38	Pre-B II	2.081F-05	8.988F-03	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	169-pP38	NK	2.109E-05	9.113E-03	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	174-pSrcFK	Mature CD8+ T	2.116E-05	9.139E-03	3.883E-05	1.677E-02	Up	Both
	PVO4	141-pPLCgamma2	Erythroblast	2.192E-05	9.471E-03	6.549E-08	2.829E-05	Up	Both
	PVO4	175-pCrkL	Mature CD4+ T	2.211E-05	9.551E-03	8.297E-05	3.584E-02	Up	Both
	PVO4	164-pSLP-76	Plasmacytoid DC	2.231E-05	9.639E-03	1.796E-05	7.758E-03	Up	Both
	PVO4	175-pCrkL	CD11bhi Monocytes	2.342E-05	1.012E-02	2.612E-05	1.128E-02	Up	Both
	PVO4	154-pSHP2	Plasma cell	2.370E-05	1.024E-02	n.s.	n.s.	Up	Marrow 1 Only
	PV04	156-PZAP70/Syk	Iviature CD4+ T	2.406E-05	1.039E-02	1.852E-08	7.999E-06	Up	BOTH
		159-pCIKL	Plasma cell	2.499E-05	1.080E-02	n s	n s	Un	Marrow 1 Only
		1-22 621212		2.3136-03	1.00JL-0Z			<b>7</b> 2	

					Marrow 1		Marrow 2		
				Marrow 1	Bonferroni-	Marrow 2	Bonferroni-	Direction	Dataset(s) with
Inhibitor	Stimulus	Antibody	Population	test P value	P value	test P value	P value	of Shift	(Bonferroni P < 0.05)
	PVO4	172-pS6	Pre-B I	2.653E-05	1.146E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	141-pPLCgamma2	CD11bhi Monocytes	2.674E-05	1.155E-02	4.535E-06	1.959E-03	Up	Both
	PVO4	156-pZAP70/Syk	Mature CD8+ T	2.768E-05	1.196E-02	2.241E-07	9.681E-05	Up	Both
	PVO4	159-pSTAT3	Pre-B II	2.779E-05	1.200E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	150-pSTAT5	Mature CD8+ I	2.813E-05	1.215E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	156-PZAP70/SYK		2.901E-05	1.253E-02	4.693E-09	2.027E-06	Up Down	Both Marrow 1 Only
	PV04	171-pBtk/ttk 156-n7AP70/Svk	Naïve CD4+ T	2.943L-03	1.272L-02	4 420F-08	1 910F-05	Un	Roth
	PVO4	150-pSTAT5	Naïve CD4+ T	3.102E-05	1.340E-02	n.s.	n.s.	Un	Marrow 1 Only
	PVO4	175-pCrkL	Naïve CD4+ T	3.124E-05	1.350E-02	1.132E-04	4.888E-02	Up	Both
	PVO4	172-pS6	Naïve CD4+ T	3.247E-05	1.403E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	164-pSLP-76	HSC	3.349E-05	1.447E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	174-pSrcFK	Mature CD38lo B	3.412E-05	1.474E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	153-рМАРКАРК2	Plasma cell	3.441E-05	1.487E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	169-pP38	Plasma cell	3.453E-05	1.492E-02	9.889E-05	4.272E-02	Up	Both
	PVO4	141-pPLCgamma2	Pre-B II	3.560E-05	1.538E-02	4.256E-06	1.839E-03	Up	Both
	PVO4	150-pSTAT5	Plasmacytoid DC	3.620E-05	1.564E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	174-pSrcFK	Plasmacytoid DC	3.679E-05	1.589E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	141-pPLCgamma2	GMP	3.927E-05	1.696E-02	3.727E-05	1.610E-02	Up	Both
		154-pSHP2	Iviature CD4+ T	4.046E-05	1./48E-02	5.866E-08	2.534E-05	Up	BOTH
		150-pSTATS	CD11bbi Monocutor	4.223E-05	1.825E-02	5.0/5E-U5	2.452E-02	Up	Both
		154-µ3HP2 156-p7AP70/Syk	CD11bmid Monocytes	4.244E-05	1.835E-02	2 121E-06	9 162E-04	Un	Both
	PVO4	164-nSI P-76	Mature CD38mid B	4.543E-05	1.870E-02	8 637E-05	3 731F-02	Un	Both
	PVO4	154-pSHP2	Mature CD8+ T	4.776E-05	2.063E-02	1.561E-07	6.746E-05	Un	Both
	PVO4	153-pMAPKAPK2	Mature CD8+ T	4.877E-05	2.107E-02	5.477E-06	2.366E-03	Up	Both
	PVO4	169-pP38	Plasmacytoid DC	4.928E-05	2.129E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	151-pERK1/2	MEP	5.179E-05	2.237E-02	2.203E-06	9.518E-04	Up	Both
	PVO4	166-IkBalpha	MEP	5.197E-05	2.245E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	164-pSLP-76	MPP	5.364E-05	2.317E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	150-pSTAT5	Pre-B I	5.516E-05	2.383E-02	1.031E-04	4.453E-02	Up	Both
	PVO4	174-pSrcFK	Plasma cell	5.613E-05	2.425E-02	9.429E-06	4.073E-03	Up	Both
	PVO4	176-pCREB	MEP	6.198E-05	2.678E-02	2.175E-05	9.398E-03	Up	Both
	PVO4	154-pSHP2	Naïve CD4+ T	6.228E-05	2.690E-02	2.723E-06	1.176E-03	Up	Both
	PVO4	166-IKBalpha		6.554E-05	2.832E-02	n.s.	n.s.	Down	Marrow 1 Only
		1/4-µSICEN	нес	6.555E-05	2.652E-02 2.851E-02	11.5. 2 550F-07	1 102E-04	Un	Both
	PV04	159-nSTAT3	Mature CD4+ T	6.885E-05	2.974F-02	n.s.	n.s.	Un	Marrow 1 Only
	PVO4	172-pS6	MEP	7.401E-05	3.197E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	159-pSTAT3	нѕс	7.720E-05	3.335E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	151-pERK1/2	Pre-B I	7.891E-05	3.409E-02	1.067E-04	4.608E-02	Up	Both
	PVO4	153-рМАРКАРК2	Naïve CD8+ T	8.111E-05	3.504E-02	2.373E-05	1.025E-02	Up	Both
	PVO4	150-pSTAT5	Immature B	8.145E-05	3.518E-02	2.738E-06	1.183E-03	Up	Both
	PVO4	159-pSTAT3	Naïve CD8+ T	8.153E-05	3.522E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	153-рМАРКАРК2	Mature CD4+ T	8.308E-05	3.589E-02	5.376E-06	2.322E-03	Up	Both
	PVO4	151-pERK1/2	Myelocyte	8.862E-05	3.828E-02	2.720E-06	1.175E-03	Up	Both
		141-pPLCgamma2	IVIATURE CD38mid B	9.589E-05	4.143E-02	9.466E-05	4.089E-02	Up	BOTH
	PVO4	130-pSTATS		9.815E-05	4.240E-02	11.5. 2 200E 07	0 0000 05	Up Up	Roth
	PV04	172-nS6	HSC	3.330E-03	4.519E-02	2.209E-07	5.003E-U3	Un	Marrow 1 Only
	PVO4	172-p30 154-nSHP2	MEP	1.055E-04	4.488L-02	n.s.	n s	Un	Marrow 1 Only
	PVO4	153-pMAPKAPK2	Immature B	1.065E-04	4.601E-02	3.284E-07	1.419E-04	Up	Both
	PVO4	176-pCREB	MPP	1.079E-04	4.660E-02	2.947E-06	1.273E-03	Up	Both
	PVO4	176-pCREB	Plasma cell	1.080E-04	4.667E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	165-pNFkB	Pre-B I	1.086E-04	4.691E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	169-pP38	GMP	1.141E-04	4.928E-02	n.s.	n.s.	Up	Marrow 1 Only
	PVO4	156-pZAP70/Syk	Myelocyte	n.s.	n.s.	9.371E-09	4.048E-06	Up	Marrow 2 Only
	PVO4	151-pERK1/2	Mature CD4+ T	n.s.	n.s.	1.292E-07	5.582E-05	Up	Marrow 2 Only
	PVO4	169-pP38	Myelocyte	n.s.	n.s.	1.349E-07	5.828E-05	Up	Marrow 2 Only
	PVO4	156-pZAP70/Syk	CD11b- Monocytes	n.s.	n.s.	1.823E-07	7.876E-05	Up	Marrow 2 Only
	PV04	156-pZAP70/Syk	Pre-B I	n.s.	n.s.	1.912E-07	8.260E-05	Up	Marrow 2 Only
	PV04	141-PPLCgamma2	Findsmacytoid DC	11.5. n.c	n.s.	2.106E-07	9.098E-05	Up	Marrow 2 Only
		141-nPl Case	NK	n s	n s	3.00UE-U/	1.381E-04	Up Up	Marrow 2 Only
	PV04	168-pH3	Immature B	n.s.	n.s.	4.037E-07 5 145F-07	2.012E-04	Un	Marrow 2 Only
	PV04	154-pSHP2	Mvelocyte	n.s.	n.s.	7.221F-07	3.119F-04	Up	Marrow 2 Only
	PVO4	151-pERK1/2	Naïve CD4+ T	n.s.	n.s.	1.243E-06	5.371E-04	Up	Marrow 2 Only
	PVO4	164-pSLP-76	Erythroblast	n.s.	n.s.	1.333E-06	5.761E-04	Up	Marrow 2 Only

					Marrow 1		Marrow 2		
				Marrow 1	Bonferroni-	Marrow 2	Bonferroni-	Direction	Dataset(s) with
Inhibitor	Stimulus	Antibody	Population	test P value	P value	test P value	P value	of Shift	(Bonferroni P < 0.05)
	PVO4	174-pSrcFK	Myelocyte	n.s.	n.s.	1.638E-06	7.075E-04	Up	Marrow 2 Only
	PVO4	176-pCREB	СМР	n.s.	n.s.	2.078E-06	8.975E-04	Up	Marrow 2 Only
	PVO4	159-pSTAT3	Myelocyte	n.s.	n.s.	2.139E-06	9.243E-04	Up	Marrow 2 Only
	PVO4	151-pERK1/2	Naïve CD8+ T	n.s.	n.s.	2.996E-06	1.294E-03	Up	Marrow 2 Only
	PVO4	152-Ki67	Plasma cell	n.s.	n.s.	3.190E-06	1.378E-03	Up	Marrow 2 Only
	PVO4	154-pSHP2	Erythroblast	n.s.	n.s.	3.243E-06	1.401E-03	Up	Marrow 2 Only
	PVO4	164-pSLP-76	NK	n.s.	n.s.	3.592E-06	1.552E-03	Up	Marrow 2 Only
	PVO4	151-PERK1/2		n.s.	n.s.	6.707E-06	2.897E-03	Up	Marrow 2 Only
	PVO4	153-PIVIAPKAPK2	IVIEP Matura CD28la D	n.s.	n.s.	6.890E-06	2.976E-03	Up	Marrow 2 Only
	PVO4	164-PSLP-76	Immature CD3810 B	n.s.	n.s.	7.085E-06	3.001E-03	Up	Marrow 2 Only
		132-NI07	Myelocyte	n.s.	n s	7.122E-06	3.077E-03	Up	Marrow 2 Only
		153-pCIKE 153-pMΔΡΚΔΡΚ2	GMP	n.s.	n.s.	7.455E-00	3.452E-03	Un	Marrow 2 Only
		153-pΜΔΡΚΔΡΚ2	CD11hmid Monocytes	n.s.	n.s.	7.550E-00 8.616E-06	3 722E-03	Un	Marrow 2 Only
	PV04	168-nH3	Myelocyte	n.s.	n s	9.856F-06	4 258F-03	Un	Marrow 2 Only
	PVO4	154-nSHP2	CMP	n.s.	n.s.	9.963E-06	4.304F-03	Un	Marrow 2 Only
	PVO4	153-pMAPKAPK2	Pre-B I	n.s.	n.s.	1.076E-05	4.648F-03	Un	Marrow 2 Only
	PVO4	164-pSLP-76	Megakarvocyte	n.s.	n.s.	1.212E-05	5.234E-03	Up	Marrow 2 Only
	PVO4	176-pCREB	Mature CD4+ T	n.s.	n.s.	1.358E-05	5.868E-03	dD	Marrow 2 Only
	PVO4	156-pZAP70/Svk	MPP	n.s.	n.s.	1.512E-05	6.531E-03	Up	Marrow 2 Only
	PVO4	176-pCREB	Immature B	n.s.	n.s.	1.615E-05	6.975E-03	Up	Marrow 2 Only
	PVO4	153-рМАРКАРК2	Plasmacytoid DC	n.s.	n.s.	1.617E-05	6.987E-03	Up	Marrow 2 Only
	PVO4	153-рМАРКАРК2	CD11b- Monocytes	n.s.	n.s.	1.774E-05	7.663E-03	Up	Marrow 2 Only
	PVO4	156-pZAP70/Syk	CD11bhi Monocytes	n.s.	n.s.	1.850E-05	7.991E-03	Up	Marrow 2 Only
	PVO4	169-pP38	Immature B	n.s.	n.s.	1.995E-05	8.619E-03	Up	Marrow 2 Only
	PVO4	156-pZAP70/Syk	Pre-B II	n.s.	n.s.	2.167E-05	9.360E-03	Up	Marrow 2 Only
	PVO4	151-pERK1/2	Pre-B II	n.s.	n.s.	2.442E-05	1.055E-02	Up	Marrow 2 Only
	PVO4	153-рМАРКАРК2	CD11bhi Monocytes	n.s.	n.s.	2.471E-05	1.068E-02	Up	Marrow 2 Only
	PVO4	156-pZAP70/Syk	Mature CD38lo B	n.s.	n.s.	2.546E-05	1.100E-02	Up	Marrow 2 Only
	PVO4	164-pSLP-76	Platelet	n.s.	n.s.	2.567E-05	1.109E-02	Up	Marrow 2 Only
	PVO4	151-pERK1/2	Mature CD8+ T	n.s.	n.s.	3.121E-05	1.348E-02	Up	Marrow 2 Only
	PVO4	153-рМАРКАРК2	Myelocyte	n.s.	n.s.	3.176E-05	1.372E-02	Up	Marrow 2 Only
	PVO4	151-pERK1/2	Mature CD38lo B	n.s.	n.s.	3.225E-05	1.393E-02	Up	Marrow 2 Only
	PVO4	150-pSTAT5	Myelocyte	n.s.	n.s.	3.723E-05	1.608E-02	Up	Marrow 2 Only
	PVO4	156-pZAP70/Syk	MEP	n.s.	n.s.	3.940E-05	1.702E-02	Up	Marrow 2 Only
	PVO4	168-pH3	HSC	n.s.	n.s.	4.137E-05	1.787E-02	Up	Marrow 2 Only
	PVO4	156-pZAP70/Syk	Plasma cell	n.s.	n.s.	4.201E-05	1.815E-02	Up	Marrow 2 Only
	PVO4	168-pH3	MPP Matura CD201a D	n.s.	n.s.	4.804E-05	2.076E-02	Up	Marrow 2 Only
	PVO4	141-pPLCgammaz	Nature CD3810 B	n.s.	n.s.	6.4/3E-05	2.796E-02	Up	Marrow 2 Only
	PVO4	153-PIVIAPKAPKZ		n.s.	n.s.	0.518E-05	2.810E-02	Up	Marrow 2 Only
		169 pU2	Civip Plasmacutoid DC	n.c.	n.c.	7.209E-05	2 120E 02	Up	Marrow 2 Only
		169-nP38	Platelet	n.s.	n.s.	7.243E-05	3.125L-02	Un	Marrow 2 Only
	PV04	154-nSHP2	Pre-B II	n.s.	n s	8 312F-05	3 591F-02	Un	Marrow 2 Only
	PV04	176-pCRFB	Mature CD8+ T	n.s.	n.s.	8.805E-05	3.804F-02	Un	Marrow 2 Only
	PVO4	166-lkBalpha	Immature B	n.s.	n.s.	8,950E-05	3,866E-02	Up	Marrow 2 Only
	PVO4	153-pMAPKAPK2	Pre-B II	n.s.	n.s.	9.050E-05	3.910E-02	Up	Marrow 2 Only
	PVO4	156-pZAP70/Syk	CMP	n.s.	n.s.	9.233E-05	3.989E-02	Up	Marrow 2 Only
	PVO4	153-pMAPKAPK2	MPP	n.s.	n.s.	9.519E-05	4.112E-02	Up	Marrow 2 Only
	PVO4	156-pZAP70/Syk	Mature CD38mid B	n.s.	n.s.	9.588E-05	4.142E-02	Up	Marrow 2 Only
	PVO4	172-pS6	Plasma cell	n.s.	n.s.	1.065E-04	4.602E-02	Up	Marrow 2 Only
	PVO4	164-pSLP-76	GMP	n.s.	n.s.	1.111E-04	4.798E-02	Up	Marrow 2 Only
	SCF	159-pSTAT3	CD11bmid Monocytes	3.809E-06	1.646E-03	7.218E-06	3.118E-03	Up	Both
	SCF	159-pSTAT3	HSC	4.904E-06	2.119E-03	n.s.	n.s.	Up	Marrow 1 Only
	SCF	159-pSTAT3	CD11bhi Monocytes	5.638E-06	2.435E-03	1.816E-07	7.846E-05	Up	Both
	SCF	159-pSTAT3	CD11b- Monocytes	5.831E-06	2.519E-03	1.222E-05	5.279E-03	Up	Both
	SCF	159-pSTAT3	Plasma cell	8.590E-06	3.711E-03	n.s.	n.s.	Up	Marrow 1 Only
	SCF	159-pSTAT3	Mature CD4+ T	8.635E-06	3.730E-03	n.s.	n.s.	Up	Marrow 1 Only
	SCF	159-pSTAT3	NK	1.148E-05	4.960E-03	n.s.	n.s.	Up	Marrow 1 Only
	SCF	172-pS6	MEP	1.230E-05	5.314E-03	n.s.	n.s.	Up	Marrow 1 Only
	SCF	159-pSTAT3	Naïve CD4+ T	1.791E-05	7.739E-03	2.592E-05	1.120E-02	Up	Both
	SCF	159-pSTAT3	Naïve CD8+ T	2.172E-05	9.382E-03	n.s.	n.s.	Up	Marrow 1 Only
	SCF	174-pSrcFK	Pre-B II	2.211E-05	9.551E-03	n.s.	n.s.	Up	Marrow 1 Only
	SCF	1/2-p56	HSC	2.761E-05	1.193E-02	n.s.	n.s.	Up	Narrow 1 Only
	SCF	139-pSTAT3	Iviature CD8+ I	2.92/E-05	1.264E-02	n.s.	11.5.	Up	Narrow 1 Only
	SCE	171-pBtk/Itk	CD11bbi Moncostas	3.401E-05	1.469E-02	n.s.	n.s.	Up	Marrow 1 Only
	SCE	176-nCPEP	MED	3.703E-05	1.599E-02	n.s.	n s	Un	Marrow 1 Only
	50	110-PENEB	IVILI	4.0102-05	1.732E-02	11.3.	11.3.	96	

				Marrow	4	Marrow 1 Bonforroni-		Marrow 2	Marrow 2 Bonforroni		Datacot(c) with
				unadjus	ted t-	corrected t-tes	st เ	unadjusted t-	corrected t-test	Direction	significant shift
Inhibitor	Stimulus	Antibody	Population	test P va	alue	P value	t	test P value	P value	of Shift	(Bonferroni P < 0.05)
	SCF	159-pSTAT3	Immature B	4.3	362E-05	1.884E-0	02 r	n.s.	n.s.	Up	Marrow 1 Only
	SCF	174-pSrcFK	CD11b- Monocytes	5.1	155E-05	2.227E-(	02 r	1.5.	n.s.	Up	Marrow 1 Only
	SCF	171-pBtk/ltk	Mature CD38lo B	5.3	362E-05	2.316E-(	02 r	n.s.	n.s.	Up	Marrow 1 Only
	SCF	174-pSrcFK	CD11bmid Monocytes	7.2	256E-05	3.135E-(	02 r	n.s.	n.s.	Up	Marrow 1 Only
	SCF	172-pS6	Immature B	1.:	102E-04	4.759E-(	02 r	n.s.	n.s.	Up	Marrow 1 Only
	SCF	168-pH3	HSC	n.s.		n.s.		1.799E-05	7.771E-03	Up	Marrow 2 Only
	SCF	176-pCREB	MPP	n.s.		n.s.	_	3./26E-05	05 1.610E-0 05 2.336E-0	Up	Marrow 2 Only
	SCF	159-pSTATS	HSC	n s		n s	+	5.587E-05	2.550E-02 2.413E-02	Un	Marrow 2 Only
	TNFa	166-lkBalpha	MEP	1.5	780E-07	7.688E-0	05 r	n.s.	n.s.	Down	Marrow 1 Only
	TNFa	166-IkBalpha	CD11bhi Monocytes	2.8	835E-07	1.225E-0	04 r	n.s.	n.s.	Down	Marrow 1 Only
	TNFa	168-pH3	CD11b- Monocytes	7.8	874E-07	3.402E-0	04	3.278E-05	1.416E-02	Up	Both
	TNFa	166-IkBalpha	CD11bmid Monocytes	9.9	950E-07	4.298E-0	04	7.287E-05	3.148E-02	Down	Both
	TNFa	166-IkBalpha	Plasmacytoid DC	3.9	929E-06	1.697E-0	03 r	n.s.	n.s.	Down	Marrow 1 Only
		166-IkBalpha	CD11b- Monocytes	5.	285E-06	2.283E-0	03	3.239E-05	1.399E-02	Down	Both Marrow 1 Only
	TNFa	166-lkBalpha	CMP	5.0	233E-06	2.449E-0 2.693E-0	03 r	n.s.	n s	Down	Marrow 1 Only
	TNFa	169-pP38	CD11bmid Monocytes	1.0	064E-05	4.597E-(	03 r	n.s.	n.s.	Up	Marrow 1 Only
	TNFa	176-pCREB	Plasmacytoid DC	1.2	242E-05	5.367E-(	03 r	n.s.	n.s.	Up	Marrow 1 Only
	TNFa	168-pH3	Plasmacytoid DC	1.8	846E-05	7.974E-(	03	9.493E-06	4.101E-03	Up	Both
	TNFa	176-pCREB	Naïve CD4+ T	1.9	996E-05	8.622E-0	03 r	n.s.	n.s.	Up	Marrow 1 Only
	TNFa	171-pBtk/Itk	Mature CD38lo B	2.0	033E-05	8.784E-0	03 r	n.s.	n.s.	Up	Marrow 1 Only
	TNFa	172-pS6	MEP	2.3	385E-05	1.030E-(	02 r	n.s.	n.s.	Up	Marrow 1 Only
		172-pso 168-pH3	CD11b- Monocytes	2.0	001E-05	1.225E-0	021	7 530F-05	11.5. 2 253E-02	Up	Both
	TNFa	168-pH3	CD11bhi Monocytes	5.	167E-05	2.232E-(	02 r	1.s.	n.s.	Up	Marrow 1 Only
	TNFa	171-pBtk/Itk	HSC	5.2	297E-05	2.288E-0	02 r	n.s.	n.s.	Up	Marrow 1 Only
	TNFa	176-pCREB	MEP	5.5	567E-05	2.405E-0	02 r	n.s.	n.s.	Up	Marrow 1 Only
	TNFa	166-IkBalpha	NK	5.6	635E-05	2.434E-0	02 r	n.s.	n.s.	Down	Marrow 1 Only
	TNFa	176-pCREB	Pre-B I	5.9	987E-05	2.587E-0	02 r	n.s.	n.s.	Up	Marrow 1 Only
		169-pP38	CD11bhi Monocytes	6.4	439E-05	2.781E-0	02 r	n.s.	n.s.	Up	Marrow 1 Only
	TNFa	174-psrcFK 172-nS6	HSC	6.0	940E-05	2.948E-0 2 998E-0	02 r	n.s.	n s	Un	Marrow 1 Only
	TNFa	172-pS6	CD11bmid Monocytes	7.0	044E-05	3.043E-0	02 r	n.s.	n.s.	Up	Marrow 1 Only
	TNFa	166-IkBalpha	Pre-B I	7.	572E-05	3.271E-0	02 r	n.s.	n.s.	Down	Marrow 1 Only
	TNFa	168-pH3	MEP	7.	592E-05	3.280E-0	02 r	n.s.	n.s.	Up	Marrow 1 Only
	TNFa	176-pCREB	HSC	8.0	601E-05	3.716E-0	02	4.973E-05	2.148E-02	Up	Both
	TNFa	169-pP38	CD11b- Monocytes	9.8	802E-05	4.234E-(	02 r	n.s.	n.s.	Up	Marrow 1 Only
		171-pBtK/ltk	Mature CD8+ I	1.0	127E-04	4.653E-0	02 r	n.s.	n.s.	Up	Marrow 1 Only
	TNFa	169-pP38	Megakarvocyte	n.s.	1276-04	n.s.	021	5.881E-06	2.541E-03	Up	Marrow 2 Only
	TNFa	169-pP38	Erythroblast	n.s.		n.s.		6.402E-06	2.766E-03	Up	Marrow 2 Only
	TNFa	169-pP38	Myelocyte	n.s.		n.s.		9.237E-06	3.990E-03	Up	Marrow 2 Only
	TNFa	168-pH3	HSC	n.s.		n.s.		1.050E-05	4.537E-03	Up	Marrow 2 Only
	TNFa	168-pH3	Immature B	n.s.		n.s.		1.987E-05	8.585E-03	Up	Marrow 2 Only
		1/6-pCREB	MYPLOCUTO	n.s.		n.s.	+	2.347E-05	1.014E-02	Up	Marrow 2 Only
	TNFa	168-nH3	MPP	n s		n s	+	3.245E-05	1.402E-02 1 730F-02	Un	Marrow 2 Only
	TPO	150-pSTAT5	MEP	2.2	251E-05	9.725E-0	03 r	000E-05	n.s.	Up	Marrow 1 Only
	ТРО	169-pP38	CD11bmid Monocytes	4.1	194E-05	1.812E-0	02 r	n.s.	n.s.	Down	Marrow 1 Only
	ТРО	171-pBtk/Itk	HSC	6.8	863E-05	2.965E-0	02 r	n.s.	n.s.	Up	Marrow 1 Only
	ТРО	171-pBtk/Itk	Mature CD38lo B	8.9	917E-05	3.852E-(	02 r	n.s.	n.s.	Up	Marrow 1 Only
	ТРО	172-pS6	MEP	1.:	119E-04	4.835E-0	02 r	n.s.	n.s.	Up	Marrow 1 Only
		152-Ki67	Plasma cell	n.s.		n.s.	-	4.546E-06	1.964E-03	Up	Marrow 2 Only
	ΤΡΟ	150-pSTAT5 151-nFRK1/2	нос	n.s.		n s	-	7.000E-00 4 149E-05	3.312E-03	Un	Marrow 2 Only
	ТРО	169-pP38	Erythroblast	n.s.		n.s.		6.241E-05	2.696E-02	Down	Marrow 2 Only
DMSO	Unstim	166-IkBalpha	Myelocyte	n.s.		n.s.		3.950E-05	1.707E-02	Up	Marrow 2 Only
DMSO	Unstim	168-pH3	Immature B	n.s.		n.s.		5.184E-05	2.240E-02	Up	Marrow 2 Only
DMSO	Unstim	164-pSLP-76	Myelocyte	n.s.		n.s.	T	7.644E-05	3.302E-02	Up	Marrow 2 Only
DMSO	Unstim	152-Ki67	Plasma cell	n.s.	1045	n.s.	0.5	7.811E-05	3.374E-02	Up	Marrow 2 Only
Dasatinib	Unstim	174-pSrcFK	CD11bhi Monocytes	1.1	194E-07	5.159E-(	05	3.084E-07	1.332E-04	Down	Both
Dasatinib	Unstim	174-pSICEK	CD11b- Monocytes	1.	340E-07	5.821E-0	02	1.895E-06	8.185E-04	Down	Both
Dasatinih	Unstim	174-pSrcFK	Pre-B II	3.	968E-07	1.714F-(	04	4.139F-05	1.788F-02	Down	Both
Dasatinib	Unstim	168-pH3	CD11bhi Monocytes	7.	723E-07	3.336E-(	04	8.731E-05	3.772E-02	Down	Both
Dasatinib	Unstim	168-pH3	CD11bmid Monocytes	2.	508E-06	1.083E-0	03	6.421E-05	2.774E-02	Down	Both

Inhibitor	Stimulus	Antibody	Population	Marrow 1 unadjusted t-	N E C	Marrow 1 Bonferroni- corrected t-test	Marr unad	ow 2 Ijusted t-	Marr Boni corre	ow 2 ferroni- ected t-test	Direction	Dataset(s) with significant shift
Decetinih	Unctim		CD11hmid Monocutor		06		n c	r value	r va	lue	Down	(Bomerrow 1 Oply
Dasatinib	Unstim	169 pH 2	CD11b Monocytos	2.031E- 4 101E	00	1.130E-03	n.s.		11.S.		Down	Marrow 1 Only
Dasatinib	Unstim	150-p115	Immature B	4.101L- 9.164E-	00	2 959F-03	n c		n c		Down	Marrow 1 Only
Dasatinib	Unstim	171-nBtk/ltk	Mature CD38lo B	9.800F-	06	4 233E-03	n s		n s		Down	Marrow 1 Only
Dasatinib	Unstim	174-nSrcFK	Naïve CD8+ T	1 077F-	05	4.255E 05		5 849F-05	11.5.	2 527F-02	Down	Both
Dasatinib	Unstim	171-pBtk/ltk	HSC	1.438F-	05	6.214F-03	n.s.	5.0452 05	n.s.	2.5272.02	Down	Marrow 1 Only
Dasatinib	Unstim	169-pP38	CD11bhi Monocytes	1.740E-	05	7.519E-03	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Unstim	174-pSrcFK	Mature CD38lo B	1.854E-	05	8.008E-03	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Unstim	174-pSrcFK	Mature CD4+ T	2.659E-	05	1.149E-02	-	1.054E-04		4.552E-02	Down	Both
Dasatinib	Unstim	174-pSrcFK	Mature CD8+ T	2.669E-	05	1.153E-02		1.677E-05		7.245E-03	Down	Both
Dasatinib	Unstim	159-pSTAT3	HSC	2.773E-	05	1.198E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Unstim	174-pSrcFK	Naïve CD4+ T	3.631E-	05	1.569E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Unstim	174-pSrcFK	Plasmacytoid DC	3.782E-	05	1.634E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Unstim	159-pSTAT3	CD11bhi Monocytes	5.538E-	05	2.392E-02		2.816E-05		1.216E-02	Down	Both
Dasatinib	Unstim	174-pSrcFK	NK	5.718E-	05	2.470E-02		7.593E-06		3.280E-03	Down	Both
Dasatinib	Unstim	171-pBtk/Itk	Mature CD8+ T	6.339E-	05	2.738E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Unstim	172-pS6	CD11bmid Monocytes	7.153E-	05	3.090E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Unstim	159-pSTAT3	CD11bmid Monocytes	8.838E-	05	3.818E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Unstim	174-pSrcFK	Immature B	9.382E-	05	4.053E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Unstim	174-pSrcFK	Mature CD38mid B	9.551E-	05	4.126E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Unstim	141-pPLCgamma2	CD11b- Monocytes	9.771E-	05	4.221E-02		1.097E-04		4.737E-02	Down	Both
Dasatinib	Unstim	172-pS6	Megakaryocyte	1.062E-	04	4.588E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Unstim	169-pP38	Erythroblast	n.s.	r	n.s.		2.581E-06		1.115E-03	Down	Marrow 2 Only
Dasatinib	Unstim	156-pZAP70/Syk	CD11b- Monocytes	n.s.	r	n.s.		4.038E-06		1.744E-03	Down	Marrow 2 Only
Dasatinib	Unstim	1/4-pSrcFK	Myelocyte	n.s.	r	n.s.		1.954E-05		8.443E-03	Down	Marrow 2 Only
Dasatinib	Unstim	152-KI67	Plasma cell	n.s.	r	n.s.		3.859E-05		1.66/E-02	Up	Marrow 2 Only
Dasatinib	Unstim	156-PZAP70/SYK	CD11bmid Wonocytes	n.s.	r	n.s.		5.042E-05		2.178E-02	Down	Marrow 2 Only
Dasatinib	Unstim	104-µSLP-70	Endbroblast	n.s.	<u> </u>	n.s.		7 01 4E 0E		2.770E-02	Down	Marrow 2 Only
Dasatinib	Unstim	151_nERK1/2	HSC	n.s.		n.s.		8 365E-05		3.419E-02	Down	Marrow 2 Only
Dasatinib	Unstim	161-pLKK1/2	CD11bmid Monocytes	n.s.		n.s.		0.303L-03		3.014L-02	Down	Marrow 2 Only
Dasatinib	BCR	174-p3LF-70	CD11bhi Monocytes	1.3. 1.478F-	07	6 386E-05		1 122E-06		4 845E-04	Down	Both
Dasatinib	BCR	174-pSrcFK	CD11bmid Monocytes	1.630E-	07	7.041E-05		9.053E-06		3.911E-03	Down	Both
Dasatinib	BCR	174-pSrcFK	CD11b- Monocytes	3.388E-	07	1.464E-04		1.073E-04		4.636E-02	Down	Both
Dasatinib	BCR	168-pH3	CD11bhi Monocytes	7.394E-	07	3.194E-04		5.006E-05		2.163E-02	Down	Both
Dasatinib	BCR	174-pSrcFK	Pre-B II	8.452E-	07	3.651E-04	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	169-pP38	CD11bmid Monocytes	1.950E-	06	8.422E-04	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	168-pH3	CD11bmid Monocytes	2.367E-	06	1.023E-03		4.780E-05		2.065E-02	Down	Both
Dasatinib	BCR	168-pH3	CD11b- Monocytes	3.165E-	06	1.367E-03		1.086E-04		4.691E-02	Down	Both
Dasatinib	BCR	174-pSrcFK	Naïve CD8+ T	2.288E-	05	9.882E-03	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	171-pBtk/Itk	Mature CD38lo B	2.616E-	05	1.130E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	169-pP38	CD11bhi Monocytes	3.378E-	05	1.459E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	159-pSTAT3	CD11bhi Monocytes	5.339E-	05	2.306E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	174-pSrcFK	Mature CD4+ T	6.657E-	05	2.876E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	169-pP38	CD11b- Monocytes	6.952E-	05	3.003E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	171-pBtk/ltk	HSC	7.294E-	05	3.151E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	174-pSrcFK	Mature CD8+ T	8.073E-	05	3.488E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	174-pSrCFK	Mature CD38I0 B	8.667E-	05	3.744E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	159-pSTAT3	CD11bmid Monocytes	8.739E-	05	3.775E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	BCR	171 pD+k/l+k	Naive CD4+ 1	9.184E-	05	3.907E-02	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib		1/1-pBlk/llk	CD11b Monocytos	1.014E-	04	4.579E-02	11.S.		11.S.		Down	Marrow 1 Only
Dasatinib	BCR	171-pFLCgammaz	NK	1.037L-	04	4.738L-02	n c		n c		Down	Marrow 1 Only
Dasatinib	BCR	156-n74P70/Svk	CD11h- Monocytes	1.111L-		4.001L-02	11.3.	4 028E-05	11.5.	1 740F-02	Down	Marrow 2 Only
Dasatinib	BCR	169-nP38	Frythrohlast	n.s.		n.s.		5 609E-05		2 423F-02	Down	Marrow 2 Only
Dasatinib	BCR	151-pFRK1/2	Mature CD4+ T	n.s.		n.s.		9.935E-05		4.292F-02	Un	Marrow 2 Only
Dasatinib	Flt3L	174-pSrcFK	CD11bhi Monocytes	1.745E-	07	7.539E-05		2.787E-07		1.204E-04	Down	Both
Dasatinib	Flt3L	174-pSrcFK	CD11bmid Monocytes	1.980E-	07	8.551E-05		1.852E-06		8.000E-04	Down	Both
Dasatinib	Flt3L	174-pSrcFK	CD11b- Monocytes	5.056E-	07	2.184E-04		1.549E-05		6.691E-03	Down	Both
Dasatinib	Flt3L	174-pSrcFK	Pre-B II	5.267E-	07	2.275E-04		3.179E-05		1.373E-02	Down	Both
Dasatinib	Flt3L	168-pH3	CD11bhi Monocytes	8.431E-	07	3.642E-04		3.186E-05		1.377E-02	Down	Both
Dasatinib	Flt3L	169-pP38	CD11bmid Monocytes	2.770E-	06	1.197E-03	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Flt3L	168-pH3	CD11bmid Monocytes	3.577E-	06	1.545E-03		3.356E-05		1.450E-02	Down	Both
Dasatinib	Flt3L	174-pSrcFK	Naïve CD8+ T	8.633E-	06	3.729E-03		3.477E-05		1.502E-02	Down	Both
Dasatinib	Flt3L	168-pH3	CD11b- Monocytes	8.853E-	06	3.824E-03		1.074E-04		4.641E-02	Down	Both
Dasatinib	Flt3L	171-pBtk/Itk	Mature CD38lo B	8.930E-	06	3.858E-03	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Flt3L	169-pP38	CD11bhi Monocytes	1.589E-	05	6.864E-03	n.s.		n.s.		Down	Marrow 1 Only
Dasatinib	Flt3L	174-pSrcFK	Mature CD38lo B	1.843E-	05	7.963E-03	n.s.		n.s.		Down	Marrow 1 Only

				Marrow 1	Marrow 1 Bonferroni-	Marrow 2	Marrow 2 Bonferroni-	Direction	Dataset(s) with
Inhibitor	Stimulus	Antibody	Population	test P value	P value	test P value	P value	of Shift	(Bonferroni P < 0.05)
Dasatinib	Flt3L	171-pBtk/Itk	нѕс	1.912E-05	8.261E-03	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	Flt3L	159-pSTAT3	Immature B	1.928E-05	8.330E-03	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	Flt3L	174-pSrcFK	Mature CD4+ T	2.120E-05	9.158E-03	6.503E-05	2.809E-02	Down	Both
Dasatinib	Flt3L	174-pSrcFK	Mature CD8+ T	2.162E-05	9.341E-03	1.160E-05	5.013E-03	Down	Both
Dasatinib	Flt3L	174-pSrcFK	Naïve CD4+ T	2.723E-05	1.176E-02	9.114E-05	3.937E-02	Down	Both
Dasatinib	FIT3L FI+3I	171-pBtK/itk	Plasmacytoid DC	4.459E-05	1.926E-02 2.094E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	Flt3L	151-pERK1/2	Plasma cell	4.918E-05	2.125E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	Flt3L	159-pSTAT3	CD11bhi Monocytes	4.947E-05	2.137E-02	2.977E-06	1.286E-03	Down	Both
Dasatinib	Flt3L	174-pSrcFK	NK ,	5.489E-05	2.371E-02	6.076E-06	2.625E-03	Down	Both
Dasatinib	Flt3L	166-IkBalpha	MEP	7.770E-05	3.356E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	Flt3L	159-pSTAT3	CD11bmid Monocytes	8.325E-05	3.596E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	Flt3L	172-pS6	CD11bmid Monocytes	8.544E-05	3.691E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	FIT3L	176-PCREB	Plasmacytoid DC	9.450E-05	4.082E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	FILSL FIt3I	174-p3rcFK 159-nSTAT3		9.551E-05	4.120E-02 4.817E-02	n s	n s	Down	Marrow 1 Only
Dasatinib	Flt3L	169-pP38	Erythroblast	n.s.	n.s.	2.516E-06	1.087E-03	Down	Marrow 2 Only
Dasatinib	Flt3L	156-pZAP70/Syk	CD11b- Monocytes	n.s.	n.s.	7.938E-06	3.429E-03	Down	Marrow 2 Only
Dasatinib	Flt3L	151-pERK1/2	HSC	n.s.	n.s.	1.354E-05	5.849E-03	Up	Marrow 2 Only
Dasatinib	Flt3L	174-pSrcFK	Myelocyte	n.s.	n.s.	2.535E-05	1.095E-02	Down	Marrow 2 Only
Dasatinib	Flt3L	151-pERK1/2	MEP	n.s.	n.s.	5.428E-05	2.345E-02	Up	Marrow 2 Only
Dasatinib	Flt3L	156-pZAP70/Syk	CD11bmid Monocytes	n.s.	n.s.	6.920E-05	2.989E-02	Down	Marrow 2 Only
Dasatinib	FIt3L	176-pCREB	CD11bmid Monocytes	n.s.	n.s.	7.142E-05	3.085E-02	Down	Marrow 2 Only
Dasatinib	FIL3L FI+3I	174-pSrCFK	CD11bmid Monocytes	n.s.	n.s.	8.707E-05	3.701E-02	Down	Marrow 2 Only
Dasatinib	Flt3I	141-pPI Cgamma2	CD11b- Monocytes	n.s.	n.s.	1.031F-04	4.452F-02	Down	Marrow 2 Only
Dasatinib	IL7	174-pSrcFK	CD11bmid Monocytes	1.639E-07	7.079E-05	3.218E-06	1.390E-03	Down	Both
Dasatinib	IL7	174-pSrcFK	CD11bhi Monocytes	1.664E-07	7.189E-05	4.758E-07	2.055E-04	Down	Both
Dasatinib	IL7	174-pSrcFK	Pre-B II	3.725E-07	1.609E-04	4.139E-05	1.788E-02	Down	Both
Dasatinib	IL7	174-pSrcFK	CD11b- Monocytes	3.867E-07	1.670E-04	2.547E-05	1.100E-02	Down	Both
Dasatinib	IL7	169-pP38	CD11bmid Monocytes	5.724E-07	2.473E-04	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7	168-pH3	CD11bhi Monocytes	8.532E-07	3.686E-04	2.752E-05	1.189E-02	Down	Both
Dasatinib	11.7	150-pSTAT5	CD11bmid Monocytes	2.221E-06 3.450E-06	9.594E-04	n.s. 3 021E-05	n.s. 1 305E-02	Down	Narrow 1 Only
Dasatinib	117	166-lkBalpha	MFP	3.540E-06	1.529F-03	n.s.	n.s.	Un	Marrow 1 Only
Dasatinib	IL7	171-pBtk/Itk	Mature CD38lo B	3.668E-06	1.585E-03	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7	159-pSTAT3	Immature B	3.726E-06	1.610E-03	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7	150-pSTAT5	Naïve CD8+ T	4.010E-06	1.732E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	IL7	171-pBtk/Itk	HSC	5.205E-06	2.249E-03	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7	168-pH3	CD11b- Monocytes	5.647E-06	2.440E-03	8.078E-05	3.490E-02	Down	Both
Dasatinib	11.7	169-0P38		5.200E-06	2.678E-03	n.s.	n.s. 2 241E 02	Down	Narrow 1 Uniy
Dasatinib	11.7	150-pSTAT5	Mature CD4+ T	7.801F-06	3.370E-03	7.691F-05	2.241L-02 3.322F-02	Un	Both
Dasatinib	IL7	171-pBtk/Itk	Mature CD8+ T	1.528E-05	6.600E-03	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7	174-pSrcFK	Mature CD38lo B	1.546E-05	6.679E-03	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7	166-IkBalpha	CD11bhi Monocytes	1.715E-05	7.409E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	IL7	174-pSrcFK	Mature CD4+ T	1.773E-05	7.660E-03	9.963E-05	4.304E-02	Down	Both
Dasatinib	IL7	150-pSTAT5	Mature CD8+T	1.901E-05	8.213E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	IL/	174-pSrCFK	Mature CD8+ I	2.026E-05	8./52E-03	2.313E-05	9.992E-03	Down	Both Marrow 1 Only
Dasatinib	11.7	109-pF 38 174-nSrcFK	Naïve CD4+ T	2.287L-05	1.047F-02	n s	n s	Down	Marrow 1 Only
Dasatinib	IL7	174-pSrcFK	Plasmacytoid DC	3.340E-05	1.443E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7	171-pBtk/Itk	Mature CD4+ T	3.889E-05	1.680E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7	159-pSTAT3	CD11bhi Monocytes	4.407E-05	1.904E-02	6.739E-06	2.911E-03	Down	Both
Dasatinib	IL7	176-pCREB	Naïve CD4+ T	4.473E-05	1.932E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	IL7	174-pSrcFK	NK	5.077E-05	2.193E-02	1.043E-05	4.508E-03	Down	Both
Dasatinib	IL7	174-pSrcFK	Mature CD38mid B	5.098E-05	2.202E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinih	II 7	159-nSTAT2	HSC	5.120E-U5	2.214E-02 2.260F-02	n s	n s	Down	Marrow 1 Only
Dasatinih		166-lkBalpha	Plasmacytoid DC	6.003F-05	2.593F-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	IL7	159-pSTAT3	CD11bmid Monocytes	6.114E-05	2.641E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7	159-pSTAT3	Pre-B II	6.569E-05	2.838E-02	n.s.	n.s.	Down	, Marrow 1 Only
Dasatinib	IL7	171-pBtk/Itk	Naïve CD8+ T	7.491E-05	3.236E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7	166-IkBalpha	GMP	7.772E-05	3.357E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	L7	171-pBtk/Itk	MEP	9.523E-05	4.114E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	L/     7	172-p56	CD11bmid Monocytes	9.880E-05	4.268E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinih	11.7	172-pS6	Megakarvocvte	1.001F-04	4.272E-02 4.323F-02	n.s.	n.s.	Down	Marrow 1 Only
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				Marrow 1	Marrow 1 Bonferroni-	Marrow 2	Marrow 2 Bonferroni-		Dataset(s) with
Inhibitor	Stimulus	Antibody	Population	unadjusted t-	corrected t-test	unadjusted t-	corrected t-test	Direction	significant shift (Ronforroni R < 0.05)
Dasatinib	Sumulus II 7	166-lkBalpha	Population Pre-B I	1.030F-04	4.447F-02	n.s.	P value	Un	(Bomerrom P < 0.05) Marrow 1 Only
Dasatinib	IL7	172-pS6	Erythroblast	1.049E-04	4.533E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7	141-pPLCgamma2	CD11b- Monocytes	1.130E-04	4.880E-02	1.031E-04	4.452E-02	Down	Both
Dasatinib	IL7	174-pSrcFK	Immature B	1.153E-04	4.979E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	IL7 11.7	171-pBtk/ltk	CD11b- Monocytes	1.157E-04	4.999E-02	n.s.	n.s. 0 142E 04	Down	Marrow 1 Only
Dasatinib	IL7	156-pZAP70/Svk	CD11b- Monocytes	n.s.	n.s.	4.754E-06	2.054E-03	Down	Marrow 2 Only
Dasatinib	IL7	174-pSrcFK	GMP	n.s.	n.s.	3.740E-05	1.616E-02	Down	Marrow 2 Only
Dasatinib	IL7	156-pZAP70/Syk	CD11bmid Monocytes	n.s.	n.s.	5.467E-05	2.362E-02	Down	Marrow 2 Only
Dasatinib	IL7	174-pSrcFK	Myelocyte	n.s.	n.s.	6.282E-05	2.714E-02	Down	Marrow 2 Only
Dasatinib	IL7	152-Ki67	Plasma cell	n.s.	n.s.	6.520E-05	2.817E-02	Up	Marrow 2 Only
Dasatinib	11.7	176-pCRFB	CD11bmid Monocytes	n.s.	n.s.	9.240F-05	3.992F-02	Down	Marrow 2 Only
Dasatinib	IL7	176-pCREB	CD11b- Monocytes	n.s.	n.s.	9.676E-05	4.180E-02	Down	Marrow 2 Only
Dasatinib	IL7	175-pCrkL	GMP	n.s.	n.s.	1.064E-04	4.598E-02	Down	Marrow 2 Only
Dasatinib	PMAiono	176-pCREB	Naïve CD4+ T	5.097E-08	2.202E-05	2.578E-06	1.114E-03	Up	Both
Dasatinib	PMAiono	174-pSrcFK	CD11bhi Monocytes	1.408E-07	6.082E-05	6.552E-07	2.830E-04	Down	Both
Dasatinib	PIVIAIONO	172-p56	CD11bmid Monocytes	2.145E-07	9.267E-05	n.s. 6.421E-06	n.s. 2 774E-03	Down	Marrow 1 Only
Dasatinib	PMAiono	172-pS6	Naïve CD8+ T	3.149E-07	1.360E-04	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	151-pERK1/2	Plasma cell	6.242E-07	2.696E-04	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	174-pSrcFK	CD11b- Monocytes	6.922E-07	2.990E-04	1.052E-04	4.545E-02	Down	Both
Dasatinib	PMAiono	172-pS6	Naïve CD4+ T	9.902E-07	4.278E-04	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	174-pSrcFK	Pre-B II	1.210E-06	5.227E-04	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PMAiono	176-pCREB	Plasmacytoid DC	2.348E-06	1.014E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	172-pS6		4.107E-00	1.774L-03	0.208L-00	2.082L-03	Un	Marrow 1 Only
Dasatinib	PMAiono	172-pS6	Pre-B II	5.154E-06	2.226E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	172-pS6	Pre-B I	5.618E-06	2.427E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	172-pS6	CD11b- Monocytes	6.024E-06	2.602E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	176-pCREB	Mature CD38lo B	6.742E-06	2.912E-03	7.371E-06	3.184E-03	Up	Both
Dasatinib	PMAiono	176-PCREB	Pre-B I	8.716E-06	3.765E-03	7.219E-06	3.119E-03	Up	Both Marrow 1 Only
Dasatinib	PMAiono	172-pS6	Mature CD8+ T	8.997E-06	3.885L-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	176-pCREB	Naïve CD8+ T	9.497E-06	4.103E-03	2.480E-05	1.071E-02	Up	Both
Dasatinib	PMAiono	172-pS6	CD11bmid Monocytes	9.945E-06	4.296E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	172-pS6	GMP	1.068E-05	4.613E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	172-pS6	Mature CD38lo B	1.176E-05	5.079E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PIVIAIONO	176-PCKEB	Nature CD38mid B	1.4/1E-05 1 702E-05	0.354E-03	1.621E-05 6.841E-05	7.003E-03	Up	Both
Dasatinib	PMAiono	165-pNFkB	Naïve CD4+ T	1.832E-05	7.916E-03	5.206E-05	2.249E-02	Up	Both
Dasatinib	PMAiono	172-pS6	NK	2.033E-05	8.782E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	168-pH3	Mature CD38lo B	2.162E-05	9.340E-03	7.785E-06	3.363E-03	Up	Both
Dasatinib	PMAiono	168-pH3	Mature CD4+ T	2.981E-05	1.288E-02	2.204E-06	9.522E-04	Up	Both
Dasatinib	PMAiono	166-IkBalpha	Plasmacytoid DC	3.162E-05	1.366E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PMAiono	176-pCRFB	Pre-B II	3.539E-05	1.529E-02	3.217E-05	1.390F-02	Un	Both
Dasatinib	PMAiono	171-pBtk/ltk	Mature CD38lo B	3.849E-05	1.663E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PMAiono	168-pH3	Immature B	4.007E-05	1.731E-02	9.751E-07	4.212E-04	Up	Both
Dasatinib	PMAiono	151-pERK1/2	CD11b- Monocytes	4.029E-05	1.741E-02	7.527E-05	3.251E-02	Up	Both
Dasatinib	PMAiono	172-pS6	HSC	4.034E-05	1.743E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	176-pCREB	Mature CD4+ 1	4.330E-05	1.8/1E-02	1.048E-05	4.526E-03	Up	Both
Dasatinib	PMAiono	174-pSrcFK	Naïve CD8+ T	4.829E-05	2.086E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PMAiono	168-pH3	Plasmacytoid DC	5.402E-05	2.334E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PMAiono	171-pBtk/Itk	HSC	6.279E-05	2.713E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PMAiono	168-pH3	Naïve CD8+ T	6.611E-05	2.856E-02	5.733E-05	2.477E-02	Up	Both
Dasatinib	PMAiono	172-pS6	CD11bhi Monocytes	7.270E-05	3.141E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PIVIAIONO	151-pERK1/2	MEP	7.405E-05	3.199E-02	7.369E-05	3.183E-02	Up	Both Marrow 1 Only
Dasatinib	PMAiono	151-pERK1/2	HSC	1.059F-04	4.576F-02	7.401F-08	3.197F-05	Up	Both
Dasatinib	PMAiono	151-pERK1/2	CD11bhi Monocytes	1.119E-04	4.833E-02	1.052E-04	4.545E-02	Up	Both
Dasatinib	PMAiono	174-pSrcFK	Mature CD8+ T	1.139E-04	4.923E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PMAiono	151-pERK1/2	Mature CD4+ T	n.s.	n.s.	2.160E-08	9.331E-06	Up	Marrow 2 Only
Dasatinib	PMAiono	151-pERK1/2	Naïve CD4+ T	n.s.	n.s.	4.273E-08	1.846E-05	Up	Marrow 2 Only
Dasatinib	PMAiono	151-pERK1/2	MFP	n.s.	n.s.	2 608F-07	3.130E-05 1 127F-04	Un	Marrow 2 Only
Dasatinib	PMAiono	151-pERK1/2	Naïve CD8+ T	n.s.	n.s.	3.971E-07	1.715E-04	Up	Marrow 2 Only

				Marrow 1 unadjusted t-	Marrow 1 Bonferroni- corrected t-test	Marrow 2 unadjusted t-	Marrow 2 Bonferroni- corrected t-test	Direction	Dataset(s) with significant shift
Inhibitor	Stimulus	Antibody	Population	test P value	P value	test P value	P value	of Shift	(Bonferroni P < 0.05)
Dasatinib	PMAiono	151-pERK1/2	СМР	n.s.	n.s.	1.363E-06	5.887E-04	Up	Marrow 2 Only
Dasatinib	PMAiono	151-pERK1/2	Mature CD8+ T	n.s.	n.s.	1.364E-06	5.891E-04	Up	Marrow 2 Only
Dasatinib	PMAiono	151-pERK1/2	Mature CD38mid B	n.s.	n.s.	1.695E-06	7.320E-04	Up	Marrow 2 Only
Dasatinib	PMAiono	151-pERK1/2	Pre-B II	n.s.	n.s.	3.714E-06	1.604E-03	Up	Marrow 2 Only
Dasatinib	PMAiono	151-pERK1/2	NK	n.s.	n.s.	3.845E-06	1.661E-03	Up	Marrow 2 Only
Dasatinib	PIVIAIONO	151-PERK1/2		n.s.	n.s.	4.279E-06	1.848E-03	Up	Marrow 2 Only
Dasatinib	PIVIAIONO	151-PERK1/2 176-pCREB	CMP	n.s.	n.s.	7.979E-06	3.447E-03	Up	Marrow 2 Only
Dasatinib	PMAiono	151-nERK1/2		n s	n s	1 289E-05	4.303L-03	Un	Marrow 2 Only
Dasatinib	PMAiono	176-pCRFB	NK	n.s.	n.s.	1.357E-05	5.863E-03	Un	Marrow 2 Only
Dasatinib	PMAiono	165-pNFkB	Plasmacytoid DC	n.s.	n.s.	1.590E-05	6.870E-03	Up	Marrow 2 Only
Dasatinib	PMAiono	176-pCREB	MPP	n.s.	n.s.	1.718E-05	7.424E-03	Up	Marrow 2 Only
Dasatinib	PMAiono	165-pNFkB	Mature CD4+ T	n.s.	n.s.	2.677E-05	1.157E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	153-рМАРКАРК2	GMP	n.s.	n.s.	4.044E-05	1.747E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	174-pSrcFK	NK	n.s.	n.s.	4.381E-05	1.893E-02	Down	Marrow 2 Only
Dasatinib	PMAiono	165-pNFkB	Immature B	n.s.	n.s.	4.744E-05	2.049E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	176-pCREB	GMP	n.s.	n.s.	5.258E-05	2.272E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	176-pCREB	CD11bmid Monocytes	n.s.	n.s.	5.654E-05	2.443E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	174-pSrcFK	Myelocyte	n.s.	n.s.	5.841E-05	2.523E-02	Down	Marrow 2 Only
Dasatinib	PMAiono	176-pCREB	Mature CD8+ I	n.s.	n.s.	6.223E-05	2.688E-02	Up	Marrow 2 Only
Dasatinib	PIVIAIONO	152-KI07	CD11bni Wonocytes	n.s.	n.s.	6.269E-05	2.708E-02	Up	Marrow 2 Only
Dasatinib	PIVIAIONO	160-nD38	Myelocyte	n s	n s	6.586E-05	2.724E-02 2.845E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	153-pF 38	Immature B	n.s.	n s	6 796E-05	2.845E-02	Un	Marrow 2 Only
Dasatinib	PMAiono	176-pCREB	CD11b- Monocytes	n.s.	n.s.	6.850E-05	2.959E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	151-pERK1/2	Immature B	n.s.	n.s.	6.868E-05	2.967E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	176-pCREB	Immature B	n.s.	n.s.	7.644E-05	3.302E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	153-рМАРКАРК2	Mature CD8+ T	n.s.	n.s.	9.709E-05	4.194E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	153-рМАРКАРК2	Mature CD38lo B	n.s.	n.s.	1.049E-04	4.532E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	169-pP38	Mature CD4+ T	n.s.	n.s.	1.081E-04	4.668E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	152-Ki67	MEP	n.s.	n.s.	1.094E-04	4.728E-02	Up	Marrow 2 Only
Dasatinib	PMAiono	165-pNFkB	Pre-B I	n.s.	n.s.	1.106E-04	4.778E-02	Up	Marrow 2 Only
Dasatinib	PVO4	169-pP38	CD11bmid Monocytes	6.099E-08	2.635E-05	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PVO4	174-pSrcFK	CD11bhi Monocytes	2.512E-07	1.085E-04	4.902E-07	2.118E-04	Down	Both
Dasatinib	PVO4	1/4-pSrcFK	CD11bmid Monocytes	3.404E-07	1.4/1E-04	3.181E-06	1.3/4E-03	Down	Both
Dasatinib	PVO4	169-pP38	CD11b- Monocytes	1.150E-06	4.970E-04	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib		174-nSrcFK	CD11b- Monocytes	1.428L-00	6 188F-04	4 360F-05	1 883F-02	Down	Both
Dasatinib	PVO4	171-pBtk/ltk	Mature CD38lo B	1.625E-06	7.020E-04	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	174-pSrcFK	Pre-B II	1.820E-06	7.864E-04	4.625E-05	1.998E-02	Down	Both
Dasatinib	PVO4	168-pH3	CD11bhi Monocytes	1.879E-06	8.117E-04	3.597E-05	1.554E-02	Down	Both
Dasatinib	PVO4	171-pBtk/ltk	HSC	2.237E-06	9.663E-04	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	169-pP38	Mature CD8+ T	6.024E-06	2.603E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PVO4	171-pBtk/ltk	Mature CD8+ T	6.303E-06	2.723E-03	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	151-pERK1/2	Plasma cell	6.533E-06	2.822E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PVO4	168-pH3	CD11bmid Monocytes	9.831E-06	4.247E-03	4.562E-05	1.971E-02	Down	Both
Dasatinib	PVO4	169-pP38	MEP	1.218E-05	5.261E-03	2.130E-05	9.201E-03	Up	Both
Dasatinib	PVO4	169-pP38	Mature CD4+ T	1.524E-05	6.585E-03	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PVO4	1/4-pSrcFK	Naive CD8+ 1	1.826E-05	7.889E-03	5.485E-05	2.370E-02	Down	Both
Dasatinib	PVO4	1/1-pBtK/itk	Wature CD4+ I	1.960E-05	8.467E-03	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	160 pP28	CMD	2.554E-05	1 21/15 02	2 0005 05	1 2575 02	Up	Roth
Dasatinib		171-nBtk/ltk	CD11h- Monocytes	3.041L-03	1.314L-02	2.909L-03	1.237L-02	Down	Marrow 1 Only
Dasatinib	PV04	169-pP38	NK	3.206E-05	1.385E-02	n.s.	n.s.	Un	Marrow 1 Only
Dasatinib	PVO4	168-pH3	CD11b- Monocytes	3.288E-05	1.420E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	171-pBtk/ltk	CD11bmid Monocytes	3.545E-05	1.531E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	171-pBtk/ltk	Naïve CD8+ T	3.667E-05	1.584E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	166-IkBalpha	MEP	3.762E-05	1.625E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PVO4	171-pBtk/Itk	CD11bhi Monocytes	4.299E-05	1.857E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	171-pBtk/Itk	СМР	5.015E-05	2.167E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	174-pSrcFK	Mature CD4+ T	5.187E-05	2.241E-02	1.042E-04	4.501E-02	Down	Both
Dasatinib	PVO4	174-pSrcFK	Naïve CD4+ T	5.846E-05	2.525E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	174-pSrcFK	Mature CD8+ T	7.058E-05	3.049E-02	2.174E-05	9.390E-03	Down	Both
Dasatinib	PV04	169-pP38		7.153E-05	3.090E-02	8.196E-05	3.541E-02	Up	Both
Dasatinib	PVO4	171 pP+1/1+1	Piasma Cell	7.393E-05	3.194E-02	n.s.	11.5. n.c	Down	Marrow 1 Only
Dasatinih	PV04	176-pCRFR		7.458E-05	3.222E-U2	n s	n s		Marrow 1 Only
Dasatinih	PV04	171-nBtk/I+b		21/F-05	3.2/UE-U2	n s	n s	Down	Marrow 1 Only
Basatinin	1. VO+			0.3141-03	J.JJ2L-02			DOWI	indition 1 Only

					Marrow 1		Marrow 2		
				Marrow 1	Bonferroni-	Marrow 2	Bonferroni-		Dataset(s) with
Inhibitor	Stimulus	Antibody	Population	unadjusted t-	corrected t-test	unadjusted t-	corrected t-test	Direction of Shift	Significant shift (Bonferroni P < 0.05)
Dasatinib	PVO4	172-pS6	Megakarvocyte	8.577E-0	3.705E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	176-pCREB	HSC	8.601E-0	5 3.716E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PVO4	159-pSTAT3	Immature B	9.389E-0	5 4.056E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	169-pP38	Naïve CD8+ T	9.711E-0	5 4.195E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PVO4	169-pP38	Plasmacytoid DC	1.013E-0	4.375E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PVO4	174-pSrcFK	, Mature CD38lo B	1.030E-0	4.450E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	159-pSTAT3	CD11bhi Monocytes	1.100E-0	4.753E-02	1.373E-05	5.932E-03	Down	Both
Dasatinib	PVO4	171-pBtk/ltk	Plasmacytoid DC	1.115E-0	4 4.818E-02	n.s.	n.s.	Down	Marrow 1 Only
Dasatinib	PVO4	150-pSTAT5	Plasmacytoid DC	1.130E-0	4 4.882E-02	n.s.	n.s.	Up	Marrow 1 Only
Dasatinib	PVO4	141-pPLCgamma2	MEP	n.s.	n.s.	8.643E-07	3.734E-04	Up	Marrow 2 Only
Dasatinib	PVO4	169-pP38	Myelocyte	n.s.	n.s.	1.048E-06	4.528E-04	Up	Marrow 2 Only
Dasatinib	PVO4	151-pERK1/2	MEP	n.s.	n.s.	2.382E-06	1.029E-03	Up	Marrow 2 Only
Dasatinib	PVO4	164-pSLP-76	Myelocyte	n.s.	n.s.	2.510E-06	1.084E-03	Up	Marrow 2 Only
Dasatinib	PVO4	152-Ki67	Plasma cell	n.s.	n.s.	3.819E-06	1.650E-03	Up	Marrow 2 Only
Dasatinib	PVO4	153-рМАРКАРК2	GMP	n.s.	n.s.	5.369E-06	2.319E-03	Up	Marrow 2 Only
Dasatinib	PVO4	141-pPLCgamma2	Myelocyte	n.s.	n.s.	6.451E-06	2.787E-03	Up	Marrow 2 Only
Dasatinib	PVO4	169-pP38	Erythroblast	n.s.	n.s.	9.096E-06	3.930E-03	Up	Marrow 2 Only
Dasatinib	PVO4	164-pSLP-76	Immature B	n.s.	n.s.	1.166E-05	5.036E-03	Up	Marrow 2 Only
Dasatinib	PVO4	153-рМАРКАРК2	CD11bmid Monocytes	n.s.	n.s.	1.348E-05	5.823E-03	Up	Marrow 2 Only
Dasatinib	PVO4	141-pPLCgamma2	Plasmacytoid DC	n.s.	n.s.	1.476E-05	6.377E-03	Up	Marrow 2 Only
Dasatinib	PVO4	153-рМАРКАРК2	Mature CD8+ T	n.s.	n.s.	1.775E-05	7.667E-03	Up	Marrow 2 Only
Dasatinib	PVO4	151-pERK1/2	Mature CD4+ T	n.s.	n.s.	2.207E-05	9.533E-03	Up	Marrow 2 Only
Dasatinib	PVO4	153-рМАРКАРК2	CD11b- Monocytes	n.s.	n.s.	2.386E-05	1.031E-02	Up	Marrow 2 Only
Dasatinib	PVO4	141-pPLCgamma2	СМР	n.s.	n.s.	2.412E-05	1.042E-02	Up	Marrow 2 Only
Dasatinib	PVO4	176-pCREB	MPP	n.s.	n.s.	3.013E-05	1.302E-02	Up	Marrow 2 Only
Dasatinib	PVO4	174-pSrcFK	NK	n.s.	n.s.	3.187E-05	1.377E-02	Down	Marrow 2 Only
Dasatinib	PVO4	153-рМАРКАРК2	MEP	n.s.	n.s.	3.445E-05	1.488E-02	Up	Marrow 2 Only
Dasatinib	PVO4	172-pS6	Plasma cell	n.s.	n.s.	4.995E-05	2.158E-02	Down	Marrow 2 Only
Dasatinib	PVO4	169-pP38	Megakaryocyte	n.s.	n.s.	5.314E-05	2.296E-02	Up	Marrow 2 Only
Dasatinib	PVO4	156-pZAP70/Syk	CD11b- Monocytes	n.s.	n.s.	5.943E-05	2.568E-02	Down	Marrow 2 Only
Dasatinib	PVO4	153-рМАРКАРК2	CD11bhi Monocytes	n.s.	n.s.	6.167E-05	2.664E-02	Up	Marrow 2 Only
Dasatinib	PVO4	153-рМАРКАРК2	Pre-B I	n.s.	n.s.	6.388E-05	2.759E-02	Up	Marrow 2 Only
Dasatinib	PVO4	153-рМАРКАРК2	Mature CD4+ T	n.s.	n.s.	8.364E-05	3.613E-02	Up	Marrow 2 Only
Dasatinib	PVO4	153-рМАРКАРК2	Plasmacytoid DC	n.s.	n.s.	1.048E-04	4.528E-02	Up	Marrow 2 Only
Dasatinib	PVO4	174-pSrcFK	GMP	n.s.	n.s.	1.085E-04	4.687E-02	Down	Marrow 2 Only
Dasatinib	PVO4	151-pERK1/2	CMP	n.s.	n.s.	1.148E-04	4.959E-02	Up	Marrow 2 Only
JAKi	Unstim	168-pH3	CD11bhi Monocytes	5.032E-0	5 2.174E-03	n.s.	n.s.	Down	Marrow 1 Only
JAKi	Unstim	168-pH3	CD11bmid Monocytes	1.069E-0	5 4.618E-03	n.s.	n.s.	Down	Marrow 1 Only
JAKi	Unstim	168-pH3	CD11b- Monocytes	1.464E-0	5 6.325E-03	n.s.	n.s.	Down	Marrow 1 Only
JAKi	Unstim	174-pSrcFK	CD11bhi Monocytes	n.s.	n.s.	2.195E-05	9.485E-03	Down	Marrow 2 Only
JAKi	Unstim	151-pERK1/2	Mature CD4+ T	n.s.	n.s.	4.792E-05	2.070E-02	Up	Marrow 2 Only
JAKi	Unstim	174-pSrcFK	CD11bmid Monocytes	n.s.	n.s.	8.519E-05	3.680E-02	Down	Marrow 2 Only
JAKi	GCSF	168-pH3	CD11bhi Monocytes	6.901E-0	6 2.981E-03	n.s.	n.s.	Down	Marrow 1 Only
JAKi	GCSF	168-pH3	CD11bmid Monocytes	1.628E-0	5 7.034E-03	n.s.	n.s.	Down	Marrow 1 Only
JAKi	GCSF	168-pH3	CD11b- Monocytes	3.209E-0	5 1.386E-02	n.s.	n.s.	Down	Marrow 1 Only
JAKi	GCSF	159-pSTAT3	HSC	5.612E-0	5 2.424E-02	n.s.	n.s.	Down	Marrow 1 Only
JAKi	GCSF	175-pCrkL	Megakaryocyte	9.351E-0	5 4.040E-02	n.s.	n.s.	Up	Marrow 1 Only
JAKi	GCSF	169-pP38	CD11bmid Monocytes	9.849E-0	5 4.255E-02	n.s.	n.s.	Up	Marrow 1 Only
JAKi	GCSF	152-Ki67	Plasma cell	n.s.	n.s.	6.182E-06	2.670E-03	Up	Marrow 2 Only
JAKi	GCSF	151-pERK1/2	Mature CD4+ T	n.s.	n.s.	2.512E-05	1.085E-02	Up	Marrow 2 Only
JAKi	GCSF	151-pERK1/2	Naïve CD4+ T	n.s.	n.s.	5.705E-05	2.464E-02	Up	Marrow 2 Only
JAKi	GCSF	165-pNFkB	Erythroblast	n.s.	n.s.	7.350E-05	3.175E-02	Down	Marrow 2 Only