SUPPLEMENTAL DATA

Table S1 CyTOF antibody Panel

Cell identify markers				
Antibodies	Clone	Metal lab		
Anti-mouse Ly6G	IA8	141Pr		
Anti-mouse CD11c	N418	142Nd		
Anti-mouse Ly6C	H1.4	162Dy		
Anti-mouse CD45	30-F11	147Sm		
Anti-mouse CD11b	M1/70	148Nd		
Anti-mouse B220	RA3-6B2	176Yb		
Anti-mouse CD25	3C7	151Eu		
Anti-mouse CD3	145-2C11	165Ho		
Anti-mouse F4/80	BM8	159Tb		
Anti-mouse CD45RB	C363.16A	145Nd		
Anti-mouse CD8	53-6.7	168Er		
Anti-mouse TCRβ	H57-597	169Tm		
Anti-mouse CD49b	HMa2	170Er		
Anti-mouse CD44	IM7	150Nd		
Anti-mouse CD4	RM4-5	172Yb		
DNA staining		191/193Ir		
Live/dead sating		195Pt		
Barcoding		102-110Pd		
	Phosphorylation markers			
Anti-mouse p4E-BP1	236B4	149Sm		
Anti-mouse pAKT	D9E	152Sm		
Anti-mouse pS6	S235/S236	175Lu		
Anti-mouse pPLCg2	K86-889.37	144Nd		
Anti-mouse pEGFR	D7A5	146Nd		
Anti-mouse pStat1	4a	153Eu		

Anti-mouse pStat3	4	158Gd
Anti-mouse pStat4	38	174Yb
Anti-mouse p38	T180/Y182	156Gd
Anti-mouse pERK1/2	D13.14.4E	171Yb

Table S2 Identified immune cells

Cell Types	Defined Markers
Monocyte derived macrophages	CD45 ^{hi} CD11b ⁺
(MoDMs)	
Microglia derived macrophages (MiDMs)	CD45 ^{lo} CD11b ⁺
DCs	CD45 ⁺ Cd11C ⁺
Monocytes	CD45 ⁺ Ly6G ⁻ Ly6C ⁺
B cells	CD45 ⁺ B220 ⁺
CD8 T cells	CD45 ⁺ CD3 ⁺ CD8 ⁺
CD4 T cells	CD45 ⁺ CD3 ⁺ CD4 ⁺
Neutrophils	CD45 ⁺ Ly6G ⁺ Ly6C ⁻
CD4 T _{EM}	CD3 ⁺ CD4 ⁺ CD44 ⁺
СD8 ТЕМ	CD3 ⁺ CD8 ⁺ CD44 ⁺
NK cells	CD45 ⁺ CD49 ⁺

Table S3 The changed phosphorylation status of immune cells in ischemic	;
hemisphere	

Phosphorylation Markers	Immune cell type have altered expression
	level
pEGFR	MoDMs
p4E-BP1	CD4 T cells
pStat4	CD3 T cells , CD4 T cells
pStat1	MoDMs
pERK1	CD4 T cells
pS6	CD3 T cells , CD4 T cells
pStat3	CD3 T cells , CD4 T cells
pPLcg2	MoDMs
pAKT	CD4 T cells, MoDMs, DCs
pP38	CD3 T cells , CD4 T cells

Supplemental Figure



Figure S1. Characterization of post-stroke immune cell populations. Variations in the numbers of distinct immune cells in the ischemic brain hemisphere were noted (compared to the sham group, ***P<0.001).



Figure S2. Analysis of blood immune cells using mass cytometry: SPADE representation (top), event plots (bottom left), and viSNE visualization (bottom right).



Figure S3. Analysis of spleen immune cells via mass cytometry: SPADE representation (top), event plots (bottom left), and viSNE visualization (bottom right).



Figure S4. Heterogeneity in lipid metabolism of Lcp1^{high} and Lcp1^{low} macrophages and monocytes in post-stroke brains. (A-B) UMAP visualization illustrating clusters of cells isolated from ipsilateral (IL) hemispheres at 24 h and 48 h post-MCAo. (C) UMAP

representation highlighting specific markers for each identified cell cluster. (D) Categorization of Lcp1^{high} and Lcp1^{low} macrophages and monocytes at 24 h and 48 h post-tMCAO. (E) Box plots depicting expression levels related to lipid metabolism pathways in Lcp1[^]high and Lcp1[^]low macrophages and monocytes, in reference to Fig. 1D.