

# Supplementary information

For

**Hypoxia-mediated impaired erythrocyte Lands' Cycle is pathogenic for  
sickle cell disease**

by

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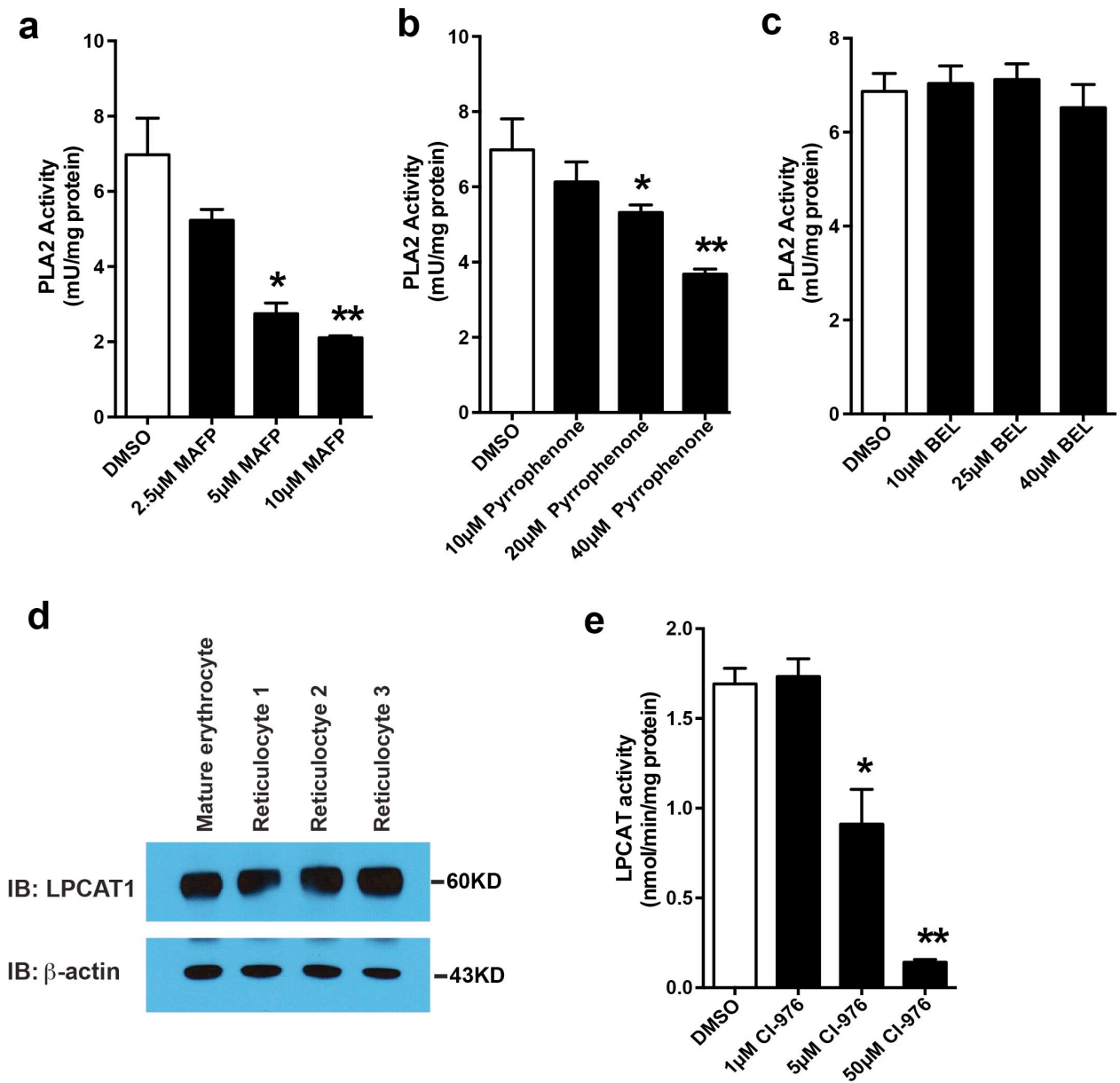
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**Running Title:** Imbalanced Lands' cycle in sickle cell disease

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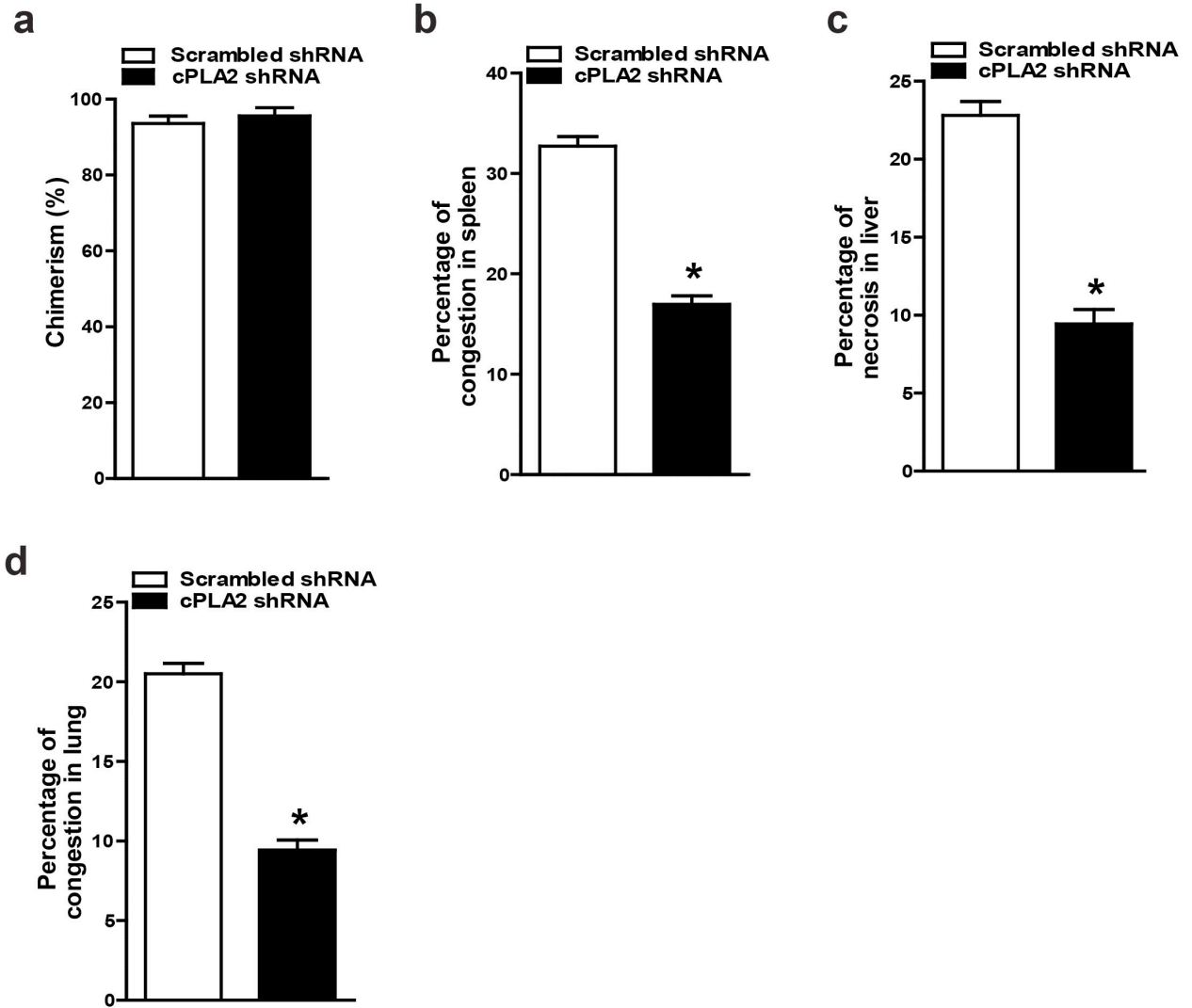
There are no conflicts of interest for any of the authors.

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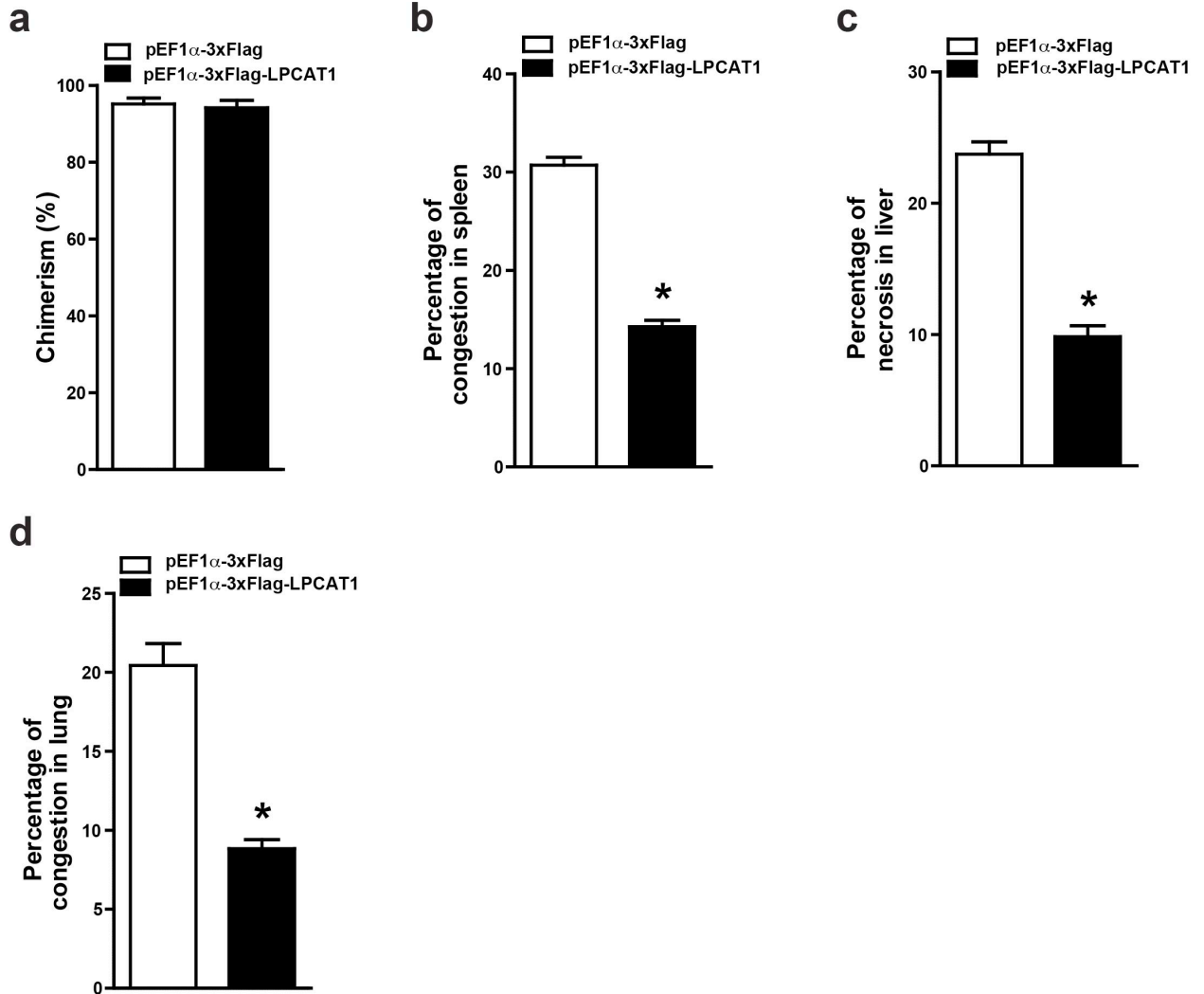


**Supplementary Figure 1. Cytosolic PLA2 and LPCAT1 are the major components of Lands' cycle for PC generation in mouse SCD erythrocytes.** (a-c) Quantification of PLA2 activity in cultured SCD Tg mice erythrocytes treated with DMSO or different dosages of cPLA2 inhibitors, MAFP (2.5µM, 5µM and 10µM) (a), Pyrrophenone (10µM, 25µM and 40µM) (b) or iPLA2 inhibitor BEL (10µM, 25µM and 40µM) (c) under 4% oxygen condition for 2 hours. Data are expressed as Mean± SEM; n = 5 in each group. (d) LPCAT protein level was measured in isolated mature erythrocytes and reticulocytes from SCD mice by Western blot. (e) LPCAT activity was measured in the cultured SCD mouse erythrocytes treated with different dosages of LPCAT inhibitor, CI-976 (1µM, 5µM and 50µM as indicated). Data are expressed as Mean± SEM; n = 4 in each group. \**P* < 0.05 and \*\**P* < 0.01 versus cultured SCD erythrocytes treated with DMSO or treated with a lower concentration of drugs,

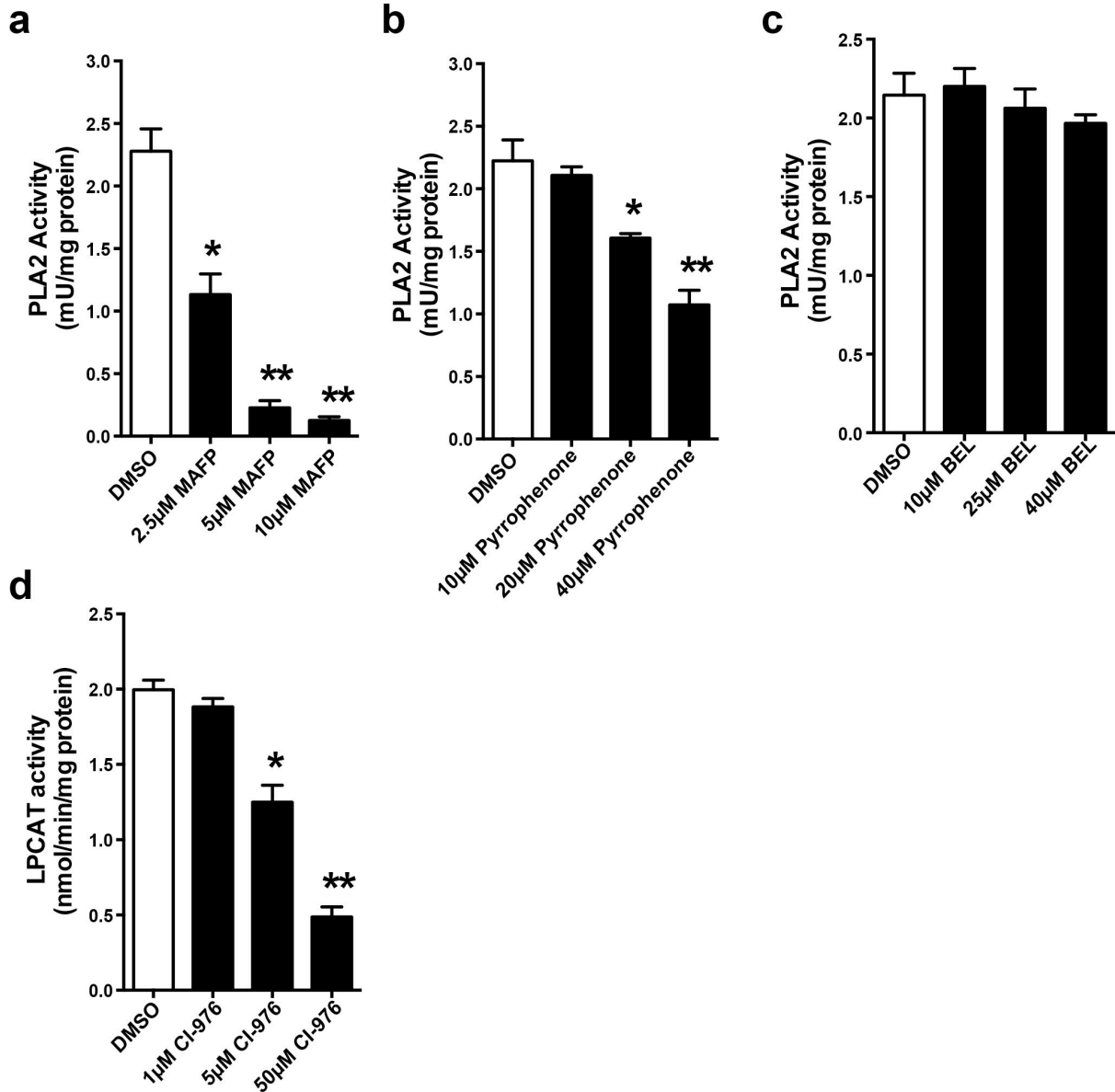
respectively.



**Supplementary Figure 2. Semi-quantitative analysis of tissue damage in BMT SCD chimeras with lentiviral shRNA specific knockdown of cPLA2 in hematopoietic stem cells (HSCs) and with scrambled shRNA. Related to Figure 3. (a)** The percentage of SCD chimeras was assessed by quantification of HbS and mouse normal Hb (HbA) in erythrocytes from BMT SCD chimeras with HSC-specific knockdown of cPLA2. Data shown represent the Mean  $\pm$  SEM. Data shown represent the Mean  $\pm$  SEM ( $n = 7$ ). **(b-d)** Semi-quantitative analysis of histological changes in spleen **(b)**, liver **(c)**, and lung **(d)** of SCD chimeras with cPLA2 knockdown in HSCs using Image-Pro Plus software. Data are expressed as Mean  $\pm$  SEM;  $n = 5-7$  in each group. \* $P < 0.05$  versus SCD chimeras with BMCs infected with recombinant lentivirus encoding scrambled shRNA.



**Supplementary Figure 3. Semi-quantitative analysis of tissue damage in BMT SCD chimeras with lentiviral overexpression of LPCAT1 and with control vector in hematopoietic stem cells (HSCs). Related to Figure 4. (a)** The percentage of SCD chimeras was assessed by quantification of HbS and mouse normal Hb (HbA) in erythrocytes from BMT SCD chimeras with overexpression of LPCAT1. Data shown represent the Mean  $\pm$  SEM. **(b-d)** Semi-quantitative analysis of histological changes in spleen **(b)**, liver **(c)**, and lung **(d)** of SCD chimeras with LPCAT1 overexpression using Image-Pro Plus software. Data are expressed as Mean  $\pm$  SEM; n = 6 in each group. \* $P$  < 0.05 versus SCD chimeras with BMCs infected with recombinant lentivirus encoding control vector.



**Supplementary Figure 4. Cytosolic PLA2 and LPCAT function as major components of Lands' cycle for PC formation in human SCD erythrocytes, Related to Figure 6.** (a-c) Quantification of PLA2 activity in isolated erythrocytes from individuals with SCD treated with DMSO or different dosages of cPLA2 inhibitors, MAFP (2.5µM, 5µM and 10µM) (a), Pyrrophenone (10µM, 25µM and 40µM) (b) or iPLA2 inhibitor BEL (10µM, 25µM and 40µM) (c) under hypoxic condition (4%). Data are expressed as Mean± SEM; n = 5 in each group. (d) LPCAT activity was measured in the cultured human SCD erythrocytes treated with different dosage of LPCAT inhibitor, CI-976 (1µM, 5µM and 50µM). Data are expressed as Mean± SEM; n = 5 in each group. \* $P < 0.05$  and \*\* $P < 0.01$  versus cultured SCD human erythrocytes treated with DMSO or treated with lower concentration of drugs, respectively.

Supplementary Table 1 Hematological parameters of SCD chimeras with or without HSCs-specific cPLA2 knockdown

| Mice                     | RBC<br>(X10 <sup>6</sup> /ml) | Hb<br>(g/dl) | HCT<br>(%)  | MCV<br>(fl) | MCH<br>(pg) | MCHC<br>(g/dl) | RDW<br>(%)  | WBC<br>(X10 <sup>3</sup> /ml) |
|--------------------------|-------------------------------|--------------|-------------|-------------|-------------|----------------|-------------|-------------------------------|
| Scrambled shRNA<br>(n=7) | 5.70±0.14                     | 6.04±1.67    | 30.34±1.88  | 50.73±0.50  | 11.14±0.89  | 21.94±0.12     | 24.7±0.23   | 18.47±1.77                    |
| cPLA2 shRNA<br>(n=5)     | 6.52±0.22*                    | 8.95±0.23*   | 34.53±1.51* | 52.9±0.45   | 13.77±0.05* | 23.65±0.46     | 20.76±0.43* | 9.82±0.92*                    |

Control vector: C57BL/6 mice were transplanted with SCD Tg mice bone marrow cells transduced with recombinant lentiviral vectors encoding scrambled-shRNA sequence;

cPLA2-shRNA: C57BL/6 mice were transplanted with SCD Tg mouse bone marrow transduced with recombinant lentiviral vectors encoding shRNA specifically for cPLA2 mRNA.

SCD, Sickle cell disease transgenic mice; RBC, red blood cell, Hb, hemoglobin; HCT, hematocrit; MCV, mean corpuscular volume; MCH, mean corpuscular hemoglobin; MCHC, mean corpuscular hemoglobin concentration; RDW, red blood cell distribution width; WBC, white blood cell. \* $P < 0.05$  versus untreated SCD transgenic mice.

Supplementary Table 2 Complete blood count in SCD chimeras with or without overexpression of LPCAT1 in donor bone marrow cells

| Mice                                   | RBC<br>(X10 <sup>6</sup> /ml) | Hb<br>(g/dl) | HCT<br>(%)  | MCV<br>(fl) | MCH<br>(pg) | MCHC<br>(g/dl) | RDW<br>(%)  | WBC<br>(X10 <sup>3</sup> /ml) |
|--|-------------------------------|--------------|-------------|-------------|-------------|----------------|-------------|-------------------------------|
| pEF1 $\alpha$ -3xflag<br>(n=6)         | 6.04±0.09                     | 6.18±0.10    | 30.90±0.63  | 50.38±0.26  | 11.46±0.07  | 20.00±0.13     | 23.48±0.14  | 17.14±0.64                    |
| pEF1 $\alpha$ -3xflag-<br>LPCAT1 (n=6) | 6.71±0.16*                    | 8.73±0.28*   | 35.36±1.12* | 52.50±0.88  | 14.04±0.10* | 23.88±0.22*    | 20.83±0.18* | 10.20±0.55*                   |

Control vector: C57BL/6 mice were transplanted with SCD Tg mice bone marrow cells transduced with recombinant lentiviral vectors encoding control vector;

Flag-LPCAT: C57BL/6 mice were transplanted with SCD Tg mouse bone marrow transduced with recombinant lentiviral vectors encoding Flag-labeled LPCAT1 sequence.

Hb, hemoglobin; HCT, hematocrit; MCV, mean corpuscular volume; MCH, mean corpuscular hemoglobin; MCHC, mean corpuscular hemoglobin concentration; RDW, red blood cell distribution width; WBC, white blood cell. \* $P < 0.05$  versus untreated SCD Tg mice.



**Supplementary Table 3: Clinical information for control individuals and SCD patients**

|                             | Control     | SCD patient |
|-----------------------------|-------------|-------------|
| Number                      | 15          | 22          |
| Gender                      | M=5 F=10    | M=10 F=12   |
| Age (Years)                 | 39.52±11.21 | 31.82±10.88 |
| RBCs ( $10^6/\mu\text{l}$ ) | 4.73±0.32   | 2.41±0.50*  |
| Hb (g/dl)                   | 13.76±1.51  | 8.26±1.47*  |
| HCT (%)                     | 41.34±1.63  | 24.71±1.40* |
| WBC ( $10^3/\mu\text{l}$ )  | 5.64±2.01   | 11.66±4.05* |
| HbS                         | NT          | 81.74±8.30  |
| HbA                         | NT          | 0.87±2.89   |
| HbA2                        | NT          | 3.12±0.62   |
| HbF                         | NT          | 14.28±8.35  |
| Hydroxyurea                 | No          | 13          |

Control: Healthy volunteers; SCD patient: sickle cell disease patient; RBC: red blood cells; Hb: hemoglobin; HCT: hematocrit; WBC: white blood cell; HbS: hemoglobin S; HbA: hemoglobin A; HbA2: hemoglobin A2; HbF: hemoglobin F; NT: not tested. \* $P \leq 0.05$  vs. control. All subjects are of African-American decent.