

SUPPLEMENTAL MATERIAL

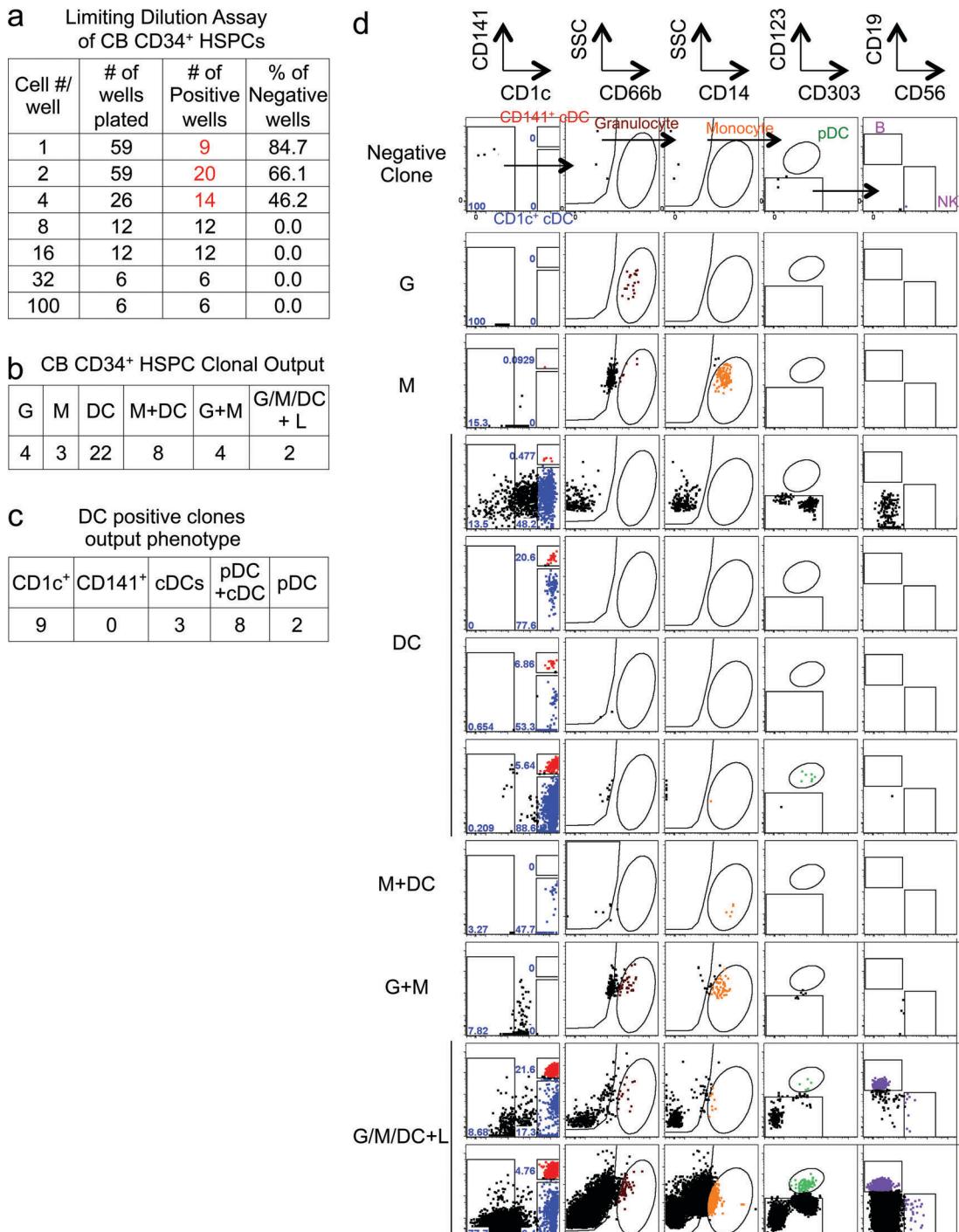
Breton et al., <http://www.jem.org/cgi/content/full/jem.20141441/DC1>

Figure S1. Clonal output of cord blood CD34⁺ HSPCs. (a) CB CD34⁺ HSPCs were cultured at 1, 2, 4, 8, or 16 cells per well for 14 d on MS5+FSG. Each well was analyzed for its positivity and lineage potential based on surface marker analysis. Positive wells were determined based on the presence of CD45⁺ cells (Materials and methods) by flow cytometry. Red font indicates wells with single clones (cell no. per well < 4.57 cells) analyzed for lineage output. (b) Table summarizes the number of clones giving rise to the specified lineages. (c) Clones generating only DCs were broken down into their respective subsets and combinations (CD1c⁺ cDCs, CD141⁺ cDCs, cDCs, pDCs+cDCs, or pDCs alone). (d) Representative flow cytometry plots of gated live human CD45⁺ cells show clones failing to generate any of the lineages (negative clones), as well as positivity for granulocytes alone (G), monocytes alone (M), DCs alone, M+DC, G+M, or Myeloid (G/M/ or DC) + Lymphoid (B or NK, L).

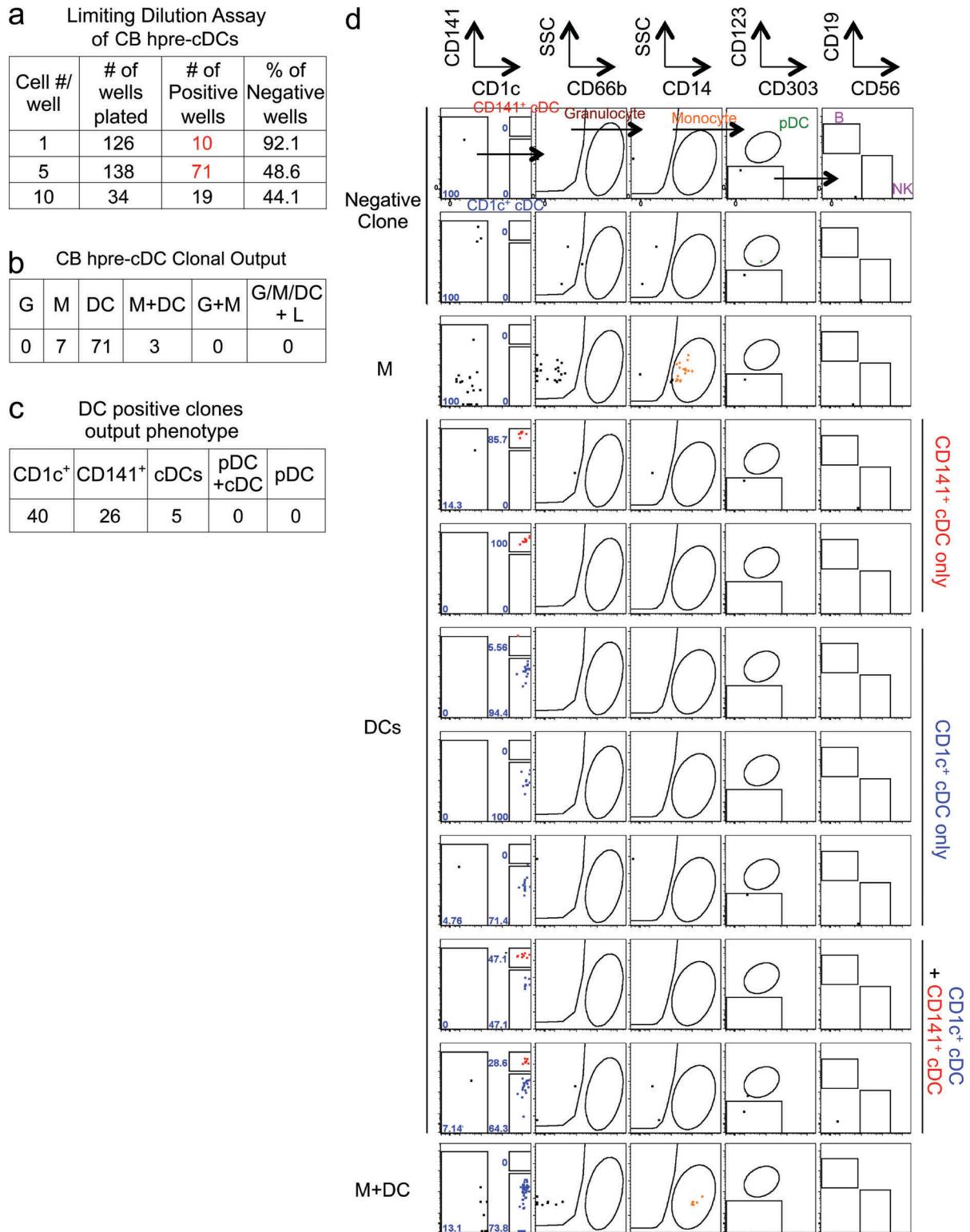


Figure S2. Clonal output of cord blood (CB) hpre-CDcs. (a) CB hpre-CDcs were cultured at 1, 5, or 10 cells per well for 5 d on MS5+FSG. Each well was analyzed for its positivity and lineage potential based on surface marker analysis. Positive wells were determined based on the presence of CD45⁺ cells by flow cytometry. Red font indicates wells with single clones (cell no. per well < 7.84 cells) analyzed for lineage output. (b) Table summarizes the number of clones giving rise to the specified lineages. (c) Clones generating only DCs were broken down into their respective subsets and combinations (CD1c⁺ cDCs, CD141⁺ cDCs, cDCs, pDCs+cDCs, or pDCs alone). (d) Representative flow cytometry plots of gated live human CD45⁺ cells show clones failing to generate any of the lineages (negative clones), as well as positivity for monocytes alone (M), DCs alone, or M+DC.

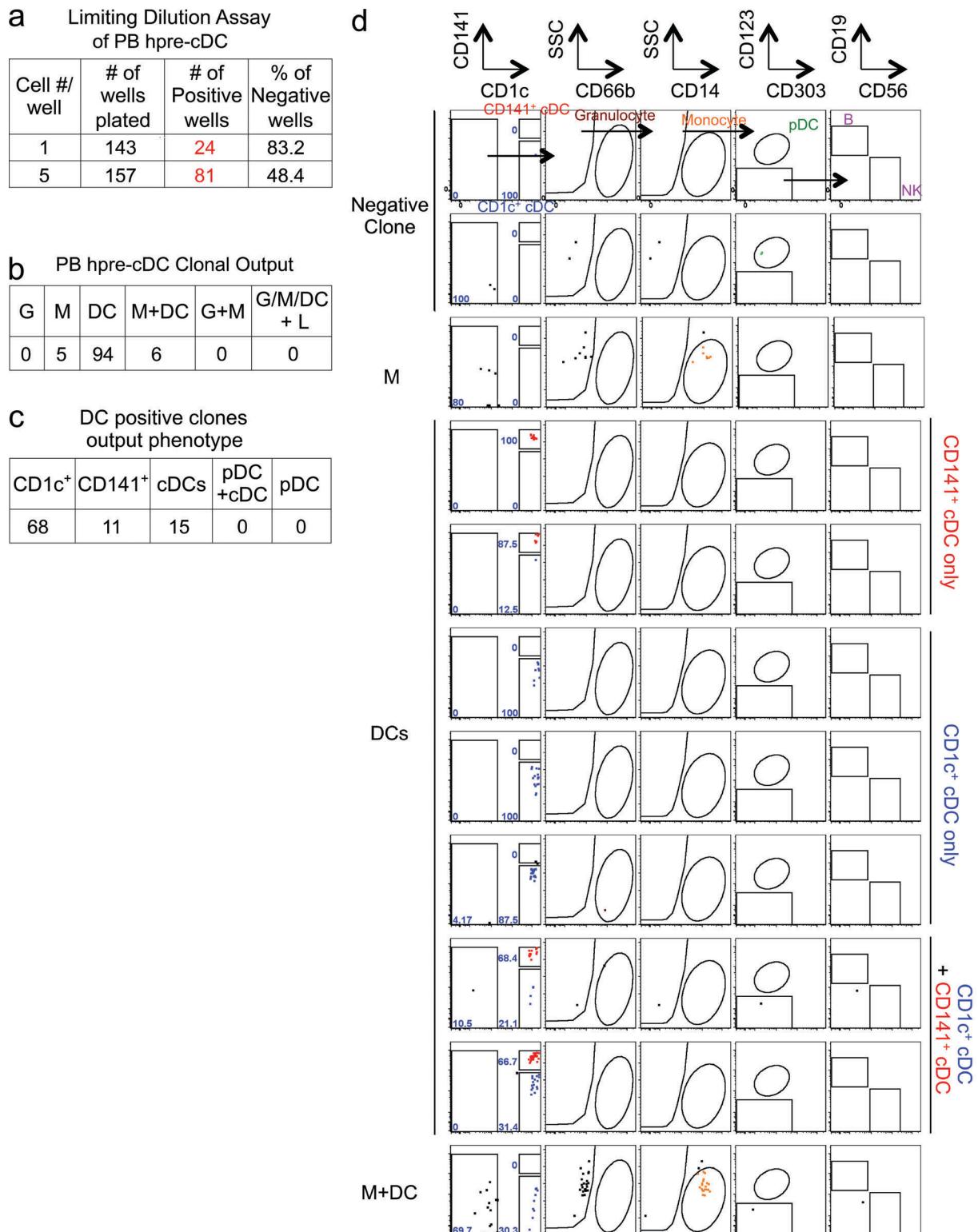


Figure S3. Clonal output of peripheral blood (PB) hpre-CDCs. (a) PB hpre-CDCs were cultured at 1 or 5 cells per well for 5 d on MS5+FSG. Each well was analyzed for its positivity and lineage potential based on surface marker analysis. Positive wells were determined based on the presence of CD45⁺ cells by flow cytometry. Red font indicates wells with single clones (cell number per well < 6.55 cells) analyzed for lineage output. (b) Table summarizes the number of clones giving rise to the specified lineages. (c) Clones generating only DCs were broken down into its respective subsets and combinations (CD1c⁺ cDCs, CD141⁺ cDCs, cDCs, pDCs+cDCs, or pDCs alone). (d) Representative flow cytometry plots of gated live human CD45⁺ cells show clones failing to generate any of the lineages (negative clones), as well as positivity for monocytes alone (M), DCs alone, or M+DC.

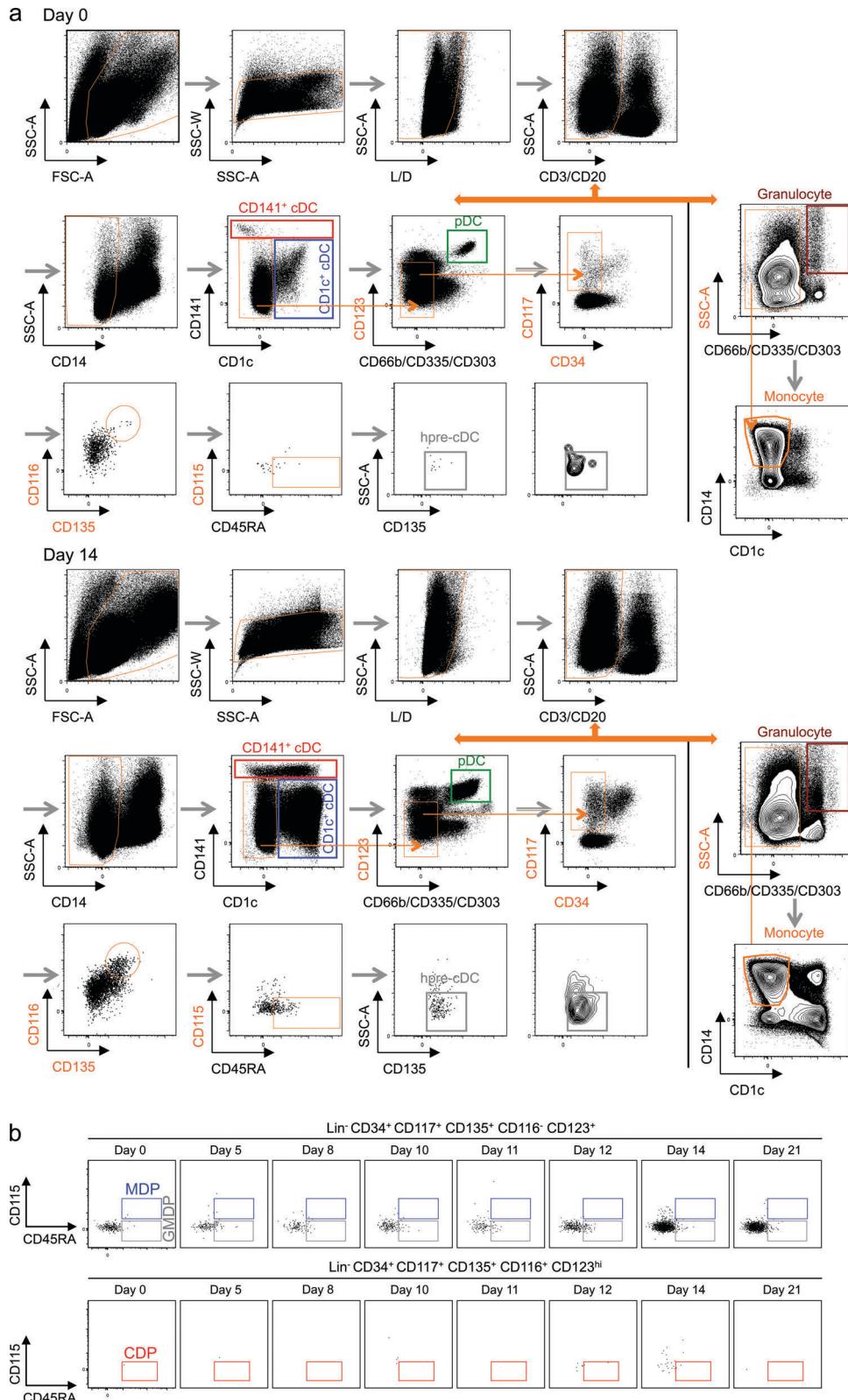


Figure S4. 14-color gating strategy for identification of human hpre-DCCs in blood. (a) PBMCs were isolated from heparinized blood using Ficoll-Hypaque. Live CD3⁻CD19⁻CD335⁻CD66b⁻CD14⁻CD1c⁻CD141⁻CD303⁻ cells, i.e., T, B, NK, neutrophils, monocytes, and DC cell-depleted, were stained for CD34, CD117, CD123, CD135, CD116, CD115, and CD45RA markers. Blood CD34⁺CD117⁺CD123⁺CD135⁺CD116⁺CD115⁺CD45RA⁺ hpre-DCCs are shown before (Day 0) and after (Day 14) Flt3L administration. (b) hCDPs, hMDPs, and hGMDPs are undetectable in the blood before and after Flt3L administration in healthy volunteers. Flow cytometry plots of gated Lin⁻CD34⁺CD117⁺CD135⁺CD116⁺CD123^{hi} cells show lack of hCDPs in blood of Flt3L-injected volunteers. Flow cytometry plots of gated Lin⁻CD34⁺CD117⁺CD135⁺CD116⁺CD123⁺ cells show lack of hMDPs and hGMDPs in blood of Flt3L injected volunteers.