	<h1>THE GOODELL LABORATORY</h1>																																																	
<p>Author</p>	<p>Nathan Boles</p>	<p>Feb.5, 2009</p>																																																
<p>Title</p>	<p>Hematopoietic Progenitor Staining</p>																																																	
<p>Introduction</p>	<p>This protocol describes the analysis or isolation of the early hematopoietic progenitors.</p>																																																	
<p>Materials</p>	<p>HBSS+ : Hanks Balanced Salt Solution (from Gibco) with 2% Fetal Calf Serum and 10 mM DMEM+ Hoechst 33342</p> <p><i>LT-HSC, ST-HSC, MPP antibody cocktail</i></p> <table border="0" style="width: 100%;"> <thead> <tr> <th style="text-align: left;"><u>Antibody</u></th> <th style="text-align: right;"><u>Concentration</u></th> </tr> </thead> <tbody> <tr> <td>Lineage- Fite</td> <td style="text-align: right;">1:100</td> </tr> <tr> <td>Gr-1</td> <td></td> </tr> <tr> <td>Mac-1</td> <td></td> </tr> <tr> <td>Ter-119</td> <td></td> </tr> <tr> <td>CD4</td> <td></td> </tr> <tr> <td>CD8</td> <td></td> </tr> <tr> <td>B220</td> <td></td> </tr> <tr> <td>Sca-1 – Pe-Cy7</td> <td style="text-align: right;">1:100</td> </tr> <tr> <td>cKit – APC-AF750</td> <td style="text-align: right;">1:100</td> </tr> <tr> <td>Flk2/Flt3 – PE</td> <td style="text-align: right;">1:50</td> </tr> <tr> <td>CD34 – AF647</td> <td style="text-align: right;">1:50</td> </tr> </tbody> </table> <p><i>CMP</i></p> <table border="0" style="width: 100%;"> <thead> <tr> <th style="text-align: left;"><u>Antibody</u></th> <th style="text-align: right;"><u>Concentration</u></th> </tr> </thead> <tbody> <tr> <td>Lineage – biotin</td> <td style="text-align: right;">1:100</td> </tr> <tr> <td>Gr-1</td> <td></td> </tr> <tr> <td>Ter-119</td> <td></td> </tr> <tr> <td>CD4</td> <td></td> </tr> <tr> <td>CD8</td> <td></td> </tr> <tr> <td>CD3</td> <td></td> </tr> <tr> <td>B220</td> <td></td> </tr> <tr> <td>CD19</td> <td></td> </tr> <tr> <td>Il7ra – Pe-Cy7</td> <td style="text-align: right;">1:50</td> </tr> <tr> <td>Sca-1 – Fite</td> <td style="text-align: right;">1:100</td> </tr> <tr> <td>cKit – Pe</td> <td style="text-align: right;">1:100</td> </tr> </tbody> </table>		<u>Antibody</u>	<u>Concentration</u>	Lineage- Fite	1:100	Gr-1		Mac-1		Ter-119		CD4		CD8		B220		Sca-1 – Pe-Cy7	1:100	cKit – APC-AF750	1:100	Flk2/Flt3 – PE	1:50	CD34 – AF647	1:50	<u>Antibody</u>	<u>Concentration</u>	Lineage – biotin	1:100	Gr-1		Ter-119		CD4		CD8		CD3		B220		CD19		Il7ra – Pe-Cy7	1:50	Sca-1 – Fite	1:100	cKit – Pe	1:100
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Protocol		<i>Notes</i>																								
1.	Prewarm DMEM+ (see below) while preparing the bone marrow.	<i>Ensure that a water bath is at precisely 37° C (check this with a thermometer!).</i>																								
2.	Using mice 8-12 weeks of age, prepare bone marrow from femurs and tibias and resuspend in HBSS+																									
3.	Aliquot bone marrow into two equal portions	<i>We find an average of 5 x 10⁷ nucleated cells per C57Bl/6 mouse. This number varies from strain to strain.</i>																								
4	Set cell concentration to 1x10 ⁶ cells/mL using 37° C DMEM+ for one aliquot (hereafter to be referred to as SP aliquot). Place other aliquot in 4° C fridge (hereafter to be referred to as Prog aliquot).	...																								
5	Add 200X Hoechst solution to the SP aliquot in the 37° C DMEM+																									
6	Incubate the SP aliquot at 37° C for 90 minutes exactly with occasional shaking.	<i>Use this time to eat lunch.</i>																								
7	Set a fraction of Prog aliquot to the side for staining controls, then split the remainder into two equal fractions (hereafter referred to as Prog1 and Prog2)	<i>For controls 5 million cells is sufficient.</i>																								

8	With 45 minutes left go ahead and make the 3 antibody cocktails outlined in the Materials section. Make enough to bring the final concentration of cells to 1×10^8 cells/mL during staining for each aliquot. After you complete the cocktails go ahead and label all the tubes you will need for the rest of your experiment.	<i>Be extra careful when making the cocktails because a mistake here can give wacky results. Also make 10% more of each cocktail than you need to ensure an adequate amount.</i>
9	Take a small fraction of your SP aliquot to use as SP controls at the end of the 90 minutes.	
10	Spin down SP aliquot for 8 min. at 2000 rpm at 4°C, and resuspend in cold Hanks+ to a volume of 1×10^8 cells/mL and keep on ice throughout the rest of the protocol.	
11	Now add the appropriate amount of antibodies to their correct aliquots (LT-HSC, ST-HSC, MPP cocktail to the SP aliquot. CMP cocktail to Prog1. CLP to Prog2)	
12	Incubate for thirty minutes on ice.	
13	Add enough Hank's+ to each control fraction to have 100µL for each color (including negative control).	
14	Stain controls for each color for both setups using Gr-1 or B220. However for the following antibodies use the actual antibody at a concentration of 1:50: Il7ra – Pe-Cy7, CD34 – AF647, CD16/32 – AF750, Flk2/Flt3 – Pe. (Stain for at least 10 minutes)	
15	Prepare a PI solution by using the lab stock of PI (1:100concentration) and Hank's+. Make enough to resuspend each of your samples in 500µL.	
16	Prepare a secondary staining solution. Put strepavidin-Pac Blue into Hank's+ at a concentration of 1:50. Make enough to stain all of your Prog1 and Prog2 samples at a cell concentration of 1×10^8 cell/mL.	
17	Wash all samples and controls with 20x Hank's+ and spin down for 8 min. at 2000 rpm at 4°C.	
18	Resuspend SP aliquots and controls in 300-500µL of PI solution, keep on ice.	

19	Incubate for 15 minutes	
20	Wash all Prog1 and Prog2 samples with 20x Hank's+ and spin down for 8 min. at 2000 rpm at 4°C.	
21	Resuspend Prog1 and Prog2 in 300-500µL of PI solution, keep on ice.	
22	Go and analyze.	<i>Colors are appropriate for a 4-laser LSRII, however if using another machine verify color scheme.</i>

References.

1. Challen G, Boles NC, Lin KY, Goodell MA. Mouse Hematopoietic Stem Cell Identification And Analysis. Cytometry A. 2009 Jan;75(1):14-24.

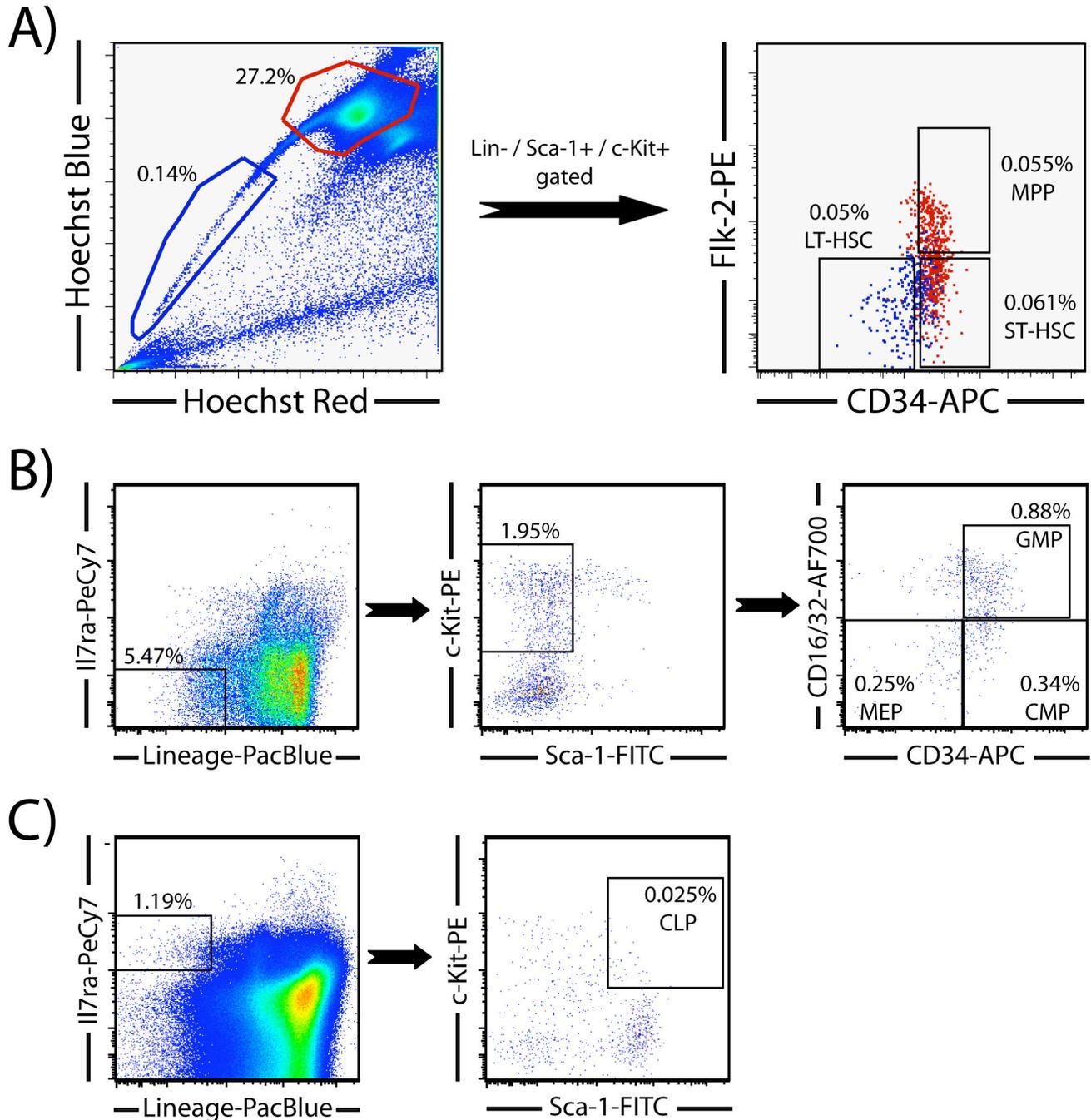


Figure 1. Separation of hematopoietic progenitor populations by flow cytometry. (A) LT-HSC, ST-HSC, and MPP gating scheme. SP cells are gated to KLS as described for Figure 1, and shown here displayed in red on a CD34 / Flk-2 plot; the SPKLS are negative for both of these markers (the Hoechst-stained cells here have been previously magnetically enriched for Sca-1 to increase the overall proportion to 0.14%). The non-SP population, also shown gated on the Hoechst plot, is also taken through a KLS selection (not shown), then displayed for CD34 and Flk2. The KLS-Flk2+CD34+ cells are multi-potential progenitors (MPP), and the Flk2-CD34+ cells are considered short-term (ST) HSC. Thus, all three of these populations can be readily sorted from one sample. (B) Gating scheme for the common myeloid progenitor (CMP), megakaryocyte-erythrocyte progenitors (MEPs) and granulocyte-macrophage progenitors (GMPs). (C) Gating scheme for the common lymphoid progenitor (CLP).

