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Correction for erythroid cell contamination in microassay for immunophenotyping of neonatal lymphocytes

E de Vries, S de Bruin-Versteeg, W M Comans-Bitter, R de Groot, G J M Boerma, F K Lotgering, J J M van Dongen

Abstract

Immunophenotyping of blood lymphocyte subpopulations in neonates and young infants is hampered by the limited amount of blood that can be collected. Contamination of the flow cytometric “lympho-gate” by normoblasts and analysed erythrocytes, and therefore the underestimation of the relative frequencies of lymphocyte subpopulations, interferes with the precise calculation of absolute counts.

A microassay was developed by adapting the lysed whole blood technique. Triple immunostaining in a single antibody staining step was used to reduce washing steps and cell loss. Introduction of a triple staining for CD71 (expressed by erythroid precursors), glycophorin A (GpA, expressed by all erythroid cells), and CD45 (expressed by all leucocytes) permitted the relative frequencies of normoblasts (CD71'/GpA'/CD45' population) and unlysed erythrocytes (CD71'/GpA'/CD45' population) to be identified and measured within the “lympho-gate” of neonatal cord blood samples. Particularly high frequencies were found (median: 31%) in cord blood samples from preterm neonates. These erythroid cells disappear rapidly by 1 week of age. The relative frequencies of erythroid cells can be used to calculate correct lymphocyte subpopulation values. Using only 0.5–0.8 ml of blood, this microassay would also be suitable for rapid prenatal immunodiagnosis of congenital immunodeficiencies.

Methods

Twenty one neonatal cord blood samples were collected by venepuncture immediately after clamping of the cord. The blood was kept at room temperature until labelling with monoclonal antibodies was performed within 12 hours of sampling. Peripheral blood was drawn by venepuncture from nine healthy adult volunteers between 20 and 40 years of age, and at 1 week of age from 14 neonates whose cord blood had been studied before. EDTA was used as an anticoagulant.

Informed consent was obtained according to the guidelines of the Medical Ethics Committee of the Erasmus University Rotterdam/University Hospital Rotterdam in all cases. The following monoclonal antibodies were used: CD3 (Leu-4a), CD4 (Leu-3a), CD8 (Leu-2a), CD16 (Leu-11c), CD19 (Leu-12), CD56 (Leu-19), CD71 (anti-transferrin receptor), CD45 (HE11), IgG1 and IgG2 (isotype controls) (Becton Dickinson, San Jose, CA), CD14 (My4), CD19 (B4) (Coulter, Hialeah, FL), CD3 (Hit3a), glycophorinA (GpA) (GA-R2), PHA, sheep E(anti-human IgG), goat (anti-human IgG) (Coulter), goat (anti-human IgM) (Coulter).
Transfusion Service, Amsterdam, The Netherlands). All monoclonal antibodies were conjugated with fluorescein isothiocyanate (FITC), phycoerythrin (PE), or the duochrome PE-Cyanin5 (PE-Cy5).

Blood samples (100, 40, 20, or 10 µl) were incubated with titrated monoclonal antibodies for 10 minutes at room temperature. After two washes with phosphate buffered saline containing 0.1% NaN₃ and 0.5% bovine serum albumin, erythrocytes were lysed with Lysing Solution (Becton Dickinson) according to the manufacturer’s instructions. Before lysis the cell pellet was loosened to prevent incomplete lysis of erythrocytes.

The samples were analysed using a FACS-cant flow cytometer (Becton Dickinson). Instrument settings were determined with cells labelled with FITC, PE, and PE-Cy5 conjugated monoclonal antibodies and unstained cells. After gating of lymphocytes on the basis of FSC and SSC (checked by CD14, CD15, and CD45 monoclonal antibody staining), 8000 events were acquired. The data were analysed using FACScan software (Becton Dickinson) in dot blots, with optimal quadrant setting checked by histogram analysis, if necessary.

The total nucleated cell count was determined on an H1 Technicon haemocytometer (Bayer, Tarrytown, NY). The relative frequency of normoblasts was carefully determined by manual differentiation of at least 400 nucleated cells. The total leucocyte count was determined according to the following formula:

\[
\frac{\text{total nucleated cell count}}{10^6} \times (100 - \% \text{ of normoblasts}) 
\]

The relative frequencies of lymphocyte subpopulations were calculated according to the following formula:

\[
\frac{\% \text{ of population within the lympho-gate}}{10^6} \times (100 - \% \text{ of normoblasts plus percentage of unlysed erythrocytes within the lympho-gate}) 
\]

The Wilcoxon rank sum test or Friedman’s test were used to compare two or more different techniques used in one blood sample, respectively (\(\alpha = 0.05\)).

Results
Development of Microassay
We incubated 100, 40, 20, and 10 µl of blood with monoclonal antibodies. When using 20 µl of blood, at least 100,000 nucleated cells remained available after completion of the staining procedure in all samples tested, as determined by use of a cell counter (Coulter Counter; Hialeah, FL). However, when using 10 µl of blood, major and variable cell loss was observed. Relative frequencies of lymphocyte subpopulations determined in parallel with either 100 µl or 20 µl of blood per test tube gave fully comparable results in five separate neonatal blood samples. Therefore, 20 µl of blood incubated with 20 µl of each monoclonal antibody per test tube was used in all further experiments.

The determination of the relative frequencies of lymphocyte subpopulations by either double or triple immunostaining was compared in one adult and six cord blood samples—for example, triple CD4/CD8/CD3 vs double CD4/CD8, CD4/CD3 and CD8/CD3 immunostaining. When using triple immunostaining, monoclonal antibodies were either incubated consecutively, with each incubation followed by two washes, or all together. No significant differences were found between the relative frequencies of lymphocyte subpopulations, as determined by the different staining methods. Therefore, in all further experiments only triple immunostaining was used with incubation of all three fluorochrome conjugated monoclonal antibodies simultaneously to speed up the microassay and minimise cell loss due to multiple washes.

Detection of Erythroid Cell Contamination
When a scatter gate is set around the lymphocyte population, it is generally not possible to exclude fully other contaminating cells without excluding lymphocytes. However, as long as the sum of the relative frequencies of B lymphocytes, T lymphocytes, and natural killer cells (the so-called “lympho-sum”) is at least 95% (or 0.95), the lympho-gate is often considered to be placed correctly. The remaining events within the lympho-gate are monocytes, granulocytes, and unidentified events, which generally consist of dead cells and debris. In our neonatal cord blood samples the lympho-sum rarely reached 95% of the lympho-gate (table 1). In preterm neonates, in particular, the lympho-sum seemed to be far too low. This was caused by a relatively high frequency of unidentified events (fig 1).

Normoblasts express CD71 and GpA, whereas erythrocytes express only GpA. In contrast, leucocytes express CD45, unlike normoblasts and erythrocytes. Therefore, CD71+/GpA+/CD45 cells within the lympho-gate can be regarded as normoblasts, whereas unlysed erythrocytes have the CD71+/GpA+/CD45 immunophenotype. By triple immunostaining with CD71-FITC, GpA-PE, and CD45-PE-Cy5, these unidentified events could be identified as contamination by normoblasts and in some samples by unlysed erythrocytes as well (fig 2) (table 1). The FSC and SSC of these normoblasts and unlysed erythrocytes was checked (“back-gating”), and was found to be relatively low, as expected for these cells (figs 2C and D).

A small percentage of events remained unidentified—that is, they were neither CD45+ leucocytes, nor CD71+/GpA+/CD45 normoblasts, nor CD71+/GpA+/CD45 unlysed erythrocytes (left lower quadrant of dot plot in fig 2B). These events may represent dead cells and debris.

Erythroid Cell Contamination in Neonatal Blood
CD71/GpA/CD45 staining was used to determine the relative counts of normoblasts and unlysed erythrocytes and to calculate the relative counts of lymphocyte subpopulations. The differences in relative frequencies of lymphocyte subpopulations in neonatal cord
blood with and without a correction for normoblasts and unlysed erythrocytes are shown in Table 2 for two samples. Normoblasts were not found by manual differentiation or flow cytometric immunophenotyping in blood samples from the same neonates at 1 week of age nor in blood samples from adults, but unlysed erythrocytes were still present in some samples at 1 week of age (Table 1).

**Discussion**

We have developed a microassay for flow cytometric determination of blood lymphocyte subpopulations in neonates and young infants by adapting the lysed whole blood technique, using one step triple immunostaining. When only double immunostaining is available, more blood is needed and less detailed results are obtained—for example, double positive CD45RA+CD45RO+ CD3+ T lymphocytes cannot be identified. This micro-assay assay needs only about 0.4 ml of blood to determine T lymphocyte subpopulations for monitoring HIV infected infants, and 0.5 to 0.8 ml of blood is sufficient for detailed evaluation of infants with suspected congenital immunodeficiencies. Extensive protocols for research purposes can also be performed, because 1.0 ml of blood is sufficient for more than 40 triple immunostainings. These are acceptable volumes in infants and even in preterm babies.

Normoblasts and unlysed erythrocytes can be present in considerable numbers in neonatal samples.

The presence of normoblasts cannot be prevented by technical measures, but strict adherence to a rigorous lysing protocol can minimise the number of unlysed erythrocytes in a sample. Therefore, identification of both cell populations is of interest. Using triple

![Image](https://example.com/image1.png)  
**Figure 1** Flow cytometric immunophenotyping of neonatal cord blood lymphocytes reveals a high frequency of unidentified events within the lympho-gate. (A) Lympho-gate setting in neonatal cord blood, based upon FSC/SSC characteristics. (B) Composition of the lympho-gate in neonatal cord blood: many unidentified events and relatively high numbers of natural killer cells were observed. (C) Lympho-gate setting in adult blood. (D) Composition of the lympho-gate in adult blood: virtually no unidentified events and relatively low numbers of natural killer cells were observed. Quadrant 1: CD14+ contaminating monocytes, and CD3+/CD16+/CD56+ natural killer cells. Quadrant 2: CD3+ T lymphocytes, CD19+ B lymphocytes, and CD15+ contaminating granulocytes. Quadrant 3: Unidentified events.

![Image](https://example.com/image2.png)  
**Figure 2** Triple staining for CD71, GpA, and CD45 identifies normoblasts and unlysed erythrocytes within the lympho-gate. (A) Lympho-gate setting. (B) CD71+/GpA+/CD45+ population of normoblasts and the CD71+/GpA+/CD45+ population of unlysed erythrocytes within the CD45 lympho-gate. (C) Back-gating of CD71+/GpA+/CD45+ normoblasts into the lympho-gate. (D) Back-gating of and CD71+GpA+/CD45+ unlysed erythrocytes into the lympho-gate.

### Table 1 Composition of “lympho-gate” in neonatal and adult blood samples

<table>
<thead>
<tr>
<th>Lympho-gate populations</th>
<th>Preterm neonates (n=6)</th>
<th>Term neonates (n=15)</th>
<th>The same neonates at one week of age (one preterm and 13 term neonates)</th>
<th>Adults (n=9)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>Median</td>
<td>Range</td>
<td>Mean</td>
</tr>
<tr>
<td>Lympho-sum†</td>
<td>57</td>
<td>61</td>
<td>24-78</td>
<td>85</td>
</tr>
<tr>
<td>CD14+ monocytes</td>
<td>2</td>
<td>2</td>
<td>1-2</td>
<td>1</td>
</tr>
<tr>
<td>CD15+ granulocytes</td>
<td>1</td>
<td>1</td>
<td>0-1</td>
<td>1</td>
</tr>
<tr>
<td>CD71+/GpA+/CD45+ normoblasts</td>
<td>31</td>
<td>29</td>
<td>8-57</td>
<td>10</td>
</tr>
<tr>
<td>CD71+/GpA+/CD45+ unlysed erythrocytes</td>
<td>4</td>
<td>3</td>
<td>2-5</td>
<td>2</td>
</tr>
</tbody>
</table>

* All numbers represent percentages.

† Lympho-sum = CD19+ + CD3+ + CD3−CD16/56+ (B+T+NK) without correction for the erythroid cell contamination of the lympho-gate.
Microassay for immunophenotyping neonatal lymphocytes

Table 2  Illustration of erythroid cell contamination bias in two cord blood samples*

<table>
<thead>
<tr>
<th>Gestational age</th>
<th>Normoblasts</th>
<th>Unlysed</th>
<th>CD5+ T lymphocytes</th>
<th>CD19+ B lymphocytes</th>
<th>CD4+CD16/56+ natural killer cells</th>
<th>Lympho-sum</th>
</tr>
</thead>
<tbody>
<tr>
<td>29 weeks</td>
<td>44</td>
<td>5</td>
<td>28</td>
<td>54</td>
<td>18</td>
<td>36</td>
</tr>
<tr>
<td>35 weeks</td>
<td>40</td>
<td>3</td>
<td>24</td>
<td>43</td>
<td>10</td>
<td>17</td>
</tr>
</tbody>
</table>

* All numbers represent percentages.
† Without correction for the erythroid contamination of the lympho-gate.
‡ With correction for the erythroid contamination of the lympho-gate.

immunostaining for CD71, GpA, and CD45, we were able to identify both normoblasts (CD71+/GpA+/CD45 cells) and unlysed erythrocytes (CD71-/GpA+/CD45 cells) within the lympho-gate, and to measure their relative frequencies. When only double immunostaining is available, GpA+/CD45- cells can be used to measure the relative frequency of erythroid cells, but normoblasts and unlysed erythrocytes cannot be identified separately. The relative frequencies of erythroid cells can be used to calculate the correct relative frequencies of blood lymphocyte subpopulations. This is especially important in preterm neonatal and prenatal blood samples, where the relative frequencies of normoblasts and unlysed erythrocytes can be high (tables 1 and 2), and the frequencies of lymphocyte subpopulations can be underestimated. This can be confusing if a congenital immunodeficiency is suspected. The erythroid cell contamination of the lympho-gate rapidly disappears after birth and is virtually absent by 1 week of age (table 1).

In conclusion, we have developed a fast and easy to use lysed whole blood microassay for immunophenotyping of neonatal and infant blood lymphocyte subpopulations. It requires only a small volume of blood (20 µl per test tube), and offers a method to correct for the presence of normoblasts and unlysed erythrocytes in neonatal samples. It would also be suitable for rapid prenatal immunodiagnosis of congenital immunodeficiencies when genetic markers are not available.7