Different MACS sorting strategies for the enrichment of Lin⁻ (CD34⁺ CD45⁻) hematopoietic progenitor cells: preliminary study

Introduction

Magnetic-activated cell sorting (MACS) has become a standard method for cell separation in many different fields. Numerous publications have demonstrated its use, from lab bench to the clinic, both small to large scale, from abundant cells to rare cells with complex phenotypes, and from human and mouse cells to many other species (Bosio et al., 2009). There are various magnetic cell separation systems currently available. They differ principally in two features: the composition and size of the magnetic particles used for cell labelling (Miltenyi et al., 1990; Ugelstad et al., 1998) and the mode (positive isolation – enrichment or negative isolation – depletion) of magnetic separation.

At present, animal stem cells are widely used in the biomedical research. Since rabbit is genetically very close to human, rabbit hematopoietic stem/progenitor cells (HSC/HPC) could be a precious animal model cell line for the study of human hematopoietic disorders. However, among the published studies, there is none available that describes either this type of rabbit cells or their isolation itself. Cryopreservation of HSC/HPC for human therapeutic use in hematopoietic stem cell banks has been carried out for more than 30 years. Three types of HSC/HPC harvests from bone marrow, mobilized peripheral blood, and umbilical cord blood (Watt et al., 2007) are now routinely obtained by the MACS enrichment of CD34 positive cells. The same principle could be used for the isolation and cryopreservation of rabbit HSC/HPC. However, currently, there is no substantial data on the isolation and/or enrichment of rabbit hematopoietic stem cells, or on the detection of rabbit CD34⁺ cells. In the few published studies, mainly anti-human CD34 antibodies (Vašíček et al., 2018) were
used, due to the unavailability of specific anti-rabbit CD34 antibodies in the market. On the other hand, HSC/HPC are defined as lineage negative cells that do not express lineage commitment markers typical for mature hematopoietic cells, including T and B cells, monocytes/macrophages, granulocytes, erythrocytes, and their committed precursors (Gallacher et al., 2000). Thus, mature cells expressing lineage commitment markers can be easily depleted from the heterogeneous mixture of hematopoietic cells (Cheng et al., 2008) in order to enrich HSC/HPC.

Therefore, the aim of this preliminary study was to isolate rabbit HSC/HPC via removing mature hematopoietic cells that express CD45 (common leukocyte antigen) from the rabbit peripheral blood and bone marrow.

**Material and methods**

**Isolation of cells from rabbit blood and bone marrow**

Mononuclear cells from rabbit peripheral blood (PBMCs) and bone marrow (BMMCs) were isolated from 3 young (5 months-old does) and clinically healthy rabbits of the NZW line as described previously (Vašíček et al., 2016; 2018).

**Indirect immunomagnetic sorting of CD45+ PBMCs and BMMCs**

Briefly, mononuclear cells were aliquoted into prepared tubes and incubated with anti-rabbit CD45 monoclonal antibody (L12/201, mouse IgG1; Biorad, UK) for 15 minutes on ice. After washing, samples were incubated with Anti-Mouse IgG1 MicroBeads (Miltenyi Biotec, Germany) according to the producer’s manual. Three different types of sorting strategy available by AutoMACS Pro Separator (Miltenyi Biotec, Germany) were used in the experiments: 1) Depletes – Depletion in sensitive mode I for the removal of cells with normal antigen expression, if purity is the highest priority or for the removal of cells with low antigen expression; 2) Depl025 – Depletion in sensitive mode III for the removal of cells with low antigen expression, special program for very sensitive depletion; and, 3) Posselds – Positive selection in sensitive mode, double-column program for the isolation of cells with frequencies lower than 5% and low antigen expression. Control samples were analysed for CD45 expression but not sorted.

**Flow cytometry**

After MACS sorting, the APC-conjugated Labelling Check Reagent (LCR; Miltenyi Biotec, Germany), which binds to the microbeads, was used to control the sorting efficiency. Moreover, control samples (unsorted cells) and both MACS fractions (negative and positive) were incubated with another clone of anti-rabbit CD45 (ISC18A, mouse IgG2a; WSU, USA) and appropriate secondary anti-mouse IgG2a-FITC antibody (eBioscience, Austria). 7-AAD (eBioscience, Austria) was used in order to ex-
clude debris and dead cells from the analysis. Labelled cells before and after sorting were evaluated using a FACSCalibur flow cytometer (BD Biosciences, USA). At least 50,000 events (cells) were analysed in each sample. Double positive cells (CD45⁺LCR⁺) in MACS fractions served as an indicator of the sorting (depletion) efficiency in negative fractions.

qPCR analysis of CD34 expression

Total RNA isolation and cDNA synthesis from the control (unsorted) samples and both CD45⁻ and CD45⁺ fractions as well as quantitative polymerase chain reaction (qPCR) were performed according to Vašíček et al. (2017). Since rabbit CD34 has predicted two transcript variants, we used universal primers that comprise both known transcripts (Vašíček et al., 2017).

Results

Flow cytometry results are summarised in table 1. According to the obtained data, isolated rabbit PBMCs showed more than 98% positivity for CD45 antigen. On the other hand, only 70% of fresh rabbit BMMCs were positive for CD45. Double staining (CD45⁺LCR⁺) of separated cells revealed the best depletion efficiency in the negative fractions that were sorted using Depl025 mode for both PBMCs and BMMCs (about 10 and 3%, respectively).

<table>
<thead>
<tr>
<th>Sample</th>
<th>Depletes</th>
<th>Depl025</th>
<th>Posselds</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>CON (CD45⁻)</td>
<td>POS (CD45⁻LCR⁻)</td>
<td>NEG (CD45⁻LCR⁻)</td>
</tr>
<tr>
<td><strong>PBMCs</strong></td>
<td>98.8</td>
<td>93.4</td>
<td>37.4</td>
</tr>
<tr>
<td><strong>BMMCs</strong></td>
<td>67.3</td>
<td>94.4</td>
<td>22.4</td>
</tr>
</tbody>
</table>

Note: CON – control (unsorted) fresh sample analysed for the initial expression of CD45, POS – positive fraction, NEG – negative fraction, LCR – Labelling Check Reagent

qPCR analyses were used to evaluate the relative expression of CD34 in the control and sorted samples. The highest CD34 expression within the negative fractions were found in the PBMCs sample sorted using Depl025 mode in comparison to the control and positive fractions or, in general, within the used sorting strategies (Fig. 1). On
the other hand, BMMCs sample sorted by Posselds mode revealed the highest CD34 expression in negative fraction in comparison to the control and positive fractions or, in general, within the used sorting strategies (Fig. 2).

**Fig. 1.** Relative expression of CD34 in control (unsorted) and sorted PBMCs normalised to B2M (beta-2-microglobulin)

**Fig. 2.** Relative expression of CD34 in control (unsorted) and sorted BMMCs normalised to B2M (beta-2-microglobulin)
Discussion

We chose three different MACS sorting strategies for the negative or positive selection of CD45+ cells in order to obtain the highest sorting efficiency or the maximum removal of mature hematopoietic cells (CD45+) from the sample. According to the producer’s manual, these strategies differed mainly in the sample loading rate and the number of used magnetic columns: 1) Depletes – one column mode with loading rate 1 mL/min; 2) Depl025 – one column mode with loading rate 0.25 mL/min; and, 3) Posselds – two column mode with loading rate 1 mL/min for each column. Thus, the sorting efficiency of the used modes could differ according to the sample loaded as well as the level of desired antigen expression.

In this preliminary study, we assessed the sorting or depletion efficiency of MACS strategies used as the percentage of CD45+LCR+ cells. The best removal of CD45+ cells in both PBMCs and BMMCs samples was obtained using Depl025 mode (Tab. 1). So, we might assume that negative fractions (CD45+) obtained by this mode can be enriched for CD34+ cells. This finding strongly agreed with qPCR analysis of rabbit PBMCs sorted by this mode, since the highest relative expression of CD34 was noticed in the negative fraction in comparison to other negative fractions or the control and positive fraction within the sample (Fig. 1). However, the highest relative expression of CD34 in BMMCs was observed in the negative fraction of the sample sorted by a Posselds mode (Fig. 2) in comparison to other negative fractions or the control and positive fractions within the sample.

Although there are presently no any published studies focusing on the enrichment of rabbit Lin- (CD34+CD45-) cells, this technique based on the depletion of cells with lineage commitment markers is widely used for the purification of mouse (Ema et al., 2006) or human (Cheng et al., 2008) hematopoietic stem and progenitor cells.

Conclusions

This preliminary study demonstrated the possibility to obtain a cell population from rabbit PBMCs or BMMCs that is enriched for CD34+ cells. The best MACS sorting strategy according to the relative expression of CD34 antigen in enriched (negative fractions) were Depl025 for rabbit PBMCs or Posselds for BMMCs. As far as we know, this is the first published study focusing on the enrichment of rabbit HSC/HPC (CD34+) cells using the removal of CD45+ cells. However, further experiments are required in order to prove these preliminary results.

Acknowledgement

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Magnetic-activated cell sorting (MACS) has become a standard method for the isolation of hematopoietic stem/progenitor cells (HSC/HPC) in human or mouse models using CD34 antibodies. However, at present, there is no useable CD34 antibody that could be successfully used for the selection of rabbit HSC/HPC. Therefore, the aim of this preliminary study was to remove all mature cells (CD45+) from the heterogeneous mixture of rabbit peripheral blood and bone marrow mononuclear cells (PBMCs and BMMCs) in order to enrich these cell populations for the CD34+ cells. Briefly, cells were incubated with a CD45 antibody and proper magnetic microbeads. Three different MACS sorting strategies were used in the experiment that differed mainly in the sample loading rate and the number of used magnetic columns. Control (unsorted) and sorted cells were assessed for the sorting efficiency (% of double positive cells for CD45 and Labeling Check Reagent – LCR) by flow cytometry and for the relative expression of CD34 antigen by qPCR. According to flow cytometry, Depl025 mode showed the best sorting efficiency in terms of the lowest percentages of CD45+LCR+ cells for rabbit PBMCs as well as BMMCs. qPCR analysis confirmed this mode as the best in terms of the relative CD34 expression for rabbit PBMCs. However, a higher relative expression of CD34 in BMMCs was obtained by another mode – Posselds. In conclusion, this study demonstrates a possible

References


Abstract
enrichment of rabbit (CD34⁺) HSC/HPC by the magnetic depletion of mature hematopoietic (CD45⁺) cells.

**Key words:** CD34, CD45, flow cytometry, hematopoietic stem cells, MACS, qPCR, rabbit

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Różne strategie sortowania MACS do wzbogacania hematopoetycznych komórek progenitorowych Lin- (CD34⁺ CD45⁻): badania wstępne

**Streszczenie**
W modelu ludzkim lub mysim, sortowanie komórek aktywowane magnetycznie (MACS) stało się standardową metodą izolacji hematopoetycznych komórek macierzystych/progenitorowych (HSC/HPC) z użyciem przeciwciała CD34. Obecnie jednak nie ma użytecznego przeciwciała CD34, które mogłoby być z powodzeniem zastosowane do selekcji króliczego HSC/HPC. Dlatego celem tego wstępnego badania było usunięcie wszystkich dojrzałych komórek (CD45⁺) z heterogenicznej mieszaniny krwi obwodowej królica i jednojądrzastych komórek szpiku kostnego (PBMC i BMMC), aby wzbogacić ich populacje dla komórek CD34⁺. Komórki inkubowano z przeciwciałem CD45 i odpowiednimi mikrokulkami magnetycznymi. W eksperymencie zastosowano trzy różne strategie sortowania MACS, które różniły się głównie szybkością ładowania próbki i liczbą użytych kolumn magnetycznych. Kontrolne (nieposortowane) i posortowane komórki oceniano pod względem wydajności sortowania (% podwójnie dodatnich komórek dla CD45 i odczynnika do sprawdzania znakowania – LCR), za pomocą cytometrii przepływowej i względnej ekspresji antygenu CD34 przez qPCR. Zgodnie z cytometrią przepływową tryb Deplo25 wykazał najlepszą skuteczność sortowania pod względem najniższego odsetka komórek CD45⁺ LCR⁺ dla PBMC królica, jak również BMMC. Analiza qPCR potwierdziła ten tryb jako najlepszy do oceny względnej ekspresji CD34 dla PBMC króliców. Najwyższą względną ekspresję CD34 w BMMC uzyskano w innym trybie – Posselds. Podsumowując, niniejsze badanie demonstruje możliwe wzbogacenie HSC/HPC królica (CD34⁺) przez magnetyczne zużożenie dojrzałych komórek krwiotwórczych (CD45⁺).

**Słowa kluczowe:** CD34, CD45, cytometria przepływowa, hematopoetyczne komórki macierzyste, MACS, qPCR, królik

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