

Research Article

Preparation of Proper Immunogen by Cloning and Stable Expression of cDNA coding for Human Hematopoietic Stem Cell Marker CD34 in NIH-3T3 Mouse Fibroblast Cell Line

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Abstract

Purpose: Transmembrane CD34 glycoprotein is the most important marker for identification, isolation and enumeration of hematopoietic stem cells (HSCs). We aimed in this study to clone the cDNA coding for human CD34 from KG1a cell line and stably express in mouse fibroblast cell line NIH-3T3. Such artificial cell line could be useful as proper immunogen for production of mouse monoclonal antibodies.

Methods: CD34 cDNA was cloned from KG1a cell line after total RNA extraction and cDNA synthesis. Pfu DNA polymerase-amplified specific band was ligated to pGEMT-easy TA-cloning vector and sub-cloned in pCMV6-Neo expression vector. After transfection of NIH-3T3 cells using 3 µg of recombinant construct and 6 µl of JetPEI transfection reagent, stable expression was obtained by selection of cells by G418 antibiotic and confirmed by surface flow cytometry.

Results: 1158 bp specific band was aligned completely to reference sequence in NCBI database corresponding to long isoform of human CD34. Transient and stable expression of human CD34 on transfected NIH-3T3 mouse fibroblast cells was achieved (25% and 95%, respectively) as shown by flow cytometry.

Conclusion: Cloning and stable expression of human CD34 cDNA was successfully performed and validated by standard flow cytometric analysis. Due to murine origin of NIH-3T3 cell line, CD34-expressing NIH-3T3 cells could be useful as immunogen in production of diagnostic monoclonal antibodies against human CD34. This approach could bypass the need for purification of recombinant proteins produced in eukaryotic expression systems.

Introduction

CD34 gene, located on long arm of chromosome 1, consists of nine exons and codes for single-chain type I transmembrane glycoprotein with molecular weight 115-120 KDa.^{1,2} cDNA coding for human CD34 was first cloned and characterized by Simmons et al.^{3,4} CD34 has two alternatively spliced full (long) and truncated (short) isoforms differ in cytoplasmic tail.^{5,6} CD34 molecule is expressed on hematopoietic stem cells (HSCs) and progenitors, and also on high endothelial venules (HEVs) of lymph nodes.⁷⁻¹⁰ This molecule belongs to sialomucin family and because of high glycosylation and several N- and O-linked sialylatedglycans, plays an important role in adhesion of hematopoietic cells to bone marrow stroma and in binding of L-selectin on naive T lymphocytes to HEVs, process plays pivotal role in homing of these cells to parafollicular region of lymph

nodes.^{1,11,12} Today, CD34 is specific selection marker for HSCs, primitive and rare cell population in bone marrow with ability to produce and differentiate to all blood cells including immune cells.¹³⁻¹⁵ This marker has been widely used for identification, enumeration and isolation of HSCs in clinical and also research areas.^{16,17} For these purposes, specific monoclonal antibodies (MAbs) against CD34 molecules are employed, and production of such useful tools are inevitable for better and more specific recognition of surface CD34.^{1,18,19}

Production of MAbs by hybridoma technology was first introduced by George Kohler and Cesar Milstein.²⁰ Up to now, huge number of investigators have employed hybridoma technology, but with some modifications including different strategies for immunization of mice. Of them, some groups have stably expressed the gene

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coding for protein of interest in mouse fibroblast cell line, NIH-3T3,²¹ and have used the cells as immunogen.²²⁻²⁴ Because of murine origin of NIH-3T3 cell line, the only immunogen part of stably transfected cells is ectopically expressed protein. Using this strategy, all problems encountered in purification of recombinant proteins in eukaryotic systems are bypassed, and intact protein with complete conformational structure is used as immunogen. In addition, transfection of cDNA coding for a specific protein in NIH-3T3 cell line has been performed for purposes other than immunization of mice, e.g. the signaling potential or functional properties of the molecule.²⁵⁻²⁸

Here, we reported cloning of long isoform of human CD34 cDNA and stable ectopic expression on mouse fibroblast cell line NIH-3T3 for further experiments to produce anti-CD34 monoclonal antibodies useful in diagnosis and even therapeutic approaches.

Materials and Methods

Cells and bacteria

KG1a and NIH-3T3 cell lines were purchased from National Cell Bank of Iran (NCBI, Tehran, Iran) and cultivated in RPMI 1640 cell culture medium (Gibco, Darmstadt, Germany) supplemented by 20% Fetal Bovine Serum (FBS) (Gibco, Darmstadt, Germany), 100 µg/ml Penicillin and 100 IU/ml Streptomycin (Gibco, Darmstadt, Germany) under humidified and 5% CO₂ conditions. E.Coli strain DH5α was purchased from Promega Inc. (WI, USA) and cultured in Luria Bertani medium.

Flow cytometry

Evaluation of surface expression of CD34 molecule on KG1a, as a source for cloning of human CD34, was performed by indirect staining of KG1a cells. 5×10^5 cells were harvested and washed by PBS 1× containing 0.1% NaN₃. Mouse monoclonal anti-human CD34 antibody (Biolegend, London, UK) was added on cells in final concentration of 5 µg/ml. In parallel, cells were stained with isotype control antibody (Biolegend, London, UK), as negative control. After 1 hour incubation at 4°C, cells were washed two times and then FITC-conjugated sheep anti-mouse immunoglobulin (Avicenna Research Institute, Tehran, Iran) was added in 1/50 dilution. Cells were incubated in a dark place for 1 hour at 4°C and after two times washing, they were scanned in flow cytometer (BD FACSCalibur flow cytometer).

Total RNA extraction and cDNA synthesis

5×10^6 KG1a cells were harvested and washed two times by RPMI 1640 culture medium. After final centrifugation, supernatant was completely discarded and the pellet was thoroughly resuspended. Cells were lysed by 1 ml RNX-plus solution (CinnaGen, Tehran, Iran) and total RNA was extracted according to manufacturer's recommendations. Briefly, after adding 200 µl chloroform (Merck, Darmstadt, Germany) and incubation for 5 minutes on ice, the solution was

centrifuged at 12000 rpm for 15 minutes at 4°C and colorless aqueous phase was transferred to other tube carefully. Isopropanol (Merck, Darmstadt, Germany) in equal volume was added and after mixing and incubation for 15 minutes on ice, the solution was centrifuged and the precipitated RNA was seen. RNA was washed in 75% ethanol (Merck, Darmstadt, Germany) and resolved in RNase-free double distilled water. Quantity and quality of RNA was evaluated by UV spectrophotometry and agarose gel electrophoresis, respectively.

Five µg RNA was employed for synthesis of first strand cDNA using MMuLV reverse transcriptase (RT) enzyme (Thermo Fisher Scientific, Inc., MA, USA) and random hexamer (N6) primer (Thermo Fisher Scientific, Inc., MA, USA). Amplification of beta actin as a house keeping gene in polymerase chain reaction (PCR), as discussed later, was performed for confirming synthesis of cDNA.

Primer design and Polymerase chain reaction

To amplify human CD34 cDNA, specific primers were designed so that forward and reverse primers contained KpnI and HindIII restriction sites, respectively. Selection of restriction enzymes was on the basis of multiple cloning sites (MCS) of expression vector will used for expression of protein in eukaryotic system. For this purpose, reference sequence for human CD34 mRNA isoform 1 (NM_001025109.1) was obtained and the sequence was imported in NEBCutter web-based software (version 2.0) to find if the sequence has restriction sites for KpnI and HindIII restriction enzymes. The sequence of forward and reverse primers were selected from the beginning and ending part of reference sequence, respectively and were analyzed in OligoCalc web-based software to check for self-complementarity and hairpin formation of primers. Also, melting temperature (T_m) of both primers was calculated for upcoming experiments and adjusted against each other. To facilitate efficient translation in eukaryotic expression system during subsequent experiments, Kozak consensus sequence (GCCACC) was considered after the sequence for KpnI restriction site (at the beginning of forward primer) and upstream of the start codon. As presence of G nucleotide at position +4 is considered as strong consensus,²⁹ i.e. it maximizes the expression level of gene of interest, nucleotide +4 was replace by G nucleotide. To terminate the translation, on the other hand, stop codon was considered at the end of sequence for CD34 cDNA and before the sequence for HindIII restriction site. Optimized PCR conditions were obtained by performing reactions in different concentrations of MgCl₂ and a spectrum of annealing temperatures. Twenty-five µl reaction mixture contained 2.5 µl 10× PCR buffer, 1.5 µl 10 mM dNTPs (Thermo Fisher Scientific, Inc., MA, USA), 1µl each primer (10 pmol/µl), 0.2 µl Taq DNA polymerase (10 U/µl) (CinnaGene, Tehran, Iran) and 1 µl cDNA. Each PCR reaction was underwent initial denaturation at 95°C for 5 minutes followed by 39 cycles of denaturation (95°C for

30 seconds), annealing (different temperatures for 30 seconds) and extension (72°C for 45 seconds), and final extension at 72°C for 10 minutes. For obtaining PCR product with high fidelity, amplification of CD34 cDNA was performed using Pfu DNA polymerase (Thermo Fisher Scientific, Inc., MA, USA) in previously optimized PCR conditions, with the exception of 2.2 minutes for extension step, and A-tailing was done at 72 °C for 7 minutes using Taq DNA polymerase. PCR products were subjected to agarose gel electrophoresis, visualized by ethidium bromide at final concentration 5 µg/ml and documented in UVP Gel Documentation System (UVP, CA, USA).

TA-cloning of CD34 cDNA

After agarose gel electrophoresis, specific band with correct size was extracted using GeneJet Gel Extraction Kit (Thermo Fisher Scientific, Inc., MA, USA) and then was ligated into pGEMT-easy vector (Thermo Fisher Scientific, Inc., MA, USA) using 3 units T4 DNA ligase (Thermo Fisher Scientific, Inc., MA, USA). After overnight incubation at 4°C, ligation mixture was transformed by heat shock method into DH5α competent bacteria, previously prepared by 0.1M CaCl₂ solution. After refreshing bacteria by adding fresh LB broth and incubation for 1 hour at 37°C and shaking in 150 rpm, they were transferred on LB agar medium containing 100 µg/ml Ampicillin (Dana, Tabriz, Iran), 40 µl X-Gal (20 µg/ml) (Thermo Fisher Scientific, Inc., MA, USA) and 40 µl IPTG (0.1mM) (Thermo Fisher Scientific, Inc., MA, USA). Plate was incubated overnight in 37°C and white colonies were evaluated by colony-PCR in conditions similar to conditions employed for amplification of CD34 cDNA. One positive colony was selected according to specific band with correct size and cultured in LB broth medium overnight at 37 °C and shaking in 250 rpm. Miniprep preparation was performed by Gene JET Plasmid miniprep kit (Thermo Fisher Scientific, Inc., MA, USA) according to manufacturer's recommendations. For initial confirmation of presence of a gene insert with correct size and proper restriction sites i.e. for KpnI and HindIII restriction enzymes (Thermo Fisher Scientific, Inc., MA, USA), double digestion was performed using 10 units of each enzyme overnight at 37°C. Digestion product along with undigested construct was run in agarose gel to visualize excised band. Consequently, selected construct was subjected for sequencing by T7 promoter and SP6 universal primers, to confirm accurate nucleotide sequence of insert. Chromatogram was analyzed first for good peaks for all nucleotides, and then entire sequence was aligned in NCBI database for checking proper sequence of inserted gene. On the other hand, flanking region of inserted gene was checked for vector-specific sequences to insure proper cloning vector and avoid cross-contamination by other commonly used vectors.

Preparation of pCMV6-Neo/CD34 recombinant construct

Double digested CD34 cDNA was sub-cloned in pCMV6-Neo digested in the same way. Ligation was done as previously described and competent DH5α bacteria were transformed by ligation mixture. Screening of colonies was performed on LB-Agar ampicillin plate and positive colonies were identified by colony-PCR reaction. Miniprep for one positive colony was prepared and direct sequencing using of V1.5 and XL39 primers was done for confirmation. To obtain high quality and quantity of finalized recombinant construct without any contamination by bacterial endotoxin, Maxiprep was prepared by EndoFree Plasmid Maxi Purification Kit (Thermo Fisher Scientific, Inc., MA, USA) according to manufacturer's recommendations.

Eukaryotic expression of human CD34 in mouse fibroblast cell line

NIH-3T3 mouse fibroblastic cells were cultured in the same conditions for KG1-a cells except of RPMI culture medium supplemented by 10% FBS. Cells were subjected to titration for resistance against G418. 10⁵ cells were seed in 6-well cell culture plate and cultured in the presence of different concentrations (300 to 800 µg/ml) of G418 (Thermo Fisher Scientific, Inc., MA, USA). They were monitored for one week and minimum concentration of G418 for selection of NIH-3T3 cells was determined. 10⁵ NIH-3T3 cells were cultured overnight in 6-well cell culture plate and transfected by the mixture of 3 µg pCMV6-Neo/CD34 recombinant construct and 6 µl JetPEI transfection reagent (Polyplus-transfection Inc. NY, USA) according to manufacturer's recommendations. In parallel, cells were transfected by expression vector devoid of CD34 cDNA (mock transfection). 48 hours after transfection, transient expression of CD34 antigen was examined in flow cytometry methods, as described earlier. To obtain stable expression, cells were cultured in complete cell culture medium containing G418. Starting concentration was determined by above-mentioned experiment on NIH-3T3 cells. Growing of transfected cells and dying of untransfected cells were monitored and live cells were subjected to gradually increasing concentrations of G418 up to 1500 µg/ml during two months.

Results

Surface Expression of CD34

Flow cytometric analysis of surface expression of human CD34 on KG1a myeloid cell line, as a source for amplification of CD34 cDNA, showed strong fluorescent detected in 99 % of cells by flow cytometer as M2. In contrast, surface staining of cells by isotype control, as negative control, showed very weak signal (M1: 1%).

Amplifying CD34 cDNA

Total RNA was purified from KG1a cells and cDNA was synthesized using 5 µg of total RNA. Processes were confirmed by agarose gel electrophoresis and beta actin gene amplification, respectively. Designed forward and reverse primers were 5'- GGTACCGCCACCATGGTG

GTCCGCAGGGGCGC-3' and 5'- AAGCTTTCACAATTCGGTATCAGCCACCACG-3'. They showed no self-complementarity and hairpin formation. On the other hand, the reference sequence had no restriction sites for selected restriction enzymes i.e. KpnI and HindIII. Using Taq DNA polymerase, conditions for amplification of CD34 cDNA were optimized. In 1 mM of MgCl₂ and annealing temperature of 62°C, specific band with 1176 bp length (1158 bp for CD34 cDNA and 18 bp for two restriction sites and Kozak consensus sequence) was obtained and the reaction was repeated using Pfu DNA polymerase, but using 2.2 minutes extension time at 72°C. After A-tailing by Taq DNA polymerase, PCR product was run in agarose gel electrophoresis (Figure 1A) and specific band was then extracted.

TA-cloning of CD34 cDNA

Extracted PCR product and pGEMT-easy TA-cloning vector were mixed in ratio 5:1 and incubated overnight. Transformation of competent bacteria was performed using heat shock method and after refreshing of bacteria, they spread on LB agar containing ampicillin, X-gal and IPTG. After overnight incubation in 37°C, 8 colonies were obtained. Of them, 5 colonies were white and selected for colony-PCR reaction. Using the same PCR conditions for amplifying CD34 cDNA, 2 colonies showed strong single bands comparable to the size for CD34 cDNA (Figure 1B). One of the colonies was cultured overnight in LB broth medium containing Ampicillin. Preparation of Miniprep was performed, and quality and quantity were measured by agarose gel electrophoresis and UV spectrophotometry, respectively. According to cloning strategy, the construct was subjected to double digestion by KpnI and HindIII restriction endonucleases and successful excision of inserted CD34 cDNA was seen (Figure 1C). Initial analysis of two-sided sequencing of the construct showed good quality peaks and alignment of the inserted sequence in NCBI data base showed complete matching of cloned CD34 cDNA to reference sequence for CD34 mRNA isoform 1 (Figure 2).

Construction of recombinant pCMV6-Neo/CD34 construction

pGEMT-easy/CD34 recombinant cloning vector was digested by KpnI and HindIII restriction enzymes. Excised CD34 cDNA with 1176 bp band was ligated into double digested pCMV6-Neo expression vector by the same enzymes and pCMV6-Neo/CD34 construction was transformed into competent DH5α bacteria. After transformation, 8 colonies were obtained from cultured DH5α bacteria in LB agar/ampicillin medium. After performing colony-PCR reaction, 3 colonies showed successful insertion and one of the colonies was selected for preparation of Miniprep, direct sequencing and preparation of Maxiprep for subsequent experiments.

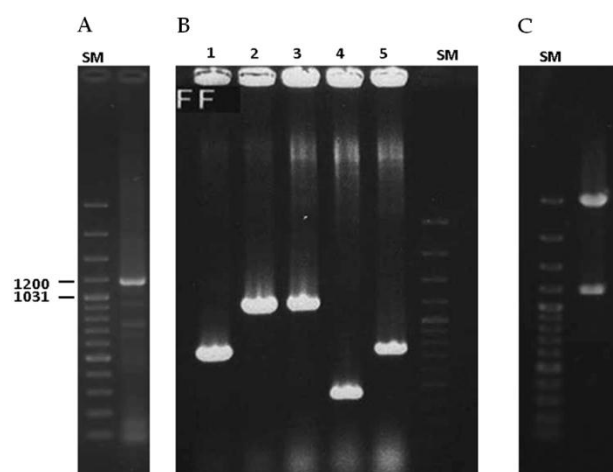


Figure 1. Cloning and subcloning of CD34 cDNA. (A) Amplification of specific band for human CD34 cDNA using Pfu DNA polymerase, (B) Colony-PCR reaction on five white colonies (1-5) after blue/white selection. (C) Excision of 1176 bp band for human CD34 cDNA after double digestion of the construct using KpnI and HindIII restriction enzymes. SM: DNA size marker (bp)

Transfection of NIH-3T3 cells and expression of human CD34 protein

NIH-3T3 mouse fibroblastic cell line was transfected by pCMV6-Neo/CD34 recombinant expression vector using JetPEI transfection reagent and transient expression of CD34 was confirmed by surface flow cytometry 48 hours after transfection (Figure 3D). After that, pCMV6-Neo/CD34 and mock-transfected and also untransfected NIH-3T3 cells were subjected to growth in complete cell culture medium containing 400 µg/ml G418, minimal dose for dying NIH-3T3 cells. Untransfected cells died and transfected cells were continued to be grown in G418 up to 1500 µg/ml during two months. Cells were finally analyzed by surface flow cytometry and significant positive reactivity of pCMV6-Neo/CD34-transfected cells (95%) (Figure 3E) was seen. On the other hand, untransfected and mock-transfected NIH-3T3 cells showed no positive signal in flow cytometry (3% and 4%, respectively) (Figure 3 B and C).

Discussion

CD34 molecule has long been used as a specific marker for hematopoietic stem cells (HSCs) in research and clinic.³⁰ For this purpose, polyclonal and monoclonal antibodies (MAbs) have been valuable tools, and several study groups tried to produce specific monoclonal antibodies against human CD34.³¹ According to modified protocol introduced in 1970s for producing mouse monoclonal antibodies,²⁰ the first and one of the most important steps of hybridoma technology is preparing a good and proper immunogen. It should be as intact as possible for preserving all potential epitopes. On the other hand, using full-length protein with proper post transcriptional modifications similar to that occurs for native protein, is helpful for obtaining functionally more active and useful MAbs.³² Additionally, immunization of animal with purified

protein is done by mixing antigen to complete (CFA) and incomplete Freund's adjuvant (IFA) usually administered subcutaneously. According to American Association for Laboratory Animal Science (IACUC) Policy on Administering Complete Freund's Adjuvant,³³ this protocol could induce unwanted local inflammation causing skin ulcerations and draining sinuses with granulomas and may lead to improper antibody response to antigen and limited number of antigen-specific clones. All above mentioned reasons force us to produce recombinant proteins in eukaryotic

expression systems, in which purification step is tedious and challenging. However, immunization of mouse using stably-transfected murine NIH-3T3 cell line needs neither adjuvant nor purification and the route of injection is intraperitoneal with minimal side-effects for animal. Stable expression of cDNA coding for protein of interest in mouse fibroblast cell line, NIH-3T3, has been used by several investigators as an approach to prepare immunogen for immunization of mouse²²⁻²⁴ and also for other research purposes.²⁵⁻²⁸

Homo sapiens CD34 molecule (CD34), transcript variant 1, mRNA

Sequence ID: [ref|NM_001025109.1|](#) Length: 2621 Number of Matches: 1

Range 1: 259 to 1413 GenBank Graphics					▼ Next Match ▲ Previous Match	
Score	Expect	Identities	Gaps	Strand		
2134 bits(1155)	0.0	1155/1155(100%)	0/1155(0%)	Plus/Plus		
Query 1	ATGCTGGTCCGCAGGGGCGCGCGCAGGGCCCAGGATGCCGCGGGGCTGGACCGCGCTT				60	
Sbjct 259	ATGCTGGTCCGCAGGGGCGCGCGCAGGGCCCAGGATGCCGCGGGGCTGGACCGCGCTT				318	
Query 61	TGCTTGCTGAGTTTGCTGCCTTCTGGGTTTCATGAGTCTTGACAACAACGGTACTGCTACC				120	
Sbjct 319	TGCTTGCTGAGTTTGCTGCCTTCTGGGTTTCATGAGTCTTGACAACAACGGTACTGCTACC				378	
Query 121	CCAGAGTTACCTACCCAGGGAACATTTTCAAATGTTTCTACAAATGTATCCTACCAAGAA				180	
Sbjct 379	CCAGAGTTACCTACCCAGGGAACATTTTCAAATGTTTCTACAAATGTATCCTACCAAGAA				438	
.....						
.....						
Query 1021	TCAGGACCTGGGACCTCCCTTGAGGCTCAGGGAAAGGCCAGTGTGAACCGAGGGGCTCAG				1080	
Sbjct 1279	TCAGGACCTGGGACCTCCCTTGAGGCTCAGGGAAAGGCCAGTGTGAACCGAGGGGCTCAG				1338	
Query 1081	GAAACGGGACCGGCCAGGCCACCTCCAGAAACGGCCATTAGCAAGACAACACGTGGTG				1140	
Sbjct 1339	GAAACGGGACCGGCCAGGCCACCTCCAGAAACGGCCATTAGCAAGACAACACGTGGTG				1398	
Query 1141	GCTGATACCGAATTG				1155	
Sbjct 1399	GCTGATACCGAATTG				1413	

Figure 2. Alignment of amplified cDNA for long isoform of human CD34 to reference sequence in NCBI database. Comparing the 1158 bp amplified sequence with reference sequence for long isoform (variant 1) of human CD34 showed complete alignment. Briefly, 5' and 3' ends of alignment have been showed

KG1 is a myeloblastic cell line with CD7⁻ CD34⁺ phenotype and has been obtained from an old AML patient. This cell line was first identified and introduced in 1978. After 35 repetitive passages, a sub-lineage named KG1a (CD7⁺ CD34⁺) was obtained with absence of some characteristics of its own parent cell line, including response to colony-stimulating factor (CSF) and expression of FCγ receptor.³⁴ To insure high expression of CD34 on KG1a cells, they were subjected to indirect immunofluorescent staining, and results showed strong surface expression of CD34. So, KG1a was used in this study as a cellular source for expression of human CD34 gene and amplifying cDNA coding for CD34 protein.

Amplification of CD34 cDNA resulted in 1176 bp amplicon, subsequently cloned in pGEMT-easy TA-

cloning vector and completely aligned to reference sequence of mRNA for long isoform of human CD34 isoform 1. cDNA was then sub-cloned in a eukaryotic expression vector for further transfection in NIH-3T3, as it has been reported to be a mouse fibroblast cell line.³⁵ For this purpose, pCMV6-Neo as an expression vector with a strong promoter for cytomegalovirus (CMV) was utilized for. pCMV6-Neo contains a resistance gene to G418 antibiotic (neomycin) and it is useable for screening of transfected cells. So transfected NIH-3T3 cells would be resistant and alive against G418 antibiotic.

Conclusion

In this study the cDNA coding for long isoform (transcript variant 1) of human CD34 was amplified and

cloned from KG1a myeloid cell line and stably expressed in mouse NIH-3T3 cell line. Stable expression of CD34 molecule in NIH-3T3 resulted in an appropriate

immunogen for production of monoclonal antibodies useful in diagnostic and research areas.

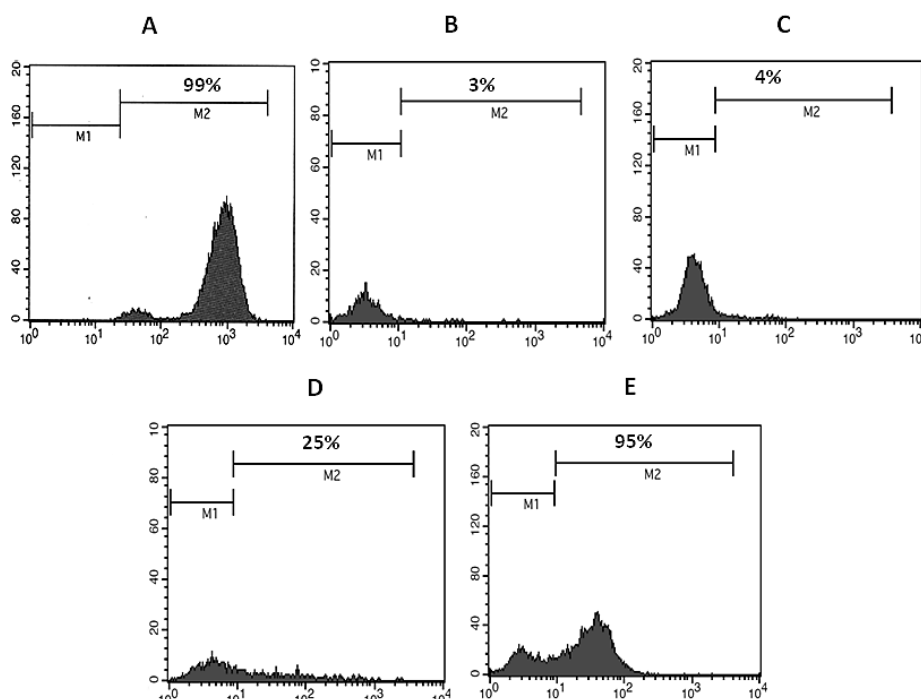


Figure 3. Flow cytometric analysis of expression of human CD34. Using specific monoclonal antibody, expression of human CD34 on KG1a (A), NIH-3T3 (B), mock-transfected NIH-3T3 (C) and pCMV6-Neo/CD34-transfected NIH-3T3 cells (D, for transient and E, for stable expression) were analyzed.

Acknowledgments

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Ethical Issues

Not applicable.

Conflict of Interest

There is no conflict of interest to be reported.

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