The CD45^{low}CD271^{high} Cell Prevalence in Bone Marrow Samples May Provide a Useful Measurement of the Bone Marrow Quality for Cartilage and Bone Regenerative Therapy

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Background: Bone marrow aspirates and concentrates are increasingly being used for musculoskeletal regenerative therapies, providing bone and cartilage progenitors. However, the quality of these bone marrow samples remains imprecise within clinical settings. As there is a need for the development of these therapies, a method of counting CD45^{low}CD271^{high} cells was optimized and tested as an indicator of bone marrow sample quality.

Methods: Bone marrow aspirates were collected from 54 donors (28 male and 26 female; median age of 48 years). The reagent concentrations were optimized for fast staining, and an acoustic-focusing flow cytometer (Attune) was used to enable automated CD45^{low}CD271^{high} cell counting in bone marrow aspirates, bone marrow concentrates, and samples loaded onto a collagen scaffold. The CD45^{low}CD271^{high} cell counts were compared with those obtained using another flow-cytometry-based method (LSR II) and with connective tissue progenitor (CTP) counts quantified using a colony forming unit-fibroblast (CFU-F) assay.

Results: The optimized method enabled the counting of CD45^{low}CD271^{high} cells within only 15 minutes. The quantified cell counts (median, 1,520; range, 96 to 20,992 cells/mL of bone marrow) were positively correlated with the CTP counts (p < 0.0001; r = 0.7237). In agreement with CFU-F and LSR II-based assays, the CD45^{low}CD271^{high} cell counts quantified using the Attune-based method decreased with age in the samples from female but not male donors (p = 0.0015 and p = 0.3877, respectively). A significant increase in CD45^{low}CD271^{high} cell counts was detected following bone marrow concentration (mean, 5-fold; 95% confidence interval [CI], 3.6 to 7.2-fold). Additionally, the number of CD45^{low}CD271^{high} cells attached to the collagen scaffold was positively correlated with the number of progenitor cells that survived on the scaffold after 2-week culture (p = 0.0348).

Conclusions: An assay for counting CD45^{low}CD271^{high} cells may provide a useful measurement of bone marrow quality. While the specificity of this measurement of CD45^{low}CD271^{high} cells remained low in our experimental conditions, CD45^{low}CD271^{high} cell counts were positively and modestly correlated with the prevalence of CTPs.

Clinical Relevance: A fast and automated assessment of bone marrow aspirate/concentrate quality using CD45^{low}CD271^{high} cell counting may be a useful tool for improving the quality of regenerative therapy.

The field of regenerative medicine is constantly evolving, with new approaches for cartilage and bone healing dominating clinical and research activities. Targeting the environment of the nonunion of fractures or joint degeneration with biological modifiers, such as progenitor cells and/or growth factors, represents a promising therapeutic strategy¹⁻³. The rationale behind such a strategy is that the repopulation of cartilage and bone defects is possible, as long as the progenitor cells are present. For example, the potential efficacy of the microfracture technique for cartilage repair in osteoarthritis could

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be related to the effect of the subchondral bone progenitor cells that produce growth factors and tissue matrix⁴. Furthermore, the use of bone marrow progenitors with or without platelet-rich plasma has been reported to aid bone repair in preclinical and clinical studies of osteochondral defects, metaphyseal bone defects, and femoral head osteonecrosis⁵⁻⁹.

Authors of previous studies have reported the clinical value of bone marrow aspirates or concentrates, showing a positive correlation between the number of applied bone marrow progenitors and favorable clinical outcomes in tibial fracture nonunion¹⁰, osteoarthritis¹¹, and osteonecrosis therapy¹²⁻¹⁵. Despite the advantages of using bone marrow aspirates or concentrates, the quality of these samples remains difficult to assess and is poorly controlled. Furthermore, the number of progenitor cells in bone marrow aspirates is widely variable, depending on the aspiration site, volume, and surgical technique^{16,17} as well as on donor-related factors such as age and sex¹⁸. The determination of the quality of bone marrow samples is crucial in order to optimize clinical outcomes, cost, and time associated with cell-based therapies. The colony forming unitfibroblast (CFU-F) assay facilitates the counting of connective tissue progenitors (CTPs) and is commonly used as an indicator for bone marrow sample quality^{19,20}; however, it usually takes several days to be informative. CTPs represent the progenitors in native tissues that are able to form colonies in vitro. However, CTP concentration and prevalence can be influenced by bone marrow processing methods. The efficiency of colony formation (the likelihood that a viable CTP will form a colony when placed into a CFU-F assay) is also dependent on culture conditions^{17,21}.

The aim of the current study was to introduce a fast and automated method, with minimum sample processing, that helps to indicate the quality of bone marrow aspirates and concentrates. Bone marrow cells isolated on the basis of the CD45^{low}CD271^{high} phenotype are known to express CD73, CD90, and CD105, but not hematopoietic lineage markers, and on culture, generate multipotential stromal cells fully consistent with the International Society for Cellular Therapy (ISCT) criteria²²⁻²⁴. Importantly, several groups have reported that no colony-forming cells are present in the CD271-negative fraction of bone marrow cells, and bone marrow colony-forming activity is completely confined to the CD45^{low}CD271^{high} cells²⁴⁻³⁰. Therefore, we chose to quantify, by a flow-cytometry-based assay, CD45^{low}CD271^{high} cell counts, as an indicator of the quality of bone marrow aspirates and concentrates. We hope that our work will contribute to the standardization of therapies and the setting of thresholds between the "success" and "failure" of musculoskeletal regenerative therapies.

Materials and Methods

Bone Marrow Aspirates

Bone marrow samples from 54 donors (28 male and 26 female) were used for this study under ethical approval (06/Q1206/127, National Research Ethics Committee Yorkshire & Humber-Leeds East). The donors were admitted at Leeds General Infirmary for orthopaedic surgery; they did not have any systemic illness, cancer, or metabolic diseases. Donor age ranged from 22 to 80 years (median, 48 years). Two groups of donor samples were used, as described in Table I. All bone marrow aspirates were consistently harvested from the same location (zone 6) of the posterior region of the iliac crest as previously described^{16,25,31}. Each sample analysis was carried out on a single bone marrow sample harvested from a single individual at a single time point.

Using Flow Cytometry for Counting CD45^{low}CD271^{high} Cells A 100-µL volume of whole bone marrow was stained using a 3-marker panel containing Vybrant DyeCycle Ruby 2.5-mM solution in dimethyl sulfoxide (DMSO) (Thermo Fisher Scientific), a DNA-selective dye that only labels the nucleated cells, enabling the gating out of red blood cells (RBCs) and platelets. Additionally, the panel contained anti-CD45 antibody (V450, clone HI30, mouse IgG1k; concentration, 100 µg/mL; BD Biosciences) and anti-CD271 antibody (PE, clone ME20.4-1.H4, mouse IgG1; concentration, 0.75 µg/mL; Miltenyi Biotec). The phenotype indicating bone marrow progenitor cells (CD45^{low}CD271^{high} cells) was applied as previously described¹⁷. The manufacturer recommendation for CD45 and CD271 antibodies was 15 minutes at room temperature and for Vybrant DyeCycle Ruby dye, 15 minutes at 37°. However, the antibody/dye staining was optimized to count CD45^{low}CD271^{high} cells within the shortest time (see Results). We used the Attune acoustic-focusing flow cytometer (Thermo Fisher Scientific), which allowed automated cell counting. For some experiments, the CD45^{low}CD271^{high} cell counts were quantified using our previously published flow cytometry-based method¹⁷. Briefly, this method involved bone marrow sample staining (using CD90, CD271, and CD45), RBC lysis, and then adding CountBright absolute counting beads (Thermo Fisher Scientific). The data acquisition for this part of the study was performed using an LSR II flow cytometer (BD Biosciences).

CFU-F Assay

The CFU-F assay was employed as previously described¹⁷ to count CTPs, whereby the bone marrow samples were added to StemMACS MSC expansion media (Miltenyi Biotec) and then cultured for 14 days. The colonies were visualized using methylene blue and counted manually. Each colony was defined as having at least 50 cells³².

TABLE I Donor Sample Groups		
	Group 1	Group 2
No. of donors	39	15
Bone marrow aspirate volume (mL)	8	60
Aspiration location	Posterior region of the iliac crest	Posterior region of the iliac crest
Assessments performed	Optimization of marker concentration (Fig. 1) Comparison of assays (Figs. 2 and 3) Assessment of CD45 ^{low} CD271 ^{high} cells attached onto collagen scaffold (Fig. 5)	Assessment of CD45 ^{low} CD271 ^{high} cells ir bone marrow concentrates (Fig. 4)

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Bone Marrow Concentration

Bone marrow samples (n = 15) were concentrated by means of the gradient centrifugation using the BioCUE device (Zimmer Biomet). The bone marrow aspirates were collected into syringes, washed with anticoagulant acid citrate dextrose (ACD), and loaded into the BioCUE device. From both pre-concentration and post-concentration fractions, aliquots were analyzed for CD45^{low}CD271^{high} cell and CTP counts using the Attune-based method and CFU-F assays, respectively. The counting of platelets was performed for some samples (n = 10) using an automated hematopoietic cell counter (Sysmex; Sysmex Ltd).

Loading of Bone Marrow Samples on a Collagen Scaffold

The bone marrow aspirates were used to load a collagen scaffold (Bio-Gide; Geistlich Pharma). Samples from pre-loading and post-loading fractions of the bone marrow aspirates were processed to count CD45^{low}CD271^{high} cells. The bone marrow-loaded scaffolds were cultured for 2 weeks and subsequently processed to quantify bone marrow progenitors that survived on the scaffolds, as previously described²¹. Briefly, the scaffolds were digested using 0.25% collagenase (Stem Cell Technologies). As the surface expression of CD271 can be reduced on cultured cells²⁴, the extracted cells were stained using CD45, CD90





Optimization of the staining and counting of CD45^{low}CD271^{high} cells using the Attune flow cytometer. **Figs. 1-A, 1-B, and 1-C** Comparison of CD45^{low}CD271^{high} cell counts quantified using 3 different volumes of anti-CD45 and CD271 antibodies and Vybrant DyeCycle Ruby (VDR) dye (Student paired t test; n = 10 samples). **Figs. 1-D, 1-E, and 1-F** Comparison of CD45^{low}CD271^{high} cell counts quantified using 1-step and 2-step staining (Student paired t test; n = 6 samples), at different staining temperatures (Student paired t test; n = 7 samples; RT = room temperature), and after 5, 10, and 15-minute staining (Student paired t test, n = 10 samples). **Fig. 1-G** Comparison of CD45^{low}CD271^{high} cell counts quantified using automated counting or using counting beads, on the Attune flow cytometer (Wilcoxon matched-pairs signed-rank test; n = 9 samples). **Fig. 1-H** The CD45^{low}CD271^{high} cell counts were quantified in bone marrow aspirates that were stained undiluted or diluted ×5 or ×10 (n = 5 samples; S = sample).

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 $(\mbox{BioLegend}),$ and CD73 (Miltenyi Biotec) antibodies and counted using counting beads.

Statistical Analysis

The statistical analysis and graph preparation were performed using Prism software (version 7.0a; GraphPad). The normal distribution of the data was assessed using the Shapiro-Wilk normality test, and the appropriate test for the data analysis was applied accordingly (as specified in the figure legends). A p value of <0.05 was considered significant.

Results

Optimization of Marker Concentration

The fast staining of the bone marrow samples was initially optimized. With respect to the anti-CD45 antibody, the

number of CD45^{low}CD271^{high} cells counted using a volume of 10 μ L of the antibody was significantly higher compared with the use of 5 μ L (p = 0.0486) but did not differ significantly from the use of 20 μ L (p = 0.3969) (Fig. 1-A). Regarding the anti-CD271 antibody, the number of CD45^{low}CD271^{high} cells quantified using a volume of 20 μ L of the antibody was significantly higher compared with the use of 10 μ L (p = 0.0450) but similar to the number quantified using 40 μ L (p = 0.5443) (Fig. 1-B). When 3 different volumes of the Vybrant DyeCycle Ruby dye were used, the CD45^{low}CD271^{high} cell counts were similar (p = 0.1901 for 3 compared with 5 μ L, and p = 0.1140 for 10 compared with 5 μ L) (Fig. 1-C).

We next tested the use of CD45 and CD271 antibodies followed by Vybrant DyeCycle Ruby (2-step staining) compared



Fig. 2

The CD45^{low}CD271^{high} cell counts quantified using the Attune-based method were compared with those obtained by the LSR II-based method and with CTP counts assessed by CFU-F assay. Examples of high-quantity (**Fig. 2-A**) and low-quantity (**Fig. 2-B**) bone marrow progenitor samples are shown. The correlation between CD45^{low}CD271^{high} cell counts quantified using the Attune-based method and the LSR II-based method (n = 33 samples) (**Fig. 2-D**) was analyzed. A Spearman r test was used for the correlation analysis.

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with the addition of all markers in 1 step. The cell counts did not differ significantly between 1-step and 2-step staining methods (p = 0.6581) (Fig. 1-D). The CD45^{low}CD271^{high} cell counts were also similar using different staining temperatures (p = 0.7237 for 4° C compared with room temperature, p = 0.1261 for 37° C compared with room temperature, and p = 0.3558 for 4°C compared with 37°C) (Fig. 1-E). The CD45^{low}CD271^{high} cell counts also did not differ significantly for 5-minute staining compared with 10-minute staining (p = 0.1981), 5-minute staining compared with 15-minute staining (p = 0.5028), or 10-minute staining compared with 15-minute staining (p = 0.7870)(Fig. 1-F).

For each bone marrow sample, the acquisition time on the Attune flow cytometer was completed within 10 minutes. An internal control for automated counting was used (the use of counting beads), and automated and bead-dependent quan-

Females

p=0.0015

r = -0.6900

Α

CD45lowCD271high cells (Attune)

102

20

60 80

AGE (years)

40

tification were comparable (p = 0.3750) (Fig. 1-G). The data also showed similar cell counts quantified when bone marrow samples were diluted ×5 or ×10 compared with undiluted samples (Fig. 1-H). Overall, we optimized an automated and simple assay of CD45^{low}CD271^{high} cells within only 15 minutes.

Comparison of Assays

CTPs (CFU-F assay)

10

102

10

10⁰

20 40 60 80

AGE (years)

Using the Attune-based method, the median percentage of CD45^{low}CD271^{high} cells per total bone marrow cells was 0.016% (95% confidence interval [CI], 0.009% to 0.032%). The absolute counts of CD45^{low}CD271^{high} cells showed a median of 1,520 cells/mL of bone marrow (95% CI, 1,056 to 6,112; range, 96 to 20,992 cells/mL of bone marrow).

The results obtained by the Attune-based, LSR II-based, and CFU-F assays were consistent, indicating, for example, high

p=0.0055

r = -0.6904





Fig. 3



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p=0.0070

r = -0.6563

CD45lowCD271high cells (LSRII)

10

10

10

101

20 40 60 80 100

AGE (years)

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or low-quality bone marrow samples (Figs. 2-A and 2-B). The CD45^{low}CD271^{high} cell counts obtained using the Attune-based method were close to those of the LSR II-based method (median, 1,311; 95% CI, 900 to 5,533; range, 87 to 20,471 cells/mL of bone marrow). However, the CD45^{low}CD271^{high} cell counts obtained by the Attune-based assay were higher than the counts of CTPs (median 60; 95% CI, 45 to 190; range, 3 to 900 CTPs/mL of bone marrow). The CD45^{low}CD271^{high} cell counts measured using the Attune-based method were positively correlated with the cell counts from the LSR II-based assay (p < 0.0001; r = 0.9801) (Fig. 2-C) and CTP counts by CFU-F assay (p < 0.0001; r = 0.7237) (Fig. 2-D).

When the data were analyzed in relation to the age and sex of the donors, a clear pattern of a negative correlation

between the CD45^{low}CD271^{high} cell counts and age was observed for the samples from female donors using the Attunebased assay (p = 0.0015; r = -0.6900) (Fig. 3-A, left panel) but not for the samples from male donors (p = 0.3877; r = -0.2102) (Fig. 3-B, left panel). This pattern was also consistently detected using the LSR II-based method and CFU-F assay (LSR II-based method: female, p = 0.0070, and r = -0.6563; male, p = 0.3708, and r = -0.2577 [Figs. 3-A and 3-B, middle panels]; and CFU-F assay: female, p = 0.0055, r = -0.6904; male, p = 0.1461, r = -0.4093 [Figs. 3-A and 3-B, right panels]). Overall, CD45^{low}CD271^{high} cell counts were comparable between the Attune and LSR II flowcytometry methods and positively correlated with the CTP counts.



Fig. 4

Assessment of CD45^{low}CD271^{high} cells in bone marrow concentrates. **Fig. 4-A** The samples of bone marrow concentrates were either undiluted or diluted $\times 5$ or $\times 10$ before the counting of CD45^{low}CD271^{high} cells (n = 8 samples; S = sample). **Fig. 4-B** The CD45^{low}CD271^{high} cell counts were compared between pre-concentration (conc) and post-concentration samples (Wilcoxon matched-pairs signed-rank test; n = 15 samples). **Fig. 4-C** The fold increase of CD45^{low}CD271^{high} cell counts was compared with the fold increase in CTPs after bone marrow concentration (Student paired t test; n = 13 samples). **Fig. 4-D** The mean fold increase (and 95% CI) of CD45^{low}CD271^{high} cells (calculated using the Attune-based method) and platelets (calculated using the Sysmex cell counter) were compared after bone marrow concentration (unpaired t test; n = 15 samples for CD45^{low}CD271^{high} cells and n = 10 samples for platelets).

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Assessment of CD45^{low}CD271^{high} cells attached to the Bio-Gide scaffold. **Fig. 5-A** The number of CD45^{low}CD271^{high} cells attached to the Bio-Gide scaffold was calculated by counting the cells in samples from pre-loading and post-loading fractions of the bone marrow aspirates. **Fig. 5-B** The number of CD45^{low}CD271^{high} cells on pre-loading and attached to the Bio-Gide scaffold is shown (n = 6 samples; S = sample). **Fig. 5-C** The correlation between the number of CD45^{low}CD271^{high} cells attached to Bio-Gide scaffold and the progenitor cells (CD45⁻CD90⁺CD73⁺) surviving on the Bio-Gide scaffold after 2-week culture was analyzed (Pearson r test; n = 6 samples).

Assessment of CD45^{low}CD271^{high} Cells in Bone Marrow Concentrates

Our optimization results showed that the quantified CD45^{low} CD271^{high} cell counts in bone marrow concentrates were generally higher after ×10 dilution compared with ×5 dilution and nondilution (Fig. 4-A), and thus, ×10 dilution of bone marrow concentrates is needed to ensure accurate estimation. The CD45^{low}CD271^{high} cell counts increased significantly after bone marrow concentration (p < 0.0001) (Fig. 4-B). The fold increase of the CD45^{low}CD271^{high} cell counts (mean, 5-fold; 95% CI, 3.6 to 7.2-fold) and those of CTPs (mean, 4.6-fold; 95% CI, 3.1 to 6-fold) were comparable (p = 0.1894) (Fig. 4-C). The Sysmex results showed an increase in the platelet counts in bone marrow concentrates (p = 0.6255), with a mean increase of 4.5-fold (95% CI, 3 to 6-fold) (Fig. 4-D). In summary, we demonstrated a method of quickly assessing increased CD45^{low}CD271^{high} cell counts in the bone marrow concentrates.

Assessment of CD45^{low}CD271^{high} Cells Attached to a Collagen Scaffold

We used bone marrow aspirates to load the Bio-Gide scaffold. The number of attached CD45^{low}CD271^{high} cells was then calculated by counting these cells in the pre-loading and postloading samples (Fig. 5-A). The number of CD45^{low}CD271^{high} cells attached to the Bio-Gide scaffold varied between samples but was consistently dependent on the pre-loading cell quantities (Fig. 5-B). Furthermore, the CD45^{low}CD271^{high} cell count strongly correlated with the number of progenitor cells that survived on the Bio-Gide scaffold (p = 0.0348; r = 0.8434) (Fig. 5-C). The CD45^{low}CD271^{high} cell assessment helped to detect the donor-related differences in cell attachment onto the scaffold.

Discussion

) one marrow samples contain CTPs that are potentially Buseful in treating degenerative musculoskeletal diseases and the nonunion of bone fractures. Processing bone marrow samples helps to concentrate these CTPs. However, the concentration and prevalence of CTPs vary widely between individuals and according to different aspiration locations and techniques¹⁶⁻¹⁸. The gold standard CFU-F assay requires at least 6 days¹⁸, and thus, clinicians currently have no way of knowing, at the time of the procedure, the quality of the bone marrow sample utilized. It would be desirable, therefore, to have a rapid measurement method that could provide insight into bone marrow quality on the day of the procedure. Here, we introduce a fast and automated quantification of CD45lowCD271high cell counts in bone marrow preparations that may be used to judge the quality of bone marrow samples. This assay was compared with another more time-consuming flow-cytometry assay¹⁷ and provided a similar range of CD45^{low}CD271^{high} cells. Both assays confirmed an age-related decline in CD45lowCD271high cells in samples among females but not among males, as was also previously reported for CTPs18.

The specificity of the CD45^{low}CD271^{high} cell measurement compared with CTP counts was low (0.05 on average); we noted 20 times more CD45lowCD271high cells with the Attunebased method than CTPs quantified by CFU-F assay. This finding agrees with those of previous studies^{17,26}. However, the CD45lowCD271high counts were positively and modestly correlated with the prevalence of CTPs (r = 0.7237). This low specificity does not prevent the use of this assay for the estimating of aspirate quality; however, it is clear that it does not enable exact measurement of CTP counts. This might be related to the senescence of some CD45lowCD271high cells in culture during CFU-F assay as a result of plating at very low clonal densities. Another possible explanation for this disparity is that CTPs represent only a subset of the CD45^{low}CD271^{high} population, as suggested recently³³. It is possible that, with the addition of more markers, this subpopulation could be defined, allowing increased specificity of the assay. One previous study tested the CD146 marker, but no further enrichment in CTP counts was detected in the CD146+CD271+ fraction compared with CD146⁻CD271⁺ fraction²⁶. Subsequently, the same group showed that the majority of CTPs resided in the CD140a⁻CD271⁺ fraction²⁸. However, our group did not find such a clear subpopulation³⁴. Others have not yet devised additional, more selective markers, while there is a mutual agreement regarding the value of CD271³⁵.

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We believe that the assessment of CD45^{low}CD271^{high} cells has a very high sensitivity (close to 100%), as other studies have shown that all bone marrow colony-forming activity is confined to CD45^{low}CD271^{high} cells, and CD271-negative cells have not been shown to have any colony-forming ability²⁴⁻³⁰. The implications of the Attune-based assay with high sensitivity and relatively low specificity is that no CTPs are missed, while some progenitor cells with potentially lower colony-forming capacity than detected in our experimental conditions can be counted.

The CFU-F assay data can vary depending on patient age and bone marrow aspiration site and volume^{17,21}. This could explain why the CTP counts in this study showed some variability from previous work³⁶. Using various bone marrow processing methods could have an additional effect on the variability of CTP counts. For example, using density gradient centrifugation during bone marrow processing causes CTP loss³⁷. We ensured optimal and consistent culture conditions by using complete and batchtested media for the CFU-F assays. Thus, the possibility of the underestimation of CTPs is small, but still exists.

Bone marrow aspirates or concentrates loaded onto scaffolds have been reported to enhance cartilage repair in knee or hip osteoarthritis, focal condylar lesions of knee articular cartilage, and talar osteochondral injuries, with promising outcomes^{9,38-41}. The results presented here show that the CD45lowCD271high cell counts can be increased 5-fold after bone marrow concentration. We also demonstrated that platelets were concentrated 4.5-fold, showing an additional value of unfractionated bone marrow concentrates with respect to providing growth factors⁴². Compared with our data, Dawson et al. demonstrated a 4-fold increase of CTPs in bone marrow concentrates⁴³. Another recent study showed that 2 concentrator devices produced substantially different counts of CTPs and dissimilar levels of growth factors⁴⁴. Our data also showed that the number of attached CD45lowCD271high cells on the Bio-Gide collagen scaffold varied depending on the initial cell count in the bone marrow samples. Collectively, this further emphasizes the potential value of CD45^{low}CD271^{high} cell count assessment to indicate the quality of bone marrow samples after concentration or when loaded onto scaffolds.

In conclusion, we report a method that can help to indicate the "potency" or "quality" of the bone marrow sample applied in clinical settings. The quantitative assessment of CD45^{low}CD271^{high} cells in bone marrow aspirates was performed rapidly, and the CD45^{low}CD271^{high} cell counts positively correlated with CTP counts. While the specificity of CD45^{low}CD271^{high} cell assessment was low compared with CFU-F assay, the sensitivity of this method was very high. Since CFU-F data cannot be immediately available on the day of surgery, our findings support the view that an assay measuring CD45^{low}CD271^{high} cell counts may instead serve as a useful surrogate measure of bone marrow quality. Additional studies on the rapid measurement of CTP prevalence in bone marrow samples with the inclusion of other specific markers are desirable to further enhance the method described here.

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