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Short communication

A Novel Automated Bioreactor for Scalable Process Optimisation of Haematopoietic Stem Cell Culture

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Abstract

Proliferation and differentiation of haematopoietic stem cells (HSCs) from umbilical cord blood at large scale will potentially underpin production of a number of therapeutic cellular products in development, including erythrocytes and platelets. However, to achieve production processes that are scalable and optimised for cost and quality, scaled down development platforms that can define process parameter tolerances and consequent manufacturing controls are essential.

We have demonstrated the potential of a new, automated, 24×15mL replicate suspension bioreactor system, with online monitoring and control, to develop an HSC proliferation and differentiation process for erythroid committed cells (CD71⁺, CD235a⁺). Cell proliferation was relatively robust to cell density and oxygen levels and reached up to 6 population doublings over 10 days. The maximum suspension culture density for a 48 hour total media exchange protocol was established to be in the order of 10⁷ cells/mL. This system will be valuable for the further HSC suspension culture cost reduction and optimisation necessary before the application of conventional stirred tank technology to scaled manufacture of HSC derived products.

Keywords: Haematopoietic stem cells, Suspension culture, Bioreactor, Cell culture automation, Process development.

Abbreviations: HSCs, haematopoietic stem cells; ambrTM, advanced microscale bioreactor; rpm, revolutions per minute; μ , mean; SEM, standard error of the mean; FITC, Fluorescein isothiocyanate; dO₂, dissolved oxygen; ANOVA, analysis of variance; SCF, stem cell factor; IGF-1, insulin-like growth factor 1; IL-3, interleukin 3; EPO, erythropoietin.

Introduction

Haematopoietic stem cells (HSCs) are multipotent cells with the ability to self-renew and differentiate into all blood cell lineages (Nielsen, 1999; Rizo *et al.*, 2006). These properties confer a wide range of possible therapeutic applications for which umbilical cord blood constitutes a readily available cell source (Neildez-Nguyen *et al.*, 2002). The development of such HSC-based products, i.e. red blood cells, has made significant progress in recent years (Douay and Andreu, 2007; Fujimi *et al.*, 2008; Giarratana *et al.*, 2005; Miharada *et al.*, 2006). However, the number of cells required for a therapeutic dose will be up to the order of 10^{12} cells per patient, necessitating a cell expansion process that sets a formidable production scale challenge for conventional bioprocessing formats. Intensification of commonly used static expansion systems (e.g. T-flasks, gas-permeable bags) is inherently limited by nutrient and gas concentration gradients, and a lack of on-line monitoring and control of key environmental conditions, limiting maximum supportable cell density (Liu *et al.*, 2006). Understanding the feasibility of a scale-up approach in conventional stirred tank systems requires the use of low volume systems for economic process design, development and optimisation, whilst retaining relevance to downstream manufacturing bioprocesses (Kirouac and Zandstra, 2008; Ratcliffe *et al.*, 2011).

Cell culture of HSCs to erythroid cells is currently costly, and scaled down processes are required for optimisation and cost reduction prior to scale up. Here we show the proliferation and erythroid commitment of HSCs at the lowest (and therefore most cost effective) suspension scale yet reported using an automated process development platform (the advanced microscale bioreactor workstation (ambrTM), TAP Biosystems, Royston, UK) (Fig. 1). The ability of this system to predict operating conditions at scale was recently demonstrated in terms of both cell productivity/growth and the physicochemical environment (Hsu *et al.*, 2012). We show that ambrTM culture offers the control required to detect different cell responses to operating parameters, such as O₂ tension and seeding density, and can support high density cell culture of HSCs. It will therefore form a valuable tool for optimisation and cost reduction of a complex and costly culture system prior to scale up.

Brief Methods

Frozen CD34⁺ enriched umbilical cord blood cells (negative selection using RoboSep[®] automated cell separator, Stem Cell Technologies, Grenoble, France) were provided courtesy of Celgene Cellular Therapeutics (New Jersey, USA) after proliferation for 7 days in 2D culture conditions using StemSpan SFEM (serum-free expansion medium) (09650, StemCell Technologies) supplemented with the following cytokines and reagents for erythroid cell proliferation and differentiation; 100ng/mL SCF, 40ng/mL IGF-1, 5ng/mL IL-3, 3IU/mL EPO (R&D Systems, Oxford, UK), 40μg/mL Lipid and 1μM Dexamethasone (Sigma-Aldrich, Poole, UK). Prior to cell defrost, ambrTM vessels were loaded with 14mL of this medium and stabilised at 37°C, experimental dO₂ level, and pH 7.4. Automated antifoam additions (20μL of 1% solution, Sigma-Aldrich) were made every 48h, or as required. Automated sodium bicarbonate additions (20μL of 1M solution, Sigma-Aldrich) maintained cultures at pH7.4 (trigger pH7.2, monitored every 2h). Cultures were initiated at several cell

seeding densities as stated in the results tables (viability $96.27\% \pm 1.12\%$ ($\mu \pm \text{SEM}$)). Cultures were sampled for cell counting (Countess automated cell counter, Invitrogen, Paisley, UK) and metabolite analysis (Nova Bio-Profile Flex, Nova Biomedical, Runcorn, UK). Replacement of culture medium was conducted through centrifugation and re-suspension of cells in fresh medium at intervals stated in results. Flow cytometry for erythroid lineage markers was performed after each expansion using CD235a-FITC or CD71-FITC (BD Biosciences, Oxford, UK) conjugated antibodies and appropriate isotype controls according to manufacturer's instructions. A Cell Lab Quanta SC Flow Cytometer (Beckman Coulter, High Wycombe, UK) was used for analysis. Experimental runs were performed in triplicate, results expressed as the mean value of the data \pm standard error of the mean ($\mu \pm \text{SEM}$). Significance was determined using ANOVA (Minitab).

Results and Discussion

In order to determine feasibility of HSC proliferation in ambrTM, to identify conditions for rapid cell proliferation, and to establish time to the primary media exchange, cells were cultured under different oxygen tensions (dO_2 5%, 15%) and cell densities ($2 \times 10^4/\text{mL}$, $8 \times 10^4/\text{mL}$). Slower proliferation observed after 5 days (5d) in all cultures indicated that the first media exchange should occur within this period. Both dO_2 and cell density had a statistically significant effect on proliferation at 5d ($P \leq 0.002$), and there was a statistically significant interaction between the two parameters ($P < 0.001$). Highest proliferation was achieved with 8×10^4 cells/mL seeding density and 15% dO_2 (Exp 1, Table 1). A further culture was conducted with total media exchange after 4 days, and every 2 days thereafter, to further optimise around these identified operating parameters (Seeding density 4×10^4 , 8×10^4 , $1.2 \times 10^5/\text{mL}$; dO_2 7.5%, 15%, 21%). Lower seeding density increased proliferation after 8 days culture ($P < 0.01$) whilst O_2 in this range was no longer significant (Exp 2, Table 1). The best growth conditions were continued to 10d (shown in **bold**) and achieved over 6 population doublings (all growth curves shown in Figure 2). In order to establish the potential of the bioreactor to develop reduced cost processing strategies, complete medium replacement, 50% medium dilution, and cytokine addition without medium dilution were trialled. No significant differences in proliferation were observed after 6 days (3.16, 3.27, 2.86 PDs respectively), but a small significant proliferative advantage was observed after 8 days (4.4, 3.9, 3.7 PDs respectively). All data from experiment 2 was used to calculate coefficient of variation of 12.5% for replicate vessels indicating system robustness across the experimental range. At the end of all experiments the cells strongly expressed markers CD71 (79-84%) and CD235a (51-64%) characteristic of erythroid lineage development and had viability of 90-97%, with no statistically significant differences between conditions. Cells processed in ambrTM were also capable of producing BFU-E colonies indicative of erythroid commitment (inset Figure 2).

In order to establish the maximum supported cell density for a 48h media exchange suspension process, and therefore the most efficient production density range, HSCs were cultured without cell dilution. Over four successive cultures cells achieved densities of 0.98 - 1.37×10^7 cells/mL. This was not dependent on the initial cell density (4×10^4 - $1.2 \times 10^6/\text{mL}$) indicating a density limit rather than an intrinsic cell proliferative limit. In order to determine

if cytokine concentrations or reactor stir rates were limiting cell density these were both increased throughout culture (up to 3-fold and 1000rpm respectively). This had no significant effect on the terminal density and did not adversely affect viability. Nutrient consumption, metabolite production, pH, and dO₂, were monitored during the culture to identify candidates for cell density limitations. pH and dO₂ set levels did not deviate, indicating these factors are unlikely to limit cell density or growth at currently observed levels. As shown inset in Figure 3, the dominant ammonia source was not related to cell culture density; pre-passage cell concentration increased by more than an order of magnitude, but the ammonia concentration range remained comparable and is therefore also unlikely to be the limiting factor for cell density at the levels observed.

Glucose consumption per cell was calculated to identify required feed rates. Average early consumption rate (0.0166ng/cell/h, range 0.007-0.025ng/cell/h, calculated from first 4 hours of culture at day 6 in all experimental conditions) indicated complete depletion would occur in approximately 24 hours at a density at the order of 10⁷ cells/mL assuming constant metabolic activity per cell. However, glucose (and glutamine) concentrations in high density culture did not reduce beneath 3.6g/L (and 2.5mmol) from fresh media levels of 4.5g/L (and 4mmol). This indicates a substantial change in glucose consumption through the culture period and a requirement for more extensive metabolic studies. Lactate concentration in culture was usually beneath the limits of detection. However, it was measured at 0.1 to 1g/L in some of the higher density cultures (all >1×10⁶ cells/mL) but showed no relationship to cell density beyond this. This lack of relationship indicates that lactate production per cell is highly variable at different points in culture. Lactate has been reported inhibitory to cell growth at the concentrations observed so it could be a factor in the reduced proliferation at higher densities or at the density limit (Hassell *et al.*, 1991).

Conclusion

HSCs are good candidates for suspension culture production systems as many of the relevant cell types are naturally non-adherent. This should facilitate their clinical production due to the easy availability of large biopharmaceutical suspension bioreactors. The data reported here shows that HSCs can be cultured in the ambr™ platform at 10mL suspension scale, the lowest yet reported, and this can generate useful data for analysing parameters influencing cell growth, critical for necessary cost reduction studies. Metabolic data indicates that achieving densities in excess of those reported will be likely to require a perfused approach to cell produced factor removal and nutrient supply. This scaled down system of stirred tank operation can successfully support HSC proliferation and differentiation and therefore provide a valuable tool for further exploring the limits of HSC bioprocessing.

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Table 1 The average population doublings achieved under several seeding density and oxygen tension culture conditions in ambrTM ($\mu \pm \text{SEM}$ of triplicate cultures) identify the levels at which these parameters support rapid cell growth.

Experiment/Culture duration	Cell Density / mL	Population Doublings @ dO ₂			
		dO ₂ 5%	dO ₂ 7.5%	dO ₂ 15%	dO ₂ 21%
Exp 1: 5d	2×10 ⁴	2.29 ± 0.10		2.09 ± 0.21	
	8×10 ⁴	2.24 ± 0.10		3.76 ± 0.16	
Exp 2: 8d	4×10 ⁴		4.98 ± 0.07	4.97 ± 0.54	4.43 ± 0.40
	8×10 ⁴			4.36 ± 0.02	4.10 ± 0.16
	1.2×10 ⁵		4.11 ± 0.02	3.91 ± 0.07	4.30 ± 0.22
Exp 2: 10d	4×10 ⁴		6.31 ± 0.09	6.05 ± 0.23	

Fig. 1. The ambr™ automated suspension culture process development workstation: (1) Liquid handler, (2) Culture vessel control stations, (3) Pipette racks, (4) Reagent and sample plates. Culture vessel inset: (5) Vessel cap, (6) Gas supply filter, (7) Gas sparge tube, (8) Impeller, (9) pH sensor, (10) dO₂ sensor. The workstation comprises 24 independent suspension culture bioreactors and offers precise control of the culture environment through online monitoring and automated control of temperature, pH, gassing, stirring, cell density and liquid handling.

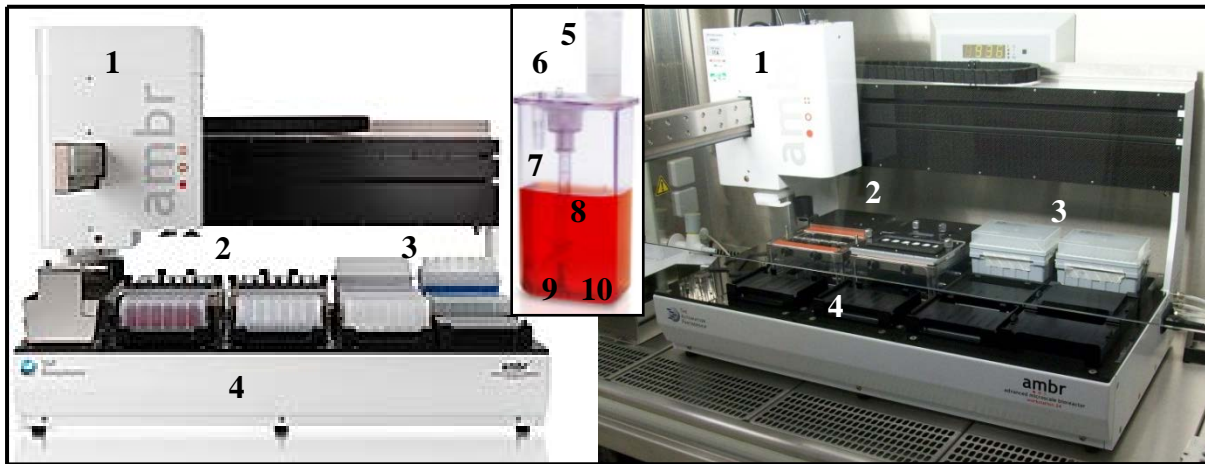


Figure 2 Growth curves for variable cell seeding density (SD) and oxygen tension (O₂) culture in ambr™. Cells were cultured under the conditions described in Table 1 (Exp 2, 8 days); Density: Low = 4×10⁴/mL, Mid = 8×10⁴/mL, High = 1.2×10⁵/mL; dO₂: Low = 7.5%, Mid =15%, High =21%. Inset graphs; cells express CD235a and CD71 after proliferation, indicators of erythroid lineage development, as shown by these typical flow cytometry profiles. Cells from the bioreactor also produced typical BFU-E colonies (inset) confirming erythroid lineage.

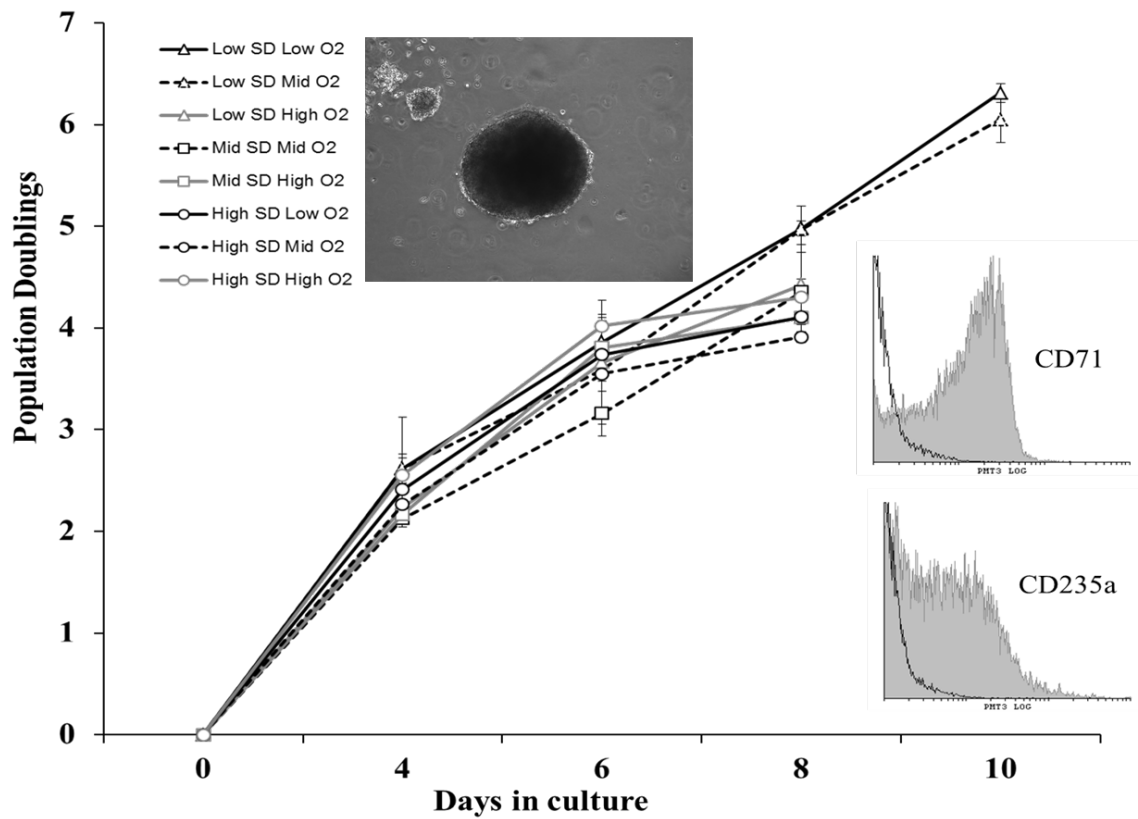


Figure 3 Growth curves for HSC culture to high density in ambr™ without cell dilution. Cells were cultured under various conditions to high density. Start and End cell density of each culture are annotated. All conditions progressively increase to 3 fold normal cytokine supplementation and 1000 rpm stir rates. Higher seeding densities reduce growth rates but do not alter terminal density. The inset graph shows a lack of strong correlation between cell density and ammonia concentration (in this range) measured at the end of each culture period.

