## Scale-up and comparability study of T-cell depletion in single inlet – single outlet QMS

Thomas Schneider<sup>1</sup>\*, Lee R. Moore<sup>1</sup>, Jeffrey J. Chalmers<sup>2</sup>, Maciej Zborowski<sup>1</sup>

<sup>1</sup>Department of Biomedical Engineering, Cleveland Clinic Lerner Research Institute, 9500 Euclid Ave., Cleveland, Ohio 44195, USA

<sup>2</sup>Department of Chemical and Biomolecular Engineering, The Ohio State University, Columbus, Ohio, USA

\*corresponding author: Thomas Schneider (email: Thomas.Schneider.81@gmx.net, Tel: 1-216-445-9342)

Pre-clinical scale T-cell depletion  $(2 \times 10^8 \text{ processed cells})$  using a single inlet – single outlet, pump controlled quadrupole magnetic flow cell sorter (QMS) was under investigation to show reproducibility of published results in a process scale-up. Buffy coats from haemochromatosis patients were spiked with cultured Kg-1a cells to mimic peripheral progenitor cells. The target T-cell fraction was immunomagnetically labeled in a one or two step labeling protocol without washing and depleted by negative selection at sorting speeds of up to  $3.3 \times 10^6$  cells per second. A mean log<sub>10</sub> T-cell depletion of 2.9 (range 2.13 to 3.74, n=4) for a one-step labeling strategy and a mean log<sub>10</sub> T-cell depletion of 3.1 (range 2.58 to 3.59, n=2) for a two-step labeling strategy showed a reasonable agreement with published data describing 1/10 as many cells processed compared to this study. The depletion experiments show high total cell recovery (92.7 ± 7.7%) and high viability (95.0 ± 3.8%) and recovery of the spiked cell fraction (81.7 ± 14.7%). The study also revealed challenges and limitations for a scale-up into clinical scale T-cell depletions (> 1 × 10<sup>9</sup> - 1 × 10<sup>10</sup>). The current goal is a 4 log<sub>10</sub> T-cell depletion. The composition of different cell fractions for an ideal depletion is shown in the Figure.



**Figure.** Idealized 4  $log_{10}$  T-cell depletion by QMS.