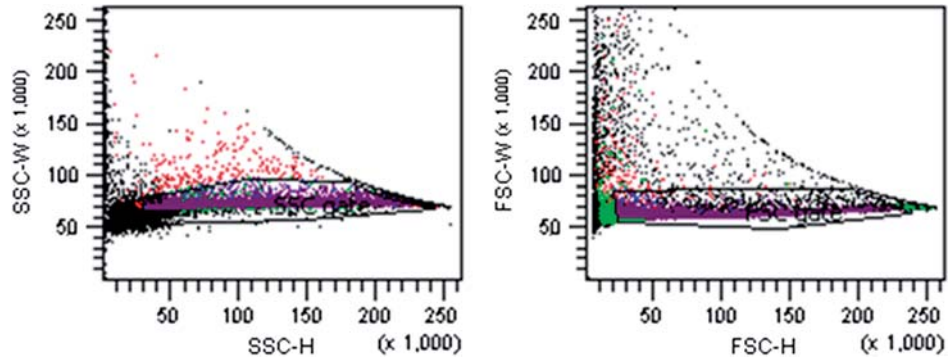
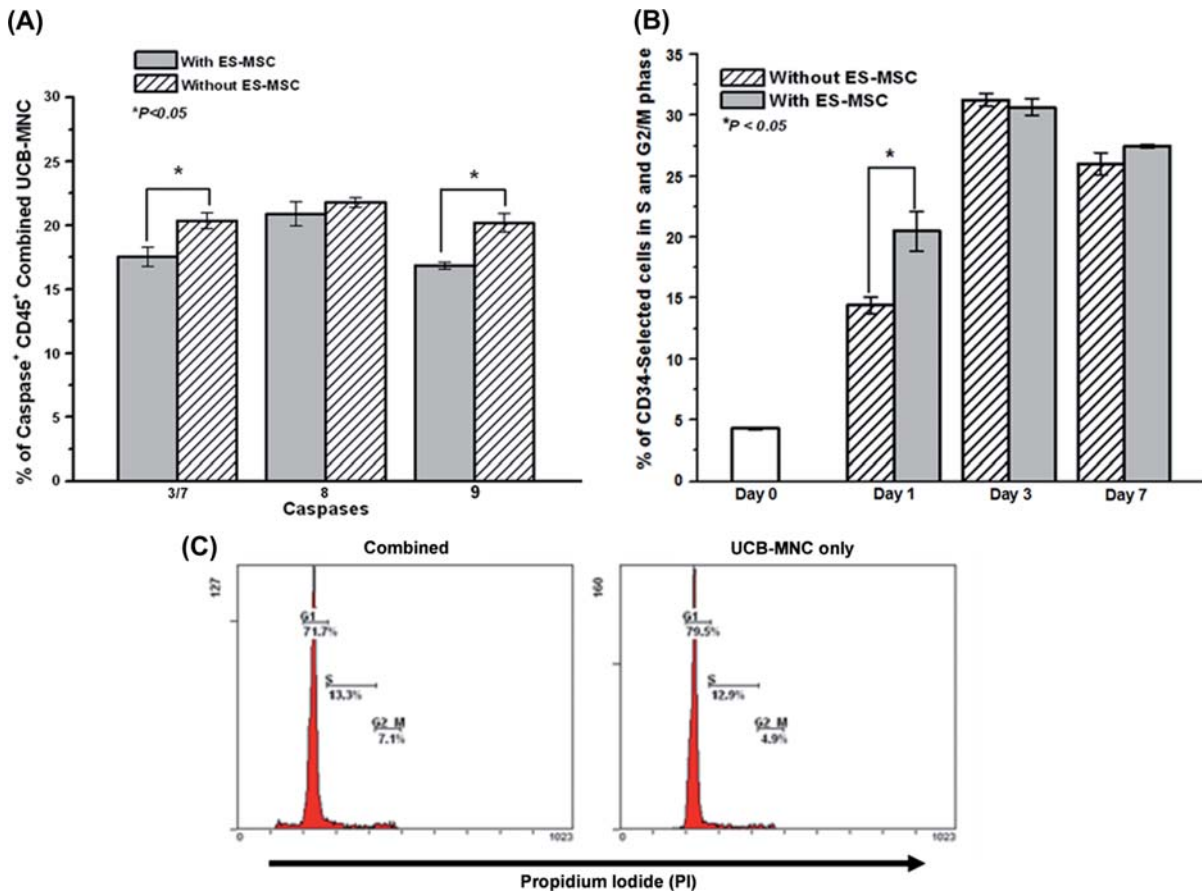


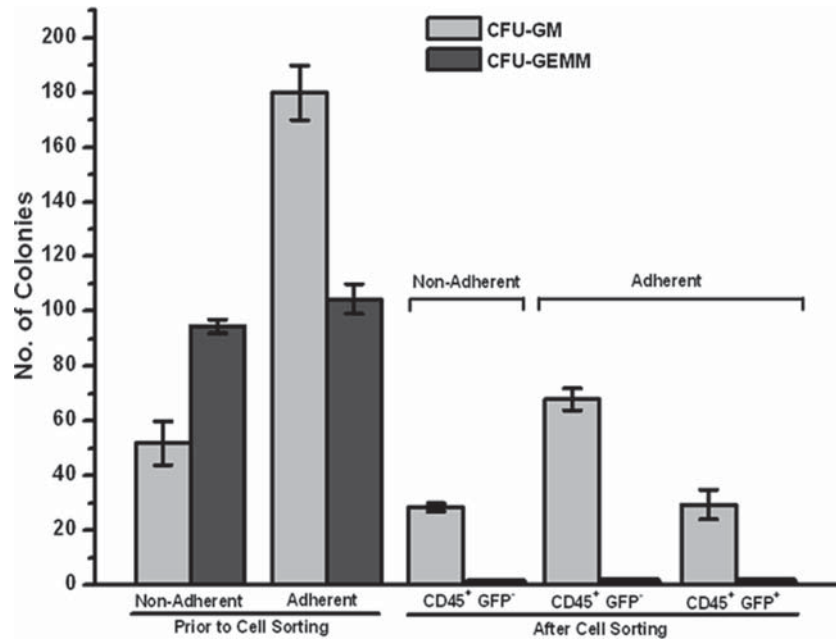
Supplementary material for Chu et al., Intercellular cytosolic transfer correlates with mesenchymal stromal cell rescue of umbilical cord blood cell viability during *ex vivo* expansion. *Cytherapy*, 2012;14:1064–1079.



Supplementary Figure 1. The height and width of the forward (FSC) and side (SSC) scatter of the CD45<sup>+</sup> GFP<sup>+</sup> cells. The parameters are used to demonstrate no cell doublets arising from UCB – MSC fusion. Data represent mean  $\pm$  S.E.M. from three independent experiments.



Supplementary Figure 2. Investigation of caspase activation and cell cycle analysis of the ES-MSC co-cultured UCB-MNC. (A) Activation of caspase 3/7, 8 and 9 (FLICA reagent) in combined CD45<sup>+</sup> UCB-MNC on Day 3 (\*represents  $p < 0.05$ ). (B) Cell cycle analysis of CD34-selected cell population in the co-culture system. The DNA contents of the cell samples were stained with propidium iodide (PI) and analyzed by flow cytometry (\*represents  $p < 0.05$ ). (C) Representative flow cytometer plots showing the cell cycle analysis of the ES-MSC co-cultured (combined fraction) and non-co-cultured CD34 selected UCB-MNC. Data represent mean  $\pm$  S.E.M. from three independent experiments.



Supplementary Figure 3. CFU-GM and CFU-GEMM assay of the adherent CD45<sup>+</sup> GFP<sup>+</sup> UCB-MNC. Fluorescence activated cell sorting (FACS) had a detrimental effect on the primitive colony forming cells. There is a complete depletion of the primitive CFU-GEMM colonies in the sorted group. CFU-GEMM is present in the control pre-sorted samples. Data represent mean  $\pm$  S.E.M. from three independent experiments.