Recombinant TAT-BMI-1 fusion protein induces ex vivo expansion of human umbilical cord blood-derived hematopoietic stem cells

Supplementary Materials



Supplementary Figure 1: Time course of the uptake of TAT-proteins by cord blood derived CD34⁺ cells. Purified TAT-GFP or TAT-BMI1 proteins were added to cultures of CD34⁺ cells at a final 100 nM concentration for 20, 40 and 120 min at 37°C. Cytospins were fixed and TAT-proteins were detected by indirect immunofluorescence with rabbit anti-HA antibody and anti-rabbit Alexafluor 488; cell nuclei were counter-stained with DAPI.



Supplementary Figure 2: TAT-BMI-1 pretreatment of CB-CD34⁺ cells enhances hematopoietic engraftment in NOG mice. Flow-cytometric peripheral blood analysis. Flow cytometric analysis of the peripheral blood of NOG mice transplanted with TAT-GFP or TAT-BMI-1-treated CD34⁺ cells 7 and 12 weeks after injection. Cytograms for each mouse show the percentage of total human leukocytes (CD45⁺), myeloid (CD33⁺) and B-lymphoid (CD19⁺) cells.



Supplementary Figure 3: TAT-BMI-1 pretreatment of CB-CD34⁺ cells enhances hematopoietic engraftment in NOG mice. Flow-cytometric bone marrow analysis. Flow cytometric analysis of the bone marrow of NOG mice transplanted with TAT-GFP or TAT-BMI-1-treated CD34⁺ cells 12 weeks after injection. Cytograms for each mouse show the percentage of total human leukocytes (CD45⁺), myeloid (CD11B⁺,CD33⁺) and lymphoid (CD19⁺, CD3⁺) cells.



Secondary Transplant PB in NOG mice - PB 9 wk

Supplementary Figure 4: TAT-BMI-1 pretreatment of CB-CD34⁺ cells enhances secondary hematopoietic engraftment in NOG mice. Flow cytometric analysis of peripheral blood (weeks 9 and 14 post-transplant) and bone marrow (week 16 post-transplant) of NOG mice transplanted with bone marrow from the primary recipients (shown in Supplementary Figure 3). Cytograms for each mouse show the percentage of total human leukocytes (CD45⁺).



Supplementary Figure 5: Purification and functional characterization of BMI-1 235 on CD34⁺ cells expansion. (A) Coomassie staining of NUPAGE 4–12% gels of the truncated, 1-235 AA form of TAT-BMI-1, called TAT-BMI-1 235 (middle lane) 34 kDa, compared with TAT-BMI-1 full length (right lane) 45 kDa and 2 μ g Bovine Serum Albumin (BSA) (left lane) 60 kDa. (B) Uptake and subcellular localization of TAT-BMI-1 235 in cord blood derived CD34⁺ cells. Purified TAT-BMI1 235 was added to cultures of CD34⁺ cells at a final 100 nM concentration for 60 min at 37°C. Cytospins were fixed and TAT-BMI1 235 was detected by immunofluorescence with rabbit anti-HA antibody and anti-rabbit Alexa Fluor 488 as described; cell nuclei were counter-stained with DAPI. (40× magnification) (C) CD34⁺ cells (5x10³/ml) were plated in StemMACS HSC Expansion Medium with StemMACS HSC Expansion Cocktail. During the first 3 days of culture, TAT-BMI-1 235, TAT-BMI-1 and the control TAT-GFP protein were added 4 times a day at a 10 nM concentration. At the time intervals indicated, the cells were counted and re-plated in fresh medium at equal cell densities. Cumulative cell expansion is shown. All assays were performed in triplicate. (D) Colony-forming cell (CFC) assays were performed on cells from the cytokine-driven cultures at day 10 and 16 × 10³ cells/well were plated in triplicate assays in semisolid StemMACS HSC-CFU medium complete with cytokines. Colonies were scored after 2 weeks and total numbers of progenitors was normalized to the total cell number in the original culture at the time of plating.