

Supplementary Figure 1

The bars in the following graphs represent the mean normalized expression of the genes that were tested by real time PCR analysis of the GFRA1-positive cell populations after immunomagnetic cell sorting for each age group (6, 21, 60 days and 8 months). The mean normalized expression and standard error values were multiplied by 1000 and presented on the Y-axis of the graphs in logarithmic scale. Standard errors represent variation between three different RNA isolations. Expression levels of all genes were normalized to beta actin.

















