Supplements

Supp. 1

Suggested scheme of hematopoiesis both *in vivo* (a) [1, 2,3] and in *in vitro* model of ES cells differentiation (b) [4, 5] in mice. The scheme of potential role of p38 kinase in hematopoiesis based on previous published results and observed in our work (c). VEGF induces expression of Etv2 in p38 kinase-dependent manner, what leads to hemangioblast formation and development. EPO/EpoR induced activation of p38 kinase and phosphorylation (green arrow) of its targets GATA1 and 2, which transcriptional activity is required for erythropoiesis. Hematopoiesis in our experiments with ES cells. The capability forming hematopoietic colony forming unit (CFU) progenitors (d), CFU-G, -GM, -M, -E, and -GEMM in differentiating wt ES cells (e). The expression of hematopoiesis specific transcripts in our model of in vitro hematopoiesis in wt ES cells (f). For details, see Materials and Methods, and Results in our manuscript.

Supp. 2

Mutated p38 α -/- cells did not express p38 α protein, in contrast to their wt counterparts. The expression of key pluripotent protein Oct4 and general abundant GAPDH was equal in both cell lines (a). When cells were differentiated by means of EB techniques, the overall level of p38 α kinase RNA did not change (b). The level of p38 α kinase protein as well as its phosphorylated form were also unchanged. This was in contrast to the level of Oct4 protein, a marker of undifferentiated pluripotent cells, that decreased continuously with differentiation time. The protein level of GAPDH, which was used as a reference gene, is also shown (c). The phosphorylated form of p38 α kinase when wt cells were treated by inhibitors of p38 kinase pathway SB203580 or SB202190 (5 μ M) for 1 hour and subsequently by 200 μ M H₂O₂ for 1 hour (d). Data are presented as representative western blot from two experiments, RNA level as mean + SEM from four independent experiments.

Supp. 3

The effect of p38 kinase pathway inhibitor (SB203580 or SB202190) on the formation of CFU and selected RNA transcripts through EB-differentiating ES cells. The number of CFU and the level of selected RNA transcripts in mutant p38 α -/- cells are also shown. For details, see Materials and Methods. The concentration of inhibitors was 5 μ M. Cells were exposed to inhibitors for the full duration of the differentiation of EBs. Inhibitors were always added to the medium when old medium was replaced by new medium, which was every two days of the culture. The number of all hematopoietical CFU and particular CFU in 6-, 10-, and 14-day-old EBs (a). The expression of RNA transcripts regulating and marking hematopoiesis (b, c) and the expression of RNA transcripts associated with erythropoiesis (d). Data are presented as mean + SEM from a minimum of three independent experiments.

Supp. 4

The table summarizes data both the effect of $p38\alpha$ -/- depletion and the effect of p38 kinase signaling inhibitors on the expression of selected RNA transcripts that are associated with hematopoiesis and that are presented on Fig. 3 (For details, see Material and Methods). It shows that the up-regulation (\uparrow), down-regulation (\downarrow) and no-change (-) of the mentioned RNA transcript expressions compare to the levels of the same RNA transcripts in wt cells. The cKit and Sca1 (both marked in bold in the table) RNA transcripts levels only have an opposite trend in expression in $p38\alpha$ -/- cells compared to wt cells treated by p38 kinase inhibitors.

	6ds old Ebs			10ds old Ebs			14ds old Ebs		
Marker	p38α-/-	SB203580	SB202190	p38α-/-	SB203580	SB202190	p38α-/-	SB203580	SB202190
CD34	\downarrow	-	-	\downarrow	\checkmark	\checkmark	\downarrow	\downarrow	\checkmark
CD38	-	\downarrow	-	\downarrow	\checkmark	\checkmark	\downarrow	\downarrow	\checkmark
cKit	\uparrow	\downarrow	\downarrow	\uparrow	\checkmark	\checkmark	\uparrow	\downarrow	\checkmark
Flk1	\downarrow	\rightarrow	\rightarrow	\rightarrow	\checkmark	\checkmark	\checkmark	-	\checkmark
Hbb-b1	\downarrow	\rightarrow	\rightarrow	\rightarrow	\checkmark	\checkmark	\downarrow	\rightarrow	\checkmark
Hbbζ	\downarrow	\downarrow	\downarrow	\downarrow	\checkmark	\checkmark	\checkmark	\downarrow	\checkmark
Hbbγ	\downarrow	\downarrow	\downarrow	\downarrow	\downarrow	\downarrow	\downarrow	\downarrow	\downarrow

HoxB4	-	-	-	-	-	-	-	-	-
GATA1	\rightarrow	\rightarrow	\rightarrow	\rightarrow	\rightarrow	\downarrow	\rightarrow	\downarrow	\rightarrow
GATA2	-	-	-	-	-	-	\rightarrow	\downarrow	\rightarrow
Klf1	-	-	-	\rightarrow	\rightarrow	\downarrow	\rightarrow	\downarrow	\rightarrow
PU.1	-	-	-	-	\downarrow	-	\downarrow	\downarrow	\downarrow
Runx1	-	-	-	-	-	-	\downarrow	\downarrow	\downarrow
Sca1	\uparrow	\rightarrow	\downarrow	\uparrow	\rightarrow	\downarrow	-	-	-
VEGF	-	-	-	-	-	-	-	\downarrow	\downarrow

References

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In vivo

In vitro





(c)

(a)

HPC/HSC

Relative number of hematopoietic colonies



Relative number of colonies in full medium





(f)

Supplement 2.







Relative number of colonies in full medium (6ds old EBs)



Type of CFU

Relative number of colonies in full medium (10ds old EBs)



Relative number of colonies in full medium (14ds old EBs)





(a)







p38α+/+ ctr

p38α-/-

p38α+/+ SB203580 **μ** p38α+/+ SB202190

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(c)



