

Thioredoxin-interacting (TXNIP) protein regulates the differentiation of erythroid precursors



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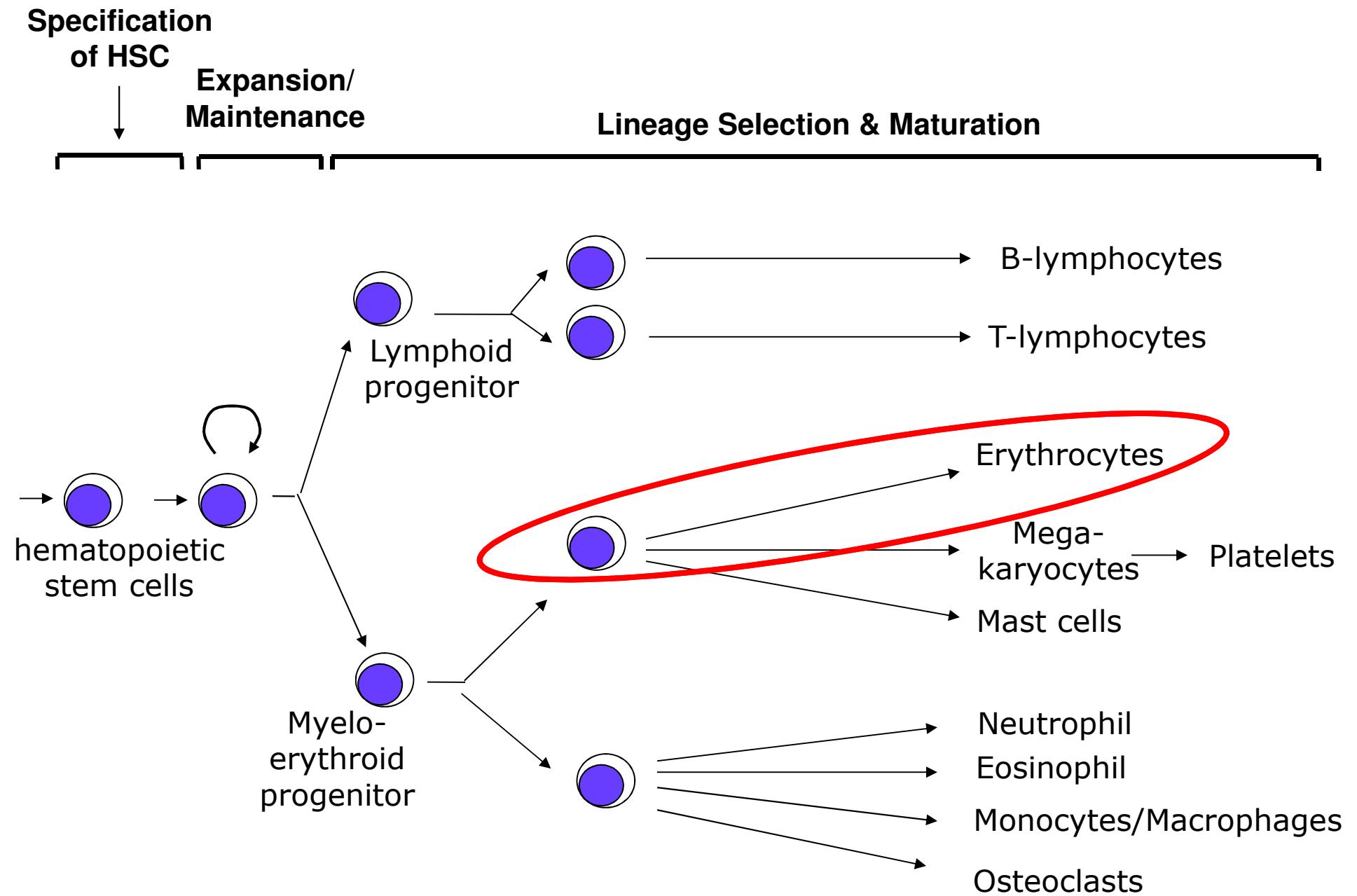
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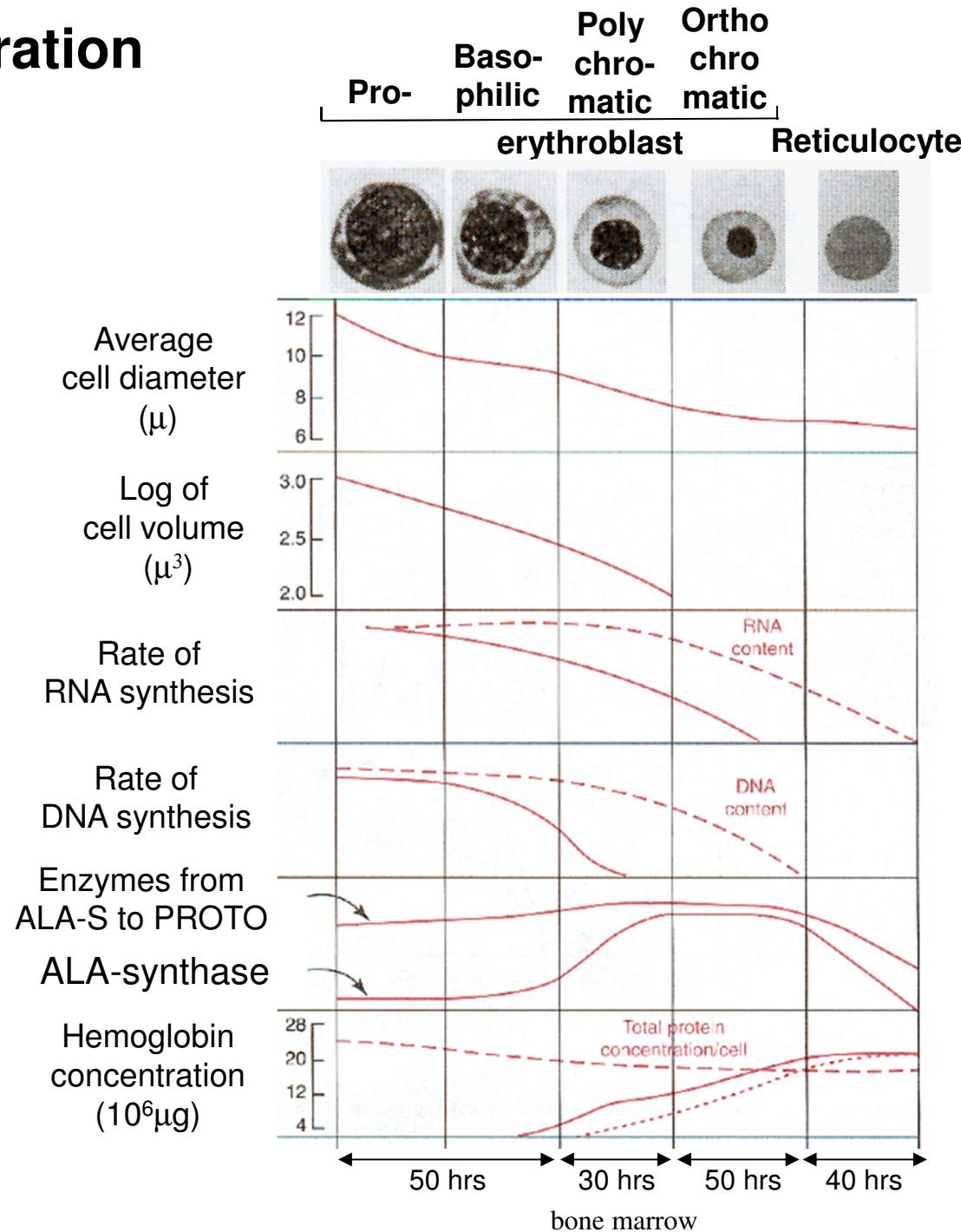


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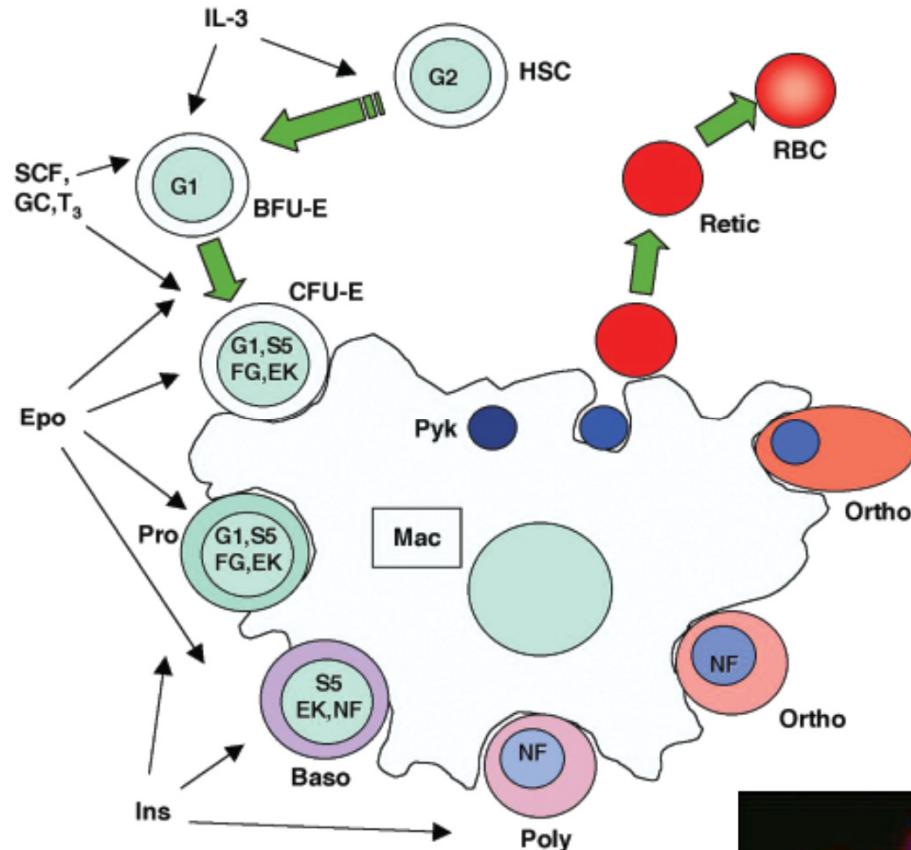
Hematopoiesis



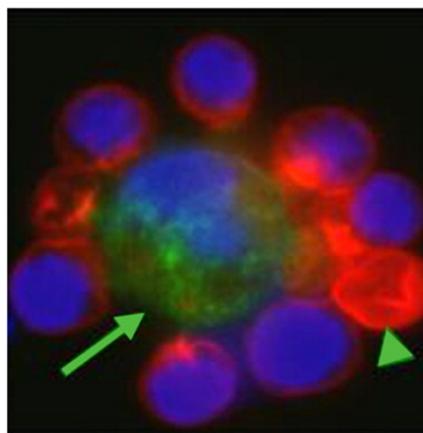
Erythroid maturation - bone marrow



Regulation of erythroid development



Ingle et al. 2004 IUBMB Life 56(4):177-184



Lee et al. 2006 Blood 108: 2064-2071

External factors

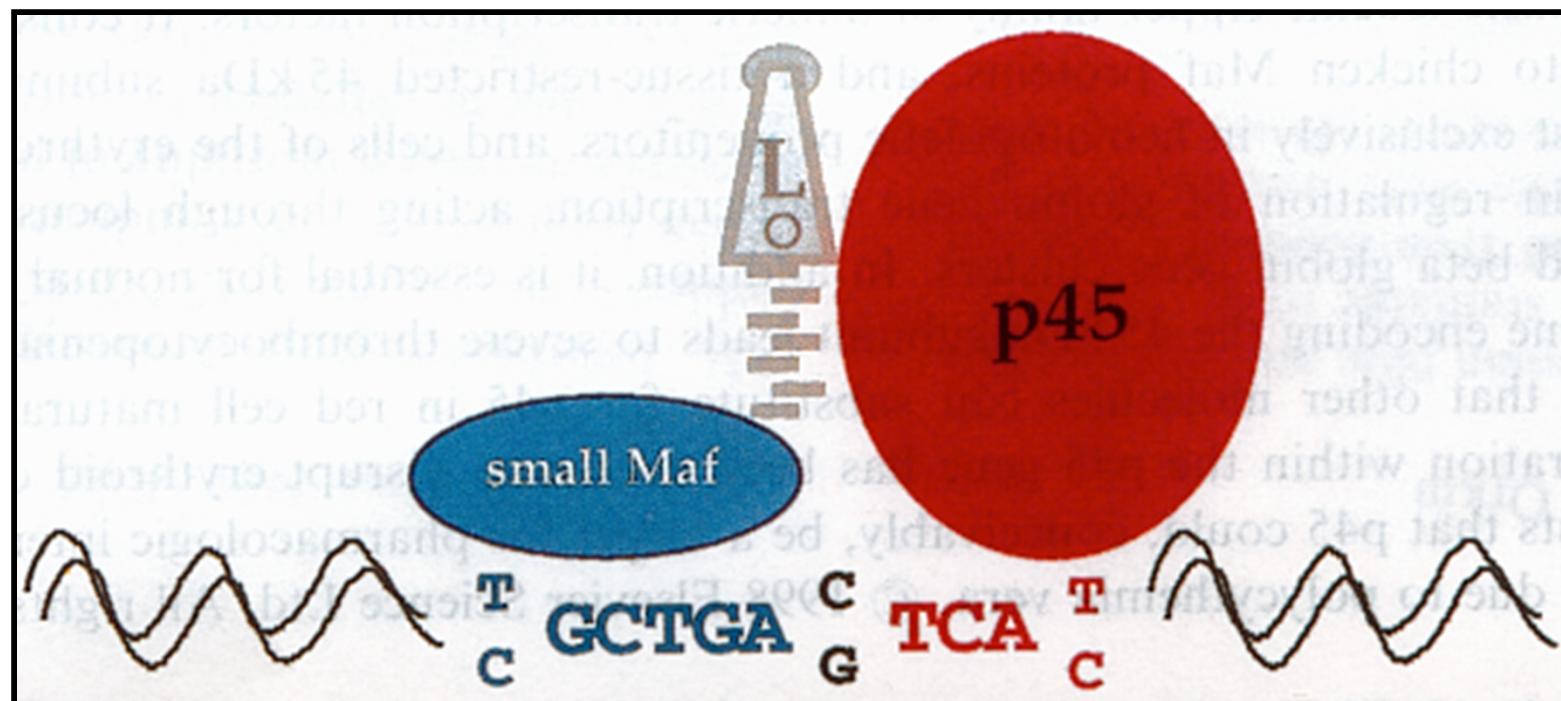
IL-3
Stem Cell Factor (SCF)
Glucocorticoids (GC)
Thyroid hormone (T₃)
Erythropoietin (Epo)
Insulin (Ins)

Transcription factors

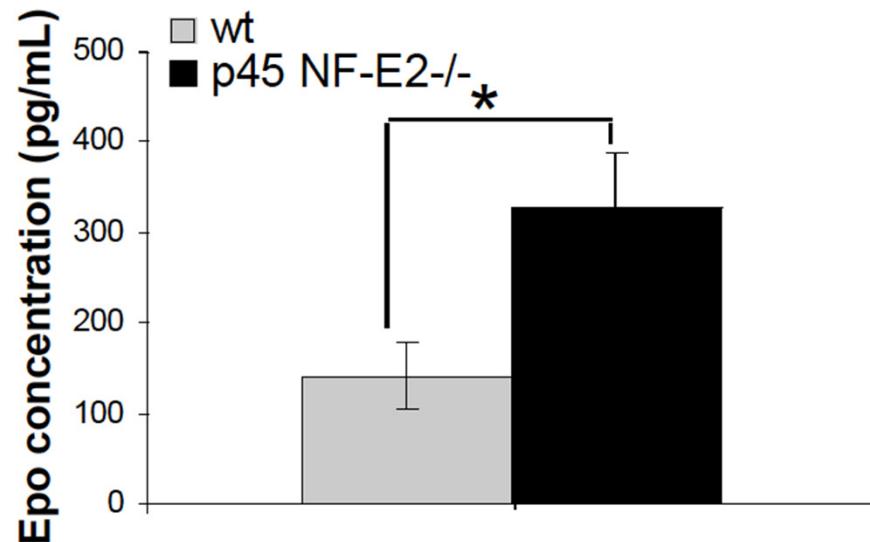
GATA-2 (G2)
GATA-1 (G1)
STAT5 (S5)
FOG (FG)
EKLF (EK)
NF-E2 (NF)

NF-E2 (nuclear factor erythroid 2)

- heterodimer (two subunits: p45 & small Maf)
- basic-leucine zipper
- large subunit specifically expressed in erythroid, megakaryocytic and mast cell lineages

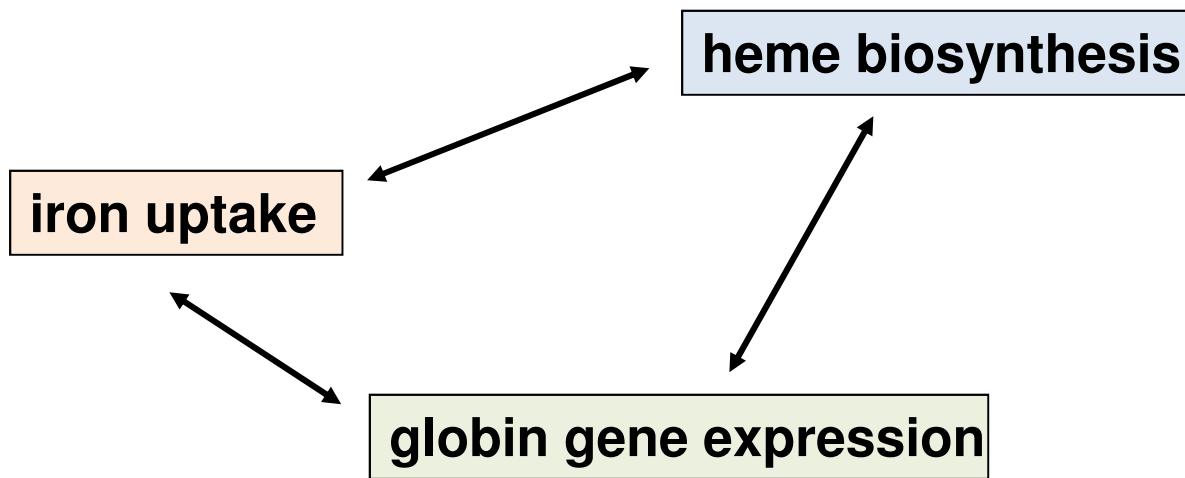


NF-E2 $^{-/-}$ mice

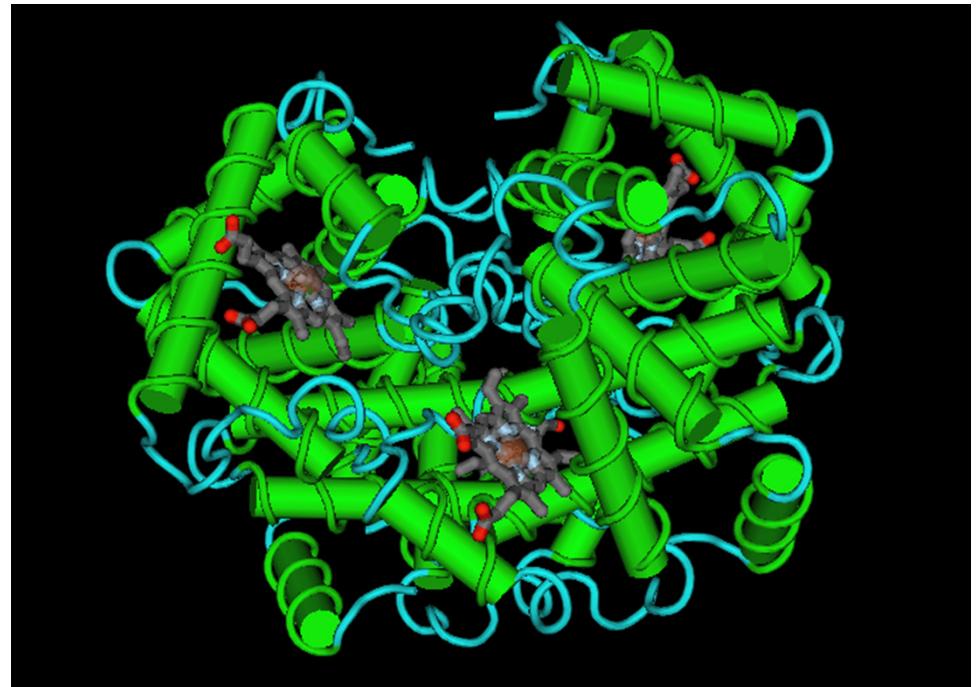


- increase in spleen erythropoiesis
- partial block in erythroid progenitor populations
- increase in GATA-1 levels
- cell cycle: increase of cells in G1

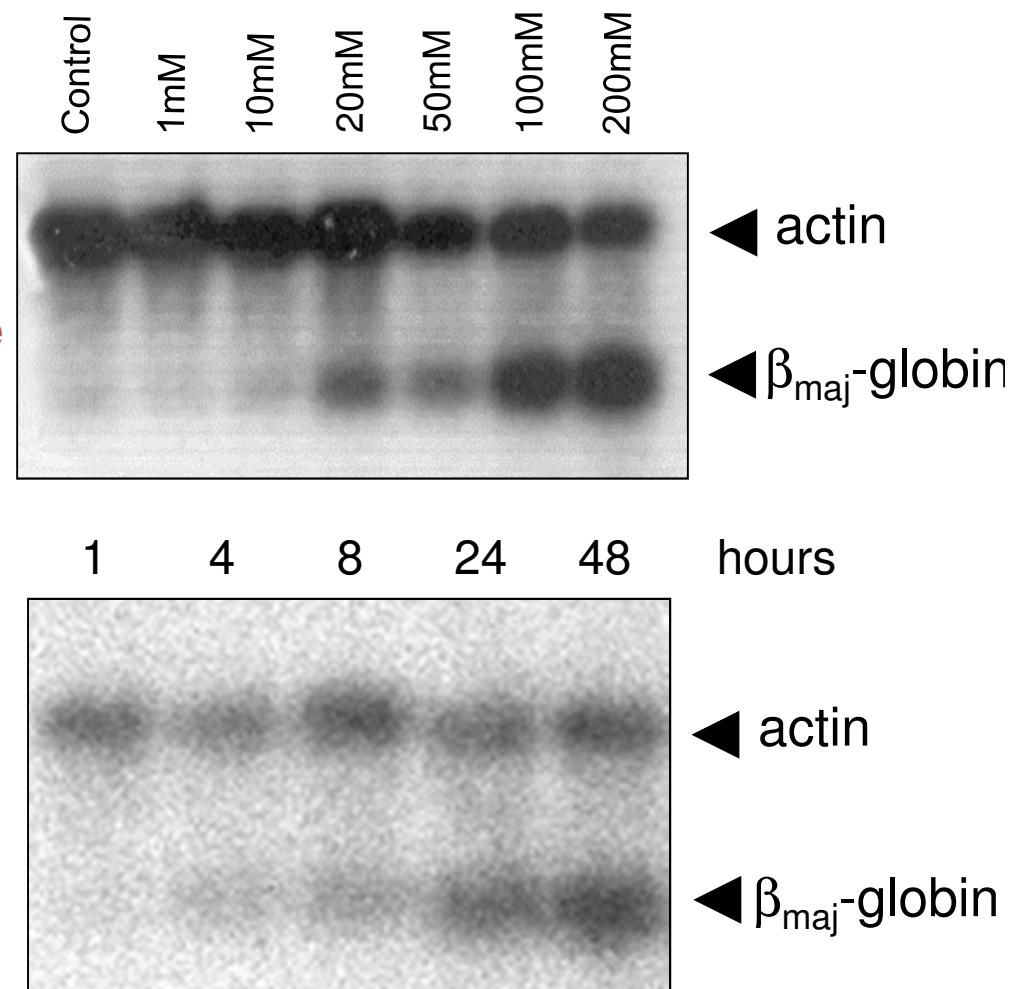
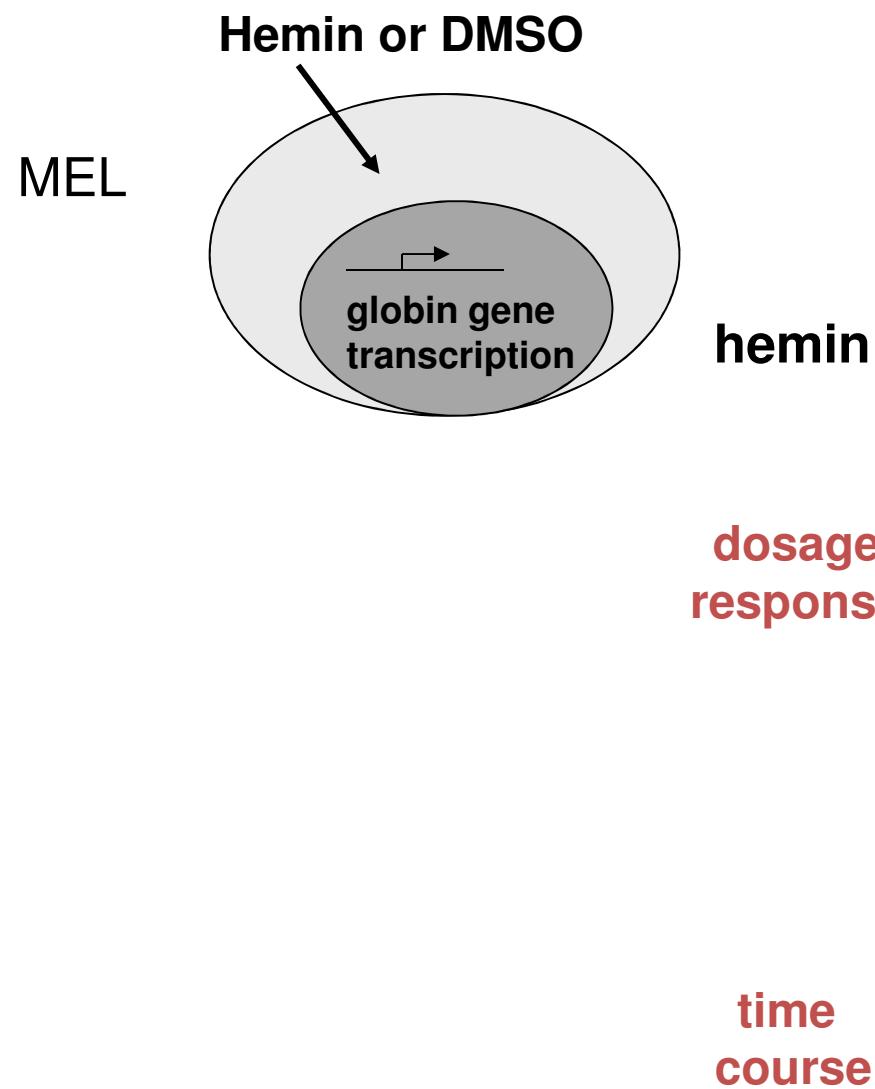
Hemoglobin synthesis



hemoglobin
 $\alpha_2\beta_2$ tetramer
α-chain 141aa
β-chain 146aa



Erythroleukemia cells



Models

MEL & G1-ER cells



Mice

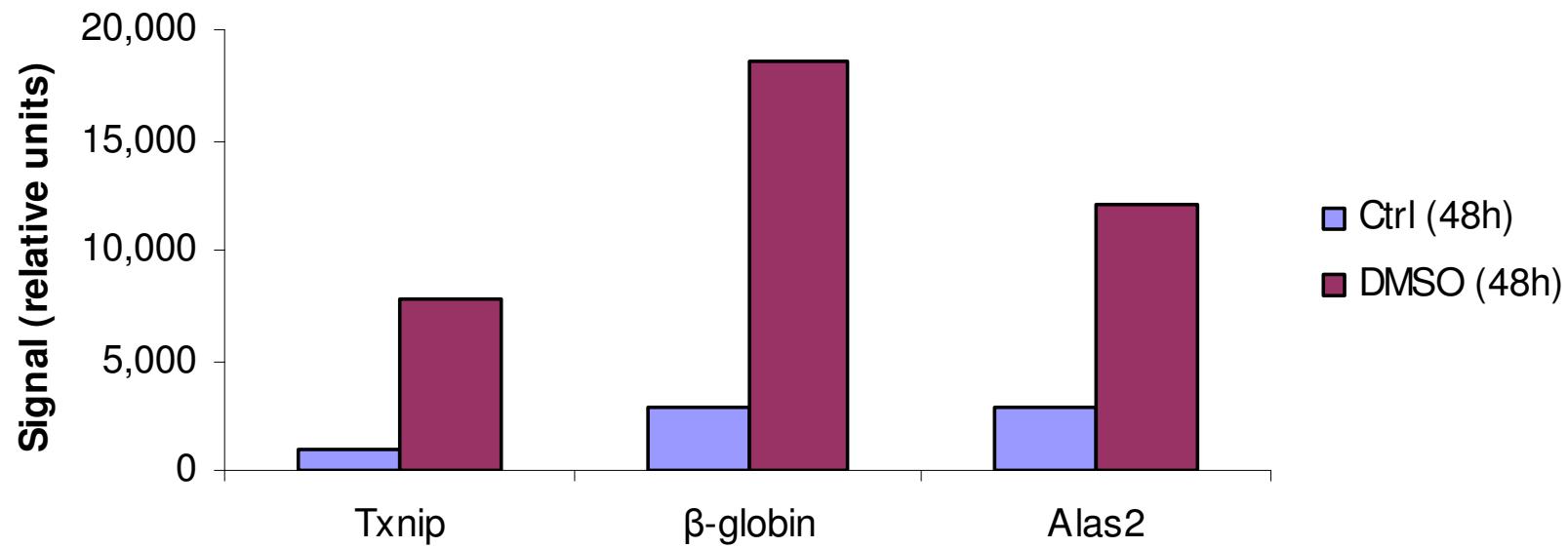


untreated

treated
(HMBA 96h)

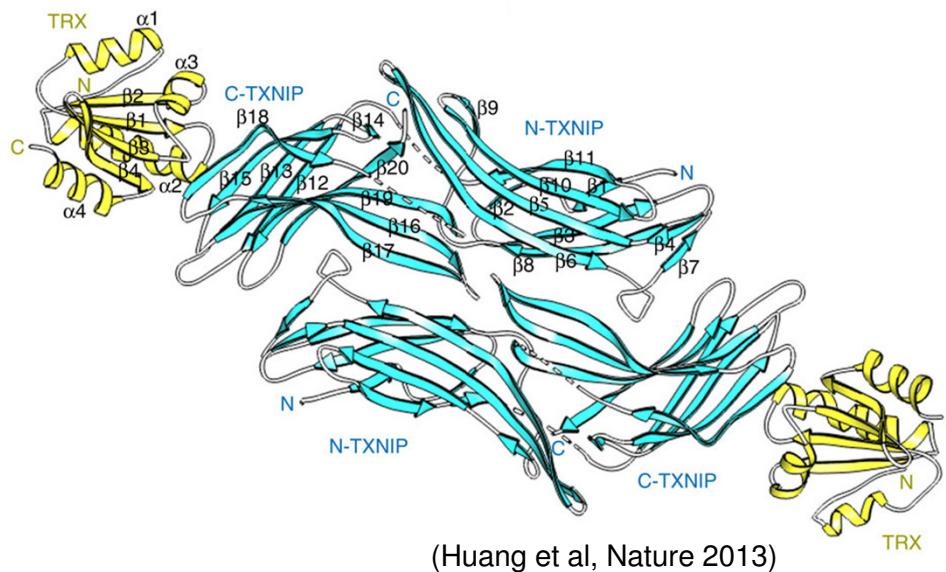
Genechip expression array (Affymetrix 430A)

- MEL cells
- differentially expressed TXNIP



TXNIP (Thioredoxin interacting protein)

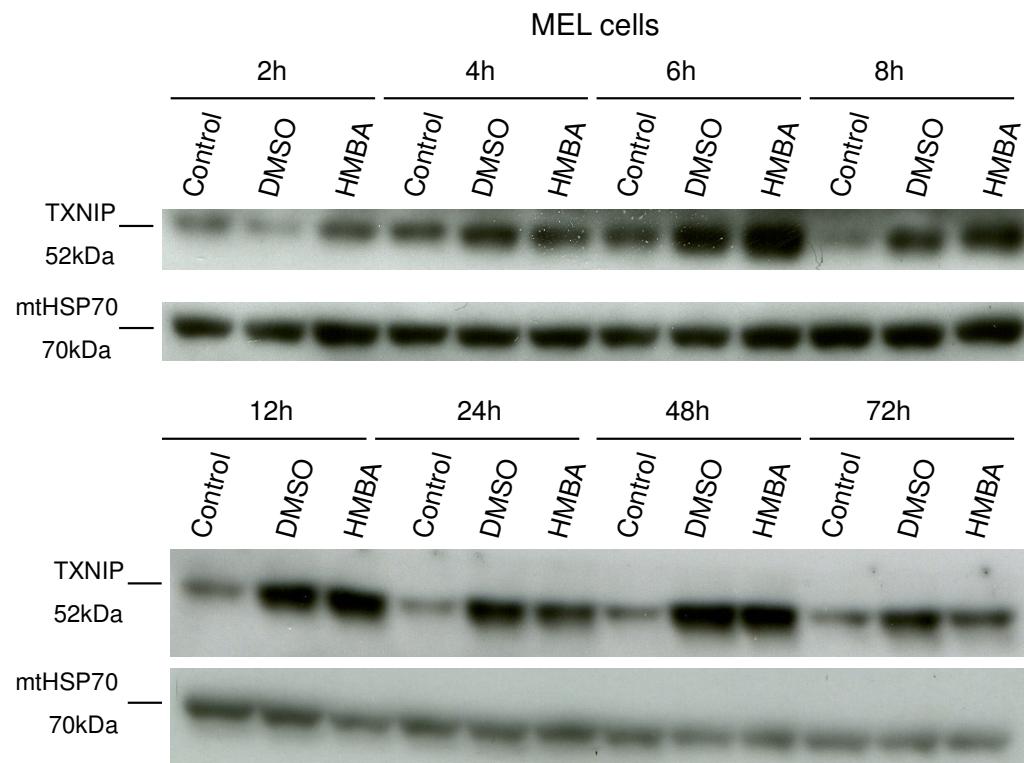
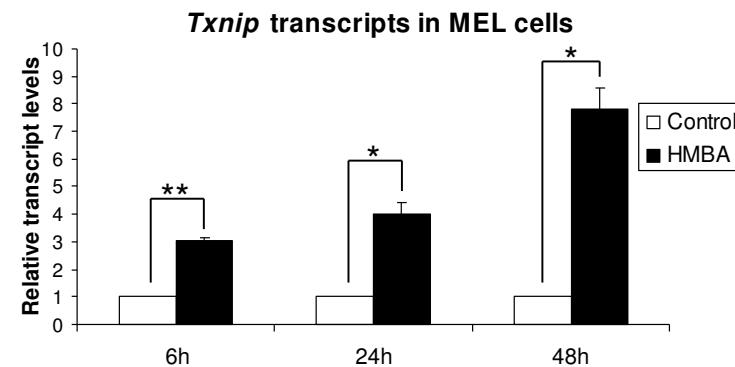
- ubiquitously expressed, ~50 kDa
- cytoplasm, nucleus, mitochondria, cell membrane
- tumor suppressor (expression reduced in many tumors)
- overexpression induces cell cycle arrest
- role in oxidative stress by inhibiting activity or bioavailability of thioredoxin



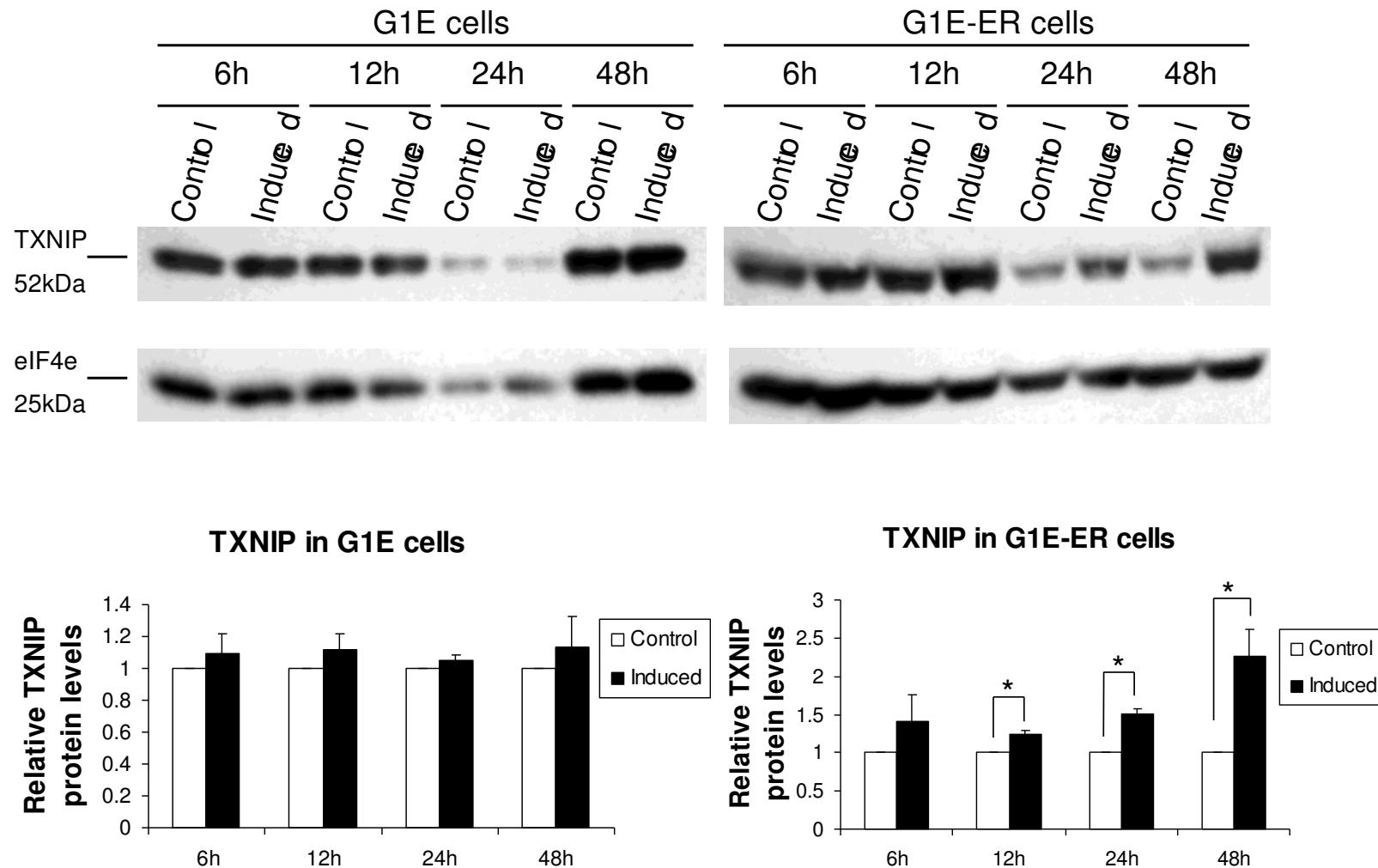
TXNIP $^{-/-}$ mice

Metabolism	In fasting conditions: hypoglycemia ↑ insulin ↑ triglycerides ↑ fatty acids in plasma (Hui et al. 2004, Donnelly et al. 2004, Oka et al. 2006)
Carcinogenesis	hepatocellular carcinoma (40% of mice) (Sheth et al. 2006) ↑ hepatocellular and bladder tumors (carcinogen-induced) (Kwon et al. 2010, Nishizawa et al. 2011)
Hematopoiesis	impaired maturation of NK cells (Lee et al. 2005) impaired function of dendritic cells (Son et al. 2008) altered HSCs (Jeong et al. 2009, Jung et al. 2013)

Induction of TXNIP in differentiating MEL cells

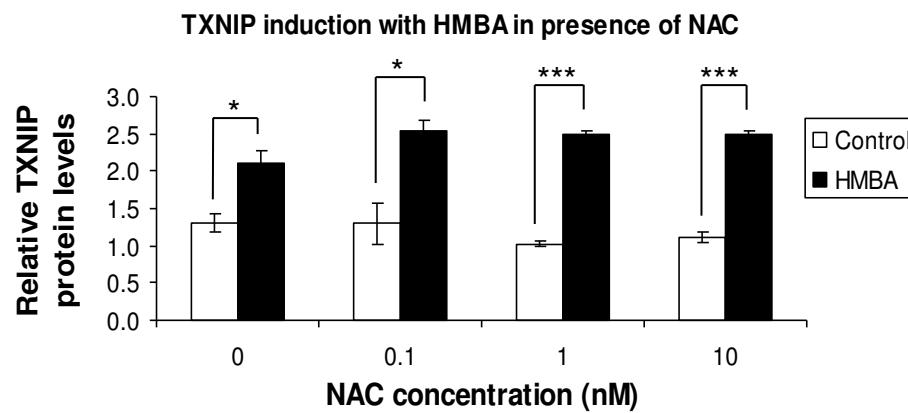
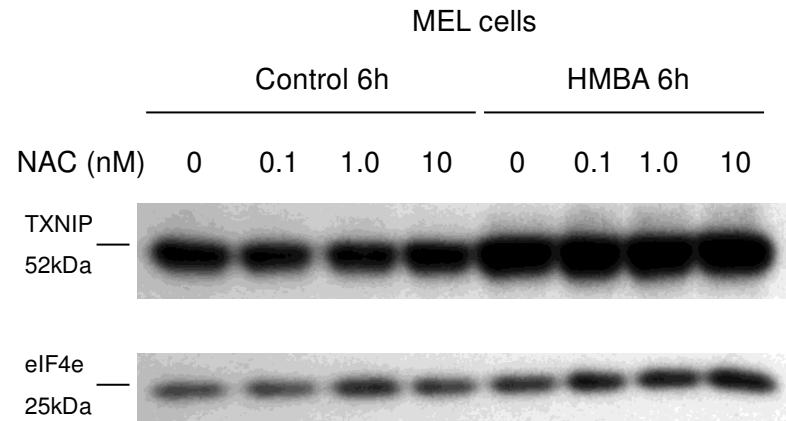


Induction of TXNIP in differentiating G1E-ER cells

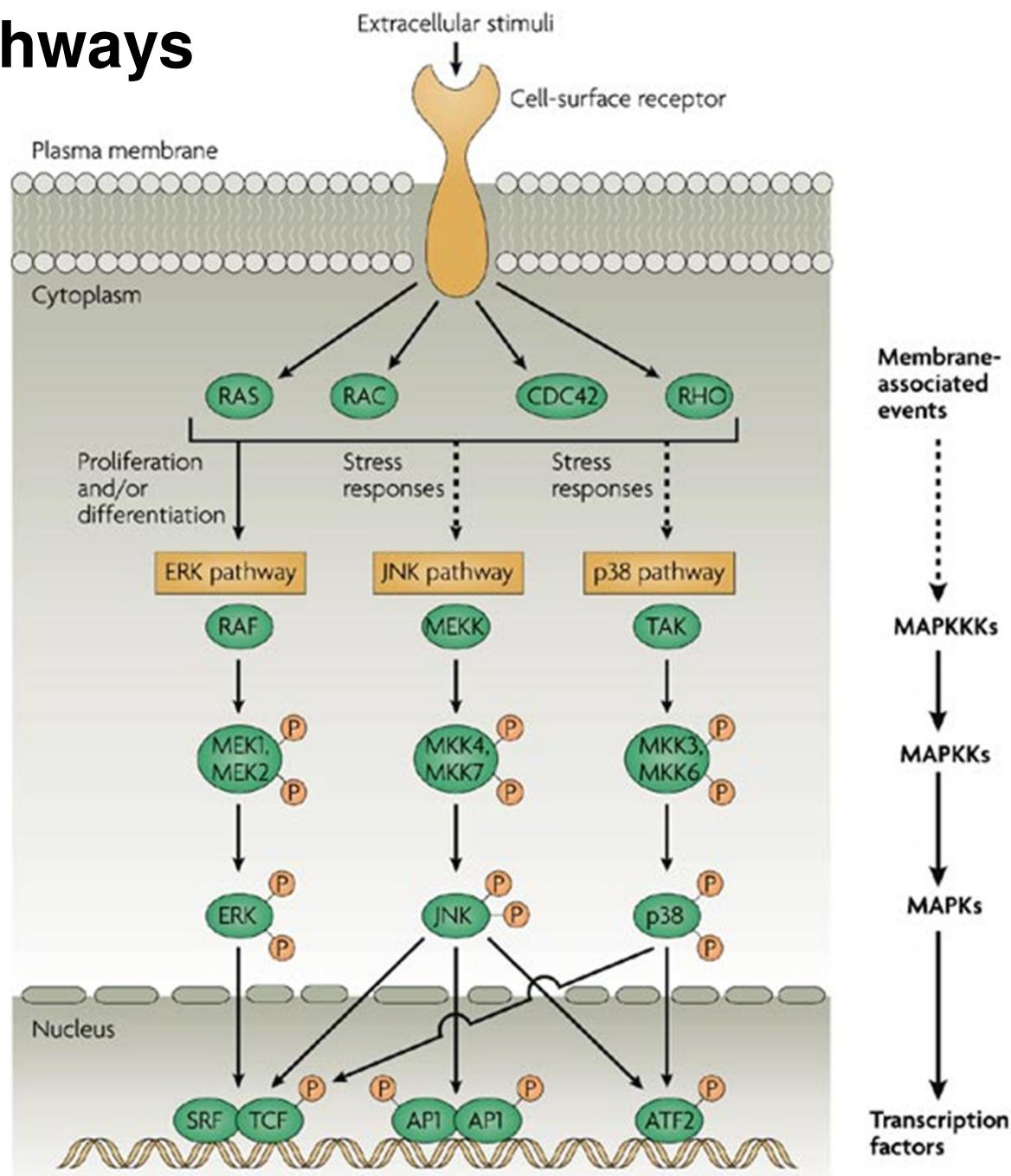


TXNIP & ROS scavenger NAC

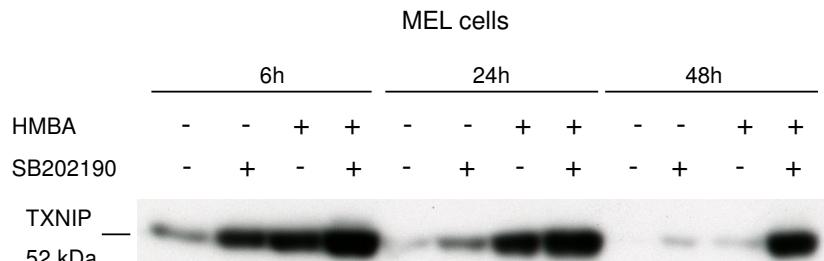
- TXNIP upregulation with HMBA unchanged



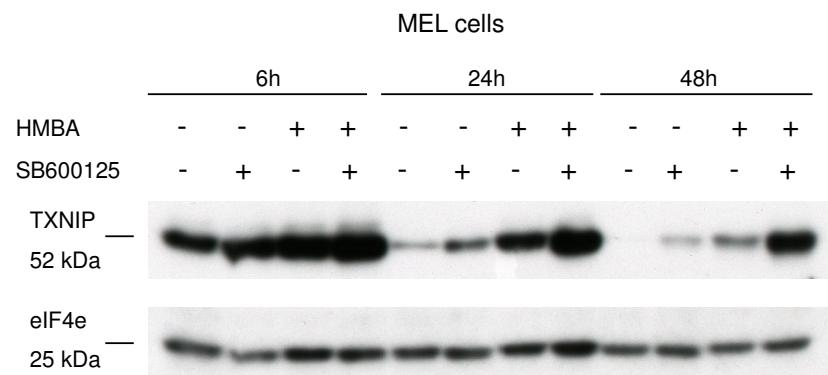
MAP kinase pathways



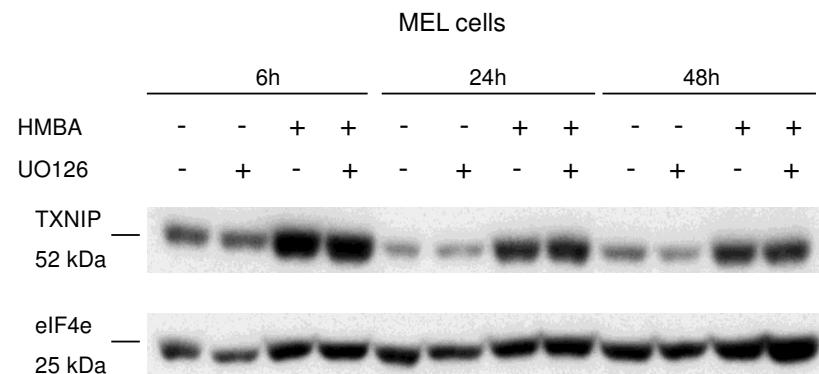
TXNIP – Regulation of MAP kinases



p38 inhibitor



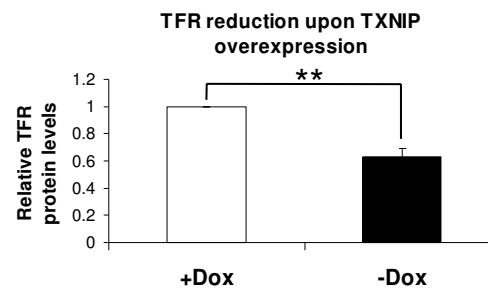
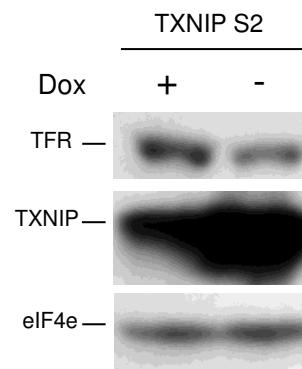
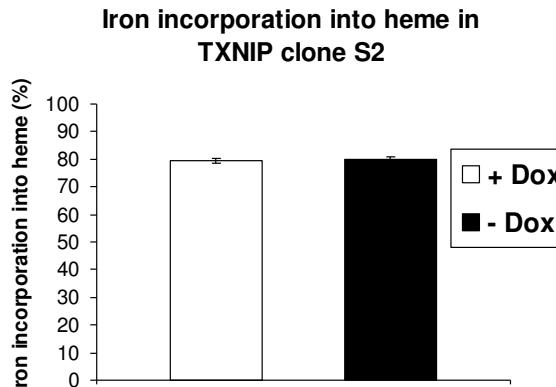
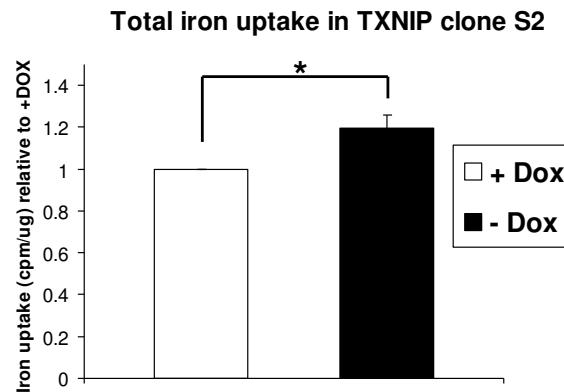
JNK inhibitor



ERK inhibitor

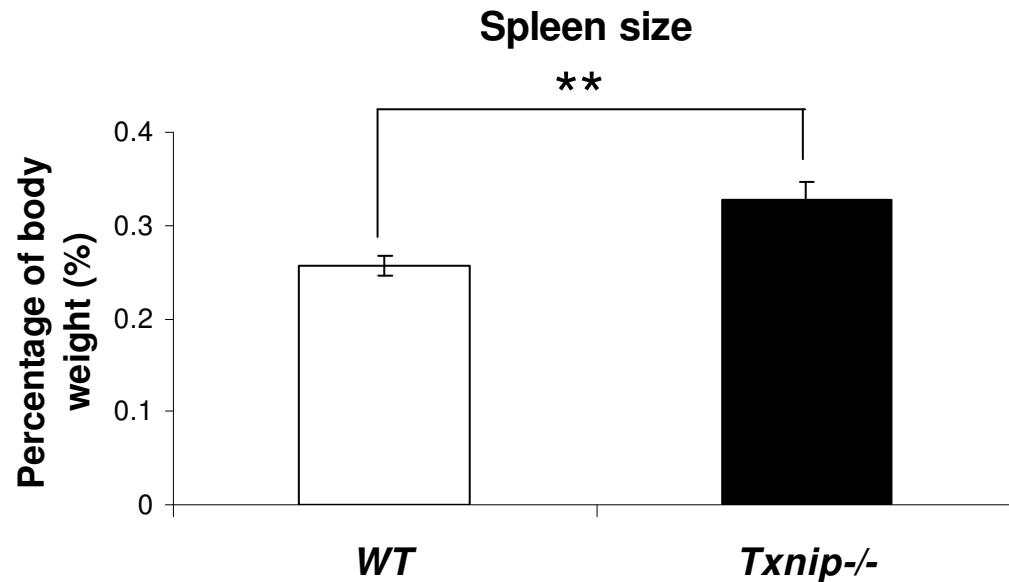
TXNIP overexpression

- increases iron uptake
- decreases TFR expression



Analysis of *Txnip*^{-/-} mice

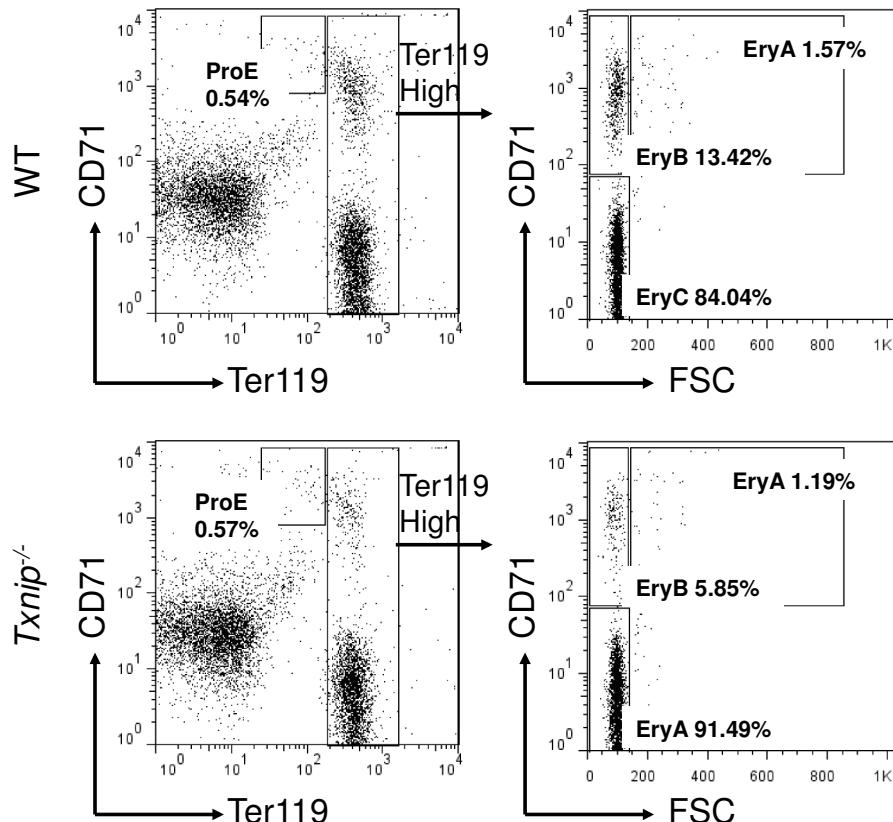
- spleen size
- blood parameters



	WT (n=5)	<i>Txnip</i> ^{-/-} (n=5)	p-value
RBC count, $\times 10^6/\mu\text{L}$	11.08 ± 0.29	11.05 ± 0.41	0.963
HGB level, g/dL	16.86 ± 0.50	17.78 ± 0.43	0.203
HCT, %	51.42 ± 1.37	53.30 ± 1.57	0.393

Erythroid precursor populations in *Txnip*^{-/-} mice

- spleen

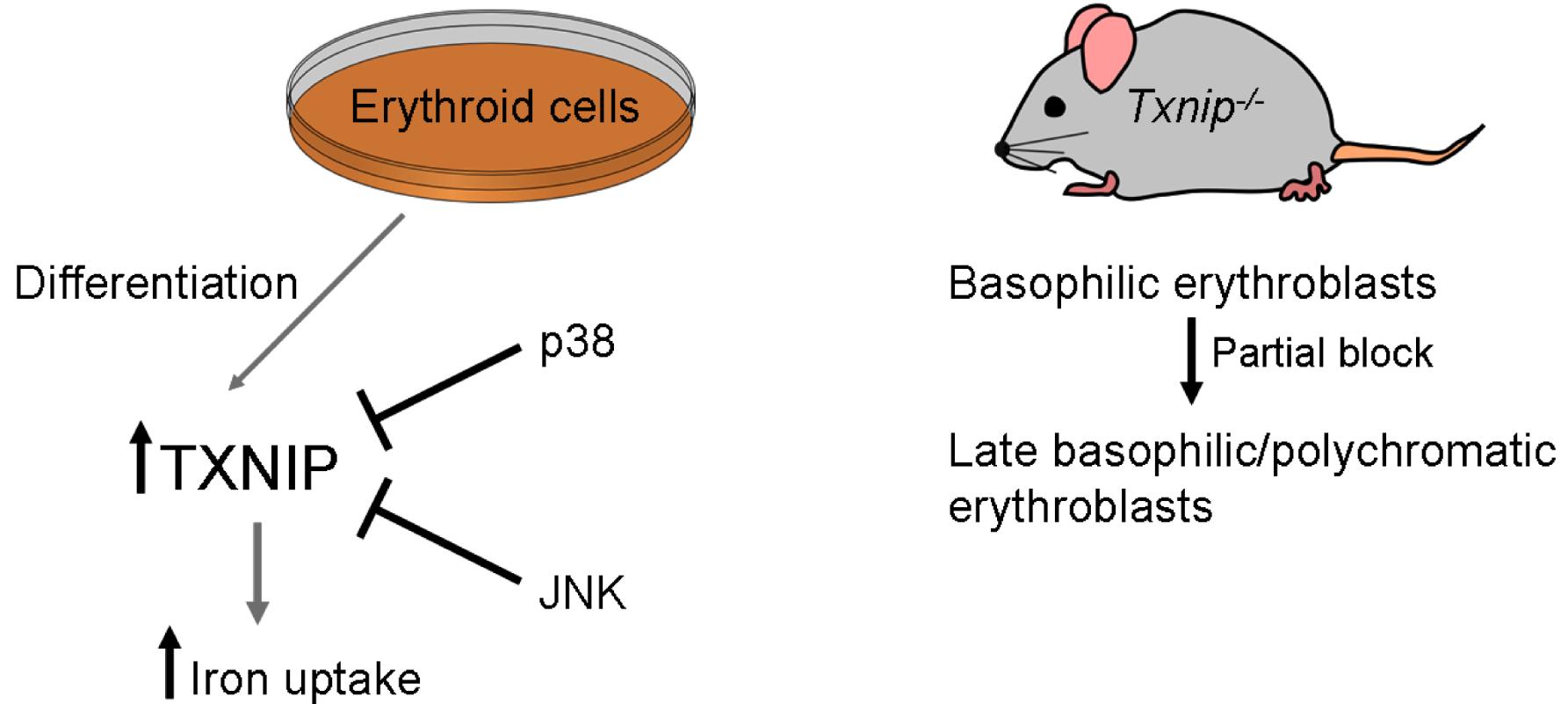


	WT (n=13)	<i>Txnip</i> ^{-/-} (n=14)	p-value
Spleen			
ProE	1.78 ± 0.35	2.37 ± 0.37	0.2580
EryA	2.99 ± 0.63	2.53 ± 0.48	0.5640
EryB	11.08 ± 1.24	6.72 ± 0.64	*0.0057
EryC	84.16 ± 1.61	88.38 ± 1.36	0.0563
Ter119+ cells	47.57 ± 1.87	45.17 ± 2.19	0.4130
Bone marrow			
ProE	6.40 ± 0.91	7.57 ± 0.69	0.315
EryA	14.90 ± 0.82	14.18 ± 0.82	0.546
EryB	25.85 ± 1.48	21.83 ± 1.44	0.063
EryC	52.85 ± 2.39	56.68 ± 2.20	0.251
Ter119+ cells	44.59 ± 2.60	42.97 ± 2.01	0.628

ProE	proerythroblasts
EryA	basophilic erythroblasts
EryB	late basophilic and polychromatic erythroblasts
EryC	orthochromatic erythroblasts and erythrocytes

Model

- Role of Txnip in erythroid differentiation



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