

YOUNG SCIENCE FORUM: IL FUTURO NASCE IN LABORATORIO



Analisi funzionale delle cellule eritroidi di pazienti con talassemia non trasfusione-dipendente trattati con luspatercept

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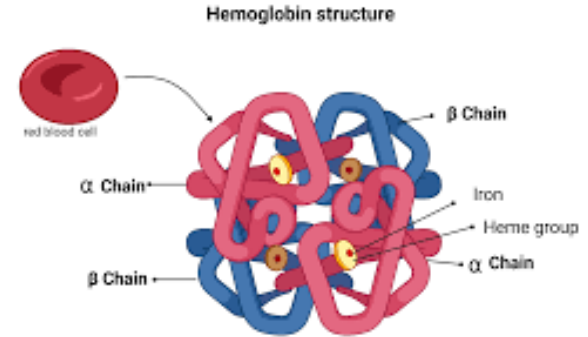
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Conflict of Interest: I declare no conflict of Interest

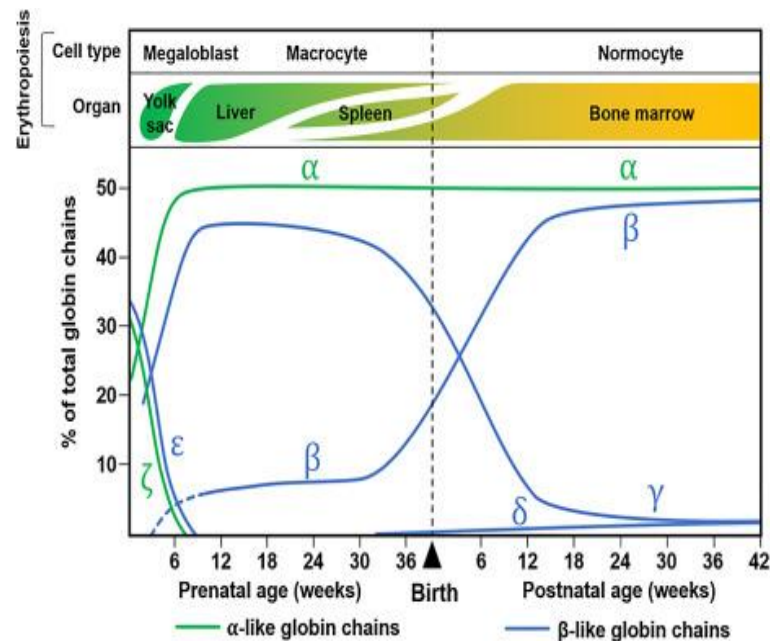
β -Thalassemia: A Disorder of Globin Synthesis

- ❖ β -thalassemia is a hereditary hemoglobin disorder characterized by reduced or absent β -globin chain synthesis.
- ❖ Reduced β -globin synthesis disrupts the α/β -globin chain balance.



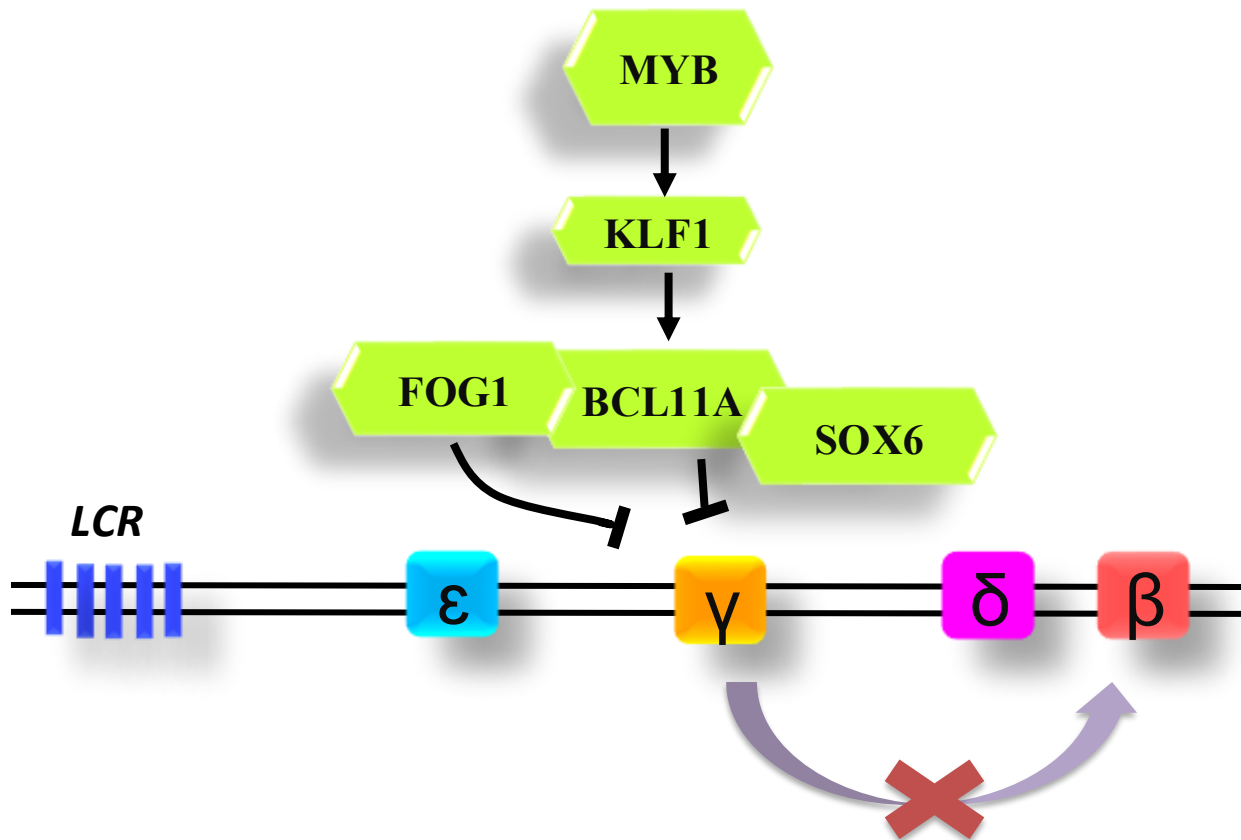
Fetal Hemoglobin (HbF) and Disease Modulation

- ❖ HbF predominates during fetal life and decreases after birth.
- ❖ β -thalassemia shows delayed γ -to- β globin switching.
- ❖ Elevated HbF reduces globin chain imbalance and ineffective erythropoiesis

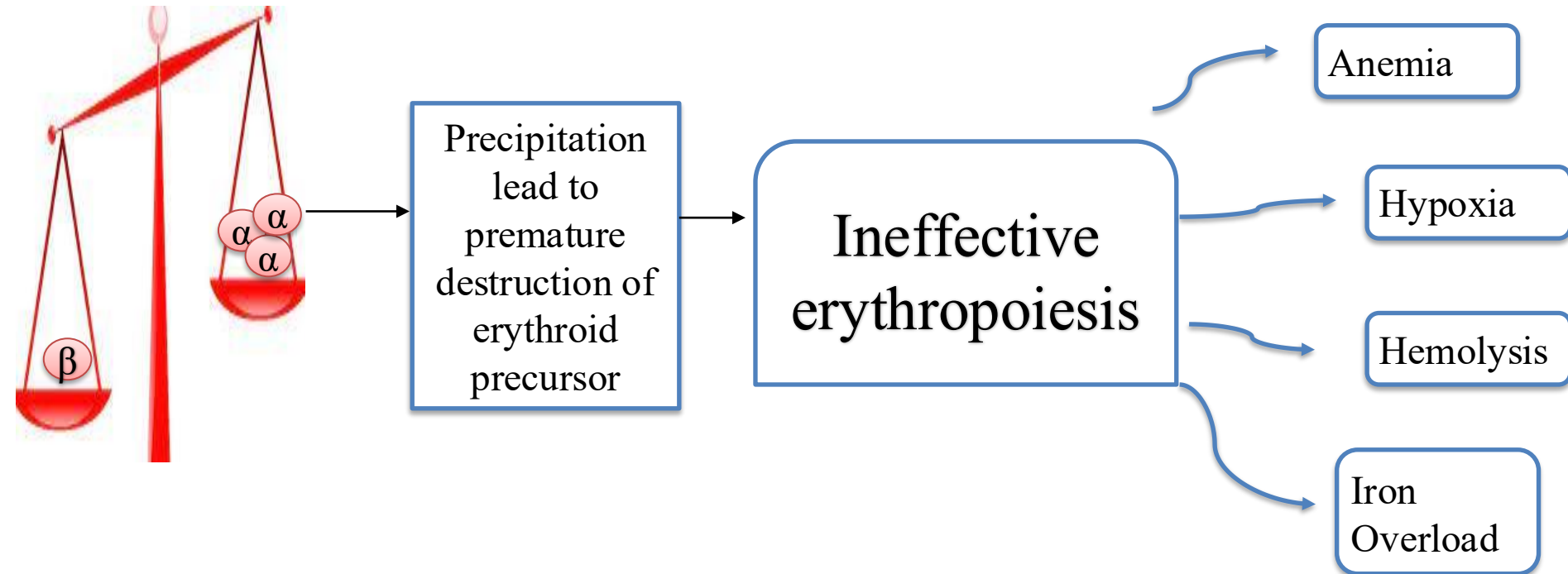


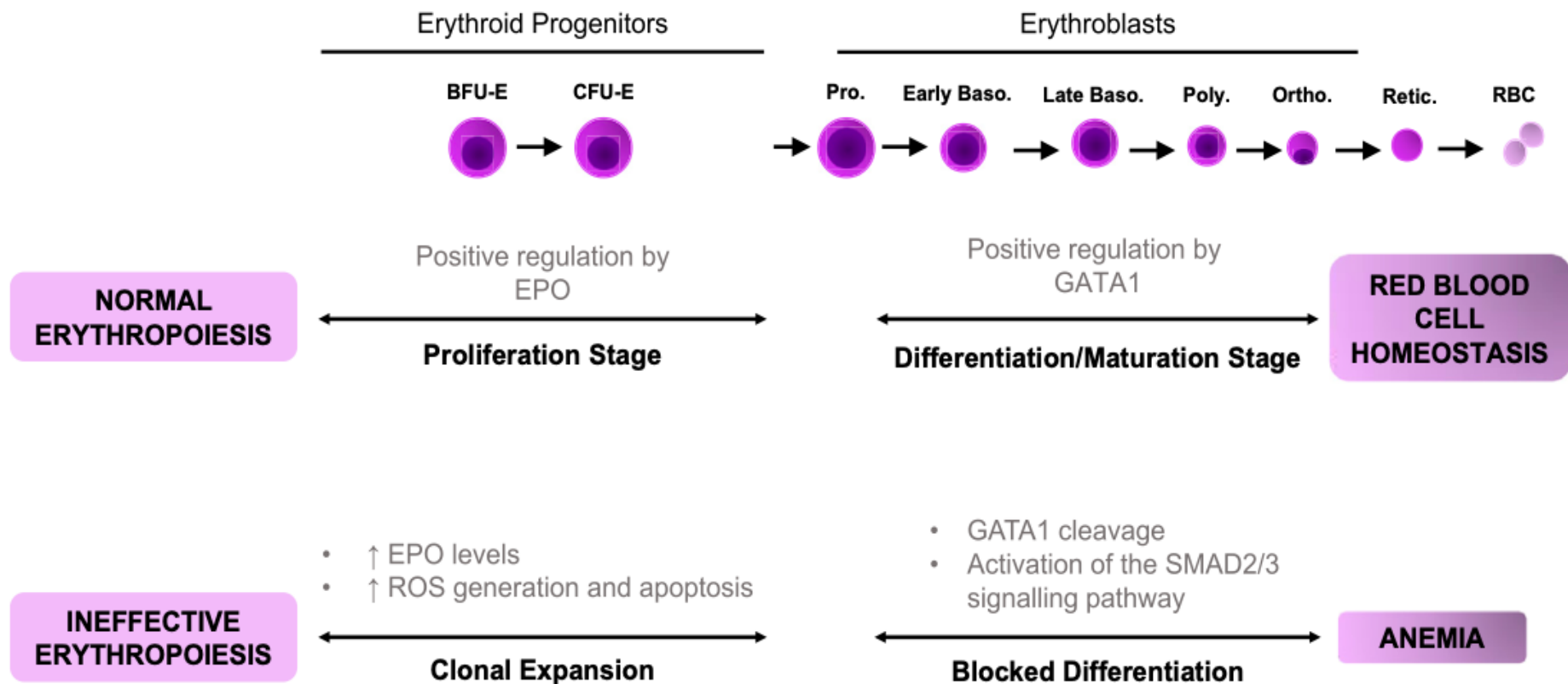
Induction of γ -globin is therapeutically beneficial in β -thalassemia.

Schematic representation of γ globin gene modulation



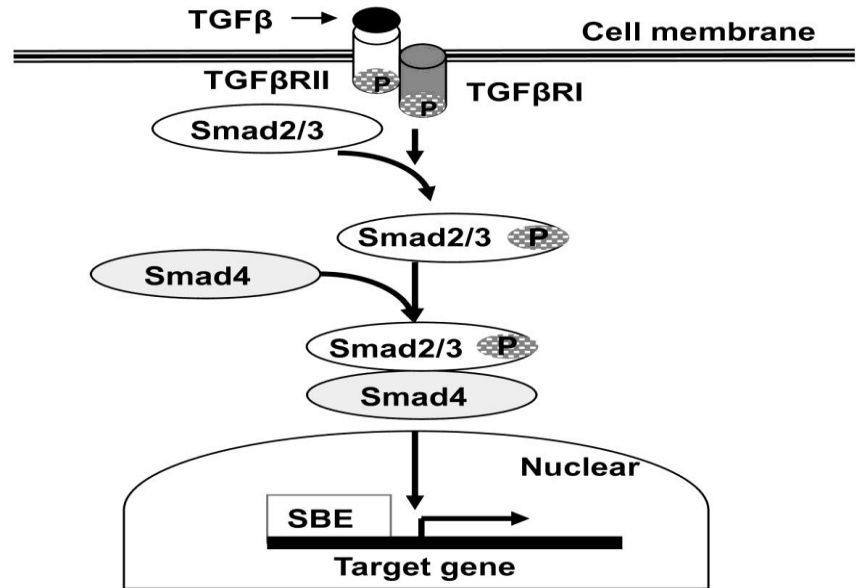
Pathophysiology of β -thalassemia





Ineffective erythropoiesis and SMAD Signalling

- TGF- β superfamily signalling contributes to ineffective erythropoiesis.
- Activin/GDF11-mediated SMAD2/3 activation suppresses late-stage erythroid maturation.

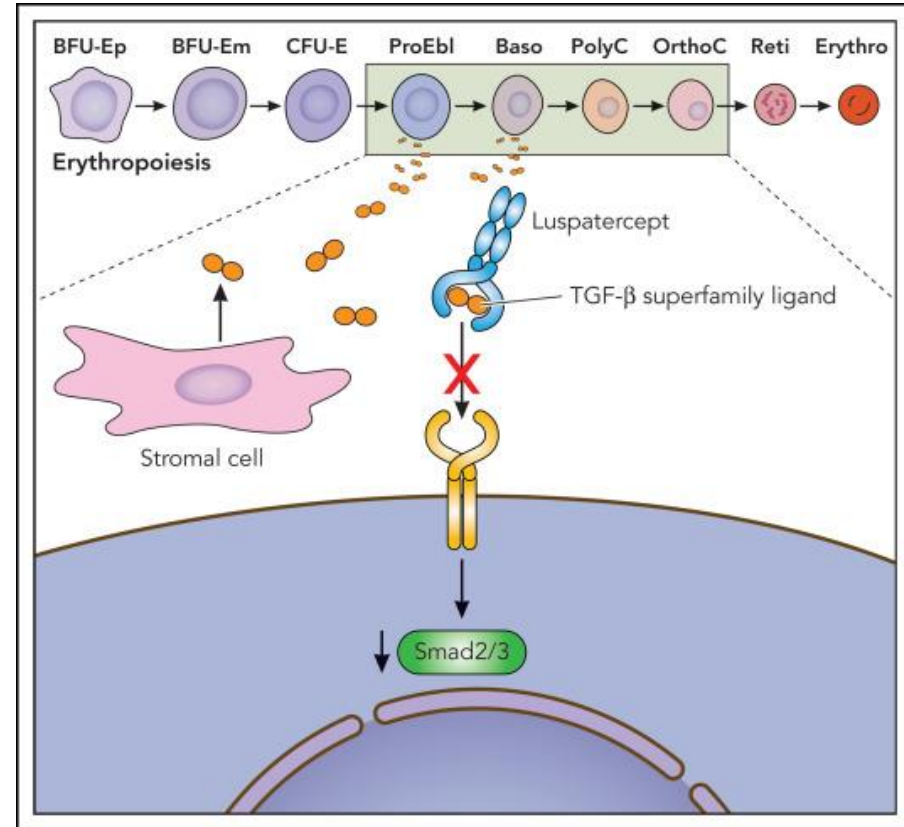


Mechanism line

GDF11 \rightarrow ActRIIB receptor \rightarrow SMAD2/3 phosphorylation \rightarrow impaired erythroid maturation

Luspatercept: Mechanism of Action

- Luspatercept (ACE-536) is an activin receptor type IIB ligand trap.
- It binds selectively to TGF- β superfamily ligands.
- This inhibits SMAD2/3 signalling.
- Result:
 - ✓ Enhanced late-stage erythroid maturation
 - ✓ Improved erythropoiesis



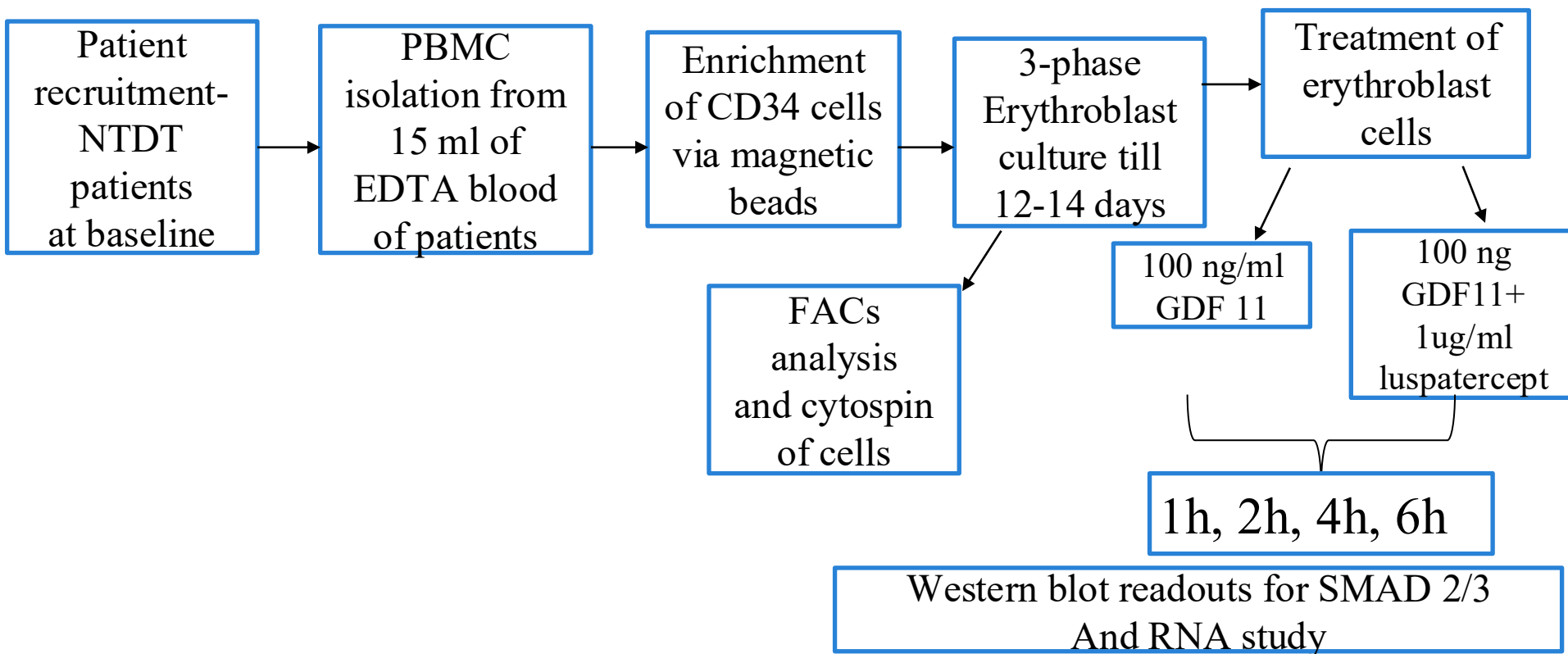
Hypothesis

Could luspatercept-mediated inhibition of SMAD2/3 signalling alter erythroid transcriptional networks and enhance γ -globin expression in β -thalassemia erythroid cells?

AIM of the Study

To investigate the effects of luspatercept on SMAD2/3 levels, erythroid transcription factors (GATA1, BCL11A) and γ globin gene expression in erythroid cells derived from non transfused dependent thalassemia patients

Workflow



Results

Isolation and Enrichment of CD34 cells

15 ml EDTA blood : PBMC: 3.2×10^7 cells

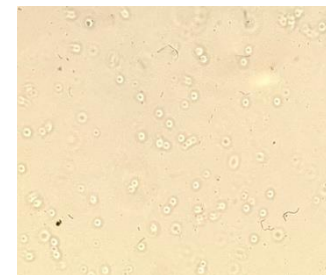


CD34+: 5.44×10^5 cells

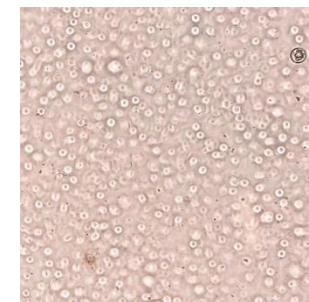


3 phase liquid culture 12 day

Erythroid cells: 3.67×10^7 cells

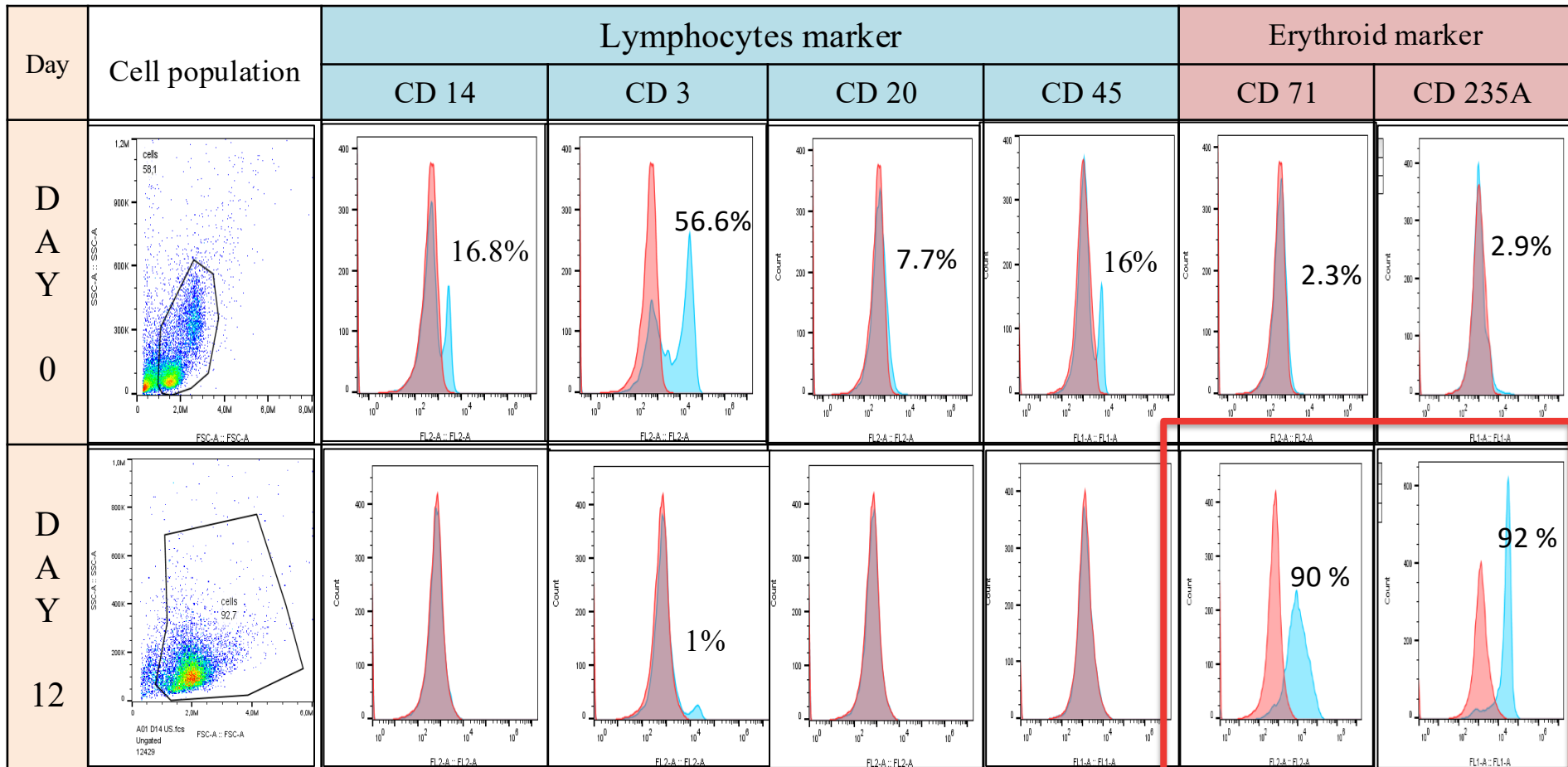


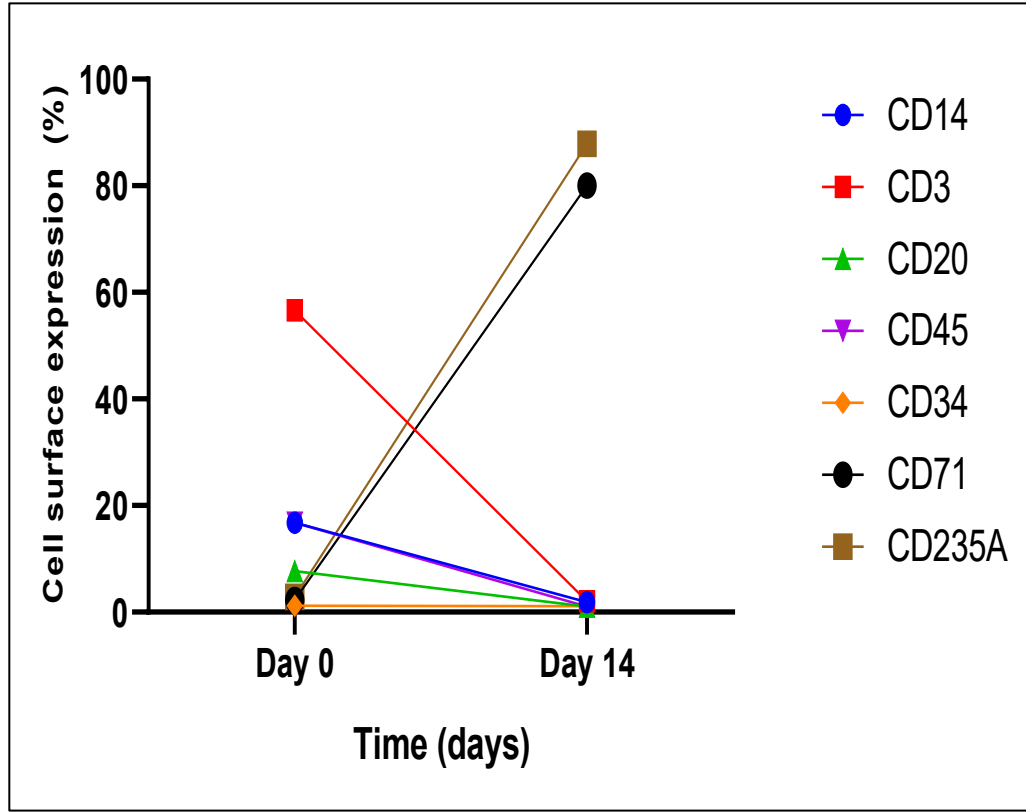
Day 0



Day 12

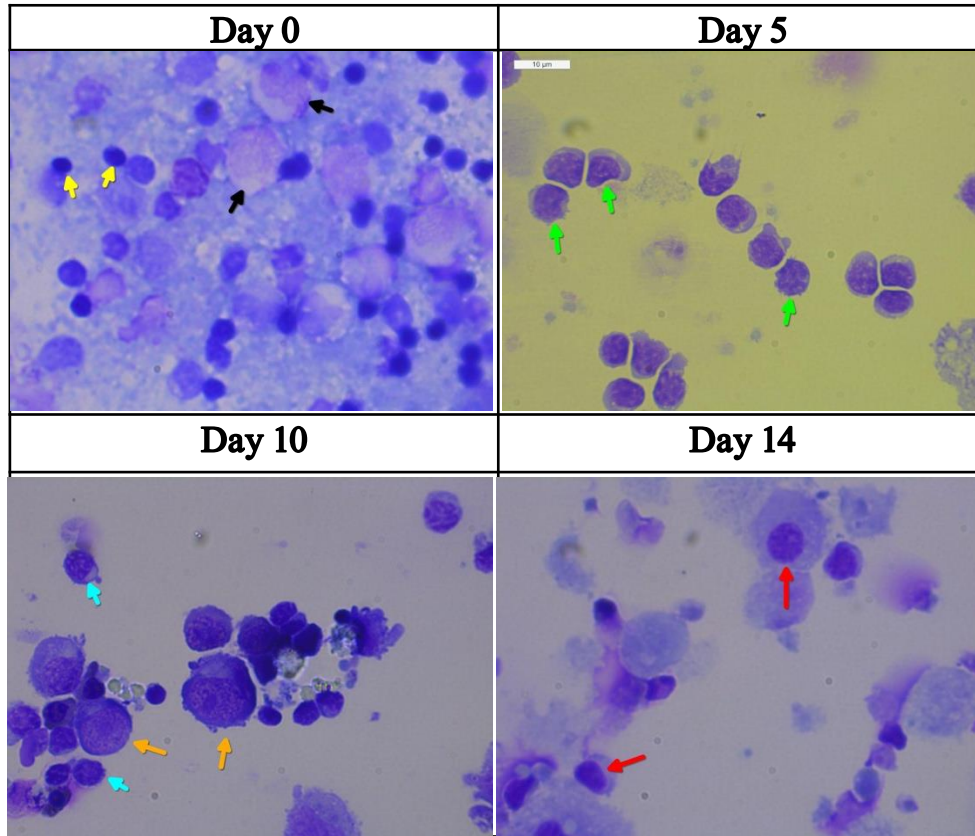
Validation of erythroid differentiation (FACS)





Efficient terminal erythroid differentiation achieved

Cell morphology assessment



Day 0: lymphocytes (yellow arrows) and monocytes (black arrows)

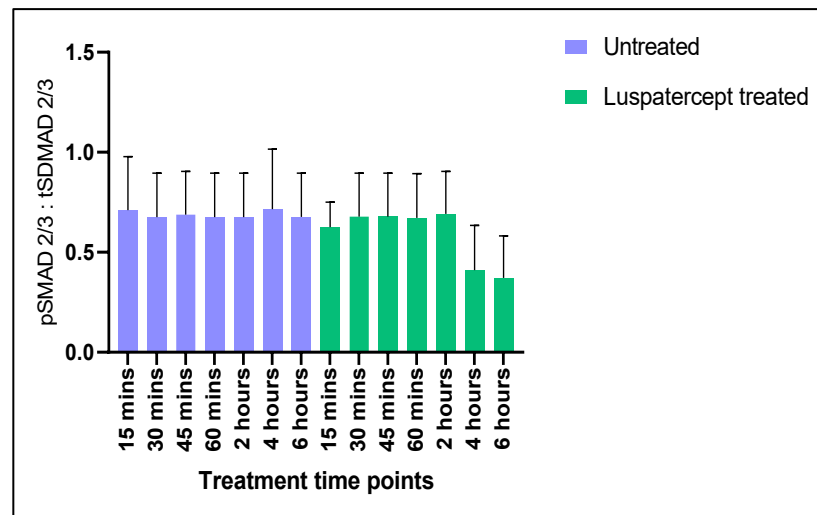
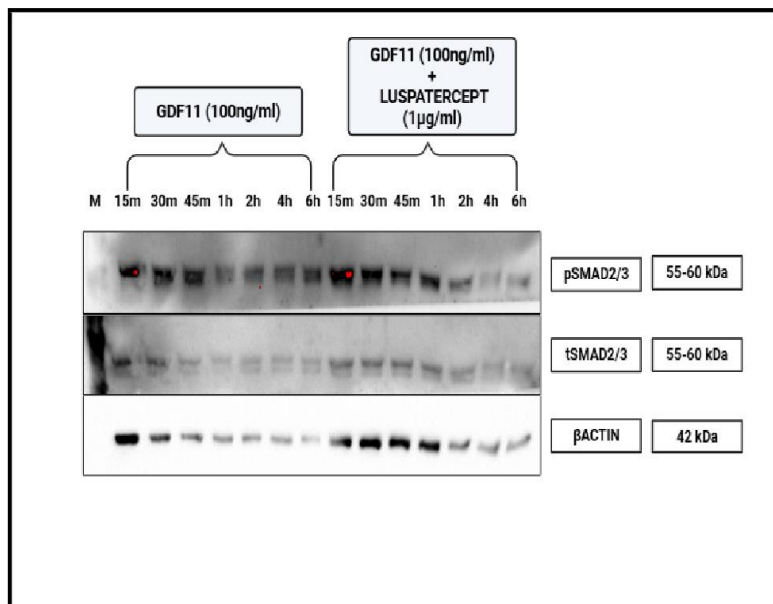
Day 5: proerythroblast stage (green arrows)

Day 10: basophilic erythroblasts, (blue arrows); polychromatic erythroblasts, (orange arrows)

Day 14: orthochromatic erythroblasts (red arrows)

Western blot Readouts SMAD2/3 signaling

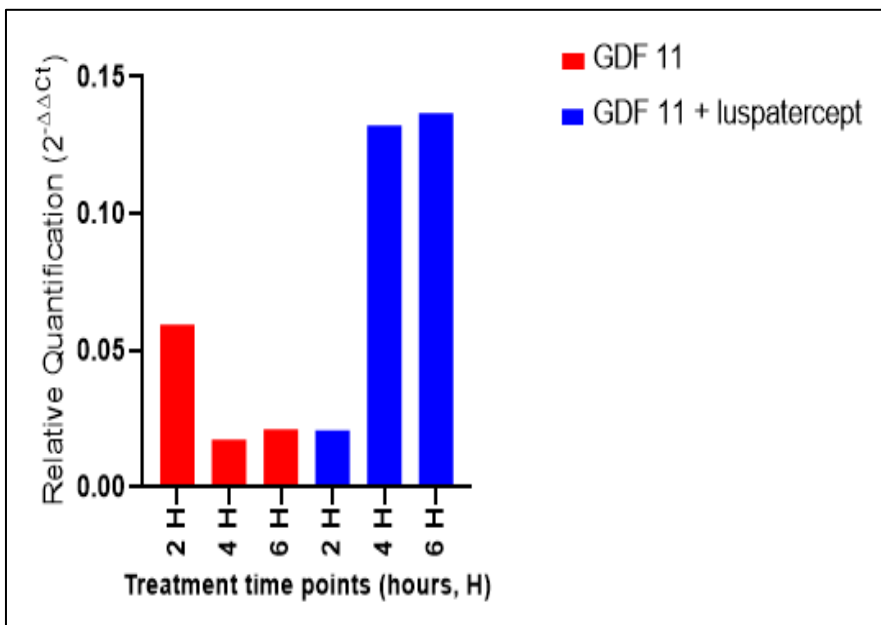
Effect of Luspatercept on pSMAD 2/3 levels



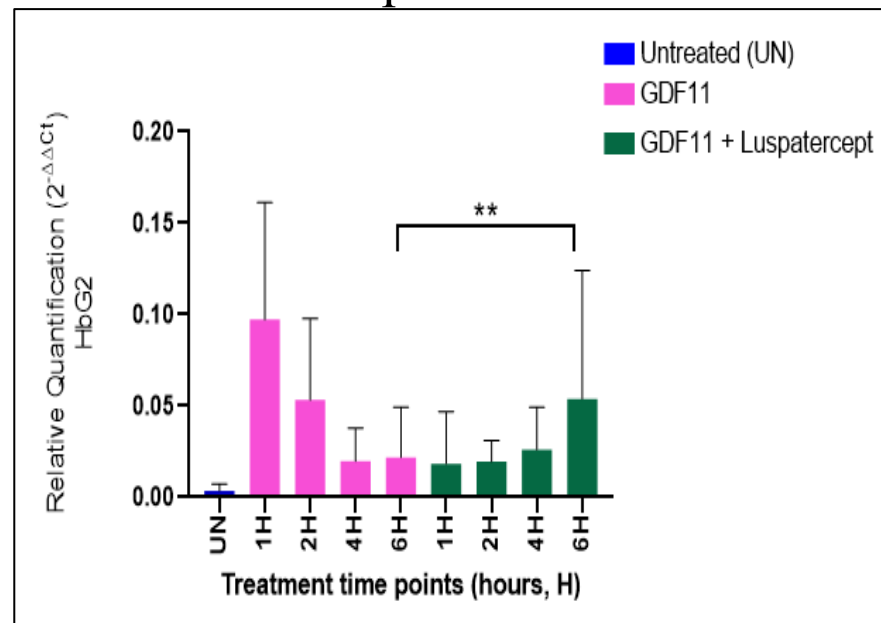
Luspatercept treatment led maximal decrease in pSMAD2/3 levels by 25% at 6 hours in erythroid cells of patients

Effect of luspatercept on HbG2 (γ) gene expression

Non transfused thalassemia

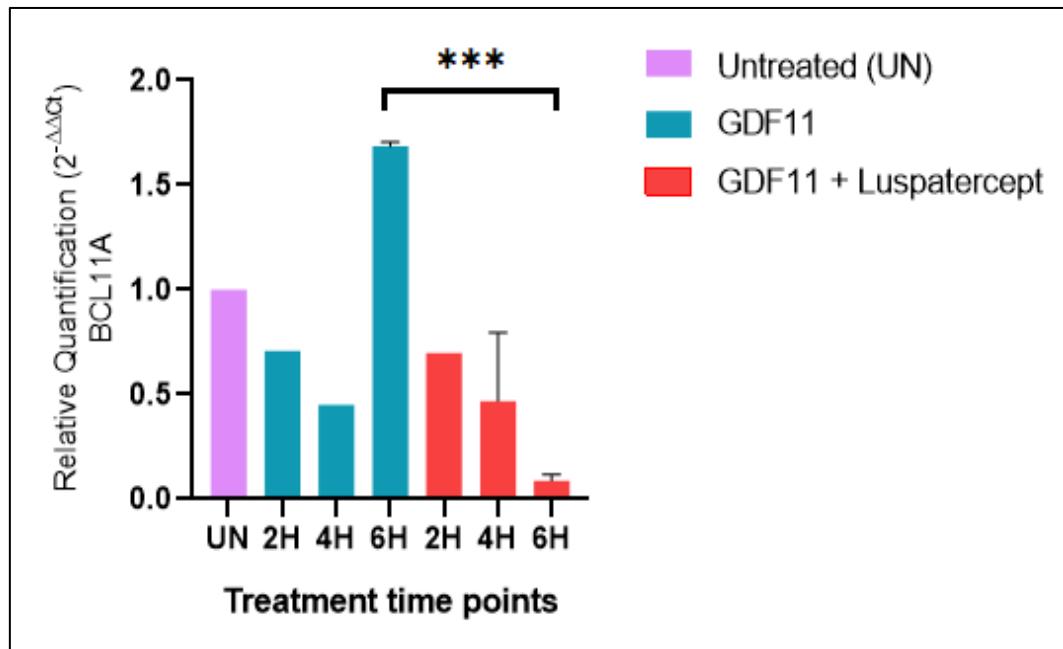


Transfused dependent thalassemia



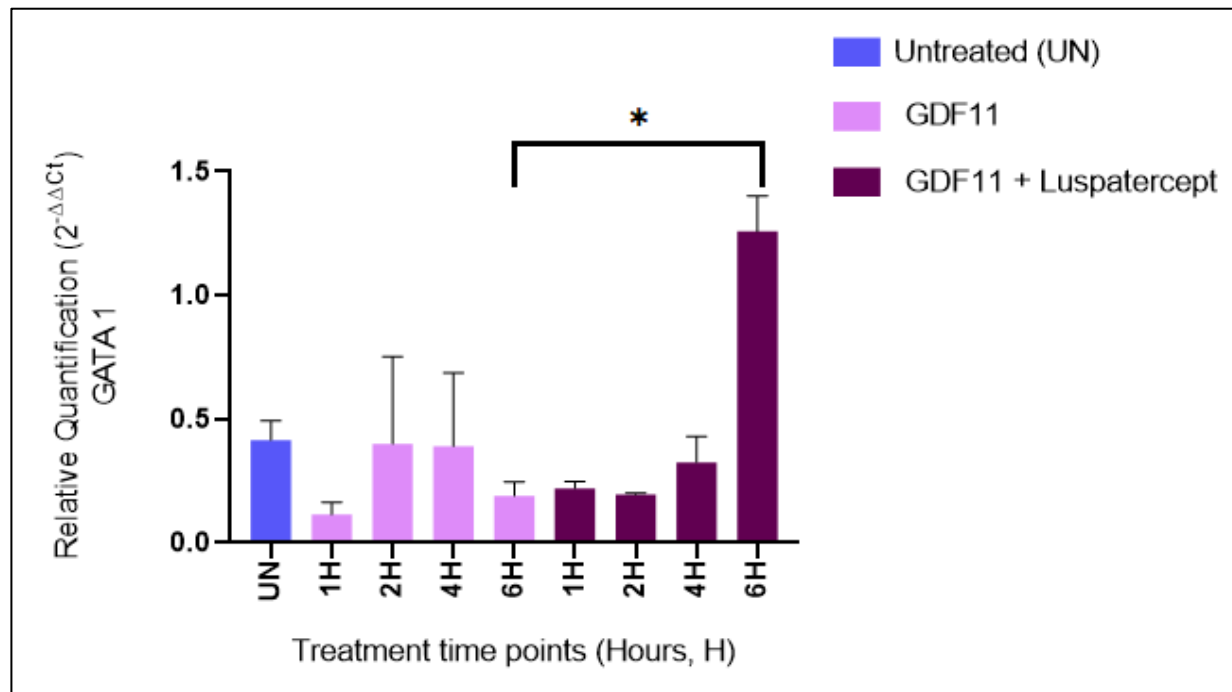
Luspatercept induced a time-dependent increase in γ -globin expression, reaching a significant 6.1-fold upregulation at 6 h compared with GDF11 alone.

Effect of Luspatercept on BCL11A expression



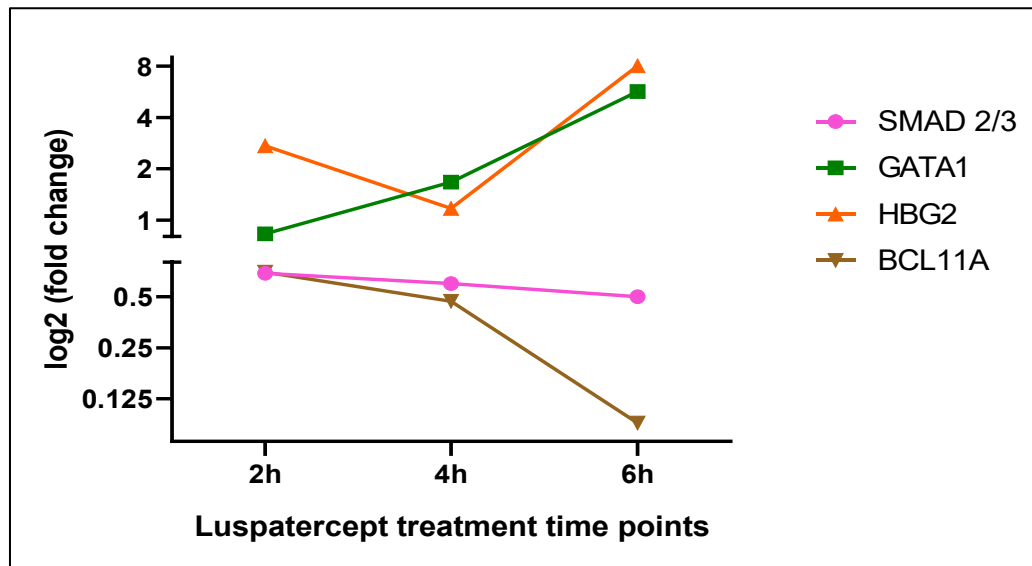
BCL11A expression decreased significantly following luspatercept treatment, reaching a 9.3-fold reduction at 6 h compared with GDF11 alone ($p < 0.0001$)

Effect of Luspatercept on GATA1 expression



GATA1 gene expression was significantly elevated (5.7-fold, $p < 0.05$) after 6 h of luspatercept treatment compared to GDF11 alone.

Integrated response of luspatercept



Luspatercept appears to coordinate signalling and transcriptional pathways associated with erythroid maturation

Conclusion

- ❖ Luspatercept reduces GDF11-induced SMAD2/3 signalling and is associated with increased GATA1 and γ -globin expression along with decreased BCL11A expression in β -thalassemia patient-derived erythroid cells.
- ❖ Overall, these findings suggest that luspatercept promotes a fetal hemoglobin–favourable erythroid transcriptional program through modulation of the SMAD–GATA1–BCL11A axis.

Plan for Future

- ❖ Future studies will use time-resolved RNA-seq (2h, 4h, 6h, 8 h) in untreated, GDF11, and GDF11 + luspatercept conditions to identify key transcripts and regulatory networks involved in its mechanism.
- ❖ Candidate genes will be functionally validated to define their role in luspatercept-mediated erythroid reprogramming.

Acknowledgment

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THANK YOU

THE **RNA WORLD** WAS
THE BEGINNING.

RNA STUDY IS
THE **FUTURE.**



ORIGIN



EVOLUTION



DISCOVERY

FUTURE

