Mitigation of Injection Site Reactions after Subcutaneous Administration of Dalcinonacog Alfa (DalcA) in Hemophilia B Using Preclinical Models

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Introduction

Prophylactic subcutaneous (SQ) administration of dalcinonacog alfa (DalcA), an engineered recombinant factor IX (FIX) variant with 22-fold enhanced potency compared with BeneFIX, provides stable and protective therapeutic FIX levels in individuals with Hemophilia B. Phase 1 and 2b studies demonstrated DalcA was well tolerated and efficacious, showing sustained FIX levels and no bleeding events from therapy start through washout. However, some subjects reported mild-to-moderate injection site reactions (ISR) consisting of pain and/or redness following initial SQ injections. To investigate the underlying ISR mechanism, we examined cutaneous microanatomic, cellular, and proteomic changes after SQ DalcA administration in the HypoSkin[®] injectable human skin biopsy platform (Genoskin, Salem, MA, USA) and Göttingen minipigs.

Reverse Translational Approach

Daily DalcA SQ injections for 8-14 days induced erythema and redness in minipigs comparable with the observed human ISR. Human skin biopsies treated with DalcA underwent proteomic analysis to assess if minipig ISR predictors could be identified in human skin 36 hours after SQ injection. ISR-like response classification in this ex vivo setting was based on observed histological modifications.



Results

Table 1: ISR-like proteomic signature

UniProt ID	Gene	Up/Down regulated	UniProt ID	Gene	Up/Down regulated
O43707	ACTN4	down	P29350	PTPN6	up
P00738	HP	down	P38606	ATP6V1A	up
P01714	IGLV3-19	down	P52597	HNRNPF	up
P35749-3	MYH11	down	P53999	SUB1	up
P35749	MYH11	down	P78371	CCT2	up
O14558	HSPB6	up	Q15181	PPA1	up
P00747	PLG	up	Q15185	PTGES3	up
P04004	VTN	up	Q7Z7G0	ABI3BP	up
P07339	CTSD	up	Q9H299	SH3BGRL3	up
P13796	LCP1	up	Q9UFN0	NIPSNAP3A	up
P29350	PTPN6	up	Q9Y3Z3	SAMHD1	up

- +From viewing the proteomic signatures in mini-pigs and human skin, 22 common proteomic regulation changes were associated with ISR-like responses from subcutaneously administered DalcA (Table 1)
- +Boolean scoring based on this signature correctly identified 2/2 ISR-like and 5/6 non-ISR-like findings in human skin with a false-positive rate of ~16% and a false-negative rate of 0%, suggesting a strong biomarker profile for ISR

Ex vivo studies in the HypoSkin ® injectable human skin biopsy platform (Genoskin, Salem, MA, USA) showed that DalcA stimulated mast cell degranulation within 4 hours. In vitro studies further demonstrated that DalcA activates the Mas-related G protein-coupled receptor X2 (MRGPRX2) expressed by mast cells and associated with dermal ISR to a similar level as wild-type recombinant FIX. Since cationic drugs can stimulate MRGPRX2, we examined MRGPRX2 activation of DalcA when presented in alternative formulations.

Figure 1 : Methodology for MRGPRX2 activation screening Table 2: Buffer compositions Formulation Code PS80 (%) pH 0.004 **A1** 0.017 **A2** 0.03 **A3 B1*** 0.004 **B2** 0.017 6.8 0.03 **B3**



+HEK cells transfected to express the G-protein coupled receptor MRGPRX2 were exposed to DalcA drug substance in formulations with varying PS80 (Tween-80) and pH levels (Table 2) +Example fluorescent signal traces for individual transfected (B) or non-transfected (A) cells demonstrate the fluorescent signal response of DalcA administration beginning at 20 seconds

Figure 2 : Buffer composition can influence MRGPRX2 activation in vitro



+Peak signal intensity from MRGPRX2 activation was reduced when stimulated with DalcA in formulation buffers with high tween content (A2 & B3) or higher pH (C1) compared to target formulation (B1) +Peak signal intensity corrected for total duration of response suggests a higher pH formulation can reduce MRGPRX2 activation *in vitro* at all tween contents (C1, C2, and C3) compared to B1 +All low pH or high tween-80 buffers (orange formulations in Table 2) demonstrated diminished stability attributes compared to the target formulation (B1) under accelerated storage conditions

* B1 matches the target specifications of the classical DalcA formulation buffer

C1

C2

C3

0.004

0.017

0.03

6.5

7.1

0.0003 0.0004 0.0018 <0.0001 0.026-< 0.0001 < 0.0001



Formulation buffers with higher pH and/or tween content reduced in vitro MRGPRX2-mediated cell activation, suggesting that optimizing the DalcA formulation buffer may reduce in vivo mast cell degranulation and downstream signaling of pain and dermal inflammation. We examined mast cell degranulation, cytokine secretion, and proteomic changes after SQ DalcA administration in Hyposkin human biopsies for a composite ISR risk profile.







+DalcA in buffer C1 performed similarly to the negative control for most cytokines for 8/10 donors +DalcA formulated in buffer C2 stimulated a different cytokine response compared to both classical Dalca

(B1) and the negative control in 4/8 donors



Conclusions

- +A biomarker signature predictive of minipig ISR was generated from the cross comparison of cutaneous proteomes of minipig and ex vivo human skin models
- +Hyposkin platform enabled evaluation and selection of optimal formulation buffer expected to lower the risk of clinical ISR upon SQ DalcA injection

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+Percent mast cells observed with moderate (A) or high (B) degranulation 4 hours after SQ injection +Relative to DalcA in B1, C1 stimulated fewer highly degranulating mast cells in most donors (insert)

icted ISR	+Consistent with clinical incidence findings, DalcA in B1 was
non-ISR	predicted to elicit an ISR-like response based on proteomic
non-B1	signature changes in 4/10 donors (D)
🗖 B1-ISR	+DalcA in C1 buffer reduced ISR-like findings to 3/10 donors
	+DalcA formulated in C2 buffer increased the incidence to 5/10
	donors, including 3 donors not predicted to experience ISR with
2	DalcA in buffer B1 ("non-B1")
-	

+Formulation buffer composition affects DalcA MRGPRX2-mediated cell activation in vitro