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## **Supplemental Data**

## **Topical Enzyme-Replacement Therapy Restores**

# **Transglutaminase 1 Activity and Corrects Architecture**

### of Transglutaminase-1-Deficient Skin Grafts

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# Figure S1. Recombinant Expression and Purification of Human Full-Length Transglutaminase 1

**A**: Coomassie stained SDS-Gel (10%) and **B**: Western Blot of different expression and purification steps. Lane 1, cell lysate before purification; lanes 2,3, washing steps with 20 mM imidazole; lane 4, washing step with 40 mM imidazole; lanes 5-10 elution steps. Lanes 6-10 in **A** show highly purified rhTG1.



### Figure S2. Intensity Scans to Record Intracellular Localization of Liposomes in *TGM1-/-*Keratinocytes

**A**: Intensity scan along the red line in the equator plane of a *TGM*-/- undifferentiated cell and **B**: of a *TGM*-/- differentiated cell. Traces of the fluorescence intensity of the lipopeptide (green), Rhod-labeled lipid (red) and cell-excluded, membrane-bound trypan blue (blue).



#### Figure S3. Immunohistochemical Detection of Differentiation Markers and TG1-Substrates after Treatment with rhTG1-LUV Preparations

In regenerated TG1-deficient grafts as well as in human TG1-deficient skin the investigated differentiation markers show a more diffuse and shifted distribution when compared to normal skin. Staining of filaggrin (A-E), involucrin (F-J), plasminogen activator inhibitor-2 (PAI-2) (K-O) and loricrin (P-T) in mice treated with rhTG1-LUV formulations reveal a normalization of the distribution patterns when compared to untreated or retinoid treated mice.

	MLVs <sup>*</sup>	LUVs <sup>#</sup>	LUVs <sup>#</sup>	SUVs <sup>¶</sup>
Diameter [nm] (Dynamic light scattering)	> 1500 nm	200 nm	100 nm	< 50 nm
Inner volume (calculated) (V=1 ml, 10 mM lipid)	> 200 µl	~ 180 µl	~ 55 µl	~ 10 µl
Amount of encapsulated protein (Dialysis, HPLC)	+++	++	++	+
Stability (Dynamic light scattering)	+	+++	+++	++
Coupling of lipopeptide vector (Fluorescence-spectroscopy)	+++	+++	+++	+++
Polydispersity-Index/Uniformity (Dynamic light scattering)	> 1.0	0.108	0.075	0.548

### Table S1. Characterisation of Liposomal Preparations of Different Diameters

\*MLVs: <u>m</u>ultilamellar <u>l</u>arge <u>v</u>esicles; <sup>#</sup>LUVs: <u>l</u>arge <u>u</u>nilamellar <u>v</u>esicles; <sup>¶</sup>SUVs: <u>s</u>mall <u>u</u>nilamellar <u>v</u>esicles; +++ high; ++ moderate; + low.