Abstract

**Background:** Recessive dystrophic epidermolysis bullosa (RDEB) and junctional EB (JEB) are inherited disorders characterised by fragility and blistering of epithelial tissues leading to pain, pruritus, and adherent scarring. The severity and chronic nature of the resultant skin wounds significantly reduces quality and length of life. Current therapies primarily consist of protective bandaging and nutritional supplementation; there is no cure for these disorders. Although the skin fragility results from a lack of C7 protein (RDEB) and laminin-332 (JEB), other serious aspects of these disorders, such as inflammation that interferes with healing and aggressive squamous cell carcinoma, have not been completely elucidated. Recent research has suggested that mitochondrial function plays a significant role in skin healing.

**Objective:** To evaluate how mitochondrial function differs in patients with RDEB and JEB.

**Method:** The energy status of RDEB and JEB patient-derived fibroblasts was determined by Seahorse analysis and metabolite production. The energetics and overall morphology of RDEB and JEB patient-derived fibroblasts were assayed as a measure of metabolic stress.

**Results:** EB patient-derived fibroblasts showed impaired oxidative phosphorylation with concomitant compensation by glycolysis. Morphological parameters were altered in RDEB and JEB fibroblasts compared with controls.

**Conclusion:** This is the first study to describe changes in mitochondrial energy metabolism, metabolic profile, and mitochondrial morphology of EB patients.
INTRODUCTION

Epidermolysis bullosa (EB) is a clinically and genetically heterogeneous group of rare inherited disorders, characterised by fragility of epithelial tissues with blistering following minimal mechanical trauma. Four major subtypes of EB have been described, each defined by a distinct plane of epidermal-dermal separation: EB simplex, junctional EB (JEB), dystrophic EB, and Kindler syndrome. The most severe forms of EB are recessive dystrophic EB (RDEB) and JEB, which currently have no curative treatment. Blister cleavage presents immediately below the lamina densa in RDEB and at the lamina lucida in JEB. The typical phenotype of these two forms involves mutilating mucocutaneous blistering, chronic cutaneous infection, and aggressive squamous cell carcinoma.

The molecular pathology of RDEB results from biallelic loss-of-function mutations in the Type VII collagen gene (COL7A1), resulting in a lack of an extracellular matrix molecule critical for skin integrity. Absent or deficient production of functional Type VII collagen (C7) protein leads to the loss of C7 homotrimer anchoring fibrils that interact with dermal and epidermal proteins to connect the skin basement membrane to the papillary dermis. One of the most severe forms of JEB is caused by loss-of-function mutations in any of the three genes (LAMA3, LAMB3, or LAMC2) composing the heterotrimeric protein laminin-332. This protein interacts with C7 at the basement membrane to attach the epidermis to the underlying layers through the integrin receptors α3β1 and α6β4.

Although genetic mutations have been identified as the origin of these diseases, the complex EB phenotype cannot be completely explained based on the adhesive functions of C7 and laminin-332. For example, the mechanisms responsible for the clinical features of EB, such as photosensitivity and cancer risks, are still poorly understood.

Several pathognomonic features not directly related to skin fragility, such as ageing, inflammation, and squamous cell carcinoma, have been strongly associated with mitochondrial dysfunction and oxidative stress. In addition, there is emerging evidence that mitochondria play an important role in skin physiology and pathophysiology. Recent research has suggested mitochondrial function promotes keratinocyte and melanocyte differentiation and pigmentation, as well as epidermal progenitor stem cell function.

Mitochondria are double-membrane enclosed organelles that generate the majority of cellular energy through the citric acid cycle and oxidative phosphorylation (OXPHOS). In addition to energy production in the form of ATP, mitochondria are involved in several metabolic pathways including fatty acid β-oxidation, the formation of iron/sulphate clusters, and haem biosynthesis. They participate in the metabolic processes of lipogenesis, gluconeogenesis, ketogenesis, steroid hormone synthesis, and ammonium detoxification. Mitochondria regulate calcium homeostasis by buffering calcium flux for the plasma membrane, endoplasmic reticulum, and apoptotic pathways.

Mitochondria are highly complex and dynamic organelles that can change in number and morphology within a cell during development, cell cycle, or when challenged by various endogenous or exogenous conditions. Defective mitochondrial architecture-function relationships are linked to many human diseases, including metabolic disorders, neurodegenerative disease, ageing, and cancer.

For these reasons, the work presented here focusses on defining the morphologic, bioenergetic, and metabolic characteristics of RDEB and JEB fibroblasts when compared to unaffected fibroblasts. This study showed altered mitochondrial morphology in both forms of EB, and its functional correlation in impaired OXPHOS and subsequent insufficient glycolytic compensation in RDEB and JEB patient-derived fibroblasts.

MATERIALS AND METHODS

Cell Line Derivation

Samples were obtained after receiving written informed consent, as approved by the University of Minnesota Institutional Review Board and in adherence with the Declaration of Helsinki. RDEB and JEB patients were screened for disease-causing mutations as part of the University of Minnesota Epidermolysis Bullosa Center standard.
of care. Primary fibroblasts were obtained from a skin punch biopsy of patients with RDEB (six patients, ranging in age from 1 to 14 years), JEB (five patients, ranging in age from 1 month to 11 years), or unaffected matched donors (two donors aged 9 and 13 years), and derived by mincing the skin tissue that had been immobilised under a sterile coverslip. Biopsies were taken from skin near blistered regions. Cells were grown in Dulbecco’s modified Eagle medium (Thermo Fisher Scientific Inc., Waltham, Massachusetts, USA), supplemented with 15% fetal bovine serum (Thermo Fisher Scientific Inc.), and 0.1 µg/mL each with penicillin and streptomycin (Thermo Fisher Scientific Inc.) at 37 °C in a humidified 5% CO₂ incubator. The culture medium was changed every third day; all experiments were conducted between passage three and passage 10.

Oxygen Consumption Rate Measurement

Oxygen consumption rate (OCR) was measured in primary fibroblasts with the XF24e Extracellular Flux Analyzer (Seahorse Bioscience, North Billerica, Massachusetts, USA). Cells were seeded at a density of 35×10³ cells per well in 24-well plates and cultured overnight at 37 °C in a humidified, 5% CO₂ incubator. For the bioenergetics profile, the fibroblasts were consecutively treated with oligomycin A (1.0 µM), carbonyl cyanide-4-(trifluoromethoxy)phenylhydrazone (FCCP) (0.7 µM), and rotenone plus antimycin A (1.5 µM). Data were normalised by cell number using the CyQuant® NF Cell Proliferation Assay kit (Thermo Fisher Scientific Inc.). Fluorescence was measured on the SpectraMax® M2 (Molecular Devices, San Jose, California, USA) with an excitation wavelength of 485 nm and an emission detection wavelength of 530 nm.

ATP/ADP Assay

Cells were seeded at a density of 1×10⁵ cells per well in 96-well plates. Following 24-hour incubation, the EnzyLight™ ATP/ADP ratio assay was performed (BioAssay Systems, Hayward, California, USA) according to the manufacturer’s protocol. The assay was normalised to the total number of cells using the CyQuant NF Cell Proliferation Assay.

Confocal Microscopy

For all experiments, cells were plated at 2.5×10⁵ cells per 6 cm dish (Thermo Fisher Scientific Inc.) and images were obtained after 48 hours on a Nikon FNI upright microscope equipped with an AIR scan head and Plan Apo LWD 25× water-immersion objective lens NA 1.1. Images of different fluorophores were acquired sequentially using the two settings: NADH 405 nm excitation, 425–475 nm emission; and flavin adenine dinucleotide (FAD) 488 nm excitation, 500–550 nm emission. The 12-bit 2048×2048-pixel images were acquired, and final magnification was adjusted by zooming with the laser to attain appropriate pixel size.

Electron Microscopy

Normal, RDEB, and JEB fibroblasts were collected using Trypsin-EDTA (Thermo Fisher Scientific Inc.) and pelleted. Cell pellets were sent to the University of Minnesota University Imaging Core for processing and imaging. The work was performed using a Philips CM12 Transmission Electron Microscope. Images were taken with the SIA L3C Digital Camera.

Live Cell Imaging

All images were obtained after 48 hours on a Nikon FNI upright microscope equipped with an AIR scan head and Plan Apo LWD 25× water-immersion objective lens with NA 1.1 in the chamber at a stable temperature of 37 °C. The cells were stained with MitoTracker® Deep Red FM (Thermo Fisher Scientific Inc.), 640 nm excitation and emission 650–720 nm. Z stacking was performed. 12-bit, 2048×2048-pixel images were acquired, and the final magnification was adjusted by zooming with the laser to attain appropriate pixel size.

Image Analysis

For all experiments testing the optical redox ratio, cells were plated at 2×10⁵ per 60 mm glass-bottomed dish (MatTek Corporation, Ashland, Massachusetts, USA). Cell images were obtained approximately 48 hours later using a Nikon FNI upright microscope (UMN Imaging Center). Images of different fluorophores were acquired sequentially using two settings: NADH 405 nm excitation, 425–475 nm emission; and FAD 488 nm excitation, 500–550 nm emission. The
pinhole, gain, and offset remained the same for every experiment. For each imaging session, there were two plates of each cell line and two fields of view for both NADH and FAD. Image acquisition took approximately 16 seconds (Z stack). Following data collection, the NADH/FAD ratio (a measure of the reduction-oxidation ratio) was calculated for every cell in each image. For each acquired NADH and FAD image, ImageJ software (Research Service Brand, National Institute of Mental Health, Bethesda, Maryland, USA) was used to obtain the integrated intensity of NADH and FAD for each cell in the image after the background fluorescence was subtracted. Box fractal dimension was calculated from two-dimensional mitochondria images, converted to binary images, then run in built-in box fractal dimension in ImageJ. Box sizes of 2, 3, 4, 6, 8, 12, 16, 32, and 64 pixels were used. For analysis of mitochondrial volume/cell volume ratio, pictures were first deconvoluted using AutoQuant X3 (Media Cybernetics, Rockville, Maryland, USA), then analysed by Imaris 7.7.0 software (Bitplane, Concord, Massachusetts, USA).22

L-lactate Assay

Cells were seeded at a density of $1 \times 10^3$ cells per well in 96-well plates. After 48 hours incubation, the L-lactate assay (ScienCell Research Laboratories, Carlsbad, California, USA) was performed according to the manufacturer’s protocol. The assay was normalised to cell number using the CyQuant NF Cell Proliferation Assay.

Statistical Analysis

Results are expressed as mean ± standard error of the mean (SEM). The significance of the difference between control and experimental conditions was analysed by unpaired Student’s t-test; p<0.05 was considered statistically significant.

RESULTS

Morphologic Changes of Mitochondria in Epidermolysis Bullosa Fibroblasts

Mitochondrial structure and function are intertwined, and the alteration of one often impacts the other. Transmission electron microscopy was used to image mitochondria in normal, RDEB, and JEB fibroblasts between five and eight matched passages after isolation (Figure 1A-C). Box fractal dimension was used to identify structural changes of the mitochondria in RDEB and JEB fibroblasts compared with control cells.22-24 Box fractal dimension is a measure of the texture of an object from two-dimensional images derived from a Z stack; it reflects a character of the outer mitochondrial membrane structure whereby a grainier structure correlates to higher fractal dimension. Box fractal dimension increased significantly: 25% in RDEB and 9% in JEB fibroblasts compared with non-EB cells (Figure 1D). These observations are consistent with the moderately swollen and more sparsely packed cristae in mitochondria from both EB groups compared with mitochondria of control fibroblasts (mean ± SEM). Live cell imaging using MitoTracker suggests a more fragmented mitochondrial network with less overall mitochondrial volume in RDEB and JEB fibroblasts compared to controls (Figure 2).

Oxidative Phosphorylation Is Impaired in Epidermolysis Bullosa Fibroblasts

To examine and quantify different components of mitochondrial function in RDEB and JEB fibroblasts during Seahorse experiments, the cells were analysed in three passages (passages six to nine). After passage 10, an overall decline in respiratory function of all cell types was observed. This may be attributed to telomere-dependent senescence.25 OCR was measured first in a basal state, then after addition of the ATP synthase (complex V) inhibitor oligomycin, the proton ionophore FCCP, and the respiratory complex I and III inhibitors rotenone and antimycin. Basal respiration was significantly decreased both in RDEB and JEB fibroblasts compared to unaffected fibroblasts derived from healthy individuals (Figure 1E). Respiration measured after treatment with oligomycin reflects decreased OCR to the extent cells are using mitochondria to generate ATP. The remaining OCR is attributed to residual electron transport and oxygen consumption independent of oxidative phosphorylation. Maximal oxygen consumption, measured after addition of FCCP (which collapses the mitochondrial membrane potential), was decreased in JEB fibroblasts (Figure 1F).
Figure 1: Mitochondria morphology and bioenergetic profile of recessive dystrophic epidermolysis bullosa and junctional epidermolysis bullosa skin fibroblasts.

TEM of mitochondria in A) normal; B) RDEB; and C) JEB fibroblasts. D) Mitochondrial morphologic changes in RDEB and JEB fibroblasts quantified using box fraction dimension. A total of 30 mitochondria were assayed for each of two normal control, six RDEB, and five JEB patient fibroblast lines (SEM error bars). E) Seahorse X24 measurement of OCR of RDEB and JEB fibroblast lines upon sequential addition of oligomycin (1.0 µM), FCCP (0.7 µM), and rotenone and antimycin (1.0 µM each). F) Seahorse X24 measurement of ECAR of normal, RDEB, and JEB fibroblast lines upon sequential addition of oligomycin (1.0 µM), FCCP (0.7 µM), and rotenone and antimycin (1.0 µM each). G) ADP/ATP ratio measured following 24-hour incubation.

Values represent the means over three passages of fibroblasts from each of two normal control, three RDEB, and three JEB fibroblast lines ±SEM.

The age ranges for the cell lines: normal (9–13 years), RDEB (1–6 years), and JEB (2–10 months).

Statistical significance was calculated with the Student’s t-test: *p<0.05; *** p<0.001.

ECAR: extracellular acidification rate; FCCP: carbonyl cyanide-4-(trifluoromethoxy) phenylhydrazone; JEB: junctional epidermolysis bullosa; OCR: oxygen consumption rate; RDEB: recessive dystrophic epidermolysis bullosa; TEM: transmission electron microscopy.
Lastly, rotenone and antimycin were injected to inhibit electron flux through complexes I and III of the electron transport chain, causing dramatic suppression of the OCR. The remaining OCR, attributable to O$_2$ consumption in non-mitochondrial organelles, was also significantly decreased in JEB fibroblasts. Moreover, the cellular energy ratio (cytosolic ATP/ADP) of RDEB and JEB fibroblasts was decreased (Figure 1G). These lower cytosolic ATP/ADP ratios determined the cell metabolism to be predominantly glycolytic.

**Epidermolysis Bullosa Fibroblasts Compensate Impaired Oxidative Phosphorylation by Glycolysis**

The normalised optical redox ratio was used to further estimate the metabolic profile of EB fibroblasts. NADH and FAD are the primary electron donor and acceptor of electron transport chain complexes I and II, respectively. Interestingly, mRNA expression profiling of RDEB fibroblasts has found significant differences in the expression of genes involved with nicotinamide metabolism compared to control fibroblasts.

The oxidation-reduction ratio is the most common optical method for measuring cell redox state, as determined by the fluorescence intensity of FAD and NADH (FAD/[FAD+NADH]). The redox ratio is strongly associated with NAD$^+$/NADH concentration and has been used in vitro and in vivo to track metabolic changes during cell differentiation and malignant transformation.

In the case of impaired OXPHOS in RDEB and JEB fibroblasts, the authors hypothesised these cells would have a significant difference in redox state compared to unaffected fibroblasts. To test this, confocal microscopy was used to examine a panel of six RDEB, five JEB, and two non-EB fibroblast lines between five and eight matched passages after isolation (Figure 3A). The optical redox ratio of RDEB and JEB fibroblasts was found to be significantly lower than the optical redox ratio of unaffected control cells (Figure 3B), consistent with abnormal respiration in these cells. To determine if glycolytic metabolism is augmented in EB cells with impaired respiration, extracellular acidification rates (ECAR) of RDEB, JEB, and control fibroblasts were measured, and it was observed that ECAR of JEB fibroblasts was significantly higher during basal respiration, but not in RDEB fibroblasts (Figure 1F). ECAR was unchanged in RDEB and JEB fibroblasts relative to normal fibroblasts during maximal and non-mitochondrial respiration. These results could indicate RDEB and JEB fibroblasts are not able to compensate for impaired OXPHOS by increasing glycolysis, although modestly elevated concentrations of L-lactate in the culture media in RDEB and JEB fibroblasts, compared to control fibroblasts, may indicate increased reliance on glycolysis for ATP production under normal oxygen conditions in these cells (Figure 3C).
DISCUSSION

New data are emerging that mitochondria have critical functions in skin physiology.\(^6\) RDEB and JEB both involve massive trauma to the skin caused by the loss of structural integrity of the basement membrane. However, the effects on bioenergetic and metabolic status of these conditions have not been studied at the cellular level. Using measurements for OCR, ATP/ADP energy ratio, and mitochondrial morphology, this study has identified a number of important variables in fibroblasts derived from RDEB or JEB patients. Critically, OXPHOS is reduced in RDEB and JEB fibroblasts. The lower basal respiration of EB fibroblasts readily indicates a lower energy output from the electron transport chain. This can reduce the amount of energy available to cells and can be detrimental for cells that carry a large metabolic load, such as fibroblasts involved in the perpetual wound healing found in patients with EB. Paired with this decreased metabolic capability appears to be an increase in the complexity of the mitochondria in EB fibroblasts, which is symptomatic of mitochondrial stress. These overall results are indicative of cell autonomous effects in RDEB and JEB fibroblasts independent of the patient wound status or extracellular environment. Future study is necessary to determine the extent

Figure 3: Metabolic profile of recessive dystrophic epidermolysis bullosa and junctional epidermolysis bullosa skin fibroblasts.

A) Representative images of NADH and FAD from RDEB and JEB fibroblasts. Redox ratio images (FAD/(NADH+FAD)) corresponding to representative NADH and FAD images (bars: 10 µM). B) Normalised optical redox ratio (FAD/(NADH+FAD)). Values represent the means over three passages of fibroblasts from each of two controls, six RDEB, and five JEB patients ±SEM. C) L-lactate production of RDEB and JEB fibroblasts. Values represent the average over three passages from each of two normal control, six RDEB, and five JEB patient cell lines ±SEM.

The age ranges for the cell lines: normal (9–13 years), RDEB (1–14 years), and JEB (2 months to 11 years).

Statistical significance was calculated with the Student’s t-test: *p<0.05; ***p<0.001.

FAD: flavin adenine dinucleotide; JEB: junctional epidermolysis bullosa; RDEB: recessive dystrophic epidermolysis bullosa.
to which these cell autonomous effects are impacted by the environmental cues present in EB skin, such as oxidative imbalance and reduced antioxidant enzymes.40

CONCLUSION

EB cells responded to reduced mitochondrial OXPHOS energy production by modestly increasing anaerobic energy production through glycolysis. This was observed through increased production of L-lactate and a reduced redox ratio, while ECAR was only significantly higher in JEB fibroblasts under basal respiratory conditions. This is significant as the cells rely upon glycolysis for energy needs, but still exhibit an energy deficit at the level of ATP availability. The reliance of EB fibroblasts upon glycolysis in a reduced energy state may contribute to the observed range of EB phenotypes. Cells that are better able to adapt or overcome changes in bioenergetic and metabolic states may lead to uneven clinical presentation among patients, or even in different areas of the same patient. One of the treatments for EB patients is bone marrow transplantation.41 Transplant outcomes have been shown to be affected by patient and donor mitochondrial haplotype.42 Haplotype differences in mitochondrial function and stress response may prove significant as this work has demonstrated the first evidence of altered function and structure of mitochondria in fibroblasts derived from RDEB and JEB patients. EB at its core is more than an extracellular matrix or wounding disorder. This work suggests that systemic response to injury may decide the outcome of cellular and subcellular defects, in turn affecting the severity and treatment response in EB patients. Skin-directed mitochondrial research opens a novel avenue toward the development of EB treatments.

References