Supplementary Appendix

This appendix has been provided by the authors to give readers additional information about their work.

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SUPPLEMENTARY APPENDIX

Integrin α 3 Mutations with Kidney, Lung and Skin Disease

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SUPPLEMENTARY METHODS

Genetic Analysis

Mutation analysis of ITGA3 was performed in Patient 1 based on a candidate gene approach, as described in Results. Thereafter, 10 patients with unclear forms of EB and eight families with focal segmental glomerulosclerosis and homozygosity at the *ITGA3* locus¹ were screened for mutations. Genomic DNA was extracted from peripheral-blood leukocytes using QIAmp® DNA mini kit (QIAGEN, Hilden, Germany). The coding exons and the exon / intron boundaries of the gene for integrin α 3 (*ITGA3*) were amplified by PCR using primers designed with Primer3 Software (version 0.4.0) (Supplementary Table 3). All PCR products were submitted to automated nucleotide sequencing in an ABI 3130XL genetic analyzer using Big Dye Terminator Chemistry (Applied Biosystems, Darmstadt, Germany). DNA sequences were compared to the reference sequence from NCBI Entrez Nucleotide database (NC 000017.10) using Mutation Surveyor[™]DNA variant analysis software (version 2.61 Softgenetics, State College, PA, USA). After written informed consent, EDTA-blood for DNA extraction was obtained from ethnically matched controls. Because the mutation found in Patient 1 abrogates the restriction site of Sac I, restriction length polymorphism was used for screening of 100 Southern Italian control chromosomes. The mutations found in Patient 2 and 3 were screened by direct sequencing of PCR products, and were excluded from 200 Palestinian-Arab and 136 Pakistani control chromosomes, respectively.

Mutation analysis of the *NPHS2* and *WT1*, genes was performed using heteroduplex analysis and direct sequencing as described before.²⁻⁴ *CFTR* mutations were analysed using two commercially available CF-gene mutations kits: INFINITI[®] CFTR-15 Assay (AutoGenomics, Vista, USA), which tests for 15 *CFTR* mutations, and LUMINEX xTAG[®] Cystic Fibrosis 39 kit v2 (Luminex Corporation, Austin, USA), which tests for 39 *CFTR* mutations. *ABCA3* was analysed by PCR and direct sequencing of the entire coding region.⁵

Morphological Analyses of Skin, Kidney and Lung

For histopathological examination with light microscopy, skin, kidney and lung biopsy specimens were embedded in paraffin, and the sections were stained with haematoxylin and eosin (H&E) by standard procedures.

Immunohistochemistry was performed using the AEC (3-amino-9-ethylcarbazole) system (Dako, Hamburg, Germany) and haematoxylin as the counter stain. The primary antibodies used are listed in Supplementary Table 4. Control tissues were from newborns/infants.

Indirect immunofluorescence staining of the skin was performed on 5 μ m cryosections, which were air dried and incubated with primary antibodies overnight at 4°C as described before.⁶ The primary antibodies used are listed in Supplementary Table 4. The mouse monoclonal antibody P1B5 was used to specifically detect human integrin α 3,^{7,8} which consists of 1051 amino acids and has a calculated molecular mass of 116 kDa (observed molecular mass of 130 kDa on SDS-PAGE under reducing conditions). The secondary antibodies were Alexa-488 anti-mouse or anti-rabbit IgG (both Invitrogen, Darmstadt, Germany). Nuclei were stained with DAPI (4',6-diamidino-2-phenylindole) (Millipore, Temecula, CA, USA). The stained sections were observed with a confocal laser scanning microscope (LSM510, Carl Zeiss, Jena, Germany).

For TEM analysis, the biopsies were fixed for at least 2h at room temperature in 3% glutaraldehyde solution in 0.1M cacodylate buffer pH 7.4, cut into pieces of ca. 1mm³, washed in buffer, postfixed for 1 h at 4°C in 1% osmium tetroxide, rinsed in water, dehydrated through graded ethanol solutions, transferred into propylene oxide, and embedded in epoxy resin (glycidether 100). Semithin and ultrathin sections were cut with an ultramicrotome (Reichert Ultracut E). Ultrathin sections were treated with uranyl acetate and lead citrate, and examined with a Zeiss EM 900 electron microscope.

In Vitro Studies of Integrin **a**3 Null Keratinocytes

Primary keratinocytes from control and Patient's 1 skin were isolated and cultivated in keratinocyte growth medium (Invitrogen) as described.⁶ For flow cytometry analysis to detect $\alpha 2$, $\alpha 3$, $\alpha 6$, and $\beta 1$ integrin subunits on the cell surface, $2x10^5$ cells were trypsinized, washed twice with PBS (phosphate buffered saline), and then incubated with the primary antibodies (Supplementary Table 4) for 15 minutes at room temperature. After washing with PBS containing 1% BSA (bovine serum albumin) and 0.05% sodium azide, the cells were incubated with FITC (fluoresceinisothiocyanat) conjugated F(ab')₂ goat anti-mouse of anti rat IgG(H+C) antibody (Immunotech, PN) for further 15 minutes. In a parallel experiment isotype controls were used. Flow cytometry acquisition was performed using the BD FACSCantoTMII (BD Biosciences, Oxford, UK).

SUPPLEMENTARY RESULTS

In the skin of Patient 1, the disorganized basement membrane was associated with cellular abnormalities. In the epidermis, the basal keratinocytes were irregular in shape, rather than polyhedral columnar as in the normal skin (Supplementary Fig. 7B, C). However, the cell-cell contacts were not affected, as shown by β -catenin staining (Supplementary Fig. 7C). Cell proliferation, as assessed by Ki67 staining, was comparable to controls (Supplementary Fig. 8A). There was a mild infiltration of CD3 positive inflammatory cells, but no evidence for CD4, CD8, CD20 or CD68 positive infiltrates (Supplementary Fig. 8B). These findings are very similar to the data derived from the mouse model with epidermis-specific ablation of integrin α 3.⁹

Because of the severe clinical lung involvement in our patients, transmission electron microscopy and immunohistochemistry were employed to uncover more subtle morphological changes. In agreement with the collagen IV staining, the transmission electron microscopy of the lung of Patient 1 revealed reduplications and irregularities of the alveolar basement membrane (Supplementary Fig. 9A). In Patient 2, abnormal lamellar bodies, the secretory organelles of type II alveolar cells involved in surfactant formation were noted (not shown). Immunohistochemistry for pro-surfactant protein C (pro-SPC) on the lung of Patient 1 demonstrated numerous type II alveolar pro-SPC-positive cells (Supplementary Fig. 9B). This is in accordance with the data derived from the mice with lung epithelial cell-specific loss of integrin $\alpha 3$.¹⁰ The anomalies of the basement membrane are consistent with those observed in kidney and skin and presumably directly related to the genetic defect.

SUPPLEMENTARY TABLES

Supplementary Table 1. Patients, *ITGA3* Mutations, Clinical Features and Investigations

Patients		Patient 1	Patient 2	Patient 3
ITGA3 mutations		c.1173_1174del p.Pro392ValfsX2	.1173_1174del c.1538-1G>A Pro392ValfsX2	
Clinical Investig	Origin features ations	Southern Italy	Gaza	Pakistan
Kidney Clinical and laboratory findings		Congenital nephrotic syndrome Peritoneal dialysis	Congenital nephrotic syndrome Peritoneal dialysis	Congenital nephrotic syndrome Peritoneal dialysis
	Histology	Globally atrophic glomeruli, segmental glomerulosclerosis, diffuse interstitial fibrosis and tubular atrophy	Right: enlarged proximal tubules with cyst formation, focal segmental sclerosis; left: renal dysplasia	Focal segmental glomerulosclerosis
	Transmission electron microscopy	NA	Focal mesangial hyperplasia, podocyte effacement	Lamellation of the basement membranes
Lung	Clinical features	Respiratory distress, oxygen-dependent, aspiration pneumonia, recurrent respiratory infections	Respiratory distress, oxygen-dependent	Respiratory distress, oxygen-dependent, aspiration pneumonia, recurrent respiratory infections
	Chest radiograph	Interstitial reticulo- nodular changes	Bilateral infiltrates	Right upper and middle lobe pneumonia
	СТ	Non-specific diffuse distortion of the pulmonary architecture consistent with interstitial lung disease	Diffuse interstitial changes consistent with diffuse lung disease	Diffuse interstitial changes consistent with interstitial lung disease
	Histology	Overinflation and mild- to-moderate simplification of airspaces ¹	Consistent with chronic pneumonitis of infancy probably superimposed on lung growth abnormality and associated with recent thromboembolism ²	Interstitial fibrosis
	Transmission electron microscopy	Reduplicated and irregular alveolar basement membrane	Abnormal lamellar bodies in the alveolar cells	NA

Skin Clinical features Immunomapping Transmission electron microscopy		Blisters, nail dystrophy	Persistent erosions Blisters, nail dystrophy	
		Focal disruption of the dermal-epidermal junction, cleavage within the plane of the basement membrane	NA	Focal disruption of the dermal-epidermal junction
		Thin lamina densa, discontinuous between the hemidesmosomes	NA	NA
Mutations were excluded in the following genes		NPHS2, WT1, ABCA3, frequent CFTR mutations	NPHS1, NPHS2, WT1 (exons 8 and 9)	NPHS2, WT1 (exons 8 and 9)
Other investigations		Laryngoscopy: normal findings	Sweat test: normal	NA
		Bronchoscopy: unremarkable	Bronchoscopy: unremarkable	
		Gastrointestinal tract biopsies: no significant abnormalities of the stomach and oesophagus, but the small intestine showed mild flattening of the villous epithelium	Barium swallowing: unremarkable	
		Cerebral MRI at the age of one month: normal anatomy	Brain ultrasound and CT: normal	

Legend: ¹an open lung biopsy was performed; ²histopathology was performed postmortem; CT, computed tomography; NA, not available; MRI, magnetic resonance imaging.

Birth n	Birth measurements:			
•	Weight (g)	4090 (90 th perce	ntile)	
•	Height (cm)	53.5 (50-75 th percentile)		
•	Head circumference (cm)	34.5 (10-25 th percentile)		
•	Apgar	8/8/8	,	
•	Umbilical arterial pH	7.25 (normal ran	ge 7.20-7.38)	
<u> </u>				
Clinica	al laboratory at day 13 of life befo	re initiation of dial	VSIS:	
BIOOD		value	Normai range	
•		5.8	< 0.6	
•	Urea (mg/dl)	186	7.2-42.0	
•	Albumin (g/dl)	1.6	> 3.5	
•	IgA (mg/dl)	</td <td>39 - 71</td> <td></td>	39 - 71	
•	lgG (mg/dl)	257	280 - 710	
•	IgM (mg/dl)	38	54 - 144	
•	Blood pH	7.21	7.35 - 7.45	
•	Base excess (mmol/l)	-15	-3 + 3	
•	Cholesterol (mg/dl)	387	62 - 189	
•	Triglyceride (mg/dl)	1314	19 - 85	
Urine	parameters			
•	Protein/creatinine ratio (mg/mg) 12.5	< 0.9	
) 12.0	0.0	
GFR ¹	ml/min/1.73 m ²	<10	40 - 60	
Respii	ratory parameters with oxygen su	pplementation:		
•	Venous CO ₂ (kPa)	6.5 - 9.0	5.5 - 6.8	
•	Respiratory rate (/minute)	60 - 120	30 - 60	
Other	features			
•	Round face with prominent for	head hypertelori	sm. antimongoloid slant, promin	ent nose with
	long saddle small mouth retro	anathism large e	ars	ient nobe with
•	Narrow chest gynecomastia			
•	Discrete clinodactyly deen furrows of the soles			
•	Shrill crv			
•	Swallowing and sucking reflexes present although swallowing was repeatedly followed by			
	couch and vomiting			
	 Mild muscular hypotonia 			
	 Microcentaly 			
	 Nicrocephaly Delayed neurological development 			
	Normal nevcho-social interactic	ons and cognitive	development	
•	 Adenoviral enteritis and subsequent marked hepatosplenomegaly at the age of six months 			

Supplementary Table 2. Associated Features and Investigations in Patient 1

Legend: ¹ Glomerular filtration rate (GFR) was estimated at day 13 of life by the Schwartz formula, using serum creatinine, height and a k constant of 0.4.

Exon	Primers	Product size (bp)
1	5'- gagagcgcagctgtgaaact	497
	5'- cttaatcctggctcccaagg	
2	5'- tgcctgccttaccctacact	397
	5'- gaagcgactagaagccgtgt	
3	5'- gctggatgggattggtagag	388
	5'- cacaggaagggacatgtgtg	
4		495
	5'- actogcatgaggtcctgttt	
5	5'- atggcaaaatgctcaccaat	392
-	5'- atcttgggagcttggaacag	
6	5'- gctggccatctggagtctac	492
	5'- ctgcaaacctctgcaaacaa	
7	5'- ggtgggtcatattggcatct	500
	5'- ccccatcctqtctcatctqt	
8	5'- catcaatcaggccaagatca	369
	5'- caqtcccaqcttctctccat	
9	5'- ccccagcaggtacagagaga	387
-	5'- ccaggettccaaggageta	
10	5'- ccatctgtgtccatgtttgc	337
	5'- ttagatagagagagaaatg	
11	5'- gtcagggacaccacagacct	392
	5'- agaagaagccgtggaagaca	
12	5'- agtatacaagtagagcttata	396
	5'- totccccaacctctotctct	
13	5'- ggcgggtccagctcttct	377
-	5'- gcactccttctggaactgga	-
14	5'- caggetetggagaaccacac	367
	5'- ttctgaccatcccacctgtt	
15	5'- caagatgggcttttctcacc	379
	5'- aagaaggcccagaccagag	
16	5'- ccttagcgtctctgctgctt	383
	5'- ctagcagccccatttgtctt	
17+18	5'- aggaggagaaggccagaaag	490
	5'- gatgatccagggaaagtgga	
19	5'- tcacagcagccctgtgatac	490
	5'- gtgggcacaagatgtgtcat	
20	5'- aggaagccccctcaagtatg	479
	5'- gcaggaaaggaaagggtagg	
21	5'- gcagaaaggccagtgtcttc	471
	5'- cctttggtcccttgtcttca	
22	5'- tatccatgtgcctggcatta	482
	5'- atgcaaagacacgcaaatca	
23	5'- aatgatgcgcatttgtgtgt	388
	5'- cttaccgtgtggccaagatt	
24	5'- cgctcctctggagtcaagtc	483
	5'- gctggtggtagaaggtggag	
25	5'- cagagagatgggatgcgttt	427
	5'- gacgaggtgtatgtgcctga	

Supplementary Table 3. Primers Used for *ITGA3* Mutation Analysis

Antigen protein	IF / IHC	FC	Clone	Source / Company
β-catenin	+		polyclonal	Abcam
CD3	+		F7.2.38	DAKO
CD4	+		4B12	DAKO
CD8	+		C8/144B	DAKO
CD20	+		L26	DAKO
CD68	+		KP1	DAKO
Collagen IV	+		C IV 22	Abcam
Collagen VII	+		LH7.2	Calbiochem
Collagen XVII	+		polyclonal	see Ref ¹¹
Fibronectin	+		ab2413	Abcam
Integrin β1		+	4B7R	Abcam
Integriņ β1	+		B3B11	Chemicon
Integrin a3	+	+	P1B5	Chemicon
Integrin α2		+	16B4	Abcam
Integrin α 6	+	+	GoH3	Progen
Integrin β4	+		3E1	Millipore
Keratin 14	+		LL002	DCS Hamburg
Keratin 10	+		LH2	Santa Cruz Biotechnology
Ki67	+		MIB-1	DAKO
Laminin α 3	+		BM165	gift of B. Burgeson, Boston
Laminin β3	+		6F12	Santa Cruz Biotechnology
Laminin y2	+		GB3	gift of G. Meneguzzi, Nice
Laminin α 5	+		So4C7	gift of L. Sorokin, Münster
Pro-surfactant	+		polyclonal	Millipore
protein C				
Tenascin C	+		578	R&D Systems

Supplementary Table 4. Primary Antibodies Used in this Study

Legend: +, used for the application; IF, immunofluorescence; IHC, immunohistochemistry; FC, flow cytometry.

SUPPLEMENTARY FIGURES



Supplementary Figure 1. Chest Radiograph and CT of Patient 2.

Panel A shows chest radiograph at 6 weeks, and panel B shows CT of the chest. Both

demonstrate diffuse interstitial changes with inter- and intralobular septal thickening.





Supplementary Figure 2. Morphologic Features of the Skin of Patient 1. In panel A, semithin sections of a skin specimen of the patient are stained with methylene blue. Light microscopy of non-blistered skin reveals a flattened epidermis and microblisters (left panel, arrows). In blistered skin, subepidermal cleavage is seen (right panel, asterisk). Panels B show transmission electron microscopy of the dermal-epidermal junction in the patient and a control. In control skin, the lamina densa appears as a strong continuous line. In the patient's skin, the lamina densa is very thin and discontinuous, and basal keratinocytes seem to detach from it. The lamina densa is present beneath the hemidesmosomes (black arrows), but interrupted in between (open arrows). The right panel shows occasional hemidesmosomes (arrow) on the blister floor, indicating intracellular rupture. The asterisk denotes the blister cavity.



Supplementary Figure 3. Identification of *ITGA3* Mutations in Patients 2 and 3.

In both panels chromatograms reveal partial sequences of the *ITGA3* gene in controls and patients. In panel A, the homozygous mutation c.1538-1G>A (arrow) found in Patient 2 affects the position -1 of the obligatory acceptor splice site of exon 12. Panel B shows the missense mutation c.1883G>C, p. Arg628Pro (arrow) identified in Patient 3 in a homozygous state.



Supplementary Figure 4. Expression of Integrin α 3 in Kidney and Lung.

Immunohistochemistry of integrin α 3 in kidney (upper panels) and lung (lower panels) of a control subject and Patient 1, demonstrating loss of integrin α 3 expression in the patient's samples.



Supplementary Figure 5. Consequences of the *ITGA3* Deletion Mutation in Keratinocytes of Patient 1. Flow cytometry of keratinocytes from the Patient 1 (red curves), and a control (green curves), with antibodies to integrins $\alpha 3$, $\alpha 2$, $\beta 1$ and $\alpha 6$ (continuous lines), or with secondary antibodies alone as a control (interrupted lines). It demonstrates complete absence of integrin $\alpha 3$ on the surface of keratinocytes derived from the patient, whereas integrins $\alpha 2$, $\beta 1$ and $\alpha 6$ are not significantly altered.



Supplementary Figure 6. Extracellular Matrix Proteins in the Skin of Patient 1.

Immunofluorescence staining is shown. Panel A shows a linear signal of collagen VII at the dermal-epidermal junction in control skin, but irregular distribution in the papillary dermis at the blister (asterisk) base in patient 1. In panels B and C, strong deposition of tenascin C and fibronectin is shown in the skin of the patient. Nuclei are visualized in blue with DAPI.



Supplementary Figure 7. Integrins and Cell-Cell Contacts in the Skin of Patient 1.

Immunofluorescence staining is shown. Panel A shows integrin α 6 in the skin of a control and Patient 1. Loss of integrin α 3 leads to discontinuous and irregular distribution of α 6 integrin (arrow). Panel B shows similar integrin β 1 patterns in control and the patient's skin. In panel C, β -catenin is used to delineate cell membranes. Note the regular, columnar shape of basal keratinocytes in normal skin. In contrast, in the patient's skin the morphology of basal keratinocytes is clearly irregular.



Supplementary Figure 8. Keratinocyte Proliferation and CD3 Positive Cells in the Skin of Patient 1.

Panels A show immunohistochemistry of Ki67 in the skin of an age-matched control and the patient. Note comparable staining patterns, demonstrating that keratinocyte proliferation is not hampered in the absence of integrin α 3. In panel B, immunohistochemistry of CD3 reveals few inflammatory cells in the skin of the patient (arrows), whereas control skin remains negative.



Supplementary Figure 9. Alveolar Basement Membrane and Pro-SPC in the Lung of Patient 1.

Panel A shows an electron micrograph representing the blood-air barrier with a capillary filled with erythrocytes, the basement membrane (BM) and pneumocytes. The lower panel is a 3-

fold magnification of the marked area in the upper panel. Note the irregular thickness (arrows) and splitting of the alveolar basement membrane (asterisk). In panel B, immunohistochemistry demonstrates abundant pro-SPC-positive type II alveolar cells.

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