A novel dermal matrix generated from burned skin as a promising substitute for deep-degree burns therapy

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Abstract. The extensive skin defects induced by severe burns are dangerous and can be fatal. Currently, the most common therapy is tangential excision to remove the necrotic or denatured areas of skin, followed by skin grafting. Xenogeneic dermal substitutes, such as porcine acellular dermal matrix (ADM), are typically used to cover the burn wounds, and may accelerate wound healing. It is assumed that burned skin that still maintains partial biological activity may be recycled to construct an autologous acellular dermal matrix, termed 'deep-degree burned dermal matrix (DDBDM)'. In theory, DDBDM may avoid the histoincompatibility issues associated with foreign or xenogeneic dermal matrices, and reduce therapy costs by making full use of discarded skin. In the present study, the collagens within prepared DDBDM were thickened, disorganized and partially fractured, however, they still maintained their reticular structure and tensile strength (P<0.01). Through microarray analysis of the cytokines present in ADM and DDBDM, it was determined that the DDBDM did not produce excessive levels of harmful burn toxins. Following 4 weeks of subcutaneous implantation, ADM and DDBDM were incompletely degraded and maintained good integrity. No significant inflammatory reaction or rejection were observed, which indicated that ADM and DDBDM have good histocompatibility. Therefore, DDBDM may be a useful material for the treatment of deep-degree burns.

Introduction

First-degree burns and superficial wounds undergo a similar wound repair process involving reconstruction of the epithelial barrier, the underlying dermis and extracellular matrix (ECM), organized by granulation tissue (1). However, second- and third-degree burns present a broad range of additional challenges for restoration of normal dermal function (2,3). The extensive skin defects induced by severe burns are dangerous and can be fatal to patients. A common therapy used to treat extensive burns is skin grafting following tangential excision to remove the necrotic or denatured skin. However, sources of skin transplants are often limited. Xenogeneic dermal substitutes, such as porcine acellular dermal matrix (ADM), can be used to cover the burn wounds, prevent infection, facilitate skin renewal and vascularization, and accelerate wound healing (4-7). However, foreign or xenogeneic tissues have partial histoincompatibility, which can induce rejection and allergic reaction (8,9). Engineered human skin matrices, including Dermagraft, DermACELL and Integra, have exhibited satisfactory effects in the clinic (10-12). However, these skin products are expensive and rare, therefore, their clinical application is restricted. It is important for the clinical therapy of patients with deep second- and third-degree burns to investigate novel dermal substitutes that avoid the limitations associated with currently used dermal matrices.

The necrotic or denatured skin of burn wounds is typically discarded during treatment due to the formation of cutaneous burn toxins (13). Burn toxins, which include free radicals, lipoproteins, lipopolysaccharides and lipid peroxides, are formed as a result of thermal injury to skin and are released into the interstitial fluid and serum during the early stages of burns (14-17). These hazardous substances have been shown to exert various effects, including immunosuppression, mitochondrial destruction, cellular energy metabolic disorders and increasing the permeability of the cell membrane, which in turn have been implicated in the various complications associated with burns, including sepsis and multiple organ dysfunction syndrome (16,18-25). In addition, bacteria can rapidly proliferate in necrotic tissue resulting in wound infection (13). It has been previously observed that sections of the skin removed from deep burn wounds partially maintained the integrity of their collagen structure, and may have the potential to be repaired under the appropriate conditions. Therefore, we hypothesized that burned skin with partial biological activity may be recycled to produce an autologous acellular dermal matrix, termed 'deep-degree burned dermal matrix (DDBDM)', and applied for the treatment of deep burn wounds (26). In theory,

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DDBDM, which originates from human skin, may avoid the histoincompatibility associated with foreign and xenogeneic dermal matrices. Furthermore, DDBDM may reduce therapeutic costs by making full use of the discarded skin. Thus, successful clinical application of DDBDM may be an improved method of treatment for patients with deep burn wounds.

In the current study, DDBDM was successfully prepared using a mouse model of a deep-degree burn wound. To determine if DDBDM and ADM produced burn toxin with different compositions, the protein expression levels of 308 cytokines were analyzed. The DDBDM was subcutaneously implanted in mice to observe whether an inflammatory reaction was induced. The aim of the present study was to investigate the use of DDBDM as a dermal matrix and evaluate its potential clinical significance for the treatment of patients with deep-degree burns.

Materials and methods

Animals and ethics statement. A total of 150 healthy male Balb/c mice (age, 12 weeks; weight, 27-33 g) were obtained from the Animal Center of Shandong University (Jinan, China). The mice were maintained under a 12 h light/dark cycle with *ad libitum* access to animal chow and water in the animal quarter at the Animal Laboratory of the Second Hospital of Shandong University (Jinan, China) at 20-24°C and 50-60% humidity. All experimental procedures were conducted according to the criteria outlined in the Guide for the Care and Use of Laboratory Animals published by the National Institutes of Health (NIH Publication no. 85-23, revised 1996). All experimental protocols were approved by Animal Care and Use Committee of the Second Hospital of Shandong University.

Establishment of burn animal model. A total of 60 healthy male Balb/c mice were used to establish the burn animal model. Following anesthesia by intraperitoneal injection of 10% chloral hydrate (0.3 ml/kg; Qilu Hospital, Jinan, China), the fur on the dorsum of each mouse was shaved. The shaved edges were protected by plastic wrap and a thin foam board. A deep-degree burn wound of 5x4 cm was created by hot water-bath burn, as follows: Using a water bath (Shanghai Jing Hong Experimental Equipment Co. Ltd, Shanghai, China) maintained at a constant temperature of 80°C, the dorsum of each mouse was bathed in the hot water for 8 sec. All the burned mice received anti-shock therapy with Lactated Ringer's solution (Baxter International Inc., Deerfield, IL, USA) by intraperitoneal injection (40 ml/kg) and were treated with 1% povidone iodine solution (Lircon Disinfection Science Technology Inc., Dezhou, China) to protect the wound. The mice were resuscitated in a warm environment by subcutaneous injection of physiological saline solution (10 ml/kg; Baxter International, Inc.) until they were fully conscious. After 72 h, the mice were anesthetized and sacrificed by decapitation, after which burned skin was immediately harvested for further experimentation.

ADM preparation method. Following anesthesia and shaving of the fur on the dorsum of 40 healthy Balb/c mice, normal skin specimens were removed and washed with sterile saline solution (Baxter International Inc.). The subcutaneous tissue was removed leaving skin sections of 0.05-1.00 mm thickness. The skin sections were placed into a mixed solution of 0.25% trypsin (Sigma-Aldrich, St. Louis, MO, USA) and Triton X-100 (Sigma-Aldrich), and shaken 100 times/min for 2 h at 37°C. Samples were repeatedly washed and shaken with phosphate-buffered saline (PBS) until the cells and trypsin/Triton X-100 solution were removed. The ADM was maintained in saline solution with 800 U/ml gentamicin (Shandong Lukang Chenxin Pharmaceuticals Co., Ltd., Jining, China) at 4°C. The whole preparation process was conducted under aseptic conditions.

DDBDM preparation method. Burned mouse skin specimens were obtained from 60 burned mice and washed with sterile saline solution. The subcutaneous tissues were removed leaving skin sections of 0.05-1.00 mm thickness. The skin sections were placed into a mixed solution of 0.25% trypsin (Sigma-Aldrich) and Triton X-100 (Sigma-Aldrich) and shaken 100 times/min for 1 h at 37°C. Samples were repeatedly washed and shaken with PBS until the cells and trypsin/Triton X-100 solution were removed. The DDBDM was maintained in saline solution with 800 U/ml gentamicin at 4°C. The whole preparation process was conducted under aseptic conditions.

Physical evaluation of ADM and DDBDM. At room temperature and in a humid environment, the prepared ADM and DDBDM were trimmed to 1x1-cm sections and measured with a Benchtop Tester (H10K-T; Tinius Olsen Testing Machine Company, Horsham, PA, USA). The samples were stretched at a rate of 1 mm/min until the actual load reached 3 MPa, then released at the same rate until the actual load was 0 MPa. After repeating the process three times, the stress-strain curve became stable. The strain value at 3 MPa in the fourth stretch was calculated using the following formula: Strain (%) = $[(L - L_0)/L_0] \times 100 = (L - 1) \times 100$, where L was the length of the dermal matrix during stress in cm and L₀ was the initial length of the dermal matrix, which was 1 cm in the present study. Subsequently, new samples were stretched at a rate of 1 mm/min until broken, recording the values of ultimate tensile strength, maximum tension and elongation at break. All the data were recorded and calculated using a JBK Measure-Control System (Beijing Jincun Electromechanical Technology Research Institute, Beijing, China).

Biotinylated antibody-based cytokine microarray assay. Total protein was extracted from skin samples using RIPA buffer (Beyotime Institute of Biotechnology, Haimen, China), after which the supernatants were obtained by centrifugation at 12,000 x g for 5 min at 4°C. The protein concentrations of the supernatants were detected using a BCA kit (Thermo Fisher Scientific Inc., Waltham, MA, USA), according to the manufacturer's protocol. A total of 308 different mouse proteins, including cytokines, chemokines, adipokines, growth factors, angiogenic factors, proteases, soluble receptors and soluble adhesion molecules, were detected using a mouse cytokine array kit (AAM-BLM-1-4; RayBiotech, Inc., Norcross, GA, USA), according to the manufacturer's protocol. Following subtraction of local background signals, the microarray signals were recorded using a GenePix 4000B microarray scanner (Molecular Devices, LLC, Sunnyvale, CA, USA) using the Cy3 channel and with a scanning

Dermal matrix	Ultimate tensile strength, MPa	Maximum tension, N	Elongation at break, %	Strain at stress of 3 MPa, %
Acellular dermal matrix	15.0±2.1	84±4	200±7	60.1±2.7
Deep-degree burned dermal matrix	10.2±1.8	57±3	182±5	41.6±1.4
P-value	<0.05	<0.01	< 0.05	< 0.01

Table I.	Physical	properties	of acellular	and deep-degr	ee burned dern	nal matrix.
				• • • • • • • • • • • • • • • • • • • •		

ADM

DDBDM



Figure 1. ADM and DDBDM under general observation. ADM, acellular dermal matrix; DDBDM, deep-degree burned dermal matrix.

wavelength of 532 nm. The positive control value was the mean fluorescent signal intensity minus the background of all the positive control spots. Following background subtraction, negative signal intensities were assigned a value of 1. If the value of all the samples tested was 1, those cytokines were removed from further analysis. The mean and the standard error of each cytokine were calculated separately.

Histological staining. The skin specimens were embedded in paraffin blocks after they had been fixed in 4% paraformaldehyde solution (Beyotime Institute of Biotechnology). Sections of $5 \,\mu$ m were deparaffinized and stained using the hematoxylin and eosin (H&E; Beyotime Institute of Biotechnology) staining method. The skin specimens were examined and evaluated in a blinded manner under a standard light microscope (CX31; Olympus Corporation, Tokyo, Japan).

DAPI fluorescence staining. Sections (5-µm thickness) were obtained, deparaffinized and stained with DAPI solution (Beyotime Institute of Biotechnology) at room temperature for 3-5 min, washed three times using PBS and observed with a fluorescence microscope.

Subcutaneous implantation of skin matrix. ADM and DDBDM were trimmed to 1x1-cm sections and implanted subcutaneously into 48 mice. Briefly, following anesthetization by intraperitoneal injection with 10% chloral hydrate (0.3 ml/kg),

the fur on the dorsum of the mice was shaved, a vertical incision was made in the middle of the dorsum and the skin was separated from the subcutaneous tissue by a blunt dissection in order to create two 15 mm-deep subcutaneous bursae. Subsequently, the ADM was implanted in the left bursa and the DDBDM in the right. The incision was closed and the gap in the subcutaneous tissue was sutured to prevent the ADM and DDBDM from touching. The wound was disinfected using 1% povidone iodine solution every day. At 1, 2, 3 and 4 weeks of subcutaneous implantation, three mice were randomly selected for sacrifice by decapitation, prior to immersion in 75% alcohol for 3 min. Subsequently, the dorsal skins of the mice were cut open and separated from the subcutaneous tissue in order to expose the implanted skin matrices, which were then stained using H&E and DAPI. Images of the skin matrices were captured under a microscope (BX53; Olympus Corporation).

Statistical analysis. All data are presented as the mean \pm standard deviation. Dual comparisons between groups were evaluated with the Student's t-test, using the SPSS software, version 19.0 (IBM SPSS, Armonk, NY, USA). P<0.05 was considered to indicate a statistically significant difference.

Results

DDBDM preparation and its biological properties. Images of ADM and DDBDM prior to implantation are presented in

	Normal	Burned		Fold				Fold
Cytokine	skin ^a	skin ^a	P-value	change	ADM ^a	DDBDM ^a	P-value	change
Activin βC	393.5±61.5	1.0±0.0	< 0.01	393.500	76.5±0.5	34.0±3.0	<0.01	2.250
6Ckine	193.5±18.5	1.0±0.0	<0.01	193.500	49.0±12.0	5.5±0.5	< 0.01	8.909
Decorin	49.0±11.0	1.0±0.0	< 0.01	49.000	913.5±76.5	180.0±6.0	< 0.01	5.075
IL-10	1,460.0±94.0	336.0±8.0	<0.01	4.345	221.0±7.0	98.0±1.0	< 0.01	2.255
IL-17	1,057.5±46.5	268.5±60.5	< 0.01	3.939	180.0±8.0	107.5±3.5	< 0.01	1.674
MMP-14	929.0±37.0	372.0±15.0	< 0.01	2.497	162.0±2.0	81.0±7.0	< 0.01	2.000
HVEM	1,193.5±118.5	499.0±212.0	< 0.01	2.392	219.5±1.5	68.5±1.5	< 0.01	3.204
TNF-α	1,327.0±32.0	637.5±18.5	< 0.01	2.082	184.0±17.0	107.0±4.0	< 0.01	1.720
IL-3	2,519.0±33.0	1,356.5±56.5	< 0.01	1.857	303.0±15.0	145.5±3.5	< 0.01	2.082
TGF-β1	1,205.5±111.5	688.5±39.5	< 0.01	1.751	212.0±22.0	115.5±4.5	< 0.01	1.835
RELM β	139.0±3.0	562.0±28.0	< 0.01	0.247	41.5±0.5	75.0±1.0	< 0.01	0.553
IL-9	148.5 ± 60.5	921.5±267.5	< 0.01	0.161	65.5±9.5	258.0±32.0	< 0.01	0.254
IL-23 R	12.8±13.2	157.5±46.5	< 0.01	0.081	2.5±2.2	14.5±4.5	< 0.05	0.172
Thrombospondin	20.5±26.0	907.0±46.0	<0.01	0.023	25.5±14.5	65.5±9.5	< 0.05	0.389

Table II. Expression values of cytokines with significant differences between burned and normal skin samples.

^aValues are presented as mean \pm standard deviation. ADM, acellular dermal matrix; DDBDM, deep-degree burned dermal matrix; 6Ckine, Chemokine (C-C motif) ligand 21; IL, interleukin; MMP, matrix matelloproteinase; HVEM, herpesvirus entry mediator; TNF- α , tumor necrosis factor- α ; TGF- β 1, transforming growth factor- β 1; RELM β , resistin-like molecule β ; R, receptor.



Figure 2. Hematoxylin and eosin and DAPI staining of normal skin, burned skin, ADM and DDBDM sections at various magnifications. ADM, acellular dermal matrix; DDBDM, deep-degree burned dermal matrix.

A		Normal skin Burned s			ned ski	in ADM			DDBDM						
						1995									
B		1	2	3	4	5	6	7	8	9	10	11	12	13	14
	1	Positive 1	Positive 1	Positive 2	Positive 2	Positive 3	Positive 3	neg	neg	6Ckine	6Ckine	Activin A	Activin A	Activin C	Activin C
	2	Artemin	Artemin	Axl	Axl	b FGF	b FGF	CD80	CD80	BAFF R	BAFF R	BCMA	BCMA	β-Catenin	β-Catenin
	3	CCL8	CCL8	CCR10	CCR10	CCR3	CCR3	CCR4	CCR4	CCR6	CCR6	CCR7	CCR7	CCR9	CCR9
	4	CD40	CD40	CD40 Ligand	CD40 Ligand	Cerberus 1	Cerberus 1	Chordin- Like 2	Chordin- Like 2	Tissue Factor	Tissue Factor	IL-2 Ry	IL-2 Rγ	CRG-2	CRG-2
	5	CXCL16	CXCL16	CXCR2 / IL-8 RB	CXCR2 / IL-8 RB	CXCR3	CXCR3	CXCR4	CXCR4	CXCR6	CXCR6	DAN	DAN	Decorin	Decorin
	6	EGF R	EGF R	EG- VEGF	EG- VEGE	Endocan	Endocan	Endoglin	Endoglin	Endostatin	Endostatin	Eotaxin	Eotaxin	Eotaxin-2	Eotaxin-2
	7	Fas	Fas	CD32b	CD32b	FGF R3	FGF R3	FGF R4	FGF R4	FGF R5β	FGF R5β	FGF-21	FGF-21	Fit-3	Fit-3
	8	G-CSF	G-CSF	GDF-1	GDF-1	GDF-3	GDF-3	GDF-5	GDF-5	GDF-8	GDF-8	GDF-9	GDF-9	GFR α-2	GFR α-2
	9	Granzyme D	Granzyme D	Granzyme G	Granzyme G	Gremlin	Gremlin	Growth Hormone R	Growth Hormone R	HGF R	HGF R	HGF	HGF	TNFRSF 14	TNFRSF 14
	10	IFN-γ	IFN-γ	IFN-γ R1	IFN-y R1	IGFBP-1	IGFBP-1	IGFBP-2	IGFBP-2	IGFBP-3	IGFBP-3	IGFBP-5	IGFBP-5	IGFBP-6	IGFBP-6
	11	IL-1 R9	IL-1 R9	IL-1 R I	IL-1 R I	IL-1 R II	IL-1 R II	IL-2	IL-2	IL-2 Rα	IL-2 Rα	IL-2 Rβ	IL-2 Rβ	IL-3	IL-3
С		15	16	17	18	19	20	21	22	23	24	25	26	27	28
	1	Activin RIB	Activin RIB	Acrp30	Acrp30	AgRP	AgRP	ALCAM	ALCAM	Angiopoietin- like 2	Angiopoietin- like 2	Angiopoietin- like 3	Angiopoietin- like 3	AR	AR
	2	BLC	BLC	втс	BTC	Cardiotro	Cardiotro	CCL1	CCL1	CCL28	CCL28	CCL4	CCL4	CCL7	CCL7
	3	CD11b	CD11b	CD14	CD14	CRP	CRP	CD27	CD27	CD27	CD27	CD30	CD30	CD30 L	CD30 L
	4	Cripto	Cripto	Crossvein	Crossvein	Cryptic	Cryptic	Csk	Csk	CTACK	CTACK	CTLA-4	CTLA-4	CXCL14	CXCL14
	5	DKK-1	DKK-1	DKK-3	DKK-3	DKK-4	DKK-4	DPPIV	DPPIV	DR3	DR3	Dtk	Dtk	EDAR	EDAR
	6	Epigen	Epigen	Epireg-	Epireg-	EPO	EPO	E-Selectin	E-Selectin	FADD	FADD	FAM3B	FAM3B	Fas	Fas
	7	FLRG	FLRG	Follistatin	Follistatin	Fractal-	Fractal-	Frizzled-1	Frizzled-1	Frizzled-6	Frizzled-6	Frizzled-7	Frizzled-7	Galectin-3	Galectin-3
	8	GER d-3	GER a-3	-like 1	-like 1	GITR	kine GITR	GITR	GITR	Glut2	Glut2	GM-CSE	GM-CSE	Granzym	Granzym
	0	ICANA 4		JOAN C				Ligand	Ligand	IFN-α/β	IFN-α/β	IFN-α/β	IFN-α/β	e B	e B
	9	ICAM-1	ICAM-1	ICAM-2	ICAM-2	ICAM-5	ICAM-5	ICK	ICK	R1	R1	R2	R2	IFN-β	i⊢n-β
	10	IGFBP-7	IGFBP-7	IGF- I	IGF- I	IGF-II	IGF-II	IL-1α	IL-1α	IL-1β	IL-1β	IL-1 R4	IL-1 R4	IL-1 R6	IL-1 R6
	11	IL-3 Rα	IL-3 Rα	IL-3 Rβ	IL-3 Rβ	IL-4	IL-4	IL-4 R	IL-4 R	IL-5	IL-5	IL-5 Rα	IL-5 Rα	IL-6	IL-6

Figure 3. Biotinylated antibody cytokine microarray assay. (A) Images of the biotinylated antibody arrays. (B and C) Grid map of cytokines from the biotinylated antibody arrays.

Fig. 1. The ADM and DDBDM were white in color, and had good elasticity and tenacity. However, the DDBDM was softer and thinner than ADM. The physical properties of ADM and DDBDM are presented in Table I. There was a statistically significant difference between ADM and DDBDM in each of the properties measured (P<0.05).

Under microscopic observation, normal skin samples exhibited an integrated structure, including epidermis, dermis and appendages. DAPI staining demonstrated the presence of high DNA levels in the nuclei of the normal skin. By contrast, collagen fibers in the burned skin samples were thickened and swollen, coagulated necrosis was observed, and the number of integrated nuclei decreased. DAPI staining demonstrated that the DNA was diffusely distributed due to karyorrhexis and karyolysis in the burned skin. No DNA was observed in ADM and DDBDM samples as they underwent acellular disposal.

D	1	2	3	4	5	6	7	8	9	10	11	12	13	14
1	2 Positiv	Positive	Positive 2	Positive 2	Positive 3	Positive 3	neg	neg	IL-6 R	IL-6 R	IL-7	IL-7	IL-7 Rα	IL-7 Rα
1	3 IL-12 R	β1 IL-12 Rβ1	IL-13	IL-13	IL-13 Rα2	IL-13 Rα2	IL-15	IL-15	IL-15 Rα	IL-15 Rα	IL-16	IL-16	IL-17	IL-17
1	4 IL-17R	D IL-17RD	IL-18 Rα	IL-18 Rα	IL-20	IL-20	IL-20 Rα	IL-20 Rα	IL-21	IL-21	IL-21 R	IL-21 R	IL-22	IL-22
1	5 IL-31 F	A IL-31 RA	Insulin	Insulin	CD18	CD18	I-TAC	I-TAC	кс	кс	Kremen-1	Kremen-1	Kremen-2	Kremen-2
1	6 L-Selec	tin L-Selectir	Lungkine	Lungkine	Lymphot actin	Lymphot actin	Lymphot oxin βR	Lymphot oxin βR	MAdCAM-1	MAdCAM-1	MCP-1	MCP-1	MCP-5	MCP-5
1	7 MIP-2	MIP-2	MIP-3α	MIP-3α	MIP-3β	MIP-3β	MMP-2	MMP-2	MMP-3	MMP-3	MMP-9	MMP-9	MMP-12	MMP-12
1	8 Osteo ntin	oo Osteopo ntin	Osteopor otegerin	Osteopor otegerin	OX40 Ligand	OX40 Ligand	PDGF C	PDGF C	PDGF Rα	PDGF Rα	PDGF Rβ	PDGF Rβ	Pentraxin3	Pentraxin3
1	9 RELM	β RELM β	Resistin	Resistin	S100A10	S100A10	SCF	SCF	SCF R	SCF R	SDF-1	SDF-1	Serum Amyloid A1	Serum Amyloid A1
2	0 TAR	TARC	TCA-3	TCA-3	TCCR	TCCR	TECK	TECK	TFPI	TFPI	TGF-β1	TGF-β1	TGF-β2	TGF-β2
2	1 TIMP	2 TIMP-2	TIMP-4	TIMP-4	TL1A	TL1A	TLR1	TLR1	TLR2	TLR2	TLR3	TLR3	TLR4	TLR4
2	2 TRAI R2	- TRAIL R2	TRANCE	TRANCE	TREM-1	TREM-1	TROY	TROY	TSLP	TSLP	TSLP R	TSLP R	TWEAK	TWEAK
2	3 VEGF	R1 VEGF R1	VEGF R2	VEGF R2	VEGF R3	VEGF R3	VEGF-B	VEGF-B	VEGFC	VEGFC	VEGF-D	VEGF-D	WIF-1	WIF-1
Е	15	16	17	18	19	20	21	22	23	24	25	26	27	28
1	2 IL-9	IL-9	IL-9 R	IL-9 R	IL-10	IL-10	IL-10 Rα	IL-10 Rα	IL-11	IL-11	IL-12 p40/p70	IL-12 p40/p70	IL-12 p70	IL-12 p70
1	3 IL-17B	R IL-17BR	IL-17C	IL-17C	IL-17D	IL-17D	IL-17E	IL-17E	IL-17F	IL-17F	IL-17R	IL-17R	IL-17RC	IL-17RC
1	4 IL-22B	P IL-22BP	IL-23	IL-23	IL-23 R	IL-23 R	IL-24	IL-24	IL-27	IL-27	IL-28	IL-28	IL-31	IL-31
1	5 Lefty-	Lefty-1	Leptin R	Leptin R	LEPTIN (OB)	LEPTIN (OB)	LIF	LIF	LIGHT	LIGHT	LIX	LIX	LRP-6	LRP-6
1	6 M-CSI	M-CSF	MDC	MDC	MFG-E8	MFG-E8	MFRP	MFRP	MIG	MIG	MIP-1α	MIP-1α	MIP-1y	MIP-1y
1	7 MMP-1	4 MMP-14	MMP-24	MMP-24	Neuregul in-3	Neuregul in-3	Neurturin	Neurturin	NGF R	NGF R	NOV	NOV	GPNMB	GPNMB
1	8 PF-4	PF-4	PIGF-2	PIGF-2	Progranu lin	Progranu lin	Prolactin	Prolactin	P-Selectin	P-Selectin	RAGE	RAGE	RANTES	RANTES
1	9 Shh-N	Shh-N	SIGIRR	SIGIRR	SLPI	SLPI	Soggy-1	Soggy-1	SPARC	SPARC	Spinesin Ectodomain	Spinesin Ectodomain	TACI	TACI
2	0 TGF-β	3 TGF-β3	TGF-βR I	TGF-βR I	TGF-β R II	TGF-β R II	Thrombo spondin	Thrombo spondin	Thymus Chemokine-1	Thymus Chemokine-1	Tie-2	Tie-2	TIMP-1	TIMP-1
2		1 TMEFF1	TNF R I	TNF R I	TNF R II	TNF R II	TNF-α	TNF-α	TNF-β	TNF-β	TPO	ТРО	TRAIL	TRAIL
2	2 TWEAK	R TWEAK R	Ubiqultin	Ubiqultin	uPAR	uPAR	Urokinase	Urokinase	VCAM-1	VCAM-1	VE- Cadherin	VE- Cadherin	VEGF	VEGF
2	3 WISP-	1 WISP-1	Neg	Neg	Neg	Neg	Neg	Neg	Positive 3	Positive 3	Positive 2	Positive 2	Positive 1	Positive 1

Figure 3. Continued. (D and E) Grid map of cytokine names from the biotinylated antibody microarrays.

H&E staining demonstrated that the collagens in the ADM were loose but well-organized, and no residual cellular nuclei were observed. In the DDBDM, the collagens were thickened, disorganized and fractured; however, the reticulate structure was maintained (Fig. 2).

Cytokine analysis using biotinylated antibody microarrays. Images of the biotinylated antibody arrays are presented in Fig. 3A, and the names of all cytokines analyzed are presented in Fig. 3B-E. If expression of a cytokine was negative in all samples, it was eliminated from further analysis. Subsequently, 227 of the 308 cytokines remained for analysis. The values from the two groups were compared. The following conditions were considered to be significant: i) A fold change in cytokine level of <0.66 or >1.5; ii) at least one value >200 and iii) P<0.05. *Effects of burn on the expression level of cytokines.* To explore whether the expression of cytokines is affected by burn, the values of cytokine expression were compared between normal and burned skin, and between ADM and DDBDM. If the results demonstrated a significant increase or decrease in both comparisons of the same cytokine, the cytokine was considered to be affected by burn. Results showed that the expression of 14 cytokines were altered by burn (Table II).

Effects of cell extraction on the expression level of cytokines. The expression levels of cytokines were compared between the normal skin and ADM, and between burned skin and DDBDM. If the results demonstrated a significant increase or decrease in both comparisons of the same cytokine, the cytokine was then considered to be affected by acellular treatment. Results



1 week

Α



4 weeks



1 week



2 weeks



3 weeks



4 weeks



Figure 4. General observations following subcutaneous implantation of the ADM and DDBDM. (A) The healed wound 1 and 4 weeks after subcutaneous implantation. (B) Images of the subcutaneously implanted ADM (left) and DDBDM (right) after 1, 2, 3 and 4 weeks. (C) Comparison of the ADM and DDBDM after they were separated from the skin. ADM, acellular dermal matrix; DDBDM, deep-degree burned dermal matrix.

showed that the levels of 23 cytokines were affected by cell extraction (Table III).

Differences in cytokine expression between ADM and DDBDM. The expression levels of cytokines in ADM and DDBDM were compared. The results are indicated in Table IV. The levels of 17 cytokines were decreased in DDBDM compared with ADM, however IL-9 and MMP-9 levels were increased.

General observation of the ADM and DDBDM following subcutaneous implantation. Following subcutaneous implantation, the skin wounds of the mice healed well (Fig. 4A). Four weeks after implantation, ADM and DDBDM degraded incompletely and maintained the some integrity (Fig. 4B and C). No marked inflammatory reaction or rejection was observed, demonstrating histocompatibility of ADM and DDBDM. After one week, the ADM and DDBDM could be easily separated from the subcutaneous tissue (Fig. 4B and C). Two weeks

Cytokine	Normal skin ^a	ADM ^a	P-value	Fold change	Burned skin ^a	DDBDM ^a	P-value	Fold change
RANTES	115.0±7.0	9.0±7.0	<0.01	12.778	208.5±102.5	19.0±10.0	<0.05	10.974
IL-6	2,195.0±32.0	226.0±20.0	<0.01	9.712	3,129.0±4.0	131.5±9.5	<0.01	23.795
IL-3	2,519.0±33.0	303.0±15.0	<0.01	8.314	1,356.5±56.5	145.5±3.5	<0.01	9.323
TNF-α	1,327.0±32.0	184.0±17.0	<0.01	7.212	637.5±18.5	107.0±4.0	<0.01	5.958
IL-10	1,460.0±94.0	221.0±7.0	<0.01	6.606	336.0±8.0	98.0±1.0	< 0.01	3.429
GDF-1	316.5±6.5	50.5±14.5	<0.01	6.267	155.5±23.5	53.0±1.0	<0.01	2.934
IL-12 p70	2,000.5±120.5	322.0±11.0	<0.01	6.213	1,984.5±21.5	150.0±16.0	< 0.01	13.230
IFN-β	1,117.5±6.5	182.0±9.0	<0.01	6.140	1,252.0±85.0	136.0±5.0	<0.01	9.206
MCP-1	1,125.0±20.0	187.5±12.5	<0.01	6.000	829.5±89.5	89.5±0.5	<0.01	9.268
IFN-γ	1,990.5±81.5	332.5±6.5	<0.01	5.986	1,357.5±75.5	163.0±1.0	<0.01	8.328
CD11b	2,094.0±231.0	355.5±4.5	<0.01	5.890	1,527.0±80.0	152.0±6.0	<0.01	10.046
IL-17	1,057.5±46.5	180.0 ± 8.0	<0.01	5.875	268.5±60.5	107.5±3.5	< 0.05	2.498
MMP-14	929.0±37.0	162.0±2.0	<0.01	5.735	372.0±15.0	81.0±7.0	<0.01	4.593
TGF-β1	1,205.5±111.5	212.0±22.0	<0.01	5.686	688.5±39.5	115.5±4.5	<0.01	5.961
IL-5	1,112.0±44.0	202.0 ± 4.0	<0.01	5.505	1,073.5±90.5	118.0±4.0	<0.01	9.097
IL-12 p40/p70	1,288.0±44.0	235.5±3.5	<0.01	5.469	1,111.5±80.5	118.5±12.5	<0.01	9.380
HVEM	1,193.5±118.5	219.5±1.5	<0.01	5.437	499.0±212.0	68.5±1.5	< 0.05	7.285
IL-5 R α	339.0±65.0	81.5±0.5	<0.01	4.160	370.0±37.0	50.0±3.0	<0.01	7.400
RELM β	139.0±3.0	41.5±0.5	<0.01	3.349	562.0±28.0	75.0±1.0	<0.01	7.493
IL-1 R9	48.5±49.6	465.0±5.0	<0.01	0.104	1.0 ± 0.0	29.0±3.0	<0.01	0.034
MMP-9	1.0±0.0	15.0±3.0	<0.01	0.067	1.0 ± 0.0	219.5±21.5	<0.01	0.005
Decorin	49.0±11.0	913.5±76.5	<0.01	0.054	1.0±0.0	180.0±6.0	<0.01	0.006
IL-2 R γ	1.0±0.0	164.5±3.5	<0.01	0.006	1.0 ± 0.0	109.0±3.0	<0.01	0.009

Table III. Expression values of the cytokines with significant differences following cell extraction.

^aValues are presented as mean \pm standard deviation. ADM, acellular dermal matrix; DDBDM, deep-degree burned dermal matrix; RANTES, regulated on activation normal T cell and secreted; IL, interleukin; TNF- α , tumor necrosis factor- α ; GDF-1, embryonic growth/differentiation factor 1; IFN- β , interferon; MCP-1, monocyte chemotactic protein-1; MMP, matrix metalloproteinase; TGF- β 1, transforming growth factor- β 1; HVEM, herpesvirus entry mediator; R, receptor; RELM β , resistin-like molecule β .

after implantation, the ADM adhered more tightly to the subcutaneous tissue than the DDBDM, due to the formation of a fibrous coat. By the third week, the ADM and DDBDM were both covered by the dense fibrous coats and were difficult to separate from the surrounding tissues (Fig. 4B). At week four, the dense fibrous coats became thinner and the ADM and DDBDM began to be absorbed (Fig. 4B and C); the ratio of absorption area between them was not significantly different (P=0.615; Fig. 5).

Histological observation of the ADM and DDBDM following subcutaneous implantation. After one week of subcutaneous implantation, infiltration of inflammatory cells from the surface to the space among the collagen was observed in both tissues; however, the DDBDM exhibited more inflammatory cell infiltration than ADM near the surface. There was no obvious change in collagen distribution between the ADM and the DDBDM. The growth of fibroblasts and neovascularization were not observed in ADM or DDBDM (Fig 6A). After two weeks, the number of infiltrating inflammatory cells was markedly increased in the ADM but decreased in the DDBDM. A small number of new blood vessels were observed, which suggested neovascularization of the dermal matrix. Meanwhile, the fibroblasts began to infiltrate and produce new collagen (Fig. 6B). In the third week, granulation tissue had formed on the ADM surface and more new blood vessels were observed. The number of fibroblasts and newly formed collagen increased while the old collagen began to degrade. In comparison, the DDBDM had thinner granulation tissue on the surface, and the process of neovascularization appeared to be slower, with a smaller number and smaller diameter of new blood vessels than in the ADM samples. Notably, numerous fibroblasts and the emergence of ordered new collagen fibers were observed in the DDBDM (Fig. 6C). Four weeks after subcutaneous implantation, more new collagen fibers emerged in place of the old ones, the majority of which had been degraded, with some forming cavities. It was observed that the DDBDM had more new collagen and less undegraded old collagen than the ADM (Fig. 6D).

Discussion

Deep-degree burn injuries can result in serious disability due to hypertrophic scar formation and skin contracture (27,28). The eschar and denatured skin are typically removed following burn injury due to adverse reactions that may result in further

Cytokine	ADM ^a	DDBDM ^a	Fold change	P-value	
IL-1 R9	465.0±5.0	29.0±3.0	16.034	< 0.01	
Decorin	913.5±76.5	180.0±6.0	5.075	< 0.01	
HVEM	219.5±1.5	68.5±1.5	3.204	< 0.01	
CD11b	355.5±4.5	152.0±6.0	2.339	< 0.01	
IL-10	221.0±7.0	98.0±1.0	2.255	< 0.01	
IL-12 p70	322.0±11.0	150.0±16.0	2.147	< 0.01	
MCP-1	187.5±12.5	89.5±0.5	2.095	< 0.01	
IL-3	303.0±15.0	145.5±3.5	2.082	< 0.01	
IFN-γ	332.5±6.5	163.0±1.0	2.040	< 0.01	
MMP-14	162.0±2.0	81.0±7.0	2.000	< 0.01	
IL-12 p40/p70	235.5±3.5	118.5±12.5	1.987	< 0.01	
TGF-β1	212.0±22.0	115.5±4.5	1.835	< 0.01	
TNF-α	184.0±17.0	107.0 ± 4.0	1.720	< 0.01	
IL-6	226.0±20.0	131.5±9.5	1.719	< 0.01	
IL-5	202.0±4.0	118.0±4.0	1.712	< 0.01	
IL-17	180.0±8.0	107.5±3.5	1.674	< 0.01	
IL-2 R γ	164.5±3.5	109.0±3.0	1.509	< 0.01	
IL-9	65.5±9.5	258.0±32.0	0.254	< 0.01	
MMP-9	15.0±3.0	219.5±21.5	0.068	< 0.01	

Table IV. Expression values of the cytokines with significant differences between ADM and DDBDM.

^aValues are presented as mean \pm standard deviation. ADM, acellular dermal matrix; DDBDM, deep-degree burned dermal matrix; IL, interleukin; R, receptor; HVEM, herpesvirus entry mediator; MCP-1, monocyte chemotactic protein-1; INF- γ , interferon- γ ; MMP, matrix metalloproteinase; TGF- β 1, transforming growth factor- β 1; TNF- α , tumor necrosis factor- α .



Figure 5. Area of ADM and DDBDM following subcutaneous implantation. Values are presented as mean ± standard deviation. ADM, acellular dermal matrix; DDBDM, deep-degree burned dermal matrix.

loss of dermis. Tissue-engineered skin substitutes are generally applied to cover the burn wound following removal of the eschar, improving wound healing (12,29,30). The clinical application of tissue-engineered skin substitutes or an ADM may avoid damage caused by multiple wound dressing changes and thus, accelerate wound healing and reduce scar formation (31,32).

The biological mechanical properties of the dermal matrix are very important. The common treatment method of the dermal matrix is to cover the burn wound following tangential excision. As a barrier between the wound and the external environment, the dermal matrix should have adequate intensity to bear pressure and tensile. In addition, it must possess suitable elasticity such that it can return to its normal length following some degree of elongation. At present, there is no standard method for evaluating the mechanical properties of tissue-engineered skin substitutes (33-36). In the present study, there was a significant difference in the mechanical intensities of ADM and DDBDM. As compared with DDBDM, ADM exhibited enhanced tensile strength and maximum tension. Furthermore, ADM had better elasticity, which was reflected by its elongation at break and strain under stress of 3 MPa. This may have occurred due to the thermal denaturation of collagen, which may have decreased its tensile strength and extensibility. In addition, the thermal injury may have damaged the network structure of the collagen. Despite this, the gaps of these parameters between ADM and DDBDM were acceptable and it was demonstrated that DDBDM prepared in the right conditions may still possess good mechanical properties.

Tissue-engineered skin substitutes are predominantly composed of seed cells and cellular supporting structures, termed the cytoskeleton. The seed cells of the skin are fibroblasts, which produce collagen to form connective tissue, and keratinocytes, which form the epithelium. The cytoskeleton is essential to the growth of seed cells and the regeneration of tissues. It is known that the collagen is one of the most important components of the ECM. ADM is a collagen skeleton without any cells. ADM can protect burn wounds from





Figure 6. Histopathological observations of the ADM and DDBDM 1, 2, 3 and 4 weeks after subcutaneous implantation. Hematoxylin and eosin staining. ADM, acellular dermal matrix; DDBDM, deep-degree burned dermal matrix.

bacterial infection and maintain the appropriate environments to accelerate the dermis reconstruction (7,37).

100×

ADM

DDBDM

When the skin is deeply burned, the entire epithelium and parts of the dermis are damaged. Pathological changes to the dermis include hyaline degeneration, swelling, enlargement and disordered arrangement of collagen. When burned, the hydrogen bonds between collagen fibers break, however, the intermolecular cross-linking is not disrupted due to its low thermal sensitivity. Thus, the collagen is shortened, while the integrity of its structure is maintained (38,39). When the skin is burned unevenly, there may be certain areas in which the cells are necrotic, however, the collagens are merely denatured and maintain a complete structure. Under certain conditions, these collagens may be repaired to become the ideal dermal matrix (40).

In the present study, it was observed that the collagen of DDBDM exhibited swelling and degradation, however, its structure was partially maintained. Therefore, it was hypothesized that the DDBDM may act as a satisfactory cytoskeleton with good tension and ductility when the digestion time of trypsin is sufficiently shortened. When observed microscopically, certain areas of collagen were enlarged, partially fractured and loosely arranged, but predominantly remained continuous in the DDBDM. No cells remained in the ADM and DDBDM following their removal with trypsin and Triton-X, which may explain why neither ADM nor DDBDM showed marked histoincompatibility. Almost all antigens able to induce allograft rejection were removed in the process of removing cells, and the main cells involved in allograft rejection, including T-cells and NK-cells (41,42), were destroyed completely. Good histocompatibility is a critical feature of tissue-engineered tissues or organs, since they are transplanted into the human body (43-46). It has previously been demonstrated that the ECM possesses almost all the features of an ideal tissue-engineered biological material, including histocompatibility, degradability, non-toxicity and mechanical properties that match those of the original tissue (47,48). In the present study, the key components of ADM or DDBDM were collagens, which are a key component of the ECM produced by fibroblasts.

A key characteristic of good histocompatibility is an inability to induce significant immune rejection responses (49-51). In the subcutaneous implantation experiment, no significant rejection response was observed for either ADM nor DDBDM, although the ADM induced a greater inflammatory reaction than DDBDM, which suggested that the so-called burn toxin did not induce adverse reactions to the DDBDM, as previously presumed. Although neovascularization was not clearly observed in the DDBDM, it exhibited a stronger ability to promote fibroblast chemotaxis and the production of new collagen. Four weeks after subcutaneous implantation with DDBDM, almost all old degenerated collagens were degraded and replaced by new collagen fibers with normal structure. Therefore, the DDBDM may be considered as a substitute to ADM to promote wound healing of patients with burns.

Burn wound healing is a complicated process involving inflammatory responses, neovascularization, granulation tissue formation, and epithelium and connective tissue remodeling (52,53). When severe burns occur, a series of harmful substances, termed burn toxins are released. Burn toxins are typically products of ECM degradation that are considered to induce adverse effects in patients with burns, including methyl guanidine, histamine, putrescine, indoles and inflammatory factors. However, beneficial burn healing factors may also be produced by degradation of the ECM, possibly resulting in chemotaxis, angiogenesis, growth factor signaling and antimicrobial activities (14,54). To investigate whether DDBDM, constructed from burned skin, contains reduced burn toxins and beneficial factors compared with ADM, the present study analyzed the expression of 308 cytokines in the mouse skin models.

The current study demonstrated that the expression level of certain cytokines in the skin tissue decreased significantly following burn injury, including interleukin (IL)-3, IL-10, IL-17 and decorin. By contrast, resistin-like molecule β (RELM β), IL-9, IL-23 R and thrombospondin levels were increased. These cytokines are closely associated with inflammatory responses. For example, IL-10 is a known anti-inflammatory cytokine (55) and tumor necrosis factor- α (TNF- α) is a primary mediator of the host response to inflammation (56). Decorin is expressed by fibroblasts (57) and has important biological functions, including regulating collagen formation, maintaining collagen arrangement and inhibiting the activity of transforming growth factor- β (TGF- β). Reducing the level of decroin in the deep dermis may lead to hypertrophic scarring (58). A previous study demonstrated that suppressors of cytokine signaling-3 (SOCS-3) may be important in regulating the balance between immunosuppression and inflammation following thermal injury (59). IL-9 induced the expression of SOCS-3 and SOCS-3 overexpression suppressed IL-9 signaling (60). In addition, IL-9 inhibits TNF- α release in lipopolysaccharide-stimulated human monocytes through TGF- β (61). TGF- β is involved in a number of wound healing processes, including inflammation, stimulation of angiogenesis, fibroblast proliferation, collagen synthesis, and deposition and remodeling of new ECM (62,63). Thrombospondin-1 and 2 are best known for their anti-angiogenic properties and their ability to modulate cell-matrix interactions (64). Thrombospondin-1 suppresses wound healing and granulation tissue formation (65), and blocks thrombospondin-1 binding to CD47, markedly increasing skin graft survival (66).

A previous study demonstrated that the expression of certain cytokines, including TNF- α , IL-1 β and IL-6, were significantly increased in the serum of animal models (67) and human patients with burns (68). Thus, the serum levels of TNF- α , IL-1, IL-6 and IL-10 may indicate the severity of this fatal condition following burn injury (69). However, in local skin tissue, the levels of these cytokines were decreased. This may be because the deep burn destroyed a large number of cells resulting in necrosis and degeneration, and the subsequent loss of cytokine expression. However, it is still unclear

why RELM β , IL-9, IL-23 R and thrombospondin levels were increased in the burned skin in the current study.

To avoid the immune response induced by the genetic material of the skin donator, acellularization was performed on the normal and burned skin to decrease histoincompatibility. In the present study, the trypsin digestion and shock-elution method was successfully used to remove cells from the dermal matrix. Undergoing this process significantly decreased the levels of the majority of cytokines compared with normal and burned skin. Notably, the level of decorin was increased significantly in ADM and DDBDM compared with normal and burned skin, respectively. Decorin is a component of connective tissue that binds to type 1 collagen fibrils and is important in matrix assembly (70,71). It may regulate the activity of TGF- β as recombinant human decorin inhibits TGF- β 1-induced contraction of collagen lattices by hypertrophic scar fibroblasts (70).

In comparison, the levels of cytokines in DDBDM were lower than those in ADM, with the exception of IL-9 and MMP-9, which may explain the observation that the DDBDM induced a smaller inflammatory response than ADM in the subcutaneous implantation experiments. MMP-9 is secreted by keratinocytes and inflammatory cells, and is associated with epithelialization. A previous study also observed an increased level of MMP-9 during the early inflammatory phase of wound healing (72).

During the formation of new granulation tissue, TGF-B1 stimulation induces the deposition of fibronectin extra domain A and α -smooth muscle actin expression by fibroblasts, resulting in enhanced synthesis of new ECM (73). TGF- β 1 is also a crucial factor in the regulation of myofibroblast differentiation, facilitating the contraction of granulation tissue (74). TGF-β has different temporal effects on wound healing and scarring, and any disruption in this expression pattern may result in hypertrophic scar formation (75). In the present study, the level of TGF-β1 in DDBDM was lower than in ADM, which may explain the thinner granulation tissue on the surface of DDBDM compared with ADM in the second week. Excess TGF- β 1 may be responsible for the overproduction of ECM, leading to tissue fibrosis and scar formation. Therefore, the DDBDM may decrease the formation of scar tissue compared with ADM in the treatment of burns.

In conclusion, the current study describes the preparation method of DDBDM. The differences in cytokine expression between the ADM and DDBDM indicated that DDBDM did not produce increased levels of harmful burn toxin. Therefore, the DDBDM may be useful as a dermal matrix. If this method is confirmed to decrease harmful inflammatory factors and utilize beneficial factors, it may be used to alleviate adverse responses and promote wound healing processes, including seed cell infiltration, angiogenesis and matrix remodeling. The DDBDM is derived from autologous dermal matrix, therefore, the histoincompatibility associated with xenogeneic or allogeneic dermal matrices is avoided. However, the present study has a number of limitations. For example, further analysis regarding the regulation key cytokines in DDBDM were not conducted, therefore, the effect of these cytokines on the body is unknown. Additionally, it is unclear whether burn healing would be promoted using the DDBDM as a dermal matrix instead of the ADM. Future investigation should aim to resolve these questions.

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