

Article

Novel Placenta-Derived Liquid Product Suitable for Cosmetic Application Produced by Fermentation and Digestion of Porcine or Equine Placenta Using Lactic Acid Bacterium *Enterococcus faecalis* PR31

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Abstract: Since ancient times, the placenta has been used to produce cosmetic and health food products, whereas fermentation is a technology that has been used to produce foods and cosmetics. For application in cosmetics, traditional placental extracts produced solely by proteolysis have not had enough moisturizing properties or the ability to stimulate the proliferation of epidermal keratinocytes. We combined these two traditional approaches to produce raw materials without such drawbacks that are suitable for cosmetic applications. Using a unique lactic acid bacterial strain, *Enterococcus faecalis* PR31, to directly ferment and digest both porcine and equine placentas, we produced the following liquid products: placenta ferment filtrates. The ferment filtrates stimulated the proliferation of not only normal human dermal fibroblasts but also epidermal keratinocytes. The ferments had higher equilibrium water content properties than traditional placental extracts, and the ferment derived from the porcine placenta maintained high stratum corneum water content levels for up to 6 h after its application on the skin. Metabolome analysis revealed various molecules that were increased by fermentation, among which lactic acid was assumed to play a central role in the high moisturizing properties. To conclude, the placenta ferment filtrates developed in this study are beneficial for cosmetic applications.

Keywords: placenta; fermentation; lactic acid bacteria; cosmetics; fibroblasts; keratinocytes; equilibrium water content; stratum corneum water content; metabolome analysis



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1. Introduction

The placenta plays an important and central role in nurturing mammalian fetuses until birth. After an animal has given birth, the placenta is delivered as an afterbirth, and according to anecdotal evidence, the mother animal is said to eat it to clean the place of delivery and for nutritional purposes [1,2]. Because of the various beneficial components of the placenta, including proteins, peptides, amino acids, growth factors, hormones, vitamins, and minerals, it has been used for both nutritional and cosmetic purposes. The oldest manuscript describing the use of the placenta as an ingredient for external applications might be “Ebers papyrus”, dating back to 3000 BC, in which the use of cat placenta as an ingredient in the mixture recommended to prevent hair graying was described (Ebers 453) [3]. Several old documents describing the usage of the placenta as traditional medicines have been discovered in Asian countries; for example, a Chinese historical book, “Compendium of Materia Medica (Ben Cao Gang Mu)” mentioned the use of dried human placenta (Ze He Che) as a revitalizer [4]. Recently, placental extracts produced by digesting the placenta of several animals with proteases or acids have been used for both inner and outer beauty, especially in Asian countries, including Japan [5–9]. For application in cosmetics, although traditional placental extracts produced solely by proteolysis have been

reported to have the property to stimulate the proliferation of dermal fibroblasts [10–12], their moisturizing effects were not sufficient, and they could not stimulate the proliferation of epidermal keratinocytes [10].

Fermentation is a traditional technology used worldwide since ancient times, mainly to produce fermented food. For cosmetic applications, a wax-based remedy containing fermented plant juice for wrinkle care and the use of sour milk for producing smooth skin were described in ancient Egypt [13]. The application of a rather sticky bread-dough-based face pack was described by Juvenal, an ancient Roman poet [14]. Recently, many cosmetic products have been produced using fermentation technologies since various molecules produced by fermentation have cosmetically beneficial effects [15]. To produce cosmetic ingredients, various microbes, such as lactic acid bacteria, bifidobacteria, yeasts, fungi, and mushrooms, are combined with various materials for fermentation [16].

In this study, we attempted to combine these two traditional approaches to solve the drawbacks of traditional placental extracts mentioned above; that is, we intended to directly ferment the placenta using microbes to produce cosmetically applicable raw materials. Although several studies described the fermentation of placental extract rather than the placenta [17–19], reports on direct fermentation of the placenta are rare, probably because in such fermentation, microbes need to digest the placenta to use its components as nutrients for their growth. To achieve this, we selected the lactic acid bacterium *Enterococcus faecalis* PR31 (PR31) as a unique strain with the ability to digest the placenta. As the placentas, we used porcine and equine ones, which have been widely used in Japan for producing cosmetically applicable placental extracts. Through the fermentation and digestion of porcine and equine placentas with PR31, we produced a novel placenta-derived liquid product suitable for cosmetic applications. In this paper, we describe the characteristics of these products for cosmetic applications.

2. Materials and Methods

2.1. Lactic Acid Bacteria

The lactic acid bacterium PR31 used in this study was isolated from *Kiteck*, a traditional fermented milk in Xinjiang Uyghur Autonomous Region (China), by Dr. Taku Miyamoto, an honorary professor at Okayama University, in 1997. This strain was selected from Miyamoto's lactic acid bacteria library, which is derived from various traditional fermented foods worldwide, for its ability to directly digest the placenta. He generously permitted us to use his library and the strain PR31. This strain was identified as *Enterococcus faecalis* through sequence analysis of the DNA encoding 16S rRNA.

Regarding its virulence, PR31 is not hemolytic, it is vancomycin sensitive, and it lacks the gelatinase phenotype in standard biological assays [20,21]; however, it has the *gelE* gene in PCR analysis [22]. A PCR analysis [22] also showed that PR31 does not have the virulent genes, *esp*, *agg*, *cylA*, *cylB*, and *cylM*, although it has *efaA*, *cpd*, *cob*, *ccf*, and *cad*, which are common in strains of *E. faecalis* isolated from fermented foods [22]. These properties indicate that the practical virulence of PR31 is too low for use in industrial fermentation, although careful handling is necessary.

2.2. The Placenta

In this study, we used porcine and equine placentas obtained from a Japanese pig farmer and New Zealand ranches, respectively, which were collected for the industrial production of traditional placental extracts in Japan. In brief, placentas delivered from healthy mothers as afterbirth were collected, washed with water, and immediately frozen until use. Before usage, the placentas were thawed and minced into several centimeters in size. Because the minced pieces derived from multiple animals were pooled, each run of production uses a mixture of all parts of the placenta.

2.3. Bacterial Culture and Fermentation

For seed culture for fermentation, PR31 was grown in a medium containing 2.4% placental extract (Snowden Co., Ltd., Tokyo, Japan), 1% yeast extract, 0.1% Tween80, and 0.5% glucose. The placental extract used for seed culture was derived from the same animal species used for fermentation.

Placental fermentation was performed as follows: Minced porcine (1 kg) or equine (0.3 kg) placentas were mixed with water (2.4 kg in total), heated for pasteurization (75 °C, 1 h), and then a sterilized glucose solution was added (final glucose concentrations were 3 and 1.25% for porcine and equine, respectively). After the seed culture of PR31 (2%) was added, the mixture was fermented for 20 h at 37 °C using a 5 L jar fermenter (Bioneer-Neo, Marubishi Bioengineering, Tokyo, Japan) under pH-stat condition (pH 6.0) with a NaOH solution, and then the obtained fermented placenta suspension was treated with protease SE-4 (0.2% of the weight of placenta; SkinMedical, Tokyo, Japan) at 40 °C for 1 h for further digestion. Finally, the fermented placenta suspension was heat-sterilized (121 °C, 20 min) and filtered to obtain a clear solution of placenta ferment filtrate. For reference, the photos at each step of the production of porcine placenta ferment filtrate are depicted in Figure 1.

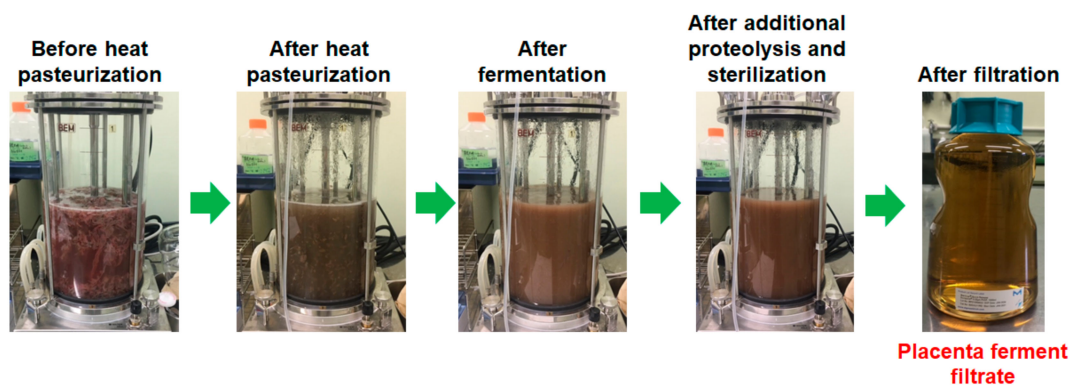


Figure 1. Photos at each step of the production of porcine placenta ferment filtrate.

2.4. Cell Proliferation Assay with Normal Human Epidermal Keratinocytes and Dermal Fibroblasts

NHEK(NB) (Kurabo, Osaka, Japan) was used as a normal human epidermal keratinocyte. The cells were cultivated in DermaLife K keratinocyte medium (Lifeline Cell Technology, Frederick, MD, USA) at 37 °C under a 5% CO₂ atmosphere in a humidified chamber. For the assay, cells were seeded in each well of a collagen-I-coated 96-well microplate at a density of 5.0×10^4 cells/mL in 100 µL. After 24 h, the medium was replaced with a medium containing 1/100 concentrations of supplements and test samples. After another 5 d of culture, the relative cell number was analyzed using the Cell Counting Kit WST-8 (Dojindo Laboratories, Kumamoto, Japan).

NHDF(NB) (Kurabo) was used as a normal human dermal fibroblast. The cells were cultivated in the FibroLife S2 Comp kit (Lifeline Cell Technology) at 37 °C under a 5% CO₂ atmosphere in a humidified chamber. For the assay, cells were seeded in each well of collagen-I-coated 96-well microplate at a density of 1.5×10^4 cells/mL in 100 µL. After 24 h of culture, the medium was replaced with Dulbecco's modified Eagle medium containing 3% fetal bovine serum and test samples. After another 3 d of culture, the relative cell number was analyzed using the Cell Counting Kit WST-8, as aforementioned.

2.5. Total Nitrogen Concentration

Total nitrogen concentrations were measured via the semi-micro Kjeldahl method using an AutoKjeldahl Unit K-370 and Digest Automat K-438 (Nihon BUCHI, Tokyo, Japan).

2.6. Size Exclusion Chromatography

Molecular weight distribution analyses of water-soluble fractions of samples were performed using an ultra-performance liquid chromatography (UPLC) system (ACQUITY UPLC H-Class, Waters, Milford, MA, USA) equipped with a size-exclusion column (ACQUITY UPLC Protein BEH SEC 125 Å, Waters). Samples were diluted in the mobile phase (100 mM phosphate buffer, pH 6.7), filtered using a 0.45 µm membrane, and injected into the system. The injection volume was 1 µL or 5 µL, the flow rate was 0.30 mL/min, and the monitoring wavelength was 220 nm. Molecular weights were calculated using a calibration curve constructed with uracil (112 Da), ribonuclease A (13.7 kDa), ovalbumin (44.2 kDa), and thyroglobulin (660 kDa).

2.7. Equilibrium Moisture Content

To determine the humectant potential of the materials, their equilibrium moisture contents (EMCs) were measured. The liquid sample was freeze-dried, and the obtained powder was placed on an aluminum plate, which was then placed in a chamber set at 40 °C and 75% relative humidity. After 48 h, the water content (WC) of the samples was determined using an infrared moisture analyzer (MA35, Sartorius, Gottingen, Germany). The EMC was calculated using the following equation:

$$\text{EMC}(\%) = \text{WC}(\%) / [100 - \text{WC}(\%)].$$

An EMC of 100% indicates that the dried material can absorb the same amount of water as its dry weight.

2.8. Metabolome Analysis

Metabolome analyses were performed at Human Metabolome Technologies Inc. (HMT, Yamagata, Japan). Before the analysis, 80 µL of the sample was mixed with 20 µL of Milli-Q water containing internal standards (H3304-1002, HMT) to attain 1000 µM. The mixture was centrifugally filtered through a Millipore 5-kDa cutoff filter (ULTRAFREE MC PLHCC, HMT) at 9100 × g, 4 °C for 60 min to remove macromolecules. The filtrate was then used for metabolome analysis according to the HMT Basic Scan package, using capillary electrophoresis time-of-flight mass spectrometry (CE-TOFMS) based on previously described methods [23,24]. Briefly, CE-TOFMS analysis was performed using an Agilent CE capillary electrophoresis system equipped with an Agilent 6210 time-of-flight mass spectrometer (Agilent Technologies, Inc., Santa Clara, CA, USA). The systems were controlled using Agilent G2201AA ChemStation software version B.03.01 (Agilent Technologies) and connected by a fused silica capillary (50 µm i.d. × 80 cm total length) with commercial electrophoresis buffer (H3301-1001 and I3302-1023 for cation and anion analyses, respectively, HMT) as the electrolyte. The spectrometer was scanned from a mass-to-charge ratio (m/z) of 50 to 1000 and peaks were extracted using the MasterHands automatic integration software (Keio University, Yamagata, Japan) to obtain peak information, including m/z , peak area, and migration time (MT) [25]. Signal peaks corresponding to isotopomers, adduct ions, and other product ions of known metabolites were excluded, and the remaining peaks were annotated according to the HMT metabolite database based on their m/z values and MTs. The areas of the annotated peaks were normalized to internal standards and sample amounts to obtain the relative levels of each metabolite. One hundred and ten primary metabolites were absolutely quantified based on one-point calibrations using their respective standard compounds.

2.9. Analysis of Stratum Corneum Water Content

To evaluate the skin-moisturizing effect of placenta ferments in humans, we employed a noninvasive method for stratum corneum water content by measuring the conductance at the skin surface at a high-frequency current of 3.5 MHz using SKICON-200EX-USB (Yayoi,

Tokyo, Japan). This method was developed based on the positive relationship between the conductance at the skin surface and hydration [26,27].

We recruited five female volunteers aged 25–55 (average = 42.4; ages of each participant were 25, 37, 41, 54, and 55) years, without skin problems such as injuries and scratches at the testing sites (inner forearm), skin sensitivity, and skin allergy, and who were not undergoing any treatment or taking medication to treat skin diseases, not pregnant or possible to be pregnant, and not breastfeeding. Before testing, the safety aspects of the test material were guaranteed using the primary skin irritation test (alternative method) and 24 h occlusive patch testing. Written informed consent was obtained from all participants beforehand. This test was performed on 27–28 November 2023.

The sites of application of the test materials were 2×2 cm square areas on both inner forearms. Before testing, the test sites were washed gently with hand soap, and the participants were acclimated for 20 min in a room set at 22 °C and 50% relative humidity. The conductance levels at test sites were determined 10 times using SKICON-200EX-USB, and the averaged values were calculated as the values for “before application”. Then, 20 µL of the test material (water or porcine placenta-derived ferment) was applied at the test site on a randomly assigned right or left inner arm and gently spread until the materials were absorbed into the skin using a finger wearing a powder-free nitril glove. At 0.5, 1, 2, 4, and 6 h after the application of the samples, the conductance at the test sites was measured as described above. Before each measurement, the participants were acclimated for 20 min as aforementioned.

2.10. Statistics

Statistical differences compared to conditions without the addition of test samples in the cell growth assays were analyzed using the Dunnett method. A statistical comparison between the test sample and water in the stratum corneum water content assay at each time point was performed using the Student’s paired *t*-test. The difference was considered statistically significant if the significance probability was <0.05.

3. Results

3.1. Digestion and Solubilization by Fermentation of the Placenta

To determine the extent of solubilization at each step of fermentation of the placentas, we measured the total nitrogen concentrations of the aqueous sample at each step after filtration with a 0.45 µm membrane filter. We employed the nitrogen concentration as a measure of the amount of variety of solubilized molecules, e.g., not only proteins but also other molecules containing nitrogen. The results are summarized in Table 1. As shown in Table 1, before heat pasteurization (Step 1 in Table 1), approximately 20% and 19% of nitrogen relative to the final products were already soluble in porcine and equine placental suspensions, respectively. By heat pasteurization (Step 2), approximately 40% and 21% of nitrogen were further solubilized, affording approximately 60% and 40% of soluble nitrogen relative to the final porcine and equine products, respectively. Fermentation with PR31 (Step 3) solubilized approximately 27% and 51% of the nitrogen, yielding approximately 87% and 91% of the final porcine and equine products, respectively. Additional proteolysis solubilized approximately 13% and 9% (Step 4). These results indicated that the contribution of the fermentation step was the largest in equine placenta solubilization, whereas that in porcine placenta solubilization was the second largest after the heat pasteurization step. Regardless of this difference, the contribution of fermentation exceeded that of additional proteolysis for both porcine (27.4% vs. 13.0%) and equine (51.2% vs. 8.7%) placental solubilization. When the same amounts of placentas for fermentation were solubilized solely by proteolysis without fermentation after heat pasteurization, the soluble nitrogen concentrations were approximately 111% and 121% relative to Step 4 for porcine and equine placentas, respectively.

Table 1. Nitrogen concentration at each step of fermentation and proteolysis of placentas.

Step for Analysis	Porcine			Equine		
	Nitrogen Concentration (mg/mL) ⁽¹⁾	Solubilized (%) ⁽²⁾	Step Solubilization (%) ⁽³⁾	Nitrogen Concentration (mg/mL) ⁽¹⁾	Solubilized (%) ⁽²⁾	Step Solubilization (%) ⁽²⁾
1. Before heat pasteurization	0.41	20.1	20.1	0.33	19.2	19.2
2. After heat pasteurization	1.23	59.6	39.5	0.69	40.1	20.9
3. After fermentation	1.79	87.0	27.4	1.57	91.3	51.2
4. After additional proteolysis and sterilization	2.06	100.0	13.0	1.72	100.0	8.7
5. Without fermentation	2.28	110.6	-	2.08	120.8	-

⁽¹⁾ Average of duplicates. ⁽²⁾ Amount of solubilized nitrogen relative to step 4. ⁽³⁾ Amount of increase in nitrogen concentration from previous step.

3.2. Molecular Weight Distribution at Each Step of the Fermentation of the Placenta

Figure 2 shows the size-exclusion UPLC chromatograms at each step of fermentation and proteolysis in the (A) porcine and (B) equine placentas, respectively. As shown in Figure 2, during fermentation, large amounts of molecules were solubilized, and the molecular weight distribution was shifted in the direction of low molecular weight in both in porcine and equine placentas. Additional proteolysis did not largely change the molecular weight distribution patterns; however, it brought about a slight shift in the low-molecular-weight direction.

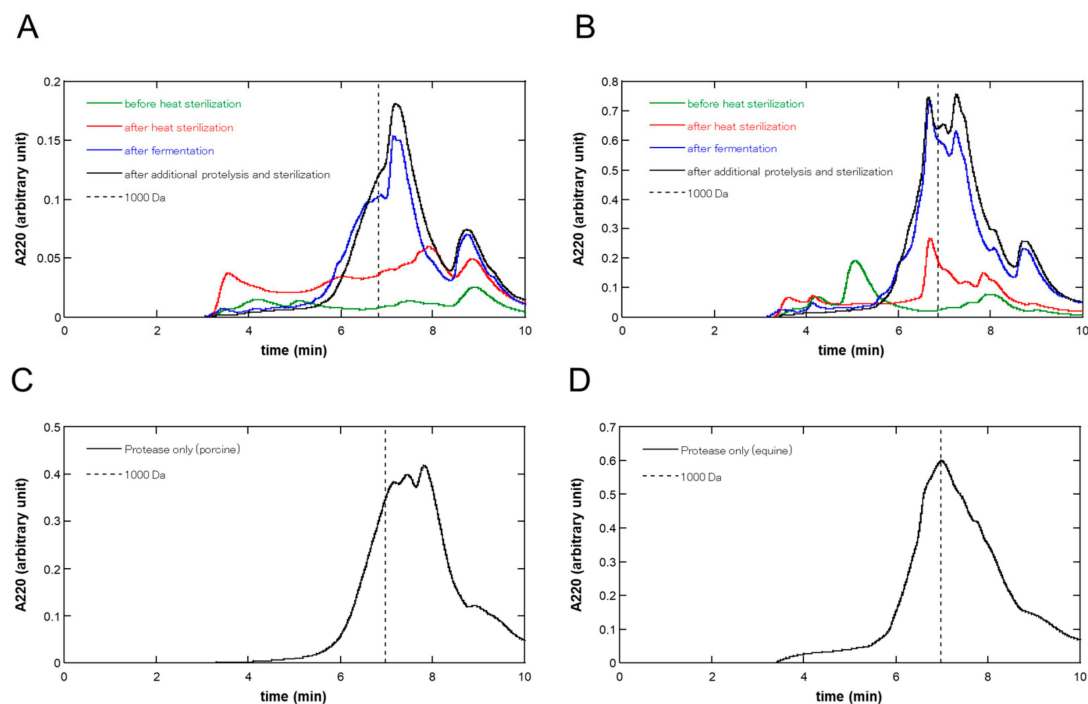


Figure 2. Size exclusion UPLC chromatograms at each step of fermentation and proteolysis of (A) porcine and (B) equine placentas. At each step, the liquid phase of the sample was filtered using a 0.45 μ m membrane filter and 1 or 5 μ L was injected into the size exclusion column for porcine or equine preparation, respectively. Absorbance at 220 nm was monitored and the chromatograms before heat pasteurization, after heat pasteurization, after fermentation, and after additional proteolysis and sterilization were shown in green, red, blue, and black lines, respectively. Size exclusion UPLC chromatograms of placental extracts produced solely by proteolysis are also shown: (C) porcine and (D) equine placental extracts. Liquid phase of the sample was filtered and 5 μ L was injected into the size exclusion column, and then analyzed as above. Each chromatogram shows the average of triplicate runs. The retention times at 1000 Da were indicated in broken lines.

Table 2 shows the molecular weight balance by discriminating at 1000 Da and the areas under the curves (AUCs) at each step of fermentation and proteolysis based on the size exclusion UPLC analysis, as shown in Figure 1. As seen in Table 2, the shifts to a low-molecular-weight direction were observed at the fermentation step (Step 3) in both porcine and equine (≤ 1000 Da: shift from 54.8% to 65.8% and 51.7% to 62.5% in porcine and equine, respectively). Additional proteolysis (Step 4) brought about a further slight low-molecular-weight shift (≤ 1000 Da: shift from 65.8% to 74.0% and 62.5% to 67.9% in porcine and equine, respectively). Although in porcine placentas, almost no low-molecular-weight shift was observed during the heat pasteurization step (Step 2), a shift from 33.9% to 51.7% was observed in the equine placenta. Regarding the amount of solubilization, as judged by the stepwise increase in AUCs, the contribution of fermentation was the largest in equine placentas (53.1%), and in porcine placentas, the contribution of heat pasteurization (43.3%) exceeded that of fermentation (26.0%), similar to the results judged by the nitrogen concentration in Table 1. Regardless of this difference, the contribution of fermentation was greater than that of additional proteolysis in both porcine and equine placentas (26.0% vs. 8.6% and 53.1% vs. 14.6%, respectively).

Table 2. Molecular weight analysis with size exclusion UPLC at each step of fermentation and proteolysis of placentas.

Step for Analysis	Porcine					Equine				
	≤ 1000 Da ⁽¹⁾ (%)	> 1000 Da ⁽²⁾ (%)	Total Area ⁽³⁾	Total area ⁽⁴⁾ (%)	Step Increase ⁽⁵⁾ (%)	≤ 1000 Da ⁽¹⁾ (%)	> 1000 Da ⁽²⁾ (%)	Total Area ⁽³⁾	Total Area ⁽⁴⁾ (%)	Step Increase ⁽⁵⁾ (%)
1. Before heat pasteurization	54.0	46.0	86.3	22.1	22.1	33.9	65.8	373.9	20.9	20.9
2. After heat pasteurization	54.8	45.2	255.1	65.4	43.3	51.7	47.7	576.6	32.3	11.3
3. After fermentation	65.8	34.2	356.6	91.4	26.0	62.5	37.0	1525.9	85.4	53.1
4. After additional proteolysis and sterilization	74.0	26.0	390.0	100.0	8.6	67.9	31.6	1786.2	100.0	14.6

⁽¹⁾ Area under the curve (AUC) of size exclusion UPLC at 1000 Da or smaller relative to the total area. ⁽²⁾ AUC of size exclusion UPLC greater than 1000 Da relative to the total area. ⁽³⁾ Total AUC of size exclusion UPLC. ⁽⁴⁾ Total AUC of size exclusion UPLC relative of the value at step 4. ⁽⁵⁾ Amount of increase in AUC from previous step.

For comparison, we analyzed the molecular weight distribution of placental extracts prepared solely with protease after heat pasteurization, which were independent runs using different placentas. The results are shown in Figure 2C,D. The ratios of AUCs corresponding to ≤ 1000 Da were 75.6% and 62.1% for porcine and equine placental extracts, respectively, which were approximately the same as those of placenta ferments, i.e., 74.0% and 67.9% for porcine and equine placenta-derived preparations (Table 2), respectively.

3.3. Changes in the Amounts of Metabolites Due to Fermentation

The results of the metabolome analyses focusing on the changes in the amounts of metabolites due to fermentation are summarized in Tables 3–6. Tables 3 and 4 present the metabolites that were 4-fold or more in porcine and equine placenta-derived ferment filtrates, respectively, compared to the corresponding proteolysate filtrates (produced solely by proteolysis), where metabolites with relative areas in fermentations larger than 1×10^{-4} were picked up. Tables 5 and 6 show the lists of metabolites that were 0.25-fold or less in porcine and equine placenta-derived ferments, respectively, compared to the corresponding proteolysates, where the metabolites with relative areas in the proteolysates larger than 1×10^{-4} were picked up. The quantification results for certain metabolites are shown in the tables.

Table 3. Compounds that are 4-fold or more in the ferment filtrate than the proteolysate filtrate of porcine placenta.

Compound Name	<i>m/z</i>	MT (min)	Relative Area ⁽⁴⁾		Concentration (μM) ⁽⁵⁾		Fold Amount in Ferment ⁽⁶⁾
			Ferment Filtrate	Proteolysate Filtrate	Ferment Filtrate	Proteolysate Filtrate	
Lactic acid	89.024	8.70	5.0×10^0	1.8×10^{-2}	412,140	1452	283.8
Ornithine	133.097	5.53	6.8×10^{-2}	8.3×10^{-3}	1002	122	8.2
Tyramine	138.091	6.69	4.9×10^{-2}	4.6×10^{-4}	538	5.06	106.3
2-Phenylethylamine	122.096	6.26	4.1×10^{-2}	N.D.	-	-	1<
2-Hydroxy-4-methylvaleric acid ⁽¹⁾	131.071	7.33	3.3×10^{-2}	2.2×10^{-4}	-	-	150.7
2-Hydroxy-3-methylvaleric acid ⁽¹⁾							
Pyruvic acid	87.009	9.90	2.0×10^{-2}	5.8×10^{-4}	2322	68.3	34.0
Citrulline	176.103	8.93	1.2×10^{-2}	2.8×10^{-3}	217	51	4.2
3-Amino-2-piperidone	115.086	6.16	9.0×10^{-3}	6.4×10^{-4}	-	-	14.1
Succinic acid	117.019	13.90	4.3×10^{-3}	1.1×10^{-4}	292	7.26	40.2
Hydroxyproline	132.065	9.72	3.7×10^{-3}	3.9×10^{-4}	61.6	6.59	9.4
Glycerophosphocholine	258.110	17.38	3.4×10^{-3}	3.9×10^{-4}	-	-	8.6
β-Hydroxyisovaleric acid ⁽²⁾							
2-Hydroxyvaleric acid ⁽²⁾	117.056	7.62	2.9×10^{-3}	2.8×10^{-5}	-	-	104.7
2-Hydroxyisovaleric acid ⁽²⁾							
N ⁵ -Ethylglutamine	175.108	9.18	2.9×10^{-3}	N.D.	-	-	1<
Propionic acid	73.029	8.83	2.4×10^{-3}	N.D.	-	-	1<
Butyric acid	87.045	8.04	1.0×10^{-3}	N.D.	-	-	1<
Isobutyric acid							
Glucose 6-phosphate	259.023	8.02	8.6×10^{-4}	N.D.	68.6	N.D.	1<
Tropic acid ⁽³⁾							
3-(2-Hydroxyphenyl)propionic acid ⁽³⁾	165.056	7.19	8.0×10^{-4}	4.3×10^{-5}	-	-	18.7
Gluconic acid	195.051	6.83	5.9×10^{-4}	1.4×10^{-4}	40	10	4.2
N-Acetylserine	148.060	18.56	5.4×10^{-4}	1.0×10^{-4}	-	-	5.3
N ¹ ,N ¹² -Diacetylspermine	287.244	6.10	4.4×10^{-4}	4.9×10^{-6}	-	-	90.3
N-Acetylhistidine	198.087	7.91	4.2×10^{-4}	1.6×10^{-5}	-	-	25.7
N-Lactoylphenylalanine	236.094	6.56	3.7×10^{-4}	N.D.	-	-	1<
N-Acetylaspartic acid	174.041	10.92	3.6×10^{-4}	2.4×10^{-5}	-	-	15.2
AMP	346.056	7.62	3.4×10^{-4}	1.7×10^{-5}	22.3	1.12	19.8
UMP	323.030	7.97	3.2×10^{-4}	2.8×10^{-5}	23.6	2.05	11.5
Terephthalic acid	165.019	12.00	3.0×10^{-4}	2.1×10^{-5}	-	-	14.3
O-Succinylhomoserine	218.067	9.53	2.8×10^{-4}	N.D.	-	-	1<
Ethanolamine phosphate	140.012	6.73	2.7×10^{-4}	5.9×10^{-6}	-	-	46.1
N ¹ ,N ⁸ -Diacetylspermidine	230.186	8.23	2.4×10^{-4}	5.1×10^{-5}	-	-	4.7
Homoserine	120.066	8.15	2.3×10^{-4}	3.3×10^{-5}	3.85	0.551	7.0
GMP	362.050	7.53	2.3×10^{-4}	N.D.	20.3	N.D.	1<
N-Acetylputrescine	131.118	6.83	2.2×10^{-4}	2.5×10^{-5}	-	-	8.8
Spermidine	146.165	3.70	2.2×10^{-4}	N.D.	2.70	N.D.	1<
Fructose 6-phosphate	259.023	8.11	2.2×10^{-4}	N.D.	15.7	N.D.	1<
CMP	322.045	7.84	2.0×10^{-4}	2.7×10^{-5}	21.5	2.92	7.4
N-Acetylthreonine	160.062	7.02	2.0×10^{-4}	3.1×10^{-5}	-	-	6.4
N ¹ -Acetylspermine	245.233	4.54	2.0×10^{-4}	N.D.	-	-	1<
Nicotinamide	123.055	6.03	1.9×10^{-4}	1.9×10^{-5}	-	-	9.9
Homocystine	269.062	7.80	1.2×10^{-4}	N.D.	-	-	1<
Mevalonic acid	147.066	7.16	1.0×10^{-4}	7.6×10^{-6}	-	-	13.6

The results obtained from a single assay of a representative sample are depicted. ^{(1)–(3)} These compounds could not be specified in this analysis. ⁽⁴⁾ Relative areas to that of internal standard are depicted. ⁽⁵⁾ Quantified values are depicted if available, where “-” denotes they are not available. ⁽⁶⁾ Ratio of relative area of ferment filtrate over that of proteolysate filtrate; N.D.: not detected.

Table 4. Compounds that are 4-fold or more in the ferment filtrate than proteolysate filtrate of equine placenta.

Compound Name	<i>m/z</i>	MT (min)	Relative Area ⁽⁴⁾		Concentration (μM) ⁽⁵⁾		Fold Amount in Ferment ⁽⁶⁾
			Ferment Filtrate	Proteolysate Filtrate	Ferment Filtrate	Proteolysate Filtrate	
Lactic acid	89.024	8.70	2.3×10^0	8.0×10^{-3}	188,642	651	289.8
Ornithine	133.097	5.53	4.3×10^{-2}	8.1×10^{-4}	637	12.0	52.9
Tyramine	138.091	6.69	3.5×10^{-2}	1.2×10^{-3}	388	12.9	30.2
2-Phenylethylamine	122.096	6.26	3.1×10^{-2}	N.D.	-	-	1<
Pyruvic acid	87.009	9.90	2.1×10^{-2}	3.4×10^{-4}	2454	40.3	61.0
2-Hydroxy-4-methylvaleric acid ⁽¹⁾	131.071	7.33	1.5×10^{-2}	1.3×10^{-5}	-	-	1114.1
2-Hydroxy-3-methylvaleric acid ⁽¹⁾							
3-Amino-2-piperidone	115.086	6.16	7.8×10^{-3}	1.4×10^{-4}	-	-	53.9
Citrulline	176.103	8.93	7.6×10^{-3}	4.0×10^{-4}	138	7.28	18.9
Hydroxyproline	132.065	9.72	3.6×10^{-3}	5.9×10^{-4}	60.1	9.8	6.1
Glyceric acid	105.019	8.37	2.8×10^{-3}	9.3×10^{-5}	-	-	30.5
N ⁵ -Ethylglutamine	175.108	9.18	2.8×10^{-3}	N.D.	-	-	1<

Table 4. Cont.

Compound Name	<i>m/z</i>	MT (min)	Relative Area ⁽⁴⁾		Concentration (μM) ⁽⁵⁾		Fold Amount in Ferment ⁽⁶⁾
			Ferment Filtrate	Proteolysate Filtrate	Ferment Filtrate	Proteolysate Filtrate	
β-Hydroxyisovaleric acid ⁽²⁾							
2-Hydroxyvaleric acid ⁽²⁾	117.056	7.62	2.4×10^{-3}	7.8×10^{-6}	-	-	312.2
2-Hydroxyisovaleric acid ⁽²⁾							
Succinic acid	117.019	13.90	1.2×10^{-3}	N.D.	79.8	N.D.	1<
Threonic acid	135.030	7.62	1.0×10^{-3}	1.8×10^{-4}	-	-	5.4
Gluconic acid	195.051	6.83	8.5×10^{-4}	7.4×10^{-5}	58.4	5.10	11.5
Tropic							
acid3-(2-Hydroxyphenyl)propionic acid	165.056	7.19	7.9×10^{-4}	4.0×10^{-5}	-	-	19.7
N-Acetylserine	148.060	18.56	7.9×10^{-4}	6.6×10^{-5}	-	-	12.0
Mevalonic acid	147.066	7.16	5.7×10^{-4}	N.D.	-	-	1<
Mevalolactone	131.070	17.84	5.1×10^{-4}	N.D.	-	-	1<
Gluconolactone	179.055	18.36	4.0×10^{-4}	4.6×10^{-5}	-	-	8.8
Terephthalic acid	165.019	12.00	3.4×10^{-4}	2.4×10^{-5}	-	-	13.8
3,4-Dihydroxyhydrocinnamic acid ⁽³⁾							
Homovanillic acid ⁽³⁾	181.051	6.93	3.3×10^{-4}	1.9×10^{-5}	-	-	17.8
Hydroxyphenyllactic acid ⁽³⁾							
N-Acetylaspartic acid	174.041	10.92	3.0×10^{-4}	N.D.	-	-	1<
N ¹ ,N ¹² -Diacetylspermine	287.244	6.10	3.0×10^{-4}	N.D.	-	-	1<
Glucose 6-phosphate	259.023	8.02	3.0×10^{-4}	8.8×10^{-6}	23.8	0.70	33.9
UMP	323.030	7.97	2.6×10^{-4}	1.8×10^{-5}	19.1	1.31	14.6
N-Acetylthreonine	160.062	7.02	1.8×10^{-4}	1.1×10^{-5}	-	-	15.6
AMP	346.056	7.62	1.8×10^{-4}	1.1×10^{-5}	11.6	0.72	16.1
Homoserine	120.066	8.15	1.6×10^{-4}	N.D.	2.64	N.D.	1<
N-Acetylhistidine	198.087	7.91	1.6×10^{-4}	N.D.	-	-	1<
Homocystine	269.062	7.80	1.4×10^{-4}	N.D.	-	-	1<
2-Aminobutyric acid	104.071	7.79	1.4×10^{-4}	N.D.	-	-	1<
N-Acetylputrescine	131.118	6.83	1.3×10^{-4}	2.1×10^{-5}	-	-	6.6
Argininic acid	176.103	7.39	1.3×10^{-4}	N.D.	-	-	1<
N-Lactoylphenylalanine	236.094	6.56	1.3×10^{-4}	6.6×10^{-6}	-	-	19.4

The results obtained from a single assay of a representative sample are depicted. ^{(1)–(3)} These compounds could not be specified in this analysis. ⁽⁴⁾ Relative areas to that of internal standard are depicted. ⁽⁵⁾ Quantified values are depicted if available, where “-” denotes they are not available. ⁽⁶⁾ Ratio of relative area of ferment filtrate over that of proteolysate filtrate; N.D.: not detected.

Table 5. Compounds that are 0.25-fold or less in the ferment filtrate than the proteolysate filtrate of porcine placenta.

Compound Name	<i>m/z</i>	MT (min)	Relative Area ⁽¹⁾		Concentration (μM) ⁽²⁾		Fold Amount in Ferment ⁽³⁾
			Ferment Filtrate	Proteolysate Filtrate	Ferment Filtrate	Proteolysate Filtrate	
Arg	175.119	5.77	2.5×10^{-2}	1.0×10^{-1}	361	1455	0.248
Tyr	182.081	9.16	6.2×10^{-3}	4.7×10^{-2}	107	808	0.133
Guanosine	284.099	10.22	5.8×10^{-4}	7.6×10^{-3}	11.3	148	0.076
Cytidine	244.093	7.80	5.3×10^{-4}	5.5×10^{-3}	10.2	106	0.096
Inosine	269.088	15.75	1.2×10^{-4}	3.6×10^{-3}	2.99	88.3	0.034
Thymidine	243.098	17.79	1.6×10^{-4}	2.1×10^{-3}	7.73	97.9	0.079
2'-Deoxycytidine	228.098	7.61	6.1×10^{-5}	2.0×10^{-3}	-	-	0.031
Uridine	245.078	17.77	4.0×10^{-4}	2.0×10^{-3}	19.9	98.4	0.202
2'-Deoxyguanosine	268.104	9.37	6.9×10^{-5}	1.2×10^{-3}	-	-	0.056
Cystine	241.031	8.90	1.6×10^{-4}	9.5×10^{-4}	-	-	0.165
2'-Deoxyinosine	253.094	14.68	N.D.	1.7×10^{-4}	-	-	<1
3'-CMP	322.045	7.98	N.D.	1.3×10^{-4}	-	-	<1
5-Methyl-2'-deoxycytidine	242.113	7.86	N.D.	1.3×10^{-4}	-	-	<1

Results obtained from a single assay of a representative sample are depicted. ⁽¹⁾ Relative areas to the internal standard are depicted. ⁽²⁾ Quantified values are depicted if available, where “-” denotes they are not available. ⁽³⁾ Ratio of relative area of ferment filtrate over that of proteolysate filtrate; N.D.: not detected.

As seen in Tables 3 and 4, the most abundant and largely increased metabolite by fermentation was lactic acid, which increased from 1452 to 412,140 μM (283.8-fold) and from 651 to 188,642 μM (289.8-fold) in the porcine and equine ferment filtrates, respectively. This was followed by pyruvic acid (68.3 to 2322 μM and 40.3 to 2453 μM, ornithine (122 to 1002 μM and 12.0 to 637 μM), and tyramine (5.06 to 538 μM and 12.9 to 388 μM) in porcine and equine placenta ferment filtrates, respectively. Other metabolites that increased

following fermentation and were quantified were citrulline, succinic acid, hydroxyproline, glucose 6-phosphate, gluconic acid, AMP, UMP, and homoserine in both porcine and equine placenta fermentations; in porcine GMP, spermidine, fructose 6-phosphate, and CMP were also quantified. Certain metabolites, such as 2-phenylethylamine, 2-hydroxy-4 (or 3)-methylvaleric acid, 3-amino-2-piperidone, and β -hydroxyisovaleric acid (or 2-hydroxyvaleric acid, or 2-hydroxyisovaleric acid), lacked quantification data.

Table 6. Compounds that are 0.25-fold or less in the ferment filtrate than the proteolysate filtrate of the equine placenta.

Compound Name	m/z	MT (min)	Relative Area ⁽¹⁾		Concentration (μ M) ⁽²⁾		Fold Amount in Ferment ⁽³⁾
			Ferment Filtrate	Proteolysate Filtrate	Ferment Filtrate	Proteolysate Filtrate	
Arg	175.119	5.77	1.4×10^{-2}	6.0×10^{-2}	195	866	0.225
Tyr	182.081	9.16	1.1×10^{-3}	1.5×10^{-2}	19.5	260	0.075
Guanosine	284.099	10.22	1.1×10^{-4}	6.3×10^{-4}	2.11	12.3	0.172
Cystine	241.031	8.90	5.0×10^{-5}	2.2×10^{-4}	-	-	0.224

Results obtained from a single assay of a representative sample are depicted. ⁽¹⁾ Relative areas to the internal standard are depicted. ⁽²⁾ Quantified values are depicted if available, where “-” denotes they are not available. ⁽³⁾ Ratio of relative area of ferment filtrate over that of proteolysate filtrate.

As seen in Tables 5 and 6, the first- and second-ranking metabolites in abundance decreased following fermentation were the two amino acids, arginine and tyrosine, respectively, in both fermented porcine and equine placentas. Arginine decreased from 1455 to 361 μ M and from 866 to 195 μ M in porcine and equine placenta ferments, respectively, and tyrosine decreased from 808 to 107 μ M and from 260 to 19.5 μ M, respectively. Other metabolites that were decreased by fermentation in the porcine placenta ferments were ten nucleic acids and one amino acid cystine, whereas in equine placenta ferments, one nucleic acid guanosine and one amino acid cystine were found to have been decreased by fermentation.

3.4. Effect on the Proliferation of Normal Human Dermal Fibroblasts

Figure 3 shows the effects on the proliferation of normal human dermal fibroblasts (NHDFs) of the placenta ferment filtrates produced by fermentation of the porcine or equine placenta followed by additional proteolysis, as described in Section 2. As shown in Figure 3, the ferment filtrates derived from both porcine and equine placentas stimulated the proliferation of NHDFs in a dose-dependent manner, with minimum effective concentrations of 1% and 0.25% for the porcine and equine placenta ferment filtrates, respectively.

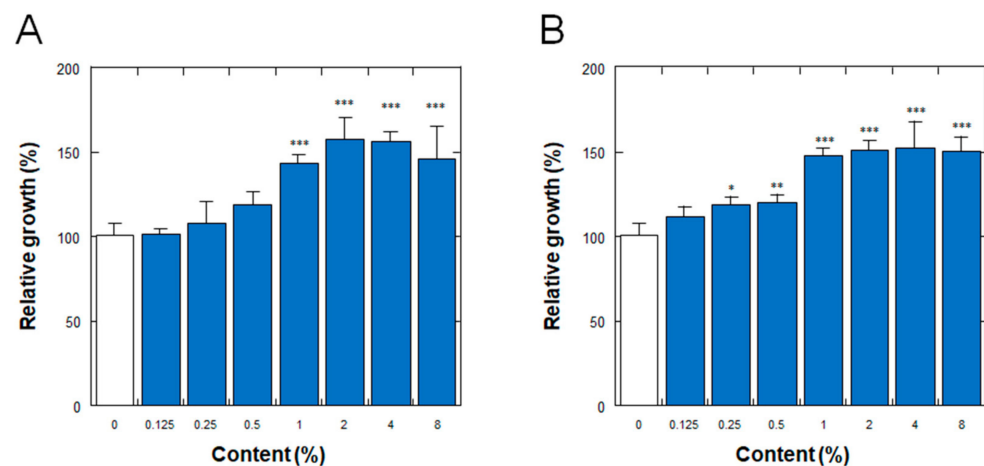


Figure 3. Effect of (A) porcine and (B) equine placenta-derived ferment filtrates on the proliferation of normal human dermal fibroblasts. Relative growths of the cells were determined by the amount

of soluble formazan from tetrazolium salt using Cell Counting Kit WST-8. The results were expressed with % values for the condition without ferments (indicated as 0%). Each bar represents the mean of 6 and 3 determinations for 0% and others, respectively, with standard deviations. * $p < 0.05$, ** $p < 0.01$, and *** $p < 0.001$ vs. 0%.

3.5. Effect on the Proliferation of Normal Human Epidermal Keratinocytes

Figure 4 shows the effects on the proliferation of normal human epidermal keratinocytes (NHEKs) of the placenta ferment filtrates as aforementioned. As shown in Figure 4, the ferment filtrates derived from both porcine and equine placentas stimulated the proliferation of NHEKs in a dose-dependent manner, with minimum effective concentrations of 0.125% and 0.25% for porcine and equine placenta ferment filtrates, respectively.

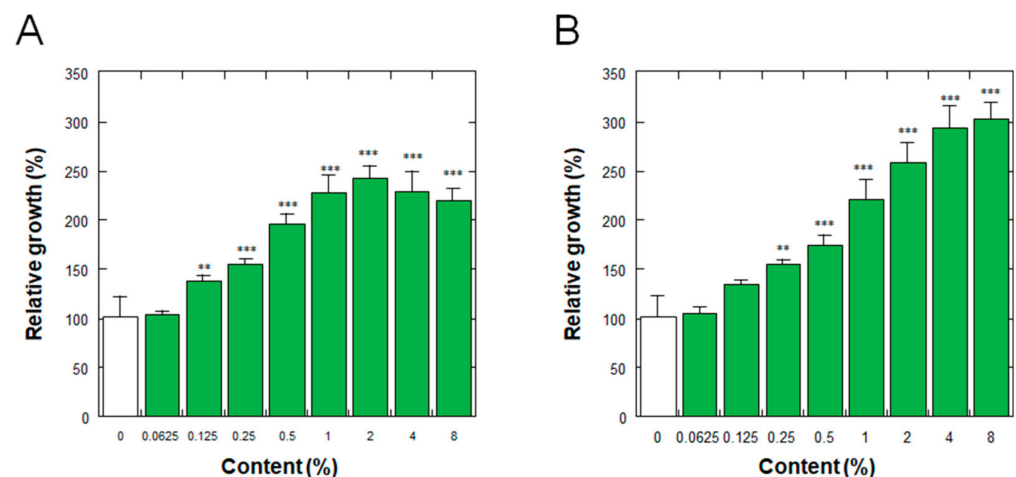


Figure 4. Effect of (A) porcine and (B) equine placenta-derived ferment filtrates on the proliferation of normal human epidermal keratinocytes. Relative growths of the cells were determined by the amount of soluble formazan from tetrazolium salt using Cell Counting Kit WST-8. The results were expressed with % values for the condition without ferments (indicated as 0%). Each bar represents the mean of 8 and 3 determinations for 0% and others, respectively, with standard deviations. ** $p < 0.01$ and *** $p < 0.001$ vs. 0%.

3.6. Equilibrium Moisture Content

Table 7 shows the equilibrium moisture contents (EMCs) at 40 °C and 75% relative humidity of placenta ferment filtrates and corresponding placenta proteolysate filtrates derived from porcine and equine placentas after their lyophilization. As shown in Table 7, the EMCs of ferment filtrates (80.3% and 79.3% for porcine and equine, respectively) were greater than those of the corresponding proteolysate filtrates (34.5% and 38.0% for porcine and equine, respectively) by 2-fold or more. As a reference, we also measured the EMCs of glycerol and sodium hyaluronate (average molecular weight = 2000 kDa) under the same conditions and observed that the values were 78% and 37%, respectively, suggesting that the EMCs of ferment filtrates and proteolysate filtrates were comparable to those of glycerol and sodium hyaluronate, respectively.

Table 7. Equilibrium moisture content of placenta-derived samples.

Sample	EMC (%)
Porcine placenta ferment filtrate	80.3 ± 2.1
Equine placenta ferment filtrate	79.3 ± 3.0
Porcine placenta proteolysate filtrate	34.5 ± 1.9
Equine placenta proteolysate filtrate	38.0 ± 0.9

The equilibrium moisture content (EMC) of each sample is shown as the mean ± standard deviation ($n = 3$).

3.7. Stratum Corneum Water Content

To determine whether the aforementioned high EMC properties of placenta ferment filtrates bring about a high moisturizing effect on human skin, we examined the effect of porcine-derived placenta fermentation on stratum corneum water content. As shown in Figure 5, at 0.5, 1, 4, and 6 h after the application of the samples, the skin surface conductance at the site of placenta ferment application was significantly higher than that of water application. This result clearly indicates that the high EMC value of placenta ferments results in their high moisturizing properties on the skin.

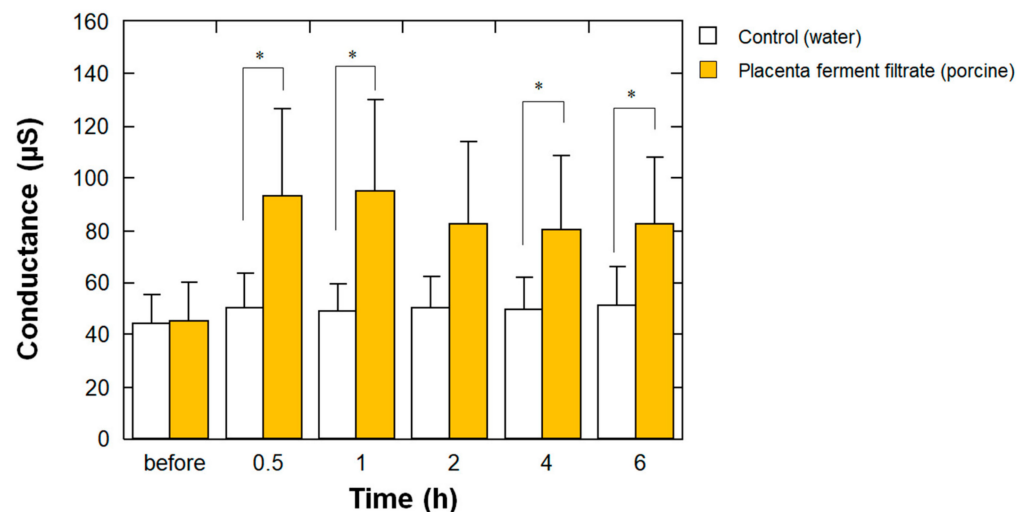


Figure 5. Effect of porcine-derived placenta ferment filtrates on stratum corneum water content. Before and after 0.5, 1, 2, 4, and 6 h from the topical application of 20 μ L of the ferment (closed bars) or water (open bars) on each forearm, the stratum corneum water content was measured using SKICON-200EX-USB as described in Materials and Methods. Each bar represents the mean with standard deviation of the measurements in five subjects. * $p < 0.05$ vs. control at each time point.

4. Discussion

In the present study, we produced a placenta ferment filtrate using a strain of lactic acid bacteria PR31 to ferment and directly digest both porcine- and equine-derived placentas. The placenta ferments produced by this method contained various molecules derived from placentas and fermentation, stimulated the proliferation of both normal human dermal fibroblasts and normal human epidermal keratinocytes, and had high EMC values. Moreover, the stratum corneum water content after topical application of the ferment derived from porcine placenta to the human skin was significantly higher than that of water until 6 h after application. These properties suggest that these placenta ferment filtrates are the materials preferred for cosmetic applications. The porcine- and equine-derived placenta ferments have been assigned the following product names as cosmetic ingredients: *Enterococcus faecalis*/Pig Placenta Ferment Filtrate and *Enterococcus faecalis*/Horse Placenta Ferment Filtrate, respectively, according to the International Nomenclature of Cosmetic Ingredients. We propose these materials as novel raw materials for cosmetic applications.

Regarding the application of fermentation to produce placenta-derived products, several reports have described methods for fermenting placental extracts (not the placenta itself) using yeast [17–19]. In other words, in these studies, yeast grew in a nutrient medium containing placental extract as a component, where the placental extract (not the placenta itself) was produced beforehand by proteolysis or other non-biological methods. An examination of the possible use of selected yeast strains for deodorizing sheep placental extract solutions has also been reported [28]. We have provided a type of porcine placental extract produced by combining fermentation and proteolysis to researchers in universities to examine its effects on lipid metabolism [29–31], where the porcine placenta was fermented

with yeast and lactic acid bacteria, and then the fraction not solubilized by the fermentation was proteolyzed [29]. However, in such a preparation, over 50% of the nitrogen was solubilized by proteolysis, indicating that the contribution of fermentation was much smaller when digesting the placenta than in the present study. In this study, a unique lactic acid bacterium strain, PR31, enabled the direct and rapid digestion of placentas to obtain liquid materials for use in cosmetics. Although we employed additional proteolysis to thoroughly digest placentas in the last step, the gains from the final proteolysis were marginal, that is, approximately 10% for the solubilization of nitrogen (Tables 1 and 2) and approximately 5% to 8% for the shift to low molecular weight (Table 2). In the preparations produced solely by proteolysis, the solubilized nitrogen concentrations were slightly higher than those in the corresponding ferments by 10–20%. This might be because, in the ferments, some components containing nitrogen were assumed to be used by PR31 as nutrients.

We found that the placenta ferment filtrates derived from both porcine and equine placentas stimulated the proliferation of both normal human dermal fibroblasts (Figure 3) and epidermal keratinocytes (Figure 4). These properties are preferable for cosmetic applications because they may promote both dermal and epidermal skin turnover. In previous studies, porcine placental extract was reported to stimulate the proliferation of dermal fibroblasts [10–12]; however, it was also reported to have no effect on the proliferation of epidermal keratinocytes [10]. Although there are some reports describing the growth-promoting activity of human placental extracts produced without heat treatment in keratinocytes [32,33], heat-labile and/or human-specific molecules may have been responsible for such an effect. The detailed molecular mechanisms underlying the effects of placenta ferment on dermal fibroblast and epidermal keratinocyte proliferation should be elucidated in future studies. In this study, since the placenta ferments were heat-sterilized at 121 °C, assuming the contribution of growth factors possibly contained in the raw placentas is unrealistic. In dermal fibroblasts, the growth-promoting effect of porcine placental extract was reported to occur via fibroblast growth factor (FGF) receptors [12], and the porcine placental extract stimulated the expression of FGF in dermal fibroblasts [10]. Therefore, assuming that certain factors in placenta ferments derived from the placenta promote FGF expression and affect dermal fibroblasts in an autocrine manner is reasonable. However, some molecules produced by fermentation might be responsible for the proliferation-promoting activity in keratinocytes of placenta ferments because placental extracts produced solely by proteolysis do not have such an effect [10]. Although detailed identification of such molecules remains a future challenge, one of the candidates might be pyruvic acid, as described below.

Regarding the differences between porcine and equine placentas, we assume that placenta ferments with the same properties were produced. However, in porcine placentas, larger amounts of materials were solubilized during heat pasteurization than in equine placentas (Tables 1 and 2), possibly owing to the difference in the placental structures. In practice, approximately 3-fold of the wet weight of porcine placenta is required to attain the same order of nitrogen concentration of placenta ferment in our laboratory. Moreover, porcine placentas are assumed to have a looser structure compared to equine placentas. Such a loose structure might have enabled us to solubilize larger amounts of materials as a kind of soup by heat pasteurization. The loose nature of the porcine placenta might also be the cause of the difference in the final molecular weight balances; that is, the ratio of low molecular weights tended to be larger in the porcine placenta-derived ferment than in the equine placenta-derived ferment (Table 2).

Basic skincare cosmetics must possess excellent moisturizing properties. As shown in the EMC analysis, placenta ferments had high EMC values comparable to those of glycerol (Table 3). Moreover, the high skin moisturizing properties of the stratum corneum were confirmed (Figure 5). As their EMC values were higher than those of the corresponding traditional placental extracts produced solely by proteolysis (Table 3), these high moisturizing properties may be assumed to have been derived from molecules produced by fermentation. One of the molecules supporting the high moisturizing properties of placental ferments

might be lactic acid, a natural moisturizing factor (NMF) [34]. In this study, lactic acid was largely increased by fermentation and was the most abundant metabolite in ferment by metabolome analysis (>100 mM in both porcine and equine placenta ferments). In addition to lactic acid, the metabolome analysis revealed that pyruvic acid, ornithine, tyramine, and citrulline largely increased to >100 μ M by fermentation in both porcine and equine placenta ferments. Ornithine and citrulline are free amino acids that are not involved in protein structure but are the central molecules in the urea cycle. In the skin, since they are thought to be NMFs derived from filaggrin by degradation and conversion [35–37], these amino acids may also act as NMFs. As both ferments and proteolysates contain various free amino acids, these molecules may also function as NMFs. Although in this study biological analyses were performed only with placenta ferments, head-to-head comparisons of the biological activities between placenta ferments and corresponding proteolysates will be helpful to elucidate the roles of metabolites derived from fermentation in future studies.

Regarding other metabolites that were found to increase during fermentation, pyruvic acid is an important molecule in the citric acid cycle. It enters the citric acid cycle after being transformed into acetyl-CoA or oxaloacetate [38]. Since the citric acid cycle contributes to energy production in cells, this molecule may participate in promoting cell growth. Tyramine is an aromatic amine derived from tyrosine via decarboxylation. Although tyramine is a possible substrate for tyrosinase, a key enzyme in melanogenesis, tyramine inhibits tyrosinase activity when measured with tyrosine as its substrate [39]. Therefore, this molecule may be beneficial for lightening the skin; however, further research is necessary to test this hypothesis. Although we found various molecules that were increased by fermentation other than those aforementioned (Tables 3 and 4), the elucidation of the biological implications of these molecules is open to future studies. Regarding the metabolites found to be decreased in the ferments (Tables 5 and 6), two amino acids, arginine and tyrosine, were the top one and two, respectively, in both porcine and equine. Cystine and several other nucleic acids are also listed. We believe that these molecules were consumed by the lactic acid bacteria during fermentation.

5. Conclusions

By directly fermenting and digesting porcine or equine placentas using a unique lactic acid bacterial strain, PR31, we produced a liquid material, placenta ferment filtrate, that is suitable for cosmetic application. Ferment filtrates derived from both porcine and equine placentas stimulated the proliferation of normal human dermal fibroblasts and epidermal keratinocytes at a concentration of 1% or higher. Such a property of ferment filtrates was superior to that of traditional placental extracts, which could not stimulate the proliferation of epidermal keratinocytes. The ferment filtrates also had 2-fold or higher equilibrium water content properties than those of the corresponding proteolysate filtrates, and the expected high skin-moisturizing properties were confirmed by stratum corneum water content analysis with the ferment filtrate derived from porcine placenta, i.e., the ferment filtrate maintained high stratum corneum water content levels for up to 6 h after its application on the skin. Using metabolome analysis, several candidate molecules that support these effects can be hypothesized. These properties of placenta ferment filtrates derived from both porcine and equine placentas support their beneficial application in cosmetics. In a future study, the molecular basis of the biological effects of the placenta ferments should be elucidated in more detail, and other biological analyses related to their cosmetic applications should be performed as well.

Author Contributions: K.M. and Y.K. performed the experiments and wrote the manuscript. All authors have read and agreed to the published version of the manuscript.

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Institutional Review Board Statement: The study on stratum corneum water content was conducted without formal approval from the institutional review board, because the test sample, ferment filtrate derived from porcine placenta, had been already commercially available from January 2023 in Japan,

and its safety aspects had been guaranteed by a primary skin irritation test (alternative method) and a 24 h occlusive patch testing. Despite the test material was thought safe as mentioned and the setting of the test was within short time-period (within 6 h), we obtained written informed consent from all participants beforehand.

Informed Consent Statement: Informed consent was obtained from all subjects involved in the study for their participation in the study on stratum corneum water content.

Data Availability Statement: The data underlying this article will be shared at the reasonable request to the corresponding author.

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Conflicts of Interest: K.M. and Y.K. are employees of Snowden Co., Ltd. This study was funded by Snowden Co., Ltd. No external funding was used.

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