Laminins in an in vitro anterior lens capsule model established using HLE B-3 cells

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Abstract. Cataracts are the most common eye disease to cause blindness in patients. The abnormal deposition of laminins (LMs) in the lens capsule and the disruption of capsular lamellar protein distribution contribute to cataract development, although the mechanism by which this occurs is currently unclear. The present study aimed to reproduce HLE B-3 basement membranes (BM) using HLE B-3 cells and to analyze the similarities of LM expression between HLE B-3 BMs and human anterior lens capsule (ALC). Immunohistochemistry (IHC), ELISA, western blot analysis and immunoprecipitation (IP)-western blot analysis were used to detect total LMs, LM trimers and 11 LM subunits in HLE B-3 cells, HLE B-3 BMs and human ALCs. In IHC staining, HLE B-3 cells and human ALCs were positive for LMs. In LM ELISA, all samples analyzed were positive for LMs. Western blot analysis detected all LM subunits except for LMα3 in HLE B-3 cell lysate, 4 subunits (LMα4, LMα2, LMα1 and LMγ1) in HLE B-3 cell culture supernatant, 5 subunits (LMα4, LMα2, LMα1, LMβ3 and LMγ1) in HLE B-3 BMs, and 3 subunits (LMα4, LMγ2 and LMγ1) in human ALCs. The results of IP-western blot analysis revealed that the LM411 trimer was detected in HLE B-3 cell culture supernatant. These results indicated that HLE B-3 BMs were similar to human ALCs in terms of LM expression. Therefore, HLE B-3 BMs could be used as an in vitro ALC model to determine the role of LMs in ALC in the pathogenesis of cataracts and to select potential anti-cataract drugs.

Introduction

A cataract is defined as any type of opacity of the eye lens that affects vision (1). Cataracts are the main cause of blindness and visual impairment worldwide, particularly in developing countries (2,3). There are a number of risk factors for the development of cataracts, including ultraviolet light exposure, diabetes, aging and other environmental factors (4-6). Despite the number of studies that have been conducted on cataracts, their pathogenesis remains unknown. Thus, effective non-surgical therapies are currently lacking, which is a substantial burden to healthcare budget (7). Dysfunction of lens epithelial cells (LECs) has been repeatedly proposed to serve a key role in cataract formation, although this mechanism is unknown (8-10).

Laminins (LMs) are large-molecular-weight trimeric glycoproteins consisting of α-, β- and γ-subunits, which integrate in almost all basement membranes (BM), functioning as scaffolds (11). It has been suggested that LM trimers are assembled inside the cell, with extracellular proteolytic processing of various subunits leading to the mature LMs (12). In addition to their function as scaffold of BMs, LMs can also interact with cell surface receptors such as integrins to control signaling events that regulate cellular proliferation, migration and differentiation (13,14).

Particular LM isoforms are present in specific tissue and cell distributions during developmental and pathological conditions, indicating that different LM isoforms have distinct functional roles (12,14). Past studies have demonstrated that LMα1, LMα5, LMβ1, LMβ2 and LMγ1 subunits exist in the
developmental human lens capsule (15,16). Another prior study demonstrated that human adult lens capsule is composed of LMα1, LMα2, LMα3, LMα4, LMβ5, LMβ1, LMβ2 and LMγ1 subunits, and the predominant LM trimer was LM521 (17).

Gene defects and the abnormal expression of LMs are closely associated with various lens diseases (18-23). LMβ2 mutations cause Pierson syndrome, which has various ocular lesions, including abnormal lens shape, posterior lenticous and cataracts (19). It has been demonstrated that development of LM-positive linear capsular densities and occurrence of lens epithelial degenerative alterations are associated with aging (20). Increased LM expression underneath lens capsules and abnormal LEC migration were identified in FE65/FE65L1 double knockout lenses, which may be involved in the formation of cortical cataracts (21). Abnormal accumulation of LMs in the lens capsule and epithelial-mesenchymal transition of LECs was also observed in patients with anterior subcapsular opacification and posterior capsule opacification (22,23).

Human primary LECs in long-term cultures could synthesize a continuous sheet of capsule-like material (24). Theoretically, human LEC lines could also produce anterior-capsule-like BM with rich LMs in vitro, which could be a suitable disease model to study the role of LMs in anterior lens capsule (ALC) in the pathogenesis of lens diseases, particularly cataracts.

In the present study, the human LEC HLE B-3 line was used to construct the BMs and further analyze the expression levels of 11 LM subunits. Similarities in LM expression between HLE B-3 BMs and human ALCs were also evaluated. The results of the present study may aid in elucidating the role of LMs in ALC and cataract pathogenesis.

**Materials and methods**

**HLE B-3 cell culture and preparation of cell lysate and culture supernatant.** The human LEC HLE B-3 line were obtained from the American Type Culture Collection (Manassas, VA, USA) and cultured in low-glucose DMEM (HyClone; GE Healthcare Life Sciences, Logan, UT, USA) containing 10% fetal bovine serum (FBS) (Gibco; Thermo Fisher Scientific, Inc., Waltham, MA, USA), 100 U/ml penicillin and 100 µg/ml streptomycin (Beyotime Institute of Biotechnology, Haimen, China) at 37°C in humidified atmosphere with 5% CO2. Cells were seeded in 10-cm dishes and grown to 95% confluence. Following discarding the culture supernatant, cells were washed twice with PBS and lysed with cold radioimmunoprecipitation assay (RIPA) buffer [50 mM Tris-HCl (pH 7.5), 150 mM NaCl, 1% Triton X-100 and 1% protease inhibitor cocktail from Sigma-Aldrich; Merck KGaA, Darmstadt, Germany] at 4°C for 30 min. Following centrifugation at 14,000 x g at 4°C for 20 min, the supernatant was harvested and kept as culture supernatant. When 95% confluent, cells were washed twice with PBS and cultured in 5 ml low-glucose DMEM without FBS for 48 h. The cell supernatant was harvested and centrifuged at 3,000 x g at 4°C for 5 min to remove debris.

**Immunocytochemical staining (ICC) of HLE B-3 cells.** For hematoxylin and eosin (H&E) staining, HLE B-3 cells grown on glass slides were stained with hematoxylin and eosin (5 and 2 min, respectively, at room temperature), and examined under a light microscope (Olympus Corporation, Tokyo, Japan; magnification, x400). For ICC of LMs, HLE B-3 cells grown on glass slides were incubated for 10 min at room temperature with 3% hydrogen peroxide, and blocked with 10% normal goat serum for 20 min at room temperature, followed by incubation for 60 min at room temperature with rabbit anti-LM antibody (1:200; cat. no. ab11575; Abcam, Cambridge, UK). Subsequently, samples were incubated for 30 min at room temperature with peroxide conjugated anti-rabbit antibody (cat. no. PV-9000; OriGene Technologies, Inc., Beijing, China) without dilution and color development was performed with 3,3′-diaminobenzidine (DAB). Staining was visualized with a light microscope (Nikon TE300, Nikon Corporation, Tokyo, Japan; magnification, x400), and images were captured with an attached digital camera and associated software (SPOT Basic™ image capture software; cat. no. SPOT53BE; SPOT Imaging, a division of Diagnostic Instruments, Inc., Sterling Heights, MI, USA).

**BM preparation using HLE B-3 cells.** When HLE B-3 cells reached 95% confluence, they were washed with PBS and cultured in fresh low-glucose DMEM medium with 5% FBS; this day was recorded as day 0. At days 0, 3, 6, 9 and 12, HLE B-3 BMs were harvested, as described previously (25,26). Briefly, HLE B-3 cells were washed with PBS four times, after which 0.03% ammonia and 0.1% Triton-PBS was added to remove HLE B-3 cells completely. Finally, HLE B-3 BM remained on the bottom of the dish, which was washed with double distilled water 4 times and harvested with 150 µl of 4X sample buffer [8% sodium dodecyl sulfate, 0.2 M Tris-Cl (pH 6.8), 40% glycerol, 0.2% bromophenol blue and 0.1 M dithiothreitol] at room temperature or 1 ml RIPA buffer on ice.

**Human ALCs and protein lysates.** Human ALCs were obtained from autopsies from 3 individuals (1 female and 2 male) aged 25-79 years old from November to December of 2016 at the Eye Bank of Heilongjiang Province (Harbin, China). Human ALCs were obtained by a single ophthalmologist using central circular capsulorhexis method from anterior surface of transparent lens (27). All experiments were performed with the approval of the Internal Review Board of Harbin Medical University and were conducted in accordance with Declaration of Helsinki Principles. Informed consent was obtained from all patients prior to mortality or from their families following mortality for inclusion of autopsy data. Samples were harvested within 4-24 h of mortality. Human ALCs were dissected from the isolated transparent lenses and were lysed with cold RIPA buffer at 4°C overnight. Finally, the supernatant was kept as human ALC lysate following centrifugation at 14,000 x g at 4°C for 20 min.

**H&E staining and immunohistochemistry (IHC) staining of human ALCs.** Following careful dissection from transparent lenses, excised human ALCs were frozen at -80°C. Frozen human ALC tissues were transversely sectioned at 5-µm thickness, mounted on glass slides, fixed with cold acetone for 15 min at 4°C, followed by H&E staining or IHC with identical steps to ICC, as aforementioned.
**Bicinchoninic acid assay.** Protein concentrations in the HLE B-3 cell lysate, HLE B-3 cell culture supernatant, human ALC lysate and various HLE B-3 BMs were measured using a Bicinchoninic Acid Assay kit (Beyotime Institute of Biotechnology) according to the manufacturer's protocol.

**ELISA.** The total LM levels were measured by using commercially available LM ELISA kit in accordance with the manufacturer's protocol (E-EL-H0128c; Elabscience, Houston, TX, USA). The antibodies used in this kit were polyclonal antibodies (pAbs) against all types of LMα, LMβ and LMγ subunits.

**Antibodies.** Goat pAbs against LMα5 and LMγ3 (1:200, cat nos. sc-16592 and sc-133178; Santa Cruz Biotechnology, Inc., Dallas, TX, USA), rabbit pAbs against LMα4, LMβ3 and LMβ2 (1:1,000, cat nos. C13067, C13071 and C30224; Assay Biotechnology Company, San Francisco, California), rabbit pAbs against LMγ1 (1:1,000, cat. no. ab69632; Abcam, Cambridge, UK), mouse monoclonal antibodies (mAb) against LMα4, LMα3, LMα2, LMα1, LMβ2 and LMβ1 (1:200, cat nos. sc-130540, SC-13586, sc-55605, sc-74418, sc-133241 and sc-17763; Santa Cruz Biotechnology, Inc.) were used in the present study.

**Western blotting analysis.** Five known LMα chains, three known LMβ chains and three known LMγ chains in HLE B-3 cell lysate, HLE B-3 cell culture supernatant and HLE B-3 BMs harvested at days 0, 3, 6, 9 and 12, and human ALC lysate were analyzed by western blotting, as described previously (28). Briefly, antigen sources including HLE B-3 cell lysate, culture supernatant or protein lysate of human ALCs were mixed with 2× sample buffer, boiled for 2 min and separated by SDS-PAGE. The separated proteins were transferred to a polyvinylidene difluoride membrane (EMD Millipore, Billerica, MA, USA). Following blocking with 5% skimmed milk in Tris-buffered saline containing 0.05% Triton X-100, and were eluted from protein G agarose precipitates contained LMα1 (1:200, cat no. 31433; Thermo Fisher Scientific, Inc.) or HRP-conjugated goat anti-rabbit IgG (1:5,000, cat no. 31433; Thermo Fisher Scientific, Inc.) diluted in solution 2 (Toyobo Life Science, Osaka, Japan) at 4°C overnight. Following washes with TBS-T, membranes were incubated with the aforementioned primary antibodies diluted in solution 1 (Toyobo Life Science, Osaka, Japan) at 4°C overnight. Following washes with TBS-T, membranes were incubated with HRP-conjugated goat anti-mouse IgG (1:5,000; cat no. 31437; Thermo Fisher Scientific, Inc.), HRP-conjugated rabbit anti-goat IgG (1:5,000; cat no. 31433; Thermo Fisher Scientific, Inc.) or HRP-conjugated goat anti-rabbit IgG (1:5,000; cat no. 31463; Thermo Fisher Scientific, Inc.) diluted in solution 2 (Toyobo Life Science) for 1 h at room temperature. Under identical experimental conditions, normal rabbit IgG, normal mouse IgG or normal goat IgG (all 1:200; cat nos. sc-2027, sc-2025 and sc-2028; Santa Cruz Biotechnology, Inc.) were used as isotype controls. Finally, the antibody-antigen complex was visualized using an Enhanced Chemiluminescent kit (Beyotime Institute of Biotechnology).

**Immunoprecipitation (IP)-western blot analysis.** All subsequent procedures were performed at 4°C, unless otherwise stated. A total of 1 ml HLE B-3 cell culture supernatant was incubated overnight on a shaker with protein G agarose (Beyotime Institute of Biotechnology) and anti-LMα4 rabbit pAb or anti-LMγ1 rabbit pAb. The same concentrations of normal rabbit IgG were used as an isotype control for IP. Precipitates, collected by centrifugation at 3,000 x g at 4°C for 2 min, were washed three times with PBS containing 0.5% Triton X-100, and were eluted from protein G agarose by boiling with 4X sample buffer for 2 min. Proteins in the supernatant were separated by SDS-PAGE and detected by western blotting, as aforementioned.

**Results**

**LMs in HLE B-3 cells.** To investigate whether HLE B-3 cells were positive for LMs, H&E staining and ICC for LMs were performed. As depicted in Fig. 1A and B, LMs were widely expressed in the cytoplasm of HLE B-3 cells. To investigate the expression of LMs in HLE B-3 cell lysate and culture supernatant, ELISA and western blot analysis was performed. ELISA for the presence of LMs generated positive results in HLE B-3 cell lysate and culture supernatant (Fig. 1C). By western blotting, all 11 LM subunits were probed in the HLE B-3 cell lysate and culture supernatant. The HLE B-3 cell lysate was positive for LMα5, LMα4, LMα3, LMα2, LMα1, LMβ3, LMβ2, LMβ1, LMγ2 and LMγ1 subunits, whereas HLE B-3 cell culture supernatant was positive for LMα4, LMα2, LMα1 and LMγ1 subunits (Fig. 1D). The 10-fold concentrated HLE B-3 cell culture supernatant was also used for detection of 11 LM subunits by western blotting, and the results were identical to those obtained using the non-concentrated supernatant (data not shown).

**LM trimer in HLE B-3 cell culture supernatant.** To investigate what type of LM trimers appeared in HLE B-3 cell culture supernatant, IP experiments with antibodies against LMα1-LMα5 for IP, and antibodies against LMβ1-LMβ3 and LMγ1-LMγ3 for western blotting were performed. The immunoprecipitate obtained with antibodies for LMα1, LMα2, LMα3 and LMα5 were negative in western blotting with antibodies for all LMβ and LMγ subunits (data not shown). The immunoprecipitate obtained with antibodies for LMα4 was positive in western blotting with antibodies for LMβ1 and LMγ1 (Fig. 2A), indicating the presence of the LM411 trimer in the HLE B-3 cell culture supernatant. To confirm this result, the immunoprecipitate obtained with antibodies for LMγ1 was also analyzed by western blotting using antibodies against LMα4 and LMβ1, which proved that LMγ1 immunoprecipitates contained LMα4 and LMβ1 (Fig. 2B). These results confirmed that the LM411 trimer existed in the HLE B-3 cell culture supernatant.

**LMs in HLE B-3 BMs.** HLE B-3 BMs were harvested at days 0, 3, 6, 9 and 12, and were analyzed for the expression of LMs by ELISA and western blotting. In LM ELISA, total LM levels in HLE B-3 BMs were similar at days 0, 3 and 6, but decreased to 70% at days 9 and 12 (Fig. 3A). Western blotting of HLE B-3 BMs detected the presence of LMα4, LMα2, LMα1, LMβ3 and LMγ1 subunits (Fig. 3B). Expression levels of LMα4, LMα2 and LMα1 in HLE B-3 BMs were relatively higher at days 3 and 6. LMβ3 expression levels increased with culture time. LMγ1 expression levels fluctuated with time irregularly, with the lowest levels at day 12.
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Figure 1. LMs in HLE B-3 cells. (A) H&E staining of HLE B-3 cells (magnification, x400). (B) ICC of LMs in HLE B-3 cells (magnification, x400). (C) LM ELISA in HLE B-3 cells. (D) Detection of 11 LM subunits in HLE B-3 cells by western blotting. 1, HLE B-3 cell lysate; 2, culture supernatant; 3, culture medium. Arrows and braces indicate the target proteins. H&E, hematoxylin and eosin; ICC, immunocytochemistry; LM, laminin.

Figure 2. LM trimers in HLE B-3 cell culture supernatant. (A) Detection of LM trimers in HLE B-3 cell culture supernatant by LMα4 IP. An antibody against LMα4 was used to precipitate LM trimers and the products were analyzed by western blotting using antibodies against LMβ1-LMβ3 and LMγ1. (B) Detection of LM trimers in HLE B-3 cell culture supernatant by LMγ1 IP. An antibody against LMγ1 was used to precipitate LM trimers and the products were analyzed by western blotting using antibodies against LMβ1-LMβ3 and LMα4. Arrows and braces indicate the target proteins. IP, immunoprecipitation; LM, laminin.
LMs in human ALCs. To investigate LM distribution in human ALCs, H&E staining and IHC for LMs was performed. As depicted in Fig. 4A and B, LMs were primarily distributed in the ALC layer, close to the lens epithelial cells. To evaluate LM expression levels in human ALC, extracts from 2 human ALCs were examined by ELISA and western blotting with antibodies against all 11 LM subunits. In LM ELISA, total LM levels in two human ALCs (N1 and N2 samples) differed by ~3-fold (Fig. 4C), which may have resulted from the differences on age or gender. In western blotting, the N1 sample was weakly positive for LMγ2 and negative for the remaining 10 LM subunits, whereas the N2 sample was positive for LMα4, LMγ2 and LMγ1 subunits and negative for the remaining eight LM subunits (Fig. 4D).

Discussion

The present study provided novel information on the LM composition in HLE B-3 cells, HLE B-3 BMs and human ALCs. The LM411 trimer existed in the HLE B-3 cell culture supernatant. Furthermore, the results of the present study clearly demonstrated that the expression pattern of LM in ALC samples was similar to the expression of BM in HLE B-3.

It has been reported that the LMγ1 subunit was present in the HLE B-3 cell lysate (29). However, whether other LM subunits exist in HLE B-3 cells remains unknown. The results of the present study demonstrated that other than the LMγ1 subunit, nine other LM subunits, including LMα5, LMα4, LMα3, LMα2, LMα1, LMβ3, LMβ2, LMβ1 and LMγ2 subunits, were also present in the HLE B-3 cell lysate. Compared with that in the cell lysate, the total LM level in the HLE B-3 cell culture supernatant was lower, and only 4 LM subunits, LMα4, LMα2, LMα1 and LMγ1, were detected by western blot analysis. The presence of the LM411 trimer in the HLE B-3 cell culture supernatant indicated that LMβ1 should also be expressed in the HLE B-3 cell culture supernatant. In addition, all LMs in the culture
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In the present study, it was also demonstrated that HLE B-3 cells could synthesize BMs rich in LMs for a short period (0-6 days following 95% confluence). HLE B-3 BMs contained LMα4, LMα2, LMα1, LMβ3 and LMγ1 subunits, which were similar to LM subunits in the HLE B-3 cell culture supernatant, except for LMβ3 and LMβ1. Considering the results of the various detection methods for the LM subunits in HLE B-3 lysate, culture supernatant and BMs, the LM411 trimer seems to be the major LM trimer in HLE B-3 BMs, although LMβ1 was not detected by western blot analysis.

Previous ICC studies have demonstrated that LMs are a major component of LECs and ALCs (22,32), results that are consistent with the findings of the present study. Studies focusing on LM subunits in human ALCs are rare; the present study provided evidence that 3 LM subunits (LMα4, LMγ2 and LMγ1) could be detected in human ALCs by western blotting. In the present study, HLE B-3 BM was positive for LMα4, LMα2, LMα1, LMβ3 and LMγ1 subunits, which contained two LM subunits (LMα4 and LMγ1) of the three that were expressed in human ALCs. Therefore, HLE B-3 BM constructed in vitro may represent a substitute for human ALCs for future studies into LM. It is a limitation that only three human ALCs were used in the present study. Further studies using more human ALC samples for evaluation of LM expression levels are necessary to confirm the results of the present study and their similarities with the established BM model of HLE B-3.

The LM411 trimer may be the predominant LM trimer in human ALCs. It is well known that LM411 can promote cell adhesion, cell migration, angiogenesis, tumor invasion and the differentiation of stem cells (33-35). Therefore, LM411 may be involved in the progression of human LEC development, migration and transformation into lens fiber cells, and that abnormal expression of LM411 may contribute to the pathogenesis of cataracts; however, this requires further confirmation.

In previous studies, LECs were mainly used as an in vitro cell model to study the possible pathogenesis of cataracts, with the results indicating that dysfunction of LECs might serve a key role in cataract formation (8-10). To the best of our knowledge, the present study is the first to report the construction of an in vitro ALC model using human LEC cell lines. The in vitro ALC model is advantageous as it allows for the study of the contribution of BM proteins, such as LMs, on the pathogenesis of cataracts. Further studies are required to confirm the present results, validate HLE B-3 BMs as an in vitro ALC model and investigate the possible molecular pathways involving LMs in the pathogenesis of cataracts in vitro and in vivo.

In conclusion, an in vitro human ALC model rich in LMs was successfully constructed using the HLE B-3 cell line, which could be valuable for the study of the molecular biological mechanisms of cataract development and for seeking novel effective drugs for cataract treatment.

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Availability of data and materials
The analyzed data sets generated during the study are available from the corresponding author on reasonable request.

Authors’ contributions
The paper is an original study presenting novel work that has not been published or accepted elsewhere. All authors have seen the manuscript and approved to submit to your journal.

Project design: LX, TX, HT and LP; experimental and data analysis: YY, QH, JH, YH, HT, SL, WX, GH, ZH, LX and TX; manuscript collection: YY, SY and TX; project guidance: QH and LX; experiment and data analysis: YY, QH, JH, YH, HT, SL, WX, GH, ZH, LX and TX; manuscript revision: HT.

Ethics approval and consent to participate
All experiments were performed with the approval of the Internal Review Board of Harbin Medical University and were conducted in accordance with Declaration of Helsinki Principles. Informed consent was obtained from all patients prior to mortality or from their families following mortality for inclusion of autopsy data.

Consent for publication
Written informed consent was obtained for all patients.

Competing interests
The authors declare that they have no competing interests.

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