Original Article 3D visualization of cellular and molecular distributions in human crystalline lenses at different ages

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Abstract: Background: The human lens is a highly organized tissue, and it is constructed of delicate inner architectures that ensure its transparency. However, the pattern of cell distribution in the intact lens has rarely been observed or traced in a three-dimensional (3D) perspective. Methods: Here, we modified and compared three different kinds of tissue transparency methods to investigate the cellular and molecular changes in the human lens at different ages from a 3D perspective. Results: First, we analyzed the general 3D parameters of cleared human lenses from 6 months to 72 years of age and found that the equator proportion remained constant with age ($23.05\% \pm 0.36$). Next, we visualized the cellular distribution patterns in the anterior capsule and equator, as well as the distribution of cortical fiber cells. Interestingly, we observed the accumulation of equatorial epithelium in adolescents and the asymmetrical denucleation of cortical fiber cells in the elderly. Zonula occludens-1 and tropomyosin receptor kinase A were also identified in the pre-equatorial germinative zone, and its presence decreased when comparing lenses of a 17-year-old to those of a 49-year-old. Conclusion: We present a 3D cellular and molecular reconstruction of the human lens, illustrating the observed alterations in human lens epithelial cells across different ages.

Keywords: Lens, optical transparency technology, 3D visualization, cataract

Introduction

The lens is one of the most crucial optical components of human eyes, and it consists of the capsule, epithelial layers, and cortical fibers [1]. The transparency, morphology, and hardness of the lens change along with aging, and under pathologic conditions such changes can lead to diseases like presbyopia and cataract [2-5]. Thus, a particular focus on lens-related research has been how the lens changes over time [6].

Much effort has been made to further understand the morphology and architecture of the lens because these form the basis for maintaining the transparency of the lens [7, 8]. The lens is composed primarily of the outer capsule and inner fibers and undergoes dynamic changes over the lifetime. Epithelial cells are closely packed into the capsule, and these cells denucleate to form fibers. It is widely recognized that the pre-equatorial zone is where new cells germinate [9, 10]. However, direct imaging proof and the detailed cellular distribution patterns have remained elusive as have been the cellular and molecular changes that occur during aging. Various gross geometric features of the lens can be quantified and applied in clinical diagnoses, such as optical coherence tomogra-

phy (OCT), which measures the volume, equatorial diameter, and plane position of the lens [11]. Three-dimensional (3D) in vivo modeling of the lens can be achieved with magnetic resonance imaging [12], and the delicate microscopic and higher dimensional features of the lens are current topics of lens research. Traditional methods, including flat-mounting and confocal microscopy techniques, have been applied to observe the epithelial cells, fiber cells, and capsule as well as to provide more quantitative details like epithelial cell density and fiber shapes and diameters [13, 14]. However, these micro-level observation techniques are incompatible with large-scale 3D views, thus impeding the creation of 3D panoramic views of the cellular arrangement of the lens.

Recently, tissue clearing techniques have been developed to obtain 3D views of the molecular and cellular components of large-scale tissue specimens [15]. The main advantage of these techniques is that they combine clearing, labeling, and imaging and thus allow the specific molecular visualization of the whole organ and the quantitative analysis of its 3D parameters [16]. Previously developed techniques rely on fixation using paraformaldehyde or formalin, but this can cause opacity of the lens, resulting in light scattering [17]. Tissue clearing is the initial process in obtaining 3D images of the entire lens, and the entire mouse eye globe has been effectively rendered transparent in previous studies [18-20]. However, when isolated lenses are removed from their neighboring eye tissues differences in the expansion capacity between fibers and the capsule make it challenging to clear such lenses [21]. In addition, none of the tissue clearing methods have been applied in human eyes, thus emphasizing the possible research value of the current work.

In the present study, we compared the modified tissue clearing methods for human lenses and obtained a 3D map of the lens's cellular characteristics. We observed changes in the expression of zonula occludens-1 (ZO-1) and tropomyosin receptor kinase A (TrkA) across different ages, thus demonstrating that optical transparency technology is useful in constructing a comprehensive molecular and cellular map of the human lens.

Materials and methods

The reagents (including antibodies), equipment, and software are shown in <u>Supplementary</u> Table 1.

Human lens samples

The present study followed the guidelines of the Declaration of Helsinki and was approved by the Ethics Committee of Wenzhou Medical University, Zhejiang, China (2020-132-K-117-02). Lenses from healthy humans without eve diseases prior to death and donated for corneal transplantation were obtained from the Eye Bank of Wenzhou Medical University. Four lenses were obtained from a 72-year-old (72 Y) and 49-year-old (49 Y), while one lens was obtained from a 17-year-old (17 Y) and one from a 6-month-old (6 M). All isolated intact lenses were obtained within 1 to 2 days after the donors died and were put into 4% paraformaldehyde (PFA) solution overnight at 4°C to start the clearing process (Supplementary Table 2).

e-CLARITY (enucleation-Clear Lipid-exchanged Acrylamide-hybridized Rigid Imaging/immunostaining/in situ-hybridization-compatible Tissue hYdrogel) for lens clearing

Enucleation, nucleus delivery, and hydrogel embedding were performed prior to e-CLARITY. These methods were adapted from those widely used in clinical cataract surgery, but in our case, the core of the lens was removed and only the capsule was kept [22, 23]. The posterior capsule of the isolated lens was removed as a 6 mm diameter circle. Then, the posterior cortex and nuclei could be easily sucked out of the lens through the hole left by the posterior capsule (<u>Supplementary Figure 1A</u>). The hollowed-out lens was refilled with 4°C hydrogel solution and incubated in a water bath at 37°C for about 5-6 hours until the hydrogel solidified.

The e-CLARITY process was performed according to previous work [24-26]. The entire procedure was performed on a shaker in a 37°C incubator. Briefly, after removing the extra gel around the lens the lens was soaked in the clearing solution (8% SDS solution/200 mM sodium borate buffer) for about one week until the lens turned clear. The clearing solution was

changed to fresh solution every day in order to prevent bursting of the capsule and to accelerate the clearing process. For immunostaining, the lens was first incubated with primary antibodies against ZO-1 (1:50 dilution, mouse monoclonal, catalog #33-9100, Invitrogen) and then incubated with secondary antibodies (1:100 dilution, Alexa Fluor 488 anti-mouse, AB 2633275, Thermo Fisher Scientific) and 4',6-diamidino-2-phenylindole (DAPI, 1:100 dilution, D9542, Sigma-Aldrich), all of which were diluted in 1× PBS/0.01% bovine serum albumin. These two antibody incubations lasted for 48 h, and the tubes were covered with aluminum foil. The lenses were washed three times with 1× PBS/0.01% bovine serum albumin for 24 h before adding the primary antibody, between the primary and secondary antibodies, and after adding the secondary antibody. Before imaging, the lenses were incubated in Easy Index as the refractive index matching solution for about 48 h.

CUBIC (Clear, Unobstructed Brain/body Imaging Cocktails and computational analysis) for lens clearing

The CUBIC lens clearing was performed according to the protocol of Susaki et al. and our previous work [27-29]. The entire CUBIC process was conducted on a shaker in a 37°C incubator. Briefly, the lens was first embedded in CUBIC clearing solution (50% for 3 hours and 100% for 3 days, consisting of 25% urea, 25% tetraethylenediamine, and 15% Triton X-100) supplemented with propidium iodide (PI, 1:100 dilution, P1304MP, Thermo Fisher Scientific). The lens was then treated with Scale solution (25% urea, 50% sucrose, and 10% triethanolamine) for refractive index homogenization (50% solution until the lens sank to the bottom and then 100% solution until the tissue cleared, usually about 48 h) and then immersed in a mixture of 50% mineral oil and 50% silicone oil for imaging.

iDISCO (Immunolabeling-enabled three-Dimensional Imaging of Solvent-Cleared organs) for lens clearing

The iDISCO process was performed according to the protocol of Renier et al. [30]. The lens was first dehydrated in a series of methanol in PBS (20%, 40%, 60%, 80%, 100%, for 1 h each at room temperature) and incubated in 66% dichloromethane/33% methanol with shaking at 4°C overnight. The lens was then bleached with 5% H₂O₂/95% methanol at 4°C overnight followed by rehydration in a series of methanol in PBS (80%, 60%, 40%, 20%, for 1 h each at room temperature). Before immunostaining, permeabilization and blocking were performed to improve the efficiency of staining. The immunostaining process was the same as for e-CLARITY discussed above, except for the primary antibody being TrkA (1:50 dilution, mouse monoclonal, MA5-15509, Invitrogen) and the washing solution being changed to PTwH solution. The process of methanol dehydration and dichloromethane incubation was also the same as above. Finally, the lens was incubated with dibenzyl ether until cleared, usually taking about 48 h.

Versatile tiling light sheet imaging and image analysis

A versatile tiling light sheet microscope [31] (Nuohai Life Science Co., Ltd.) was used to image the intact transparent lens, and Imaris software (Imaris 9.7, Oxford Instruments PLC, UK) was used to perform the 3D image analysis on our workstation (Intel Xeon W-2245 CPU @ 3.9 GHz, RAM 128 GB, x64 Win10). The original microscope image files contained four light sheet phase positions per layer scan, and these were first combined to make one image for each scan. Imaris File Converter and Imaris Stitcher (Imaris 9.7, Oxford Instruments PLC, UK) were then used to transfer the 2D images series to the 3D file. For quantification, basic Imaris Surpass tree items were used, including surface and spots. All of the data were autocalibrated based on the coefficient of expansion when comparing post-clearing to the original. The surface item was used to reconstruct the anterior capsule, the equator, and the whole lens surface, while the spots item was used to reconstruct the nuclei. Both the surface and spots were built automatically based on the immunofluorescent intensity, with the estimated diameter set to 6 µm to facilitate detection of the nuclei with the spots item.

Basic direct 3D parameters were semiautomatically calculated after imaging according to the restricted definition [32]. In brief, the lens and equatorial thicknesses are the perpendicular distances from the anterior to the posterior of the lens and the equator, and these were calculated as the number of layers multiplied by the distance between the layers. The equator proportion was estimated as the thickness of the equator divided by the thickness of the whole lens. The lens equatorial diameter was the diameter of the equator in the coronal section, and the equatorial plane position was the distance from the anterior pole to the equatorial plane. In addition, the lens surface area and volume were calculated for the whole lens.

Three indirect 3D parameters were automatically calculated after reconstruction in this study, including the number of *spots*, the area and sphericity of the *surface*, and the centroid distance between *spots* and *surface* (<u>Supplementary Figure 1B</u>). The sphericity was defined as the ratio between the surface area (A_1) of a sphere with the same volume as the reconstructed object and the surface area (A_2) of the reconstructed sphere [33], the centroid was defined as the geometric center of an assumed homogeneous mass, and the thickness was calculated semi-automatically as the distance between the sharpest signal edges under the same fluorescence threshold.

Statistical analysis

We evenly divided each sample into four regions for statistical analysis. All 3D parameters were analyzed by GraphPad Prism (v 8.0.2, GraphPad Software, Inc., San Diego, CA, USA) and are presented as means \pm standard errors. Using the Shapiro-Wilk test, the distributions were shown to be normally distributed. One-way ANOVA and Dunnett's comparison were performed to calculate the significance between the groups. Fit spline analysis was applied to fit the trend changes. P < 0.05 was set as the limit for statistical significance.

Results

Human lenses cleared by e-CLARITY, CUBIC, and iDISCO

For the clearing of human lenses, we used modified versions of three different methods of lens clearing based on reported experiences in clearing the entire mouse eye globe, including e-CLARITY, CUBIC, and iDISCO [18-20]. All three methods resulted in the clearing and immunostaining of lenses in less than 17 days (**Figure 1**, Supplementary Figure 2A), among which e-CLARITY took the longest time (17 days) and CUBIC took the shortest (about 11 days). We modified the e-CLARITY process by adding the enucleation step prior to hydrogel embedding and by increasing the concentration of SDS to 8% (Figure 1). For the CUBIC method, we added PI in the CUBIC clearing solution to shorten the time needed for immunostaining. Lastly, in the modification of iDISCO we changed the immunostaining time to 48 h for antibody embedding.

To compare the clearing efficiency of these three methods, we recorded the transparency, deformation, and pigmentation and compared the gross lens diameter after PFA fixation and after clearing. As shown in Figure 2A, the younger the lens, the more transparent it was after being isolated. The lens turned opaque after PFA fixation, and this could be reversed by our optical clearing methods. As far as the transparency was concerned, there was not much difference among the three methods. However, e-CLARITY seemed to cause a slight deformation of the lens, while CUBIC and iDIS-CO caused a small amount of pigmentation. Although the process of fixation and clearing could cause a certain degree of swelling or shrinkage when comparing the post-clearing to original diameter of the lens, both CUBIC and iDISCO showed no significant difference while e-CLARITY underwent severe expansion by about 150% (Figure 2B).

Basic parameter measurements of the lens after 3D reconstruction

To explore the basic measurement efficacy of our lens-clearing methods, we first analyzed the gross geometric measures and compared them with other methods, including OCT and digital shadow photogrammetry. Based on the even expansion and calibration of the 3D reconstruction, we could calculate several 3D parameters and could perform further microlevel analyses. Seven semi-automatically calculated parameters regarding thickness, diameter, and volume are shown in **Table 1**. Due to the high spatial resolution of the reconstruction, the sub-structure of the lens could be determined, including the equatorial region. Most of the parameters increased with age except for the equatorial proportion, which



Figure 1. Timeline for the e-CLARITY, CUBIC, and iDISCO processes. The overview of the timeline and main reagents for e-CLARITY, CUBIC, and iDISCO. One sector represents one day, so the total processing time for each method was 17, 11, and 15 days, respectively. Each method consisted of clearing (blue highlight), immunostaining (yellow highlight), and refractive index matching (purple highlight) to achieve optical clearing, followed by versatile tiling light sheet imaging.

remained constant at $23.05\% \pm 0.36\%$. When compared to other methods such as OCT and digital shadow photogrammetry, most of the parameters calculated by lens-clearing methods fit well with the other two methods [32, 34], including lens thickness, equatorial diameter, and lens column, thus supporting the feasibility of our method.

3D map of cellular changes across different ages

In order to further explore the 3D cellular architecture of the lens, we traced the cellular changes in the lens during aging by reconstructing and analyzing the nuclear signal. The nuclei of all lens cells were stained by DAPI in the iDIS-CO method and by PI in the CUBIC method (**Figure 3A**, **3B**), and the equatorial region was segmented based on the morphology characterized by the largest diameter in the coronal plane together with the equatorial front part that was defined as the anterior capsule region (<u>Supplementary Figure 2B</u>). In addition, the cortex region was defined as the subepithelial part. From a 3D view, the lens cells could be

roughly divided into three parts according to their spatial position, including the anterior capsular epithelium, the equatorial epithelium, and cortical fiber cells. Some parts of the anterior capsule were peeled off, possibly due to the shaking during the clearing process. All three groups of cells grew along with the ageincreasing diameter of the lens, especially from the 6 M to the 17 Y lenses (Figure 3A, 3B). In addition, the density of the equatorial epithelium reached its peak in the 17 Y lens. Interestingly, we found that the cortical fiber cells gathered together to form an inner ring just beneath the epithelium in the 6 M lens, and scattered in 17 Y and 49 Y lenses (Figure **3A**, **3B**). Finally, in the 72 Y lens half of the lens had lost the cortical fiber cell nuclei (Figure 3A, 3B).

Furthermore, we divided the lens into quarters to reduce sampling errors in the statistical analyses (<u>Supplementary Figure 1B</u>). Nuclei were reconstructed automatically using the Imaris *spots* item, as shown by the correlation between the reconstructed pink spots and the PI immunostained nuclei. Based on the quar-



Figure 2. Human lens clearing efficacy of e-CLARITY, CUBIC, and iDISCO. A. The gross morphology of randomlychosen lenses immediately after isolation (the top panel), after PFA fixation (the middle panel), and after clearing (the bottom panel), roughly showing the changes in size and degree of transparency. B. The coefficient of expansion of the three clearing methods after fixation and after clearing and as compared to the original. n = 3/group, ***P < 0.001 comparing within one method under different conditions; ^{###}P < 0.001 comparing among the different methods under the same conditions.

tered lens, four points were referred to as 3, 6, 9, and 12 o'clock located where the split line intersected with the edge of the lens. We quantified the number of nuclei, the surface area, and the sphericity of the anterior capsule and equator and the centroid distance between the nucleus and the capsule (Figure 3C-F). The number of nuclei in all three groups and the

Geometric parameter	Lens optical clearing methods in this experiment				OCT [32]	Digital shadow Photogrammetry [34]
	6 M	17 Y	49 Y	72 Y	0-56 Y	0-88 Y
LT (mm)	3.6	3.7	4.3	4.1	3.5-5.5	3.0-6.0
ET (mm)	0.84	0.84	1.01	0.93	NA	NA
EP (%)	23.33%	22.70%	23.49%	22.68%	NA	NA
DIA (mm)	7.3	9.2	9.3	11.0ª	5.5-10.0	6.0-11.0
EPP (mm)	1.12ª	1.68	1.67	1.98	1.4-2.5	NA
LSA (mm ²)	113	177	247ª	265ª	80-200	NA
VOL (mm ³)	93	158	250	294ª	50-250	NA

 Table 1. Comparison of basic gross geometric parameters between the lens clearing methods and
 OCT and digital shadow photogrammetry

^aBeyond the estimated range of OCT. LT, lens thickness; ET, equatorial thickness; EP, equatorial percentage; DIA, lens equatorial diameter; EPP, equatorial plane position; LSA, lens surface area; VOL, lens volume; NA, not applicable.

areas of the anterior capsule and equator changed significantly with age along with the centroid distance between the equatorial epithelium and the lens surface, especially from 6 M to 17 Y. In contrast, the centroid distance between the anterior capsular epithelium and cortical fiber cells and the lens surface, together with the sphericity, showed no significant differences among the different ages. Notably, the large decrease in anterior capsular epithelium and cortical cells (41.59% and 91.09%) occurred along with a large increase in the equatorial epithelium (150.70%), which resulted in a decrease in the total number of lens cells when reaching adolescence (**Figure 3C**).

Spatial distribution of ZO-1 in lens across different ages

To further understand the intercellular adhesion between lens cells, we determined the expression of ZO-1 in the 17 Y and 49 Y lenses (Figure 4A, 4B). ZO-1 was mainly distributed at the interface between the capsule and epithelium fibers together with some expression in the cortical fiber cells extending from the anterior capsule to the frontal portion of the posterior capsule. Focusing on its distribution in the posterior capsule, ZO-1 in the 17 Y lens extended farther than in the 49 Y lens (Figure 4a, 4b, Supplementary Figure 2C), and ZO-1 covered 66.7% of the frontal part of posterior capsule in the 17 Y lens but only 22.2% of the frontal part in the 49 Y lens. In both the 17 Y and 49 Y lenses the expression of ZO-1 in the direction of 3, 6, 9, and 12 o'clock grew thicker from the central to the peripheral zone beneath the anterior capsule and peaked just before the equatorial

region, while it gradually became thinner beneath the posterior capsule. The maximum thickness of the ZO-1 signal in the 17 Y and 49 Y lenses was 30.67 \pm 3.7 μ m and 38.40 \pm 5.94 µm, respectively, while the thickness of the nuclei was $41.58 \pm 3.62 \,\mu\text{m}$ and 51.70 ± 12.74 µm, respectively. In addition, DAPI marked the epithelium and denucleated fiber cells closely surrounding the cells expressing ZO-1 and showed a similar pattern as ZO-1 expression (Figure 4C, 4D). We also observed the changing thickness of DAPI and TrkA in the 6 M lens (Figure 4e) and found a similar pattern as ZO-1 in the 17 Y and 49 Y lenses, except that TrkA reached its thickest point just behind the equator (Figure 4E, 4F, Supplementary Figure 2D). Furthermore, we reduced the interval between the measuring planes to about 20 µm for the equatorial observation, and DAPI, TrkA, and ZO-1 all showed a slight decline from the front to the back in all lenses (Figure 4G, 4H). A similar pattern between TrkA, ZO-1, and DAPI from the anterior to the posterior indicated the close relationship between ZO-1, TrkA, and the epithelium. The slight decrease in ZO-1 and DAPI signals within the equatorial region indicated that ZO-1 in the anterior capsule peaked before the equator (Figure 4G, 4H). When comparing the three age groups, it was observed that the younger the lens, the thinner the capsule within the equatorial region (Figure 4H).

Discussion

The human lens is a highly organized structure that remains transparent throughout the lifetime, and understanding its spatial cellular and molecular features will improve our ability to

3D mapping of human lenses across different ages



3D mapping of human lenses across different ages

Figure 3. 3D cellular maps of the human lens across different ages. 3D cellular maps of (A) 6 M and 17 Y lenses stained by DAPI (dark blue) and (B) 49 Y and 72 Y lenses stained by PI (light blue). The first four columns show the frontal view, including the original signal, the reconstructed anterior capsular epithelium, the equatorial epithelium, and the cortical fiber cells. Among these, in the second column the color represents the distance from the origin with red being farther away and purple being closer. The last column shows the side view of the original signal. (C-F) Cellular 3D parameters are shown as a statistical graph comparing the different ages and different lens parameters, including the number of nuclei, the area, the centroid distance, and the sphericity. Data for the 6 M and 17 Y lenses were obtained using iDISCO, and data for 49 Y and 72 Y lenses were obtained using CUBIC. n = 4/group, *P < 0.05, **P < 0.01, ***P < 0.001 compared to the 6 M lens.



Figure 4. 3D distribution of ZO-1 and TrkA in the whole human lens. Representative images of the anterior coronal plane at a diameter of 6 mm and 9 mm showing ZO-1 (green) and DAPI (blue) in (A) the 17 Y lens and (B) the 49 Y lens, with continuous thickness changes being repeatedly calculated at 3, 6, 9, and 12 o'clock from the anterior to posterior of the lens (white, yellow, and gray represent the anterior capsule, equator, and posterior capsule region, respectively) in (C) the 17 Y lens and (D) the 49 Y lens. Representative continuous imaging from the anterior to posterior (with a measuring plane interval of 1 mm) of ZO-1 and DAPI at 9 o'clock is shown for (a) the 17 Y lens and (b) the 49 Y lens. (E) Representative images of the anterior coronal plane at diameters of 5 mm and 7 mm showing TrkA (nerve growth factor receptor, yellow) and DAPI (nucleus, blue) in the 6 M lens. (F) Continuous thickness changes of TrkA (yellow) and DAPI (blue) were repeatedly calculated at 3, 6, 9, and 12 o'clock from the anterior to posterior (white, yellow, and gray represent the anterior capsule, equator, and posterior (with a measuring the anterior capsule, equator, and posterior capsule, respectively) as shown in the 6 M lens. (e) Representative continuous imaging from the anterior capsule to the posterior (with a measuring plane interval of 1 mm) of TrkA and DAPI at 9 o'clock in the 6 M lens. Furthermore, fit spline analyses of (G) ZO-1 (green) and TrkA (orange) and (H) DAPI (blue) are shown within the equator from the anterior to posterior (with a measuring plane interval of 20 μ m). n = 4/group.

restore transparency under pathologic conditions. Our study first optimized three different methods to clear the human lens in terms of the time required and the labeling efficiency. Through 3D reconstruction and statistical analysis of the nuclear distribution within the capsule, equator, and cortex, we could observe the cellular and molecular characteristics of the whole lens. ZO-1 and TrkA had similar distribution patterns as the epithelium, indicating their close relationship.

Tissue clearing methods have recently been developed and widely applied in studies of various organs, and these methods mainly consist of clearing, immunostaining, and microscope imaging procedures [16, 35]. All of the techniques have primarily benefitted from advances in microscopy resulting in high-speed image acquisition and minimal photobleaching together with the big data computing and analysis systems that are necessary for terabyte-level image reconstructions and calculations [36, 37]. The biggest breakthrough in tissue clearing methods was by no means just the 3D visualization, but also the ability for 3D reconstruction, quantification, and modeling [38-40]. Although there are numerous mature clinical optical examination techniques such as OCT, Purkinje imaging, Scheimpflug imaging, and digital shadow photogrammetry [34, 41-43], as well as non-optical techniques such as X-ray Talbot interferometry, magnetic resonance imaging, and ultrasound biomicroscopy [44-46], their inability to support in-depth microlevel exploration limits their usefulness in understanding the micro-world buried beneath the opaque tissue. Thus, tissue clearing methods have been developed to make up for these clinical shortcomings and to more deeply explore the components of the lens.

Comparing the three different kinds of clearing methods, we found iDISCO to be the most suitable for the clearing and immunostaining of the human lens. We first applied the traditional CLARITY method based on our previous experience [24]. However, due to the weaker water absorption and expansion capacity of the capsule compared with fibers, together with the intensified expansion due to the long SDS clearing process (about 7 days), the lens tended to burst easily during the SDS clearing step (data not shown). Thus, in order to balance the inner pressure caused by fibers and the outer pressure caused by the capsule, we added the enucleation step before embedding in SDS. We modified the e-CLARITY procedure to include splitting the posterior capsule and sucking out part of the nucleus of the lens in order to balance the internal and external pressure. Our e-CLARITY data showed a great enlargement of the sample, indicating its potential use in magnifying and detecting more delicate structures, theoretically similar to the Magnified Analysis of the Proteome technique [47, 48]. Nevertheless, the splitting of the posterior capsule and enucleation of the nucleus destroys the integrity of the lens. To avoid swelling of the tissue, we next tried the organic clearing methods CUBIC and iDISCO. In the CUBIC process, we added the PI directly with the CUBIC clearing solution in order to reduce the immunostaining time, and in iDISCO we modified the antibody incubation time. Adding PI to the CUBIC clearing solution allowed us to skip the washing step for the immunostaining. iDISCO has a penetration step, so we reduced the staining time based on our lens samples as appropriate. Both PI and DAPI can bind to DNA, the only difference being the lack of cell membrane penetrability for PI [49, 50]. Owing to the membrane

penetration treatment in the whole clearing process, the entire lens nucleus could be stained. Both the CUBIC and iDISCO methods performed well in the human lens, except for slightly better transparency with the iDISCO method. Due to the specific penetration step, iDISCO was also more appropriate for immunostaining using a wide range of different markers.

Our methods could determine not only the parameters that are already being applied in clinical practice, but also several other 3D cellular parameters. In addition to well-accepted parameters like lens thickness, lens volume, lens equatorial diameter, and the radius of the curvature of the lens [32, 34, 42, 44, 51], we also defined the equatorial thickness and the equatorial proportion. We observed a constant equatorial proportion of around 22-23% in both young and aged lenses, but whether this value might change under pathologic conditions needs further clinical investigations. Moreover, the previously reported slight variation in the ratio of the equatorial height to the lens thickness across different ages, which agrees with our finding of a constant equatorial proportion [52].

We also reconstructed the position of each lens cell together with the capsule morphology to support the cell development theory of the lens. The human crystalline lens has a high degree of spatiotemporally organized tissue [53], and an increase in mass and a flattening of the anterior capsular curvature occurs during aging together with increased light scattering and spectral absorption [54]. It is believed that the anterior capsule is the only metabolically active part of the lens, with the germinative zone located just pre-equatorially where cells undergo mitosis, migrate towards the equator, and differentiate into fiber cells [55]. The newly-differentiated fiber cells elongate and become densely packed surrounding the older fiber cells, and they show upregulated expression of α -, β -, and y-crystallins along with degradation of their nuclei and organelles, thus maintaining the continuous growth and transparency of the lens [56, 57].

With regards to changes in the number of lens cells across different ages in our experiment, the following distribution patterns were observed: 1) the anterior capsular epithelium in

the 6 M lens was almost one-and-a-half times as thick as in the 72 Y lens, and the number of cortical fiber cells was almost 10 times greater, 2) the accumulation of equatorial epithelium peaked at around 17 Y, and 3) there was a slight decrease in the total number of cells in the whole lens and an increase in the number of cortical cells in the elder sample. Our results for lens cell numbers and spatiotemporal distributions from different ages (Supplementary Video 1) suggest that the infant lens is rich in cells in the anterior capsular epithelium and equatorial epithelium, especially denucleated cortical fiber cells. In the progression to adolescence, many epithelial cells in the germinative zone underwent mitosis and migrated to the equator, but only a relatively small part of the equatorial epithelium denucleated and packed into the cortex, which could explain the intense increase in equatorial epithelium and the decrease in the anterior capsular epithelium and the decrease in cortical fiber cells.

Another aspect of our study focused on ZO-1, which is associated with tight junctions and is mainly located at the epithelium-fiber interface [58]. In addition, ZO-1 can interact with various proteins and can reorganize the membrane domain and signaling pathways and thus plays a role in various cataract-related pathologic conditions like Nance-Horan syndrome [59, 60]. In our experiment, ZO-1 was mainly distributed at the epithelium-fiber interface (Supplementary Video 2). In our continuous observation of ZO-1 from the front to the back of the lens, we found that in both the adolescent and middle-aged lenses the expression of ZO-1 reached its maximum just before the equator, together with the densest nuclei, indicating the location of the germinative zone. Moreover, both the epithelium and ZO-1 became thicker in the older lenses. Another interesting thing about ZO-1 was that in the 17 Y lens ZO-1 expression extended farther towards the posterior capsule than the epithelium, partially due to the incomplete packing process of denucleated cortical fiber cells. TrkA expression in the 6 M lens followed a similar pattern and might also be partially explained by the incomplete denucleated packing process. The lack of overlap of ZO-1 and TrkA with the epithelium is an indication of the immature lens cortex during early life. Because TrkA is the receptor for nerve growth factor [61], its distribution as shown here suggests a role in the trophism and growth of the epithelium, and this needs further study at the molecular level.

There are several limitations to this study. First, due to the need to obtain fresh lenses without freezing from healthy individuals without ocular diseases, which is a rarity in young and middleaged populations, this research is constrained by a limited sample size and lack of young subjects. We specifically selected lenses representing infancy (6 months), adolescence (17 years), middle age (49 years), and the elderly (72 years). Additionally, due to our sample size limitations, staining of TrkA and ZO-1 was not performed in the aged lenses. Finally, the fibrous tissue might still cause light scattering even when it looks transparent to the naked eye, and an additional quenching step might be applied to avoid false-positive signals.

Conclusion

The three modified tissue clearing methods of e-CLARITY, CUBIC, and iDISCO were successfully applied to the human lens. Subsequent 3D cellular reconstruction revealed the cellular and molecular characteristics in the human lens across different ages together with the molecular distribution of ZO-1 and TrkA.

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Disclosure of conflict of interest

None.

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PTwH
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DMSO D8418 Sigma-Aldrich
Tween-20 P9416 Sigma-Aldrich
Heparin 375095 Milipore
Dibenzyl Ether 33630 Sigma-Aldrich
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DAPI D9542 Sigma-Aldrich
PI P1304MP Thermo Fisher Scientific
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Constant temperature incubator 7HLY-1803 Shanghai Zhichu Instrument. China
Light sheet microscope IS-18 Nuohai Life Science Co., Ltd., China
Workstation W-2245 Intel
Software
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GraphPad Prism version 8.0.2 GraphPad

Supplementary Table 1. Reagents, antibodies, equipment, and software

3D mapping of human lenses across different ages

Sample	Tissue clearing method	Immunostaining methods		
6 M	iDISCO	DAPI, TrkA		
17 Y	iDISCO	DAPI, ZO-1		
49 Y #1	CUBIC	PI		
49 Y #2	iDISCO	DAPI, ZO-1		
72 Y #1	CUBIC	PI		
72 Y #2	e-CLARITY	DAPI		

Supplementary Table 2. Tissue clearing and immunostaining methods for each individual lens



Supplementary Figure 1. Pretreatment with enucleation and subsequent e-CLARITY 3D-reconstruction. A. Pretreatment with enucleation in the e-CLARITY process. 1 - the isolated lens after removing the central 6 mm of the posterior capsule; 2 - the hollowed-out lens; 3 - the lens filled with hydrogel; 4 - sealing the lens in the Eppendorf tube with hydrogel; 5 - degassing with a vacuum desiccator; and 6 - solidified hydrogel containing the lens. B. Cellular 3D parameters are shown as a schematic diagram of definitions comparing the different ages and different lens parts, including the number of nuclei, the area, the centroid distance, and the sphericity.



D



Supplementary Figure 2. Representative 3D views of the molecular and cellular maps of the human lens. A. Randomly chosen planes magnified to show the comparison of nuclear immunostaining efficacy (dark blue for DAPI and light blue for PI) using the different clearing methods. B. Reconstructed surface of the anterior capsule and equator of the 6 M, 17 Y, 49 Y, and 72 Y lenses. C. 3D view of ZO-1 (left) together with three views from the front, side, and top of ZO-1 (middle) and DAPI (right) in addition to the top panel for the 17 Y lens and the bottom panel for the 49 Y lens. D. 3D view of TrkA (left) together with three views from the front, side, and DAPI (right) in the 6 M lens.