Local structure of human hair spatially resolved by sub-micron X-ray beam

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Human hair has three main regions, the medulla, the cortex, and the cuticle. An existing model for the cortex suggests that the α-keratin-based intermediate filaments (IFs) align with the hair’s axis, but are orientationally disordered in-plane. We found that there is a new region in the cortex near the cuticle’s boundary in which the IFs are aligned with the hair’s axis, but additionally, they are orientationally ordered in-plane due to the presence of the cuticle/hair boundary. Further into the cortex, the IF arrangement becomes disordered, eventually losing all in-plane orientation. We also find that in the cuticle, a key diffraction feature is absent, indicating the presence of the β-keratin rather than that of the α-keratin phase. This is direct structural evidence that the cuticle contains β-keratin sheets. This work highlights the importance of using a sub-micron x-ray beam to unravel the structures of poorly ordered, multi-phase systems.

While Human hair has been studied by x-ray diffraction and electron microscopy for many years1–9, a complete picture has been elusive because hair is weakly scattering, disordered, and heterogeneous. Human hair is a hierarchical structure, largely constituted of intermediate filaments (IFs) that in turn are made up of keratin molecules10. Hair has three main regions: (1) the medulla on the central axis, (2) the cortex, and, (3) the cuticle at the exterior of the hair, (Fig. 1A,B). The medulla mainly is filled with a keratin matrix partially composed of IFs that are spatially and orientationally disordered. The majority of the cortex has a dense packing of IFs aligned along the axis of the hair, but are fluid-like in two dimensions in the plane perpendicular to the hair’s axis. The cuticle is well organized in a layered 1D-ordered structure.

Fraser and colleagues proposed a molecular structural model of α-keratin and the IF in 1959 and 196410,11, but there is still controversy about the exact structure of the IFs that are a major component of hair2,3,12. In most x-ray measurements, many hair fibers were bundled together so as to assure significant signal-to-noise ratio2,7,13. The advent of high brightness synchrotron sources has enabled the usage, in a few experiments, of an x-ray micro-beam on single strands of hair. However, given the typical sizes of x-ray beams (over 2 microns) relative to the diameter of hair (60–140 microns), and the scattering geometry (x-ray perpendicular to hair’s axis), it was difficult to isolate scattering from the different regions5,12,14–16.

To spatially resolve the hair’s macro regions, we employed a sub-micron x-ray- diffraction beam of 300 nm in the high-resolution scanning direction, and 20 μ in the orthogonal direction. The samples were cross-sectional slices of hair of 30 μ thickness, with a diameter 80 μ; they were studied with the x-ray beam || and ⊥ to the hair’s axis.

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Our main discussion here will focus on the results of high-resolution SAXS/WAXS. However, we also present our STEM imaging that is complementary to, and provides a context for, the discussion of the SAXS/WAXS data. Further details on the STEM imaging is given at the end of the discussion section.

We note that hair and wool have been extensively studied by electron microscopy and one of the first high-resolution electron microscopy studies was reported by Rogers 1959\textsuperscript{17,18}, clearly showing the packing arrangement of the IFs in the cortex and cell membrane complex in the cuticle. More recent results have been obtained by TEM and tomography studies\textsuperscript{8,19}.

**Results and Discussion**

**Small angle x-ray scattering.** In Fig. 1B, we show a representation of the diffraction data obtained from the cross-section of a hair fiber, wherein we spatially positioned the Small Angle X-ray Scattering

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**Figure 1.** SEM image of a human hair fiber is shown in (A) with schematic overlays on to the cut face of the hair fiber to emphasize the relative size of the hair and the main hair regions. In (B), we spatially position the SAXS diffraction patterns relative to each other, so that the locations match the position of the x-ray beam on the sample, thus generating a SAXS “map” of human hair. This map allows us to see the spatial variation of the SAXS patterns from a 30-micron- thick cross-section of hair. The incident x-ray beam is \( \parallel \) to the hair axis. SEM zoom of selected cross section from (A) is overlapped to the SAXS map showing the real-space cross section of the cuticle and cortex. Magnified views of the characteristic SAXS patterns are shown in (C) for the medulla, (D) for the cortex, and in (N) for the cuticle. (E,F) are for the newly discovered intermediate zone. Corresponding Scanning Transmission Electron Microscopy (STEM) images shown in (G,H,J,L).
(SAXS) part of each diffraction pattern as it was taken, so allowing us to see the evolution of the IF structural arrangements within each region of hair. We superimposed the boundaries of the different regions that are identified by the SAXS pattern. The magnified images of the characteristic SAXS patterns for the medulla, cortex and cuticle are shown respectively in Fig. 1C,D and 1N. In Fig. 1E,F, we show a SAXS diffraction pattern for a previously unidentified region, an “aligned cortex” region, which we elaborate on below. Complementary Scanning Transmission Electron Microscopy (STEM) images for these regions are shown in 1G, 1H, 1J and 1L.

Both the SAXS data and the STEM images were obtained from different slices from the same hair strands, but prepared differently for the different methods, (see method section). We emphasize that we present the STEM images to give an indication of the real-space morphology of a given hair region from which the diffraction data is obtained, to help the reader interpret the observed diffraction pattern. The STEM images are discussed later.

The characteristic SAXS for the medulla region, shown in Fig. 1C, is a symmetric pattern that is monotonically decreasing in intensity with momentum transfer. The matching real-space image, 1G, shows that the medulla is spatially and orientationally disordered, viz., consistent with the SAXS pattern. The characteristic SAXS for the cortex in Fig. 1D, shows that intensity does not decrease monotonically, but instead has an isotropic ring at 94 Å, representing the averaging of randomly oriented nearest-neighbor correlations between IFs. The matching real-space electron-microscopy image, 1H, shows the cortical cells (dark grey contrast) and the random network of walls between cortical cells (light grey contrast). Electron microscopy does not have the spatial resolution to show individual IFs. The cuticle (Fig. 1N,L) is made up of macro layers of about 0.5 μ thick as is shown in the real-space electron-microscopy image in Fig. 1L, with a total thickness of cuticle, in our case, of about 3–4 μ. Here, the SAXS pattern in Fig. 1N shows an array of x-ray diffraction peaks typical of a layered structure aligned with the walls of the hair. While the layering observed in both the electron microscopy and the SAXS data are orientationally aligned, in this experiment the two techniques have mutually exclusive spatial-resolutions.

We now address the newly observed “aligned cortex” region. As we move the x-ray beam from deep inside the cortex towards the cuticle, the SAXS pattern changes from an isotropic 94 Å ring with uniform intensity around the ring (Fig. 1D), into one in which the intensity becomes anisotropic, as shown in Fig. 1E. The evolution of the diffraction pattern from Fig. 1D to Fig. 1E is a continuous function of position. Our interpretation of the intensity of the anisotropic ring is that the angular-probability distribution of next nearest neighbor bonds is influenced by their proximity to the cuticle-cortex interface. As we continued to move the x-ray beam closer to the interface, a new set of peaks appeared, consistent with the layering (210 Å and 111 Å), shown in Fig. 1F. These new peaks are precursors to the structure in the cuticle, but coexist with the 94 Å ring with an anisotropic intensity distribution.

In Fig. 2, we show diffraction images with better counting statistics for the cuticle (Fig. 2A,B) and cortex (Fig. 2D,E). We observed that, in addition to the primary ring at 94 Å, there are weaker, broader rings at 53.5 Å, 46.2 Å and 31 Å. This is consistent with 2D fluid-like disorder of close packed aligned IFs, probably with a short-range hexagonal, para-crystal arrangement.20,21

Line projections of the cuticle and cortex region are shown in Fig. 2C,F. In the cuticle, we observe up to 5 diffraction peaks with d spacings of 210.2 Å, 111.6 Å, 71.9 Å, 54.5 Å, 42.3 Å, which indicates a rearrangement of the keratin molecules not typical of the α-phase. A well-ordered, layered structure is more indicative of β keratin, as we confirm below.

To compare with previous publications, we took diffraction images with the beam ⊥ to the hair axis, as depicted in Fig. 3A–C. STEM images of the cortex and cuticle, cut in a longitudinal plane are shown respectively, in Fig. 3D,E. The SAXS from the cortex (Fig. 3B) shows two peaks at ±qα on an axis that is ⊥ to the direction of the hair’s axis. Relative to this axis, the cuticle peaks that are at the top of the hair (Fig. 3A) and bottom of the hair (Fig. 3B) are offset, respectively, by 4.6 ± 0.1 degrees and −4.5 ± 0.1 degrees. These angular offsets have been correlated with the orientation of the cuticle scales relative to the hair axis, as viewed from the ⊥ direction (SEM image in Fig. 3F). Other than these angular offsets, the layer spacing of the cuticle in Fig. 3A,C are consistent to those observed in Fig. 2B. Clearly, the SAXS data from the || and ⊥ directions show (Fig. 1N,A,C) that the cuticle is layered on the 210 Å length-scale, while electron microscopy (Fig. 1L) shows that it is layered on the 5000 Å length scale.

Wide angle x-ray scattering. As shown in Fig. 4, wide angle x-ray scattering (WAXS) with the x-ray beam ⊥ to the hair axis, reveals structural information internal to the IFs. Three features are shown; a broad anisotropic 10 Å peak, a broad isotropic 4.6 Å ring, and a sharp arc at 5.2 Å. In Fig. 2A,D where the incident x-ray beam is || to the hair’s axis, we can compare the 10 Å ring in the cuticle, and cortex. Clearly, the 10 Å feature is stronger in the cortex. The 10 Å ring reflects the correlations between keratin-coiled coils.21,22

Comparing the WAXS patterns of the cortex (Fig. 4A) and the cuticle (Fig. 4B) more carefully as is done in Fig. 4C, we find that the sharp 5.2 Å peak, which is characteristic of the α keratin, is present in the cortex but absent in the cuticle. In contrast, the broad 4.6 Å feature simply is less intense in the cuticle, but its width is unchanged between the two regions. The absence of the 5.2 Å peak, and a weakening of the 10 Å peak are characteristic of a transition of α to β keratin. These results corroborate suggestions from Raman scattering,23 viz., that the cuticle has β keratin. Considering the layered morphology of
the cuticle, and the known propensity of β keratin to form layered structures, it is reasonable to assume that the cuticle contains β keratin sheets.

An existing model on cuticle structure obtained primarily from SAXS data is reported in the literature5,12, but this model is incomplete because it does not incorporate WAXS measurements. In this model it is proposed that the layered diffraction peaks come only from the complex of the cuticle membrane and that the rest of the cuticle is amorphous5,12. The WAXS data we have presented here indicates that a significant fraction of the cuticle is constituted of β keratin. While a smaller x-ray beam and more study is needed to resolve the structure of cuticle sub-regions, such as the cuticle membrane complex, exocuticle or endocuticle, our results suggest that the previous model based only on SAXS data taken in orthogonal geometry needs to be reconsidered.

Scanning Transmission Electron Microscopy. As mentioned above, the STEM measurements were primarily included to show the real space morphology of the different macro regions of hair. Below, we discuss some details of the STEM images.

Cross sections. In Fig. 1, we show the cross sections of the cuticle, cortex, and medulla. Figure 1G shows the medulla with a visible macro porous section. This empty porous part is a result of preparing the sample. The keratin matrix liquid, that normally fills the porous part, has evaporated during the preparation of the 60-nm thick sample slice. Figure 1H shows the cortex, the main visible components of which are 500-nm macro fibrils and circular black objects which are the hair’s pigment, melanin. There also are some gray zones, with different contrast due to the different orientation of IFs in the zones8,17. In Fig. 1J we show the intermediate zone of the cortex/cuticle, wherein we can observe the alignment and accumulation of the cell walls oriented by the hair’s curvature. Finally, in Fig. 1L, we show the cuticle’s macro layers with very high contrast in the layers. Different gray contrast and stratification is visible. It is known that hair contains and accumulates minerals in the cuticle, and we suspect that the minerals are represented by some of the lighter contrast, compact shapes in Fig. 1L, but these are beyond the scope of this work.

Longitudinal sections. Figure 3D to 3E are the STEM images of slices cut longitudinally along the hair’s axis. The intermediate zone at the cuticle/cortex interface clearly is visible (Fig. 3D). We also emphasize
here that the longitudinal section (Fig. 3E) of the cortex looks quite different from its cross-section (Fig. 1H). The macro-fibrils clearly are visible in cross section (Fig. 1H), but not as obvious as in the longitudinal section.

In the longitudinally cut cortex region (Fig. 3E) we see that hair pigments (melanin) with sizes of about 1- to 2-microns are grouped in long clusters and align with the long hair’s axis.

Conclusions. By using a sub-micron x-ray beam, we were able to resolve the structure of the regions of human hair. Within the cortex, we found a new region near the cuticle where the local arrangement of the IFs are oriented by the wall. Additionally, we present direct structural evidence that the cuticle of human hair contains $\beta$-keratin sheets. This study highlights the importance of a sub-micron x-ray beam in unraveling the structures of poorly ordered, biological multi-phase systems. We anticipate that smaller x-ray beams will improve our understanding of hair, and similarly complex systems.

Methods
Sample preparation. Long hair strands were cut from the back of the head from males between 25–30 years old of Japanese origin. The hair is virgin and never had been treated with any reagents except for a normal wash. All slices for the experiments are cut far from follicles or the end of the hair, in order to get the healthy part of the hair.

For embedded hair samples, long hair fibers were placed horizontally in a mold on a flat silicon-wafer that keeps the hair centered in the middle of the mold and leveled. No stretching, pressure, or deformation is applied to the hair. Resin (Pelco Eponate 12 with BDMA) is poured over the hair and allowed to solidify. The embedded samples were left at room temperature to solidify. Just before the measurements, the hair was sliced to 60nm thickness using an ultramicrotome (model: RMC Boeckeler PowerTome X) for STEM measurements, and to 30 microns thickness by microtome (model: Leica RM 2135) for x-ray measurements. We used a MT-Diamond blade made by Pelco with a knife angle of 45 degrees. All samples are examined by optical- and SEM-microscope before taking the x-ray measurements.

X-ray measurements. X-ray measurements were performed in air, at room temperature, with the relative humidity around 48%, at the X13B beamline at the National Synchrotron Light Source.
at Brookhaven National Laboratory. This beamline is an undulator-based beamline with a gravity-fed water-cooled monochromator for enhancing the beam’s stability. The monochromatized photons are incident on a Kinoform lens, which creates a focused x-ray beam of 300 nm by 20 μ on the sample. The sample is positioned with a six degree angle of freedom stage (Smarpod 110.45), which, in turn, is placed on top of a high precision rotary air-bearing stage (MICOS UPR-120Air). Consequently, the sample can be translated in three degrees of freedom (3 DOF) with 100-nm resolution, and rotationally oriented with 0.001 degree resolution in with 3 DOF. Downstream of the sample is a beam-stop constructed of a 4-mm-long cylinder of Tungsten with diameter 500 microns. Downstream of the beam stop, there is a choice of two detectors, a visual detector that is used to position the hair in the direct x-ray beam, and a large area X-ray CCD from Princeton Instruments for quantitatively measuring the X-ray diffraction data. The visual detector system consists of a 100-micron thick YAG crystal, viewed by a microscope objective. The CCD has a diagonal dimension of 165 mm, with a 56-micron pixel size.

**Figure 4.** Comparison of the wide angle x-ray scattering of cortex shown in (A) and cuticle shown in (B) measured with x-ray beam ⊥ to the hair’s axis. (C) showing the line projections of the integrated intensities for the peak 5.2 Å and the peak 4.6 Å of the two respective regions.
Relevance of the micro-beam SAXS methodology. Hair has a heterogeneous arrangement of sub-components with slightly different structures. A traditional SAXS pattern, with a large x-ray beam, results in a sum of diffraction SAXS patterns, weighted by the volume fraction of the different regions. In the case described here, where the volume of the cortex is much larger than that of the cuticle, the traditional SAXS approach in the past has obscured the cuticle’s signal. By providing an x-ray beam that was small enough to select just the cuticle region, we were able to observe the absence of the relatively weak signature of the α-keratin phase in the cuticle region.

**STEM and SEM measurements.** The SEM and STEM imaging were performed with Scanning Electron Microscope FEI Quanta 50F. STEM images are taken on hair cross-sections (60 nm thickness) without any staining or coatings.

**Ethics approval.** Several participants were interviewed, and signed consent was obtained from the participants for the use of the samples. In total, 3 samples of hair were obtained from the men, ranging in age from 20 to 30 years old. The samples have no personal identification and the experimental protocols were approved by Brookhaven National Labs, Photon Sciences Safety Committee and the Office of Research Administration, which processes all human subject protocols. Protocols are in according with the “Resolution 199/96 on research involving human subjects” as well as the guideline of Brazilian National Health Council according to the Decree n: 93933 for research involving human subjects.

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Author Contributions
V.S. and K.E.-L. conceived and conducted X-ray experiments, analyzed results and prepared manuscript. V.S., J.B. and F.E.M. prepared the samples and conducted the SEM and TEM experiments, A.S. manufactured and supplied kinoform lenses, crucial for the X-ray experiments, All authors reviewed the manuscript.

Additional Information
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