Keratins as the main component for the mechanical integrity of keratinocytes

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Keratins are major components of the epithelial cytoskeleton and are believed to play a vital role for mechanical integrity at the cellular and tissue level. Keratinocytes as the main cell type of the epidermis express a differentiation-specific set of type I and type II keratins forming a stable network and are major contributors of keratinocyte mechanical properties. However, owing to compensatory keratin expression, the overall contribution of keratins to cell mechanics was difficult to examine in vivo on deletion of single keratin genes. To overcome this problem, we used keratinocytes lacking all keratins. The mechanical properties of these cells were analyzed by atomic force microscopy (AFM) and magnetic tweezers experiments. We found a strong and highly significant softening of keratin-deficient keratinocytes when analyzed by AFM on the cell body and above the nucleus. Magnetic tweezers experiments fully confirmed these results showing, in addition, high viscous contributions to magnetic bead displacement in keratin-lacking cells. Keratin loss neither affected actin or microtubule networks nor their overall protein concentration. Furthermore, depolymerization of actin preserves cell softening in the absence of keratin. On reexpression of the sole basal epidermal keratin pair K5/14, the keratin filament network was reestablished, and mechanical properties were restored almost to WT levels in both experimental setups. The data presented here demonstrate the importance of keratin filaments for mechanical resilience of keratinocytes and indicate that expression of a single keratin pair is sufficient for almost complete reconstitution of their mechanical properties.

Formation of a barrier capable of protecting tissue from external damage, chemical factors, and pathogens while resisting mechanical stress, external pressure, or shear force is one of the main functions of epithelial tissues. Keratinocytes represent the major cell type of mammalian epidermis and are mainly responsible for barrier functionality (1–3). On the molecular level, mechanical cell properties mainly depend on cytoskeletal fibrous structures (3), namely actin filaments, microtubules, and intermediate filaments (IFs). Although the contribution of actin filaments and microtubules to the resilience of many cell types is widely accepted (4), it has been hypothesized for many years that the resilience of epithelia against various types of deformation depends largely on keratins (5–7). These form a stable network spanning from the cell periphery to the nucleus. Peripheral filaments dynamically enlarge into thicker filaments and progressively intermingle with the preexisting network by continuously moving centripetally until, in the center of the cell, a dense network of keratin filaments encircles the nucleus (8, 9).

Keratins are encoded by a large multigene family of more than 50 genes that are specifically expressed depending on distinct developmental pathways and physiological requirements (10, 11). Based on their amino acid sequence, type I keratins display an overall acidic character and differ considerably from the more basic type II keratins (12). Filament assembly requires both keratin types because of the obligatory heterodimer composition of keratin IFs (13). Given that most epithelia express 4–10 different keratin subunits (14), the total and isotype-specific contribution of the overall keratin network to mechanical properties of epithelia remains highly challenging to analyze in vivo. An additional difficulty in experimentation is the absence of drugs to specifically disrupt the keratin IF system.

Most results on the mechanical properties and functions of IFs are based on biomimetic systems and disease models. One of the hallmarks of IFs is their low bending stiffness. In conjunction with nonlinear strain stiffening observed in networks at large deformations, these features supposedly enable IFs to serve as a mechanical buffer system protecting cells from environmental stress (15–17). High tensional loads have already been observed for keratin in vivo (18). Strain stiffening goes along with a predominantly elastic behavior in biomimetic systems (19). A similar elastic response was also found in particle-tracking microrheology experiments on epithelial cells (20). Furthermore, on the level of transgenic mice and patients with heritable fragility of the epidermis, a clear correlation between molecular integrity of keratin filaments (K5 and K14) and mechanical toughness of epithelia was shown (21, 22).

Generation of keratin KO mouse strains by deleting either the type I or type II keratin gene cluster (23) results in complete loss of keratin filaments in epidermal cells and provides the unique opportunity to accomplish accurate and exact biomechanical analyses of keratin-free keratinocytes for the very first time. Until now, these cell systems have been characterized for various keratin subunits and cell types (14) that have been used to attempt to elucidate the contribution of keratin to cell mechanics. Here, we provide the first experimental evidence for the crucial role of keratin IFs in cell mechanics and their vital contribution to epithelial tissue integrity.

Significance

For decades, researchers have been trying to unravel one of the key questions in cell biology regarding keratin intermediate filament function in protecting epithelial cells against mechanical stress. For many different reasons, however, this fundamental hypothesis was still unproven. Here we answer this pivotal question by the use of keratin KO cells lacking complete keratin gene clusters to result in total loss of keratin filaments. This lack significantly softens cells, reduces cell viscosity, and elevates plastic cell deformation on force application. Reexpression of single keratin genes facilitates biomechanical complementation of complete cluster loss. Our manuscript therefore makes a very strong case for the crucial contribution of keratins to cell mechanics, with far-reaching implications for epithelial pathophysiology.


The authors declare no conflict of interest.

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cell biological and biochemical functions (24, 25). Of note, increased cell motility and altered desmosomal and hemidesmosomal adhesion were identified in these studies.

In the current work, we analyze the keratin type I and type II mutant keratinocyte cell lines and compare them to WT controls established in parallel and rescue cell lines reexpressing keratins 5 and 14. Mechanical properties are examined in living cells from outside by atomic force microscopy (AFM) (26, 27) and from inside by magnetic tweezers (28) (Fig. S1). The data provide compelling evidence for the hypothesis that keratins determine the mechanical stability and integrity of epithelial cells by direct effects on various mechanical parameters such as elasticity, viscosity, spring constant, and relaxation time.

Results

Loss of Keratin Filaments Leads to Significant Cell Softening. Keratin filaments are believed to strengthen epithelial cells against various types of mechanical stress, mainly indentation, local deformation, and strain. To investigate this hypothesis, we characterized the mechanical properties of WT and keratin-lacking cell lines by complementary approaches, i.e., by AFM and magnetic tweezers experiments. To consolidate our data beyond doubt, both keratin type I mutant cells (KtyI−/−) and keratin type II mutant cells (KtyII−/−), as well as corresponding rescue cell lines, were examined in parallel.

In a first series of experiments, WT and KtyI−/− cell lines were characterized for their resilience against external force by AFM indentation. Immunocytochemistry stainings revealed well-developed keratin networks in WT cells with thin keratin filaments in the cell periphery and more prominent keratin bundles within the cell body surrounding the nucleus as a stable cage (Fig. 1). In contrast, keratin filaments were completely absent in KtyI−/− cells. AFM indentation experiments were performed above the nucleus and the cell body (Fig. 1 D–F and Fig. S1A). Subsequently, cell elasticities E were calculated for increasing indentation depths (Fig. S2 and SI Text, AFM Apparent Modulus Plots). Plateau regions indicating ideal indentation depths for cell elasticity analysis were identified in an indentation range of 300–600 nm above the nucleus and estimated around 200 nm for the cell body. These indentation depths were always less than 10% of the respective overall cell thickness [7.0 ± 1.4 (SD) μm for WT (n = 6) and 5.0 ± 0.8 μm for KtyI−/− cells (n = 4) on the nucleus and ~4 μm for both cell types on the cell body]. In all AFM experiments, at least three force-distance curves were recorded for each cell position. The total number of analyzed cells was >100 for WT and KtyI−/−. Superposition of the resulting raw data curves that are exemplarily given for one position in Fig. 2A indicated reliable reproducibility. The data showed that cells responded elastically with no permanent plastic deformation or active response to probing. Calculating the applied forces necessary to indent 100–600 nm directly from the raw data revealed significant differences between WT and KtyI−/− cells (KO) are indicated (P = 0.05, **P = 0.01, ***P = 0.001); error bars are SEM.

To accurately calculate cellular elasticities, force-distance curves were analyzed by the classical Hertzian spherical indenter model (Eq. 1) as implemented by the AFM manufacturer (Fig. 3A). For all areas analyzed, the resulting Young’s moduli indicated a significantly higher (Fig. 3 B and C) stiffness of WT cells compared with KtyI−/− cells [nucleus: EWT = 459 ± 31 Pa (SEM) (n = 125) and EKyI−/− = 343 ± 18 Pa (n = 109); significance level, 0.003]; cell body: EWT = 752 ± 100 Pa (n = 117) and EKyI−/− = 412 ± 75 Pa (n = 113); significance level, 0.0002]. Force maps from whole WT and KtyI−/− cells also demonstrated an overall softened cell phenotype on keratin loss, most prominently above the nucleus (Fig. 3A–D).

Using an alternative approach for force-distance curve analysis by accurately determining the indenter contact points and by describing curves via a standard power law function, the presented results were fully confirmed with an even better fit accuracy (SI Text, Alternative Force-Distance Curve Analysis and Figs. S3–S6).

Identical AFM experiments on keratin KtyII−/− mutant cells lacking the complete set of type II keratins resulted in very

Fig. 1. Keratin network characterization and AFM analysis regions. (A–C) Immunofluorescence micrographs of the keratin network in WT, KtyI−/− KO, and K14 rescue (RES) keratinocytes. The KtyI−/− cell outline is given by a dashed line. (Scale bars, 10 μm.) (D) Top view phase contrast image of an AFM tip-less cantilever aligned with the cell body of a living adherent keratinocyte. (Scale bar, 10 μm.) (E) Side view diagram of a cantilever positioned above the nucleus and (F) the cell body.
similarity data. Mutant cells were significantly softer (∼40%) than the corresponding WT cells (n ≥ 10 cells each).

Reexpression of Keratin 14 Rescues Keratin Network and Cell Elasticity. To prove that the absence of keratins was responsible for altered cell stiffness, rescue cells stably expressing keratin 14-YFP (yellow fluorescent protein) were generated (K14), leading to IF formation by pairing with the endogenously produced K5. Cells were analyzed by the above-described AFM protocol. Proper keratin network formation was assessed by fluorescence microscopy (Fig. 1C) before indentation in each of the analyzed cells (n = 46). Elastic moduli were retrieved by fitting the resulting force distance curves with the Hertz model. To minimize errors, WT and KtyI−/− cells (n = 57 and n = 51, respectively) were measured in parallel using the same cantilever (Fig. 4E and F). We found that the average cell elasticity was significantly increased on reexpression of K14. Interestingly, Young’s moduli of rescue cells showed a biphasic distribution, with ∼50% of cells presenting reduced elasticity as in KtyI−/− cells and the other ∼50% presenting WT or even more elevated stiffness.

Keratin Stiffens Cells Against Intracellular Deformation and Elevates Cell Elasticity. Although AFM analyses are based on the application of extracellular indentation forces, we sought for an alternative method to probe for intracellular force measurements of adherent cells. To this end, we adapted magnetic tweezers (for a scheme, see Fig. S1B) to examine intracellular deformation by local straining in keratin-free cells. In this set of experiments, KtyII−/− cells, corresponding WT cells and K5-YFP expressing rescue cells, were used. The KtyII−/− cells much like the KtyI−/− cells were completely devoid of keratin filaments, and reexpression of a single keratin, in this instance K5, was sufficient for restitution of an extended keratin filament network (Fig. 5A).

Cyclic force pulses applied to superparamagnetic beads resulted in viscoelastic response curves exemplified by Fig. 5B for WT and KtyII−/−. Major bead displacements, however, were noted in KtyII−/− cells that resulted in incremental bead movement toward the magnetic tweezers tip with repeated pulses, indicating a lower resistance of the keratin-free cytoplasm. The response curves were then rescaled, normalized by the amplitude of the applied magnetic force, and averaged. The resulting curves revealed a significantly different viscoelastic behavior of WT vs. KtyII−/− cells. Stable reincorporation of GFP-keratin 5 (K5) in KtyII−/− cells and subsequent direct fluorescence microscopic preselection for cells with a well-developed keratin network almost fully restored viscoelastic behavior to WT levels (Fig. 5C).

Extension portions of these processed response curves recorded on each cell phenotype were fitted via Eq. 2. After an initial quick elastic response [initial jump of amplitude 1/(k0 + η1)] in the creep compliance, the system enters a viscous regime (with a characteristic transition time equal to τ) until a flow regime dominated by η0 (viscous friction coefficient) is established.

Parameters obtained from the fits (Table 1) clearly indicate a lowering of the cytoplasmic viscosity by more than 40% in the absence of a keratin network. This reduction is comparable to the reduced cell elasticity found above by AFM experiments and is accompanied by an overall softening of KtyII−/− cells in comparison with the corresponding WT cells (n = 117, nKO = 113).

![Fig. 3](image1)

![Fig. 4](image2)
Keratin Loss Does Not Affect Actin and Microtubule Networks. To verify that reduced cell resilience and viscosity in the absence of keratins were not significantly influenced by altered expression of actin and tubulin, Western blot analysis was performed (24). Overall protein levels for all cell lines analyzed here indicated no major changes in protein concentration of actin and tubulin (Fig. 6B). In agreement with previous findings, staining for actin and tubulin demonstrated that the overall actin and microtubule organization was unaffected by the absence of keratins (Fig. 6A). To further exclude a putative influence of actin on altered mechanical properties of KO cells, actin filaments were disassembled by latrunculin A in WT and KtyI−/− cells. Subsequent AFM measurements on those cells identified a continuous, basically unchanged difference in elasticity and therefore significantly (significance level, 0.003) stiffer cell properties for WT cells compared with KtyI−/− cells, strongly supporting the dominant role of the keratin network in determining cell stiffness (Fig. 6C).

Discussion
The elasticity of cells and their resistance to external mechanical stress have been intensively analyzed for decades because of their crucial contribution to many cell functions such as differentiation (29) or cancer progression (30). Mechanical stability is especially relevant for epidermal keratinocytes, which are subjected to intense deformation in all directions. Although one of the main characteristics of epithelial cytoskeleton is the predominance of keratin filaments spanning the entire cytoplasm and surrounding the nucleus as a dense network (8), mechanical analyses have largely focused on actin filaments. On the other hand, it has been proposed that IFs serve as a major mechanical buffer system protecting cells from environmental stress (5). Seminal publications have tried to shed light on this hypothesis by use of biomimetic model systems on artificial IF networks (15), extracted keratin networks (20, 31), and analyses on whole cells or mouse models (32, 33). In essence, many of these analyses identified a very low bending stiffness for keratin filaments and a minor effect of keratin on cell elasticity. At the same time, however, rearrangements of the keratin network in various cancer cell lines were hypothesized to induce significant changes in cell elasticity (5).

Using two independent keratinocyte cell lines, both lacking the entire keratin cytoskeleton, and matched rescue cell lines reexpressing the basal cell–specific keratin pair K5/14, we conclusively demonstrate that elasticity, effective spring constants, and cell viscosity are significantly affected by keratins, leading to softened, less viscous cells in their absence. The use of two independent keratin-free cell models, the previously established keratin type II−/− mutant cells and the newly isolated keratin type I−/− keratinocyte mutant cells, in combination with matched WT controls and rescue cell lines largely excludes cell line–specific artifacts and considerably strengthens the tenacity of our findings. Both mutant cell lines lacked keratin filaments completely, showed very similar cell morphologies, and presented very similar biomechanical deficiencies as reflected by almost identical reductions in Young’s moduli that were identified in our AFM analyses. Furthermore, all alterations in mechanical cell properties were shown to be fully dependent on keratin loss. Here, neither actin nor MT protein levels nor corresponding cytoskeletal structures were affected. These results are well in line with results on protein levels found before for KtyII−/− cells (23, 24). Additionally, pharmacological actin network disruption reduced the overall cell elasticity as one would expect on removal of cortical actin but did not influence the clear biomechanical differences between WT and KtyI−/− cells. These data therefore clearly show that our AFM measurements were sufficiently deep to deform not only cortical actin but also the keratin network beneath.

Keratin material properties additionally explain the even more drastic keratin mutant phenotypes observed in magnetic tweezers stretching experiments. Although AFM-induced indentation experiments work primarily against the comparatively low IF bending stiffness of only 4−12 × 10−27 Nm2 (15), microrheological magnetic tweezers experiments locally stretch the keratin network to such an extent that they work against increasing stiffness at high strain. Keratin filament extensions of more than 2.5-fold have been observed in vitro (34), and stretched keratins have been imaged in cultured keratinocytes that were elongated by 133% (18) without filament rupture. The absence of plastic deformations on application of successive magnetic tweezers pulses in WT cells in the current study provides further compelling evidence for the effectiveness of keratin networks as mechanical buffer system. Consequently, the absence of this buffer system in keratin-deficient cells resulted in reduced viscosity, lowered effective spring constants, and plastic cell deformation. These observations are in line with reports on other IF systems. For example, vimentin-deficient fibroblasts presented a reduced stiffness on mechanical stress application by extracellular magnetic twisting cytometry (35). In another study, Brown et al. (36) noted that the rigidity of T lymphocytes depends primarily on intact vimentin filaments as determined by flow cytometry. On the other hand, AFM measurements on rat fibroblasts with a disrupted vimentin cytoskeleton identified localized stiffening (37).

Table 1. Viscoelastic cell parameters

<table>
<thead>
<tr>
<th>Parameter</th>
<th>KO</th>
<th>RES</th>
<th>WT</th>
</tr>
</thead>
<tbody>
<tr>
<td>$k_{eff}$ (pN/μm)</td>
<td>8</td>
<td>11.8</td>
<td>12.3</td>
</tr>
<tr>
<td>SD</td>
<td>1.1</td>
<td>3.2</td>
<td>1.8</td>
</tr>
<tr>
<td>$τ$ (s)</td>
<td>10.8</td>
<td>18.5</td>
<td>18.3</td>
</tr>
<tr>
<td>SD</td>
<td>1.3</td>
<td>4.2</td>
<td>3.0</td>
</tr>
<tr>
<td>$\eta$ (pN s/μm)</td>
<td>0.4</td>
<td>0.43</td>
<td>0.48</td>
</tr>
<tr>
<td>SD</td>
<td>0.11</td>
<td>0.19</td>
<td>0.17</td>
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Viscoelastic parameters obtained from fitting Eq. 2 to the averaged normalized magnetic tweezers data of WT, K5 rescue (RES), and KtyII−/− mutant cells (KO). $k_{eff}$, effective spring constant; $\eta$, viscous friction coefficient; $τ$, characteristic relaxation time.
Different type I and type II keratins are simultaneously expressed in resting keratinocytes. The most prominent keratins of basal epidermal cells are the type II K5 and the type I keratins K14 and K15 (14). K5 deletion results in pronounced cytolysis and neonatal lethality (38), whereas K14 depletion leads to epidermolysis bullosa simplex-like disease where mechanical trauma exacerbates blistering (39). Assuming a similar role of K14 and K15, we decided to test for rescue of overall type I KO by reexpression of K14 only. At the same time, K5 rescues were analyzed here to induce formation of the same keratin network by expression of K5 as sole type II keratin. Both rescue clones formed a well-detectable keratin network (Figs. 1 and 5). To our surprise, reexpression of K5 rescued basically all biomechanical phenotypes of KtyI−/− cells to WT levels when analyzed by magnetic tweezers. Similar results were found for KtyI−/− cells reexpressing K14 recuperating elasticity levels similar to the WT when analyzed by AFM indentation. A slightly enhanced spread in cell elasticity found for this rescue may be caused by different K14 expression levels.

In summary, our data clearly prove the vital character of keratin filaments for biomechanical cell properties as a buffer system to withstand mechanical stresses with far-reaching implications for epithelial pathophysiology.

Materials and Methods

Cell Culture. Keratinocyte cell line KtyI−/− lacking type II keratins and control WT cell lines were generated and described previously (23, 24). Generation of the type II rescue cell line KII KO K5-GFP reexpressing GFP-tagged keratin 5 was described in previously (25).

For generation of the cell line KtyI−/− lacking the complete set of type I keratins, homozygous C57Bl16 mice harboring two floxed alleles of the type I keratin cluster were crossed with 129S1hprt Cre mice. The resulting heterozygous mice were crossed to get 25% type I keratin cluster-deficient mice. Keratinocytes were isolated from the epidermis of E18.5 WT and KtyI−/− embryos and cultivated under conditions that permit spontaneous immortalization as described in Turksen (40).

To rescue KtyI−/− cells, human keratin 14-YFP cDNA was cloned into lentiviral pLVX-puro vector using XhoI and NotI restriction sites. Lentivirus for transduction was produced using the Lenti-X expression system (Clontech). KtyI−/− cells were incubated with lentivirus for 24 h in the presence of 8 μg/mL polybrene (Sigma-Aldrich). Subsequently, cells were selected with puromycin (8 μg/mL; PAA), and clones stably expressing hK14-YFP were generated by limiting dilution cloning. Cultivation of keratinocytes was performed as described previously (25). Cells were grown at 5% (vol/vol) CO2 and the appropriate temperature. Rescue cells were incubated in medium containing 8 μg/mL puromycin for selection purposes. For some experiments, KtyI−/− and WT cells were incubated with medium containing 0.5 μM latrunculin A for 60 min at 37 °C. Directly before analysis, medium was replaced, and AFM measurements were performed within the next 60 min. The effectiveness of actin depolymerization was verified by immunocytochemistry on identically incubated cells.

Western Blotting. SDS/PAGE on crude protein extracts of all cell lines analyzed in this work was performed as described previously (41). Separation of total protein extracts was performed by standard procedures (4-20% SDS-PAGE). Using GAPDH as constitutively expressed marker (antibody G8795; Sigma-Aldrich), overall protein levels of actin and tubulin were determined using mAB8457 (CST) and MAB1864 (Millipore) as antibodies. A standard immunoblotting protocol was performed as described previously (25) using actin and tubulin (see above) or keratin-specific antibodies (rabbit anti-pan keratin; ProGen Biotechnik). Images were recorded either on an ApoTome.2 (Zeiss) using AxioVision LE software (Zeiss) or acquired using a laser scanning confocal microscope (LSM 710; Zeiss), both equipped with a Plan-Apochromat 63×/1.4 oil immersion objective.

Immunocytochemistry. For immunocytochemical experiments, 2.5 × 10^6 cells were grown on collagen I-coated glass coverslips for 48 h. Labelings were performed as described previously (25) using fluorescently labeled phalloidin (Invitrogen) for actin and antibodies against tubulin (see above) or keratin (rabbit anti-pan keratin; ProGen Biotechnik). Images were recorded either on an ApoTome.2 (Zeiss) using AxioVision LE software (Zeiss) or acquired using a laser scanning confocal microscope (LSM 710; Zeiss), both equipped with a Plan-Apochromat 63×/1.4 oil immersion objective.

AFM Measurements. AFM imaging and force spectroscopy were performed using a JPK Nanowizard Life Science version instrument (JPK) equipped with a fluid chamber (Biocell; JPK) for live cell analysis and an inverted optical microscope (Axiovert 200; Zeiss) for sample observation. AFM imaging of fixed and live cells was performed as described in SI Text, AFM Measurements. During indentation experiments, each analyzed position was recorded by at least three force curves and a cantilever speed of v = 1.5 μm/s. Force set points of 1.48 and 0.5 nN were used for the nucleus and cell body, respectively.
AMF Data Analysis. The approaching parts of the recorded force-distance curves were analyzed using the JPK DP software (Hertz fit, sphere model) to obtain values of Young’s modulus \( E \) (Fig. 2A). With this option, the software uses the fit function described by Sneddon (42):

\[
F(a) = \frac{E}{1 - v^2} \left[ \frac{a^2 + R^2}{2} \ln \frac{R + a}{R - a} - \frac{\pi R}{2} \right].
\]

Here, \( h \) is the indentation depth, \( a \) is the contact radius of the indenter, \( R \) is the indenter radius, and \( v \) is the Poisson ratio (set to 0.5 assuming cell’s incompressibility), and \( E \) is the cell’s Young’s modulus. In the fits, Young’s modulus \( E \) and the contact point were used as fit parameters. For estimating the optimal indentation depth, so-called “apparent modulus” plots were used based on ref. 43 (Fig. S2 and SI Text, AFM Apparent Modulus Plots).

Because infinite sample thickness is assumed in the Hertz model, we limited the indentation depth at most 5–10% of the sample thickness. Accordingly, curves recorded when indenting on the nuclei were fit for a range of 0 to 500 nm after the contact point. For the cell body, comparably thinner, this distance was taken to be 200 nm. Distributions of resulting elasticities \( E \) were analyzed for differences with the parameter-free two-sided Mann-Whitney-Wilcoxon test.

Magnetic Tweezers Setup and Measurements. Construction and calibration of the self-made magnetic tweezers and measurement procedures are described in detail in Fig. S7 and in SI Text, Magnetic Tweezers Measurements.


In short, the instrument is based on a self-etched, water-cooled magnetic core of an electromagnet attached to a micromanipulator. M-270 Dyna-beads (Invitrogen) were incorporated as superparamagnetic beads using a PDS-1000/He particle delivery system (BioRad).

Magnetic Tweezers Data Analysis. Single peaks were normalized by the magnetic force applied at every time frame (calculated from the calibration curve \( F = F(d) \), \( d \) being the bead-tip distance; Fig. S8 and SI Text, Magnetic Tweezers Measurements) and averaged for every cell type. Following Bausch et al. (28), the extensional parts of these averaged creep-response curves (i.e., distance divided by force) were fitted by a four-element mechanical equivalent circuit composed of a dashpot in series with a Zener body:

\[
\frac{1}{\tau_i} = \frac{1}{\tau_0} - k_i \exp\left( -\frac{d}{\tau_i} \right),
\]

where \( k_i \) describes the effective spring constant of the system, \( \tau_0 \) indicates the viscous friction coefficient of the bead, and \( d \) is the relaxation time characteristic of the transition from an elastic to a viscous regime. Uncertainties of fit results were determined as SDs from 500 times fitting synthetic data created by adding to the fit result normally distributed random numbers with scatter taken from the experiment.

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Supporting Information

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SI Text

**AFM Measurements.** For atomic force microscopy (AFM) analyses, 12,000 keratinocytes were seeded on collagen I-coated (6 μg/cm² collagen, rat-tail; BD Bioscience) glass coverslips (0.24 mm; Menzel-Gläser) 16–24 h before the experiment. AFM imaging of fixed and live cells was performed in contact mode using silicon tips (Nanosensor point probes-CONT-50) of nominal resonance frequency \( f_0 = 10–17 \) kHz and nominal spring constant \( k = 0.07-0.4 \) N/m. Images were processed using the JPK DP software. Indenters for probing cell elasticity were prepared by mounting silica microspheres of a nominal radius of 5 μm (G. Kisker GbR, PSI-5.0, surface plain) to silicon AFM tipless cantilevers of nominal resonance frequency \( f_0 = 7 \) kHz and nominal spring constant \( k = 0.04 \) N/m (Nanoworld Arrow TLI-Au with Tl’ Au back tip coating) using two-component glue (UHU plus Endfest 300; UHU). Smooth silica beads were picked under microscopy control. After attachment, microsphere diameters were measured using a confocal microscope equipped with an LD Plan Neofluar Ph.2 (40×0.6) objective. AFM indentation curves were recorded at room temperature at sites of the nucleus and the cell body of single keratinocytes. Directly before measurements, cells on coverslips were washed once in PBS, mounted in the JPK Biocell chamber, and immersed in freshly prepared Hepes-buffered medium. Vitality of keratinocytes was checked visually. Before each experiment, the cantilevers were immersed in medium for 10 min to allow thermal equilibration and subsequent calibration using the thermal noise method. Measured spring constants ranged between 20 and 50 mN/m.

**AFM Apparent Modulus Plots.** A test to determine the optimal depth of indentation to be analyzed in AFM elasticity measurements consists of the analysis of the so-called “apparent-modulus” plots. For a material as heterogeneous and complex as a cell, Young’s modulus \( E \) is not a constant value but varies according to the depth at which the AFM tip indents all of the different layers of soft biological material. Plateau regions in these \( E(\delta) \) plots should be looked for, and the range of indentation \( \delta \) giving rise to a roughly constant value of \( E \) should be used as a fit range during data analysis. By progressively extending the fit range (sphere model, AFM manufacturer’s software) on single indentation curves, plateaus of \( E \) were identified between 200 and 600 nm (nucleus). No systematic plateau could be identified for measurements performed on the cell body. Fit ranges for further data analysis were chosen to be 500 and 200 nm for nucleus and cell body, respectively, also in light of the restrictions on the indentation depth imposed by the Hertz model (\( \delta < 5-10\% \) of total sample thickness). Although different fit ranges impede the direct comparison of elasticity values for the cell body with those above the nucleus, analysis parameters were always kept identical between all analyzed cell lines.

**Alternative Force-Distance Curve Analysis.** Data fits using the Hertz model (Fig. S3 A and B) always yielded slightly curved residuals and contact points that were consistently and significantly below the measured curves. Therefore, this description is only a first approximation to the very complicated mechanical system and should be improved. Underlying reasons for the failure of this popular model are most likely roughness of cells (Fig. S4 A and B), sphere irregularities (Fig. S4C), and, most importantly, the nonhomogeneous and nonisotropic mechanical material properties of cells. We therefore developed an alternative approach by accurate determination of contact points from the approach curves alone (Fig. S5A and see below). Curves were then fitted using a standard power law function

\[
F = A \delta^b. \tag{S1}
\]

In the following, we will call the prefactor \( A \) “apparent stiffness.” If applicable, the exponent chosen is denoted by a subscript.

This approach can be connected to the Hertz model as follows. For indentations much smaller than the radius of the spherical indenter, which is in all our experiments, the solution of the Hertz model, Eq. 1, can be simplified to yield

\[
F(\delta) = \frac{4}{3} \frac{E}{(1-v^2)} \sqrt{R} \delta^{3/2}. \tag{S2}
\]

Thus, the Hertz model predicts a power law with exponent 3/2. The general power law function Eq. S1 can be regarded as a generalization of the usual Hertz model.

Apparent stiffnesses \( A_{1/2} \) resulting from such a model again proved WT cells to be significantly stiffer than keratin-deficient cells above the nucleus and cell body (Fig. S5 B and C; significance levels of 0.015 and 0.004, respectively).

In a next step, apparent stiffness \( A \) and exponent \( b \) were both selected as free fit parameters. From fitting all data curves we obtained Gaussian distributions of the exponent \( b \) with almost identical mean exponential values of 2.09 ± 0.75 (SD) for WT cells \((n = 117)\) and 2.01 ± 0.65 for KtyI−/− cells \((n = 109)\) (Fig. S5 D and E). The exponent was therefore fixed to a value of 2 and apparent stiffnesses \( A_{2} \) were fitted for all force-distance curves. In contrast to the Hertz model the resulting fits showed minimal residuals without curvature and, most importantly, consistent contact points were used (Fig. S3B). The fit results also revealed a highly significant cell softening of ~30% on keratin loss at nuclear and cell body positions (Fig. S3 C and D). The mean values were: \( A_{2} = 1,460 \pm 1,060 \text{ N/m}^2 \) (SD) \( W.T. \), \( A_{2} = 1,120 \pm 520 \text{ N/m}^2 \) (KtyI−/−, significance level 0.008) for the nucleus, and \( A_{2} = 2,150 \pm 1,400 \text{ N/m}^2 \) (WT), \( A_{2} = 1,610 \pm 1,220 \text{ N/m}^2 \) (KtyI−/−, significance level 0.003) for the cell body.

On rescue cells, proper keratin network formation was assessed by fluorescence microscopy (Fig. 1C) before indentation in each of the analyzed cells \((n = 46)\). The resulting force distance curves were also analyzed by the standard power law function (Eq. S1) with exponent \( b \) as a free parameter. Because K14 cells were also characterized by a mean value of 2.01 ± 0.75 (SD) (Fig. S5E), all curves were analyzed for apparent stiffness \( A_{2} \) with \( b = 2 \) as before.

The Hertz model applies to the case of a smooth, spherical, and rigid indenter deforming a semi-infinite, perfectly elastic, and homogeneous substrate. To partially test these assumptions, surface roughness of the cells and of the spherical indenter were assessed. Reflection interference contrast microscopy (RICM) imaging of silica beads showed a variety of shape imperfections.

Contact-mode AFM high-resolution live cell imaging performed on small portions of the plasma membrane’s surface allowed to ascertain cell’s roughness, which had average values of ~50 nm as computed over an area of 500 × 500 nm² via

\[
\text{RMS}_{\text{rough}} = \sqrt{\frac{1}{N} \sum_{i=1}^{N} (z_i - \bar{z})^2}, \tag{S3}
\]

where \( N \) is the image pixel number, \( z_i \) is the height at that pixel, and \( \bar{z} \) is the average of \( z_i \).
Indentation curves were analyzed by a Matlab-implemented software according to the following algorithms. Raw data from force-distance approach curves were recalibrated by setting the sensitivity (nm/V) and spring constant (N/m) of the cantilever used for measuring, corrected for baseline offset and tilt and corrected for tip sample separation using the AFM manufacturer’s data processing software. For differentiation, the curves (of typically ~20,000 data points) were averaged to yield 100 final data points.

The maximum position of the third derivative of the smoothed curve was used as the x coordinate of the contact point. The corresponding offset force value was calculated by averaging over a portion of the noncontact part of the unsmoothed curve (50 values next to the contact position).

The first 500/200 nm (nucleus/cell body, respectively) following the contact point of the recalibrated indentation curve was used as a fit range. The general power law $F(\delta) = A \delta^b$ (i.e., Eq. S1) was fitted via a least-squares method. $A$ and $b$ were the only free fit parameters. In a successive step of the analysis, $b$ was user-constrained, thereby leaving the apparent stiffness $A$ as the only free parameter. The residuals were computed as difference between measurement and best fit.

**Magnetic Tweezers Measurements.** For the magnetic tweezers, solenoids were fabricated with a VACOFLUCX 17 (VACUUMSCHMELZE GmbH & Co.) core (2.28 mm in diameter), which has a saturation magnetization of 2.2 T. An electrochemical etching process described by ref. 1 was applied to produce tips with a radius of 5 μm. Initially a 10-mm section of one side of the core was mechanically reduced to a diameter of ~1 mm (Fig. S7A). This section of the core was exposed to the etching solution, which is comprised of 80 mL 85% (vol/vol) phosphoric acid, 70 mL 95.0–97.0% (vol/vol) sulfuric acid, and 50 mL distilled water, whereas the rest of the core was protected by a photoresist (AZ 4562; Microchemical GmbH) (Fig. S7B). A DC voltage of 12 V was applied (Fig. S7C). The lower part of the shield was removed after the core section exposed to the solution was etched to 40% of the original diameter. Etching was continued with a reduced voltage of 8 V (Fig. S7D). The etching was stopped when the distal part of the core dropped down (Fig. S7E and F). The coil surrounding the magnetic core was produced from 100-μm copper wire (3,000 turns). For cooling, the isolated coil was housed in a surrounding chamber, which was circulated by cold water.

To calibrate the particle movement, Stokes’ law was used. The magnetic force applied to a bead of radius $r$ moving with velocity $v$ in a fluid of known viscosity $\eta$ can be calculated via Stokes’ law

$$F = 6\pi r \eta v.$$  \[S4\]

Dynabeads M-270 with a diameter of 2.8 μm and an SD of 1.4% were used for force calibration experiments. The beads were immersed in three different fluids with viscosities of 1,000, 3,500, and 5,000 centiStokes (vinyl terminated polyDimethylsiloxanes; abcr GmbH & Co.). The current supply of the tweezers coil was kept to 215 mA, both for calibration and further measurements. The beads and the tip of the magnetic tweezers were simultaneously tracked to determine both the velocity and the distance between bead and tip.

Fig. S8A shows the force-distance dependence of three magnetic beads, each deflected in one of the three above described calibration solutions. The movements of the beads were within an angle of 5° with respect to the tweezers axis. The fitting curve

$$F(x) = a(x+c)^b,$$  \[S5\]

where $x$ denotes the bead-core-distance. Parameters $a$, $b$, and $c$ were determined by fitting Eq. S5 to the measured force distance relation displayed in Fig. S8A. The result was used for calculating forces acting in magnetic tweezers experiments on cells.

For magnetic tweezers experiments, cells were grown to confluency on collagen-coated glass slides (35 mm Ø); 250 mg of superparamagnetic M-270 Dynabeads (Invitrogen) was “shot” onto the cells using a PDS-1000/He particle delivery system (BioRad). Beads remaining outside the cells were removed by rigorous washing. After trypsinization and reseeding, single cells with single incorporated beads were selected for analyses after 12 h. An Eppendorf micromanipulator 5171 (Eppendorf) was used to place the tip of the tweezers next to the cell at bead height and at a lateral distance of 40–60 μm. Repeated rectangular pulses of 5 s in length were applied every 10 s and recorded at a frequency of 50 frames/s using an EOS 650D/Rebel T4i single-lens reflex (SLR) camera (Canon). Beads still attached to the outside of the cells had only loose contact and were removed by the first pulse. The entire setup was placed in a climate chamber heated to 37 °C. The bead and magnetic tweezers tip position were tracked over time using implemented algorithms of Adobe After Effect CS4 (Adobe Systems) and Fiji.

Fig. S2. Examples of the dependency of the Young’s modulus $E$ on the portion of indentation curve taken as fit range. For measurements performed on the nucleus a plateau is observable between 200–600 nm.

Fig. S3. Fit residuals and apparent stiffness $A$ of WT and KtyI$^{-/-}$ cells. (A) Fitted portion (500 nm) of a force-distance curve recorded on a nucleus. Gray, data; blue, AFM manufacturers software fit (Eq. 2, sphere model); blue dot, fitted contact point; green, fit of a power law (Eq. S1) with fixed exponent $b = 2$; red, same fit with $b = 1.5$ (Hertz model, cf. Eq. S2); red dot, contact point determined directly from measured curve as described in SI Text. (B) Corresponding fit residuals. (C) Cumulative histograms of the apparent stiffness $A$ obtained from fitting the power law function $F(\delta) = A\delta^b$ with $b = 2$ to the approaching portion of all force-spectroscopy curves recorded on nucleus (fit range: 500 nm, $n_{WT} = 89$, $n_{KO} = 110$) and (D) cell body (fit range: 200 nm, $n_{WT} = 117$, $n_{KO} = 113$).
Fig. S4. (A) Contact-mode AFM error signal image of a live WT keratinocyte. (Scale bar, 10 μm.) (B) 3D reconstruction of the surface topography of a 14 × 1.3-μm² portion of the plasma membrane used to estimate surface roughness. (C) Reflection interference contrast microscopy (RICM) image showing shape and surface roughness of the silica beads used for AFM cantilever modification. The arrow marks a representative bead selected for mounting. (Scale bar, 10 μm.)

Fig. S5. (A) Plot showing contact point identification on a typical AFM indentation curve recorded on a cell’s nucleus. Gray, original data; dark gray, smoothed data used for third derivative calculation (dashed line). The contact point is identified as the maximum of the curve’s third derivative. Zero of tip-sample separation axis: contact point as identified by the AFM manufacturer’s data analysis software. (B) Cumulative histograms of the apparent stiffness $A$ obtained from fitting the power law function $F(\delta) = A\delta^b$ with an exponent $b = 1.5$ (Hertz model) to force-indentation curves recorded on WT/KtyI⁻/⁻ (KO) nucleus ($n_{WT} = 89$, $n_{KO} = 110$, different on a significance level of 0.015) and (C) cell body ($n_{WT} = 117$, $n_{KO} = 113$, different on a significance level of 0.004). (D) Frequency histogram of the exponent $b$ obtained from fitting the power law function to the extended portion of all indentation curves recorded on nucleus (fit range: 500 nm) and cell body (fit range: 200 nm) of WT ($n = 118$) and KO ($n = 116$) cells. (E) Corresponding cumulative histogram, showing the almost perfect overlap of the distributions (data recorded on RES cells are included). A total of four cantilevers were used during the complete series of experiments.
Fig. S6. (A) Histogram showing the distribution of apparent stiffness $A_y$ obtained from indentation experiments performed on the nucleus of WT ($n = 57$), Ktyl$^{−/−}$ (KO; $n = 51$), and rescue (RES; $n = 46$) cells and (B) corresponding cumulative histograms.
Fig. S7. Electrochemical etching process outline. (A) A lathed core before etching. (B) Two fitted pipette tips were mounted onto the core surface. The lower one had to be sealed to protect the end of the core. The length of the tip was controlled by the exposed length of the core. (C) A 12-V DC voltage was applied on the core to etch the exposed part electrochemically. (D) Once the core diameter is etched down to 40% of the original, the lower shield was removed. The voltage was set to 8 V to avoid too large etching currents. (E) The power switched off once the distal part dropped down. (F) The upper shield was removed.

Fig. S8. (A) Magnetic tweezers force-distance calibration curve. (B) Schematics of the mechanical equivalent circuit used for fitting the viscoelastic response of the cytoplasm-incorporated magnetic beads used in magnetic tweezers experiments (1). In Eq. 2, we used the relaxation time $\tau$ that is related via $\tau = \frac{\gamma_1 k_0 + k_1}{k_0 k_1}$ according to ref. 2 with the parameters of the equivalent circuit shown here.