Stiffness heterogeneity of small viral capsids

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Nanoindentation of viral capsids provides an efficient tool in order to probe their elastic properties. We investigate in the present work the various sources of stiffness heterogeneity as observed in atomic force microscopy experiments. By combining experimental results with both numerical and analytical modeling, we first show that for small viruses, a position-dependent stiffness is observed. This effect is strong and has not been properly taken into account previously. Moreover, we show that a geometrical model is able to reproduce this effect quantitatively. Our work suggests alternative ways of measuring stiffness heterogeneities on small viral capsids. This is illustrated on two different viral capsids: Adeno associated virus serotype 8 (AAV8) and hepatitis B virus (HBV with $T = 4$). We discuss our results in light of continuous elasticity modeling.

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I. INTRODUCTION

The elastic properties of viral capsids can be inferred using nanoindentation experiments. Because viral capsids' sizes range from tens to hundreds of nanometers, these experiments are usually performed with an atomic force microscope (AFM), whose operating size range is appropriate [\[1\]](#page-10-0). Capsids of different virus families have been probed this way in several works, exhibiting various interesting features [\[2–9\]](#page-10-0). Depending on the identity of capsids and the amplitude of indenting force, one can observe elastic behavior with a linear or nonlinear response, plastic behavior, and eventually mechanical rupture. From the biological point of view, all these observations are relevant since they address the global stability of viruses, as well as their deformability, which is important physical information to fully understand the various steps of their replication cycle.

Most past works dealing with nanoindentation of capsids have focused on their linear response. This linear regime allows the *stiffness* to be defined effectively as the ratio between the force applied and the indentation. Typical stiffness ranges from 0.04 N m^{-1} for the influenza virus [\[5,10\]](#page-10-0) to 3 $N m^{-1}$ for the immature particle of HIV-1 in the presence of envelope glycoproteins, for example [\[11\]](#page-10-0). From a structural perspective, viruses are composed of mainly nucleic acids packaged inside self-assembled protein shells and eventually a surrounding lipid bilayer or membrane. For most viral capsids, the self-assembly involves multiple copies of identical proteins. Yet, the topology of a closed shell imposes that the environment at the scale of a single protein is not unique: at exactly 12 locations on the surface of the virus, proteins belong to a *capsomer* or face that is pentameric (five proteins),

while they belong to hexameric capsomers (six proteins) anywhere else. The precise location of pentamers determines the global shape of the viral capsid $[12]$. It can be either regular with icosahedral symmetry, or irregular and elongated. Focusing on icosahedral viruses, the three main symmetries (twofold, threefold, and fivefold) strongly suggest that elastic properties of capsids are not uniform [\[13\]](#page-10-0).

Heterogeneities in stiffness measurements have been observed on some viruses, such as the MVM parvovirus [\[4\]](#page-10-0), or bacteriophages ϕ 29 [\[2\]](#page-10-0). In the former case, heterogeneities have been attributed to the difference in virus orientation during the adhesion on the substrate, thus offering different capsomer or environments (with twofold, threefold, or fivefold symmetry) on its highest altitude as probed by the AFM tip. Yet, the effect was observed in the presence of viral DNA. Analysis of the AFM images allowed us to correlate mechanical properties with capsid orientation. In the case of bacteriophages, the difference in stiffness was traced back again to the orientation of the capsid during adhesion, but this was interpreted as a signature of the *asphericity* of the virus rather than inhomogeneity at the single capsomer level. However, these observations of stiffness heterogeneity are not systematic, and, for example, it is not possible to detect significant variations of stiffness for some viruses like Cowpea Chlorotic Mottle Virus (CCMV), depending on its orientation [\[3\]](#page-10-0). As a consequence, the presence or absence of stiffness heterogeneity and its origin are far from being correctly understood. The purpose of the present work is to investigate quantitatively, both experimentally and theoretically, signatures for these mechanical heterogeneities.

This paper is organized as follows. In the next section, we present our experimental approach based on AFMnanoindentation for two different viral capsids, namely Adeno-associated virus 8 (AAV8) and hepatitis B virus (HBV, with $T = 4$ capsid conformation), and then the numerical and

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analytical approach in order to predict and interpret the results of indentation experiments quantitatively. Both approaches are combined and discussed in Sec. [IV.](#page-10-0) These findings are then discussed in light of past literature. Consequences of future experiments are also expressed.

II. EXPERIMENTS OF AFM NANOINDENTATION ON VIRAL CAPSIDS

A. AFM nanoindentation experiments

1. AAV purification

Stock of AAV8 vector particles was produced and purified on CsCl gradients by calcium phosphate transfection of HEK-293 cells as described previously [\[14\]](#page-10-0). The vector plasmid AAVCMVeGFP included the enhanced green fluorescence cDNA under the control of the cytomegalovirus promoter. The total length of the vector was 3769 bases, including the AAV inverted terminal repeats. The helper plasmids used for production were pDG8. After purification and titration by qPCR, the viral stock was kept at -80° C.

2. HBV purification

HBV capsids were obtained from the assembly of Cp183 proteins produced in bacteria following a protocol detailed in [\[15\]](#page-10-0). Cp183 capsids were expressed in *E coli* and purified similarly as described previously for truncated Cp149. Plasmid of pRSFT7-HBc183opt was transformed into E coli BL21* CodonPlus (DE3) cells and grown at 37° C in LB medium culture. When OD600 reached around 2, an expression was induced with 1 mM IPTG overnight at 20° C. Cells were collected by centrifugation at 6000 g for 20 min. For 1 L of culture, cells were resuspended in 15 mL of TN300 buffer (50 mM Tris, 300 mM NaCl, 2.5 mM EDTA, 5 mM DTT, pH 7.5) and incubated on ice for 45 min with 1 mg/mL of chicken lysozyme, 1X protease inhibitor cocktail solution, and 0.5% Triton X-100. 6 μ L of benzonase nuclease were added to digest nucleic acids for 30 min at room temperature. Cells were broken by sonication and centrifuged at 8000 g for 1 h to remove cell debris. The supernatant was loaded onto a 10–60 % sucrose gradient buffered with 50 mM Tris pH 7.5, 300 mM NaCl, 5 mM DTT, and centrifuged in an SW-32Ti Beckman Coulter swinging bucket rotor at 140 000 g for 3 h at 4° C. Capsids were identified in gradient fractions by 15% SDS-polyacrylamide gel and precipitated by 40% saturated $(NH_4)_2SO_4$. After incubation on ice for 1 h and centrifugation at 20 000 g for 1 h, pellets were resuspended in 10 mL of purification buffer (50 mM Tris pH 7.5, 5% sucrose, 5 mM DTT, 1 mM EDTA). The protein solution was centrifuged again for 15 min to remove insoluble pellets. The supernatant containing soluble capsids was dialyzed overnight against purification buffer at 4 ◦C. The purified CP183 capsids obtained in such an assembly process contain random short *E Coli* nucleic acids [\[16,17\]](#page-10-0).

3. AFM sample preparation for AAV8 and HBV

The viral stock of AAV8 or HBV capsids was diluted to a concentration of approximately 6×10^8 capsids/ μ L in TN 1 mM [Tris 10 mM (pH = 7.4) and nickel(II) chloride (NiCl₂) 1 mM]. Immediately after, $5 \mu L$ of the solution was deposited

onto mica disks previously cleaned using adhesive tape. After 20 min of incubation to favor adhesion, we added 30 μ L of TN 1 mM to the sample and another 70 μ L of TN 1 mM in the AFM liquid imaging cell before imaging.

B. AFM imaging and nanoindentation protocol

The samples were imaged using a Bruker Nanoscope V Multimode 8 AFM using PeakForce Mode in liquid. We used three different cantilevers: SNL and ScanAsyst-fluid+ with a nominal tip radius of 2 nm, and ScanAsyst-fluid with a nominal tip radius of 20 nm. The SNL and the two ScanAsyst cantilevers have a nominal spring constant of 0.35 and 0.7 N/m, respectively. We use mainly ScanAsyst fluid+cantilevers to compare both AAV8 and HBV nanoindentation and AFM imaging. We used the two cantilevers with a tip radius of 2 nm (ScanAsyst fluid+) and 20 nm (ScanAsyst fluid) to study the size tip effect in the AAV8 particles (Supplemental Material Fig. 2) [\[18\]](#page-10-0). Also, we studied the effect of the stiffness of the cantilever in AAV8 nanoindentation; for this, we used the SNL and ScanAsyst fluid+cantilevers (Supplemental Material Fig. 3) [\[18\]](#page-10-0). Large-scale AFM square images of 3 μ m sides (512×512 pixels) at a 2 Hz scan rate were obtained to verify the deposition conditions on a large number of capsids. Figure [1](#page-2-0) of the Supplemental Material [\[18\]](#page-10-0) shows a zoom of those large AFM images for AAV8 and HBV capsids.

The capsids were then imaged one by one to perform nanoindentation. Images were scanned at 2 Hz over scan segments of 300–500 nm broad. The maximal force varied between 200 and 300 pN. Using the Point and Shoot feature of Bruker acquisition software, we defined a line with 15 equidistant points over the capsid and 3 points far from the capsid to get the surface reference force curve [see Fig. $1(a)$]. Additionally, a horizontal height profile that passes through the highest point of the capsid is plotted and will be used to determine the capsid radius $R_{cap,AFM}$ used to normalize the stiffness profile before alignment and averaging. Nanoindentation data have been analyzed semiautomatically using a homemade MATLAB code. Figure $1(b)$ shows the force-displacement $F(z)$ curves separately according to the indentation position over the viral capsid. The black line is the best curve among the three recorded over the surface far from the capsid, and its slope represents the cantilever spring constant k_c . All those curves were aligned with the contact point.

AFM nanoindentation data analysis

We used two methods to extract the stiffness from the capsid. In the first method, we considered the force curve as a function of the cantilever displacement $F(z)$ that contains the deformation of both the cantilever and the capsid in the first linear regime. Assuming the capsid and the cantilever behave like two springs in series, the capsid stiffness k_{cap} was calculated using the relation $k_{cap} = k(F(z)) = (k_{eff}k_c)/(k_{eff} - k_c)$. The cantilever stiffness k_c is determined using the thermal noise method, and the effective stiffness k_{eff} is the slope of the linear part of the $F(z)$ curve. In the second method, k_{cap} is extracted from the force indentation $F(\delta)$ curve as simply its slope. To obtain a $F(\delta)$ curve, we used the fact that for a given

FIG. 1. Nanoindentation on AAV8 viral particles. (a) Typical image of an AAV8 viral particle deposited on the surface. Indentations are performed at different locations represented by colored circles. Three additional indentations, shown by three black dots, were made away from the capsid to obtain a reference surface curve that contains only the cantilever deformation. The topographic profile along the black line is also shown. (b) Force-displacement curves for the different locations: left: gray points for virus-free locations; center: green points on the border of viruses; right: blue points in the center. The black curve represents an indentation made on the surface and away from the position of the virus (the best of the three *surface* curves recorded away from the viral capsid). By subtracting at a fixed force the two *z*-positions on the black curve and a curve obtained within the capsid, it is possible to extract the capsid deformation δ . (c) Single-particle stiffness extracted at different locations, either through force-displacement $F(z)$ or force-indentation $F(\delta)$ curves. The indentation positions *r* along the *x*-axis of the AFM image have been normalized using the capsid radius *R*cap, AFM obtained from the height profile in (a). (d) Ensemble-averaged stiffness at different locations ($N = 58$ viruses). For each lateral position, the symbol (square or losange) is the median of the local stiffness distribution, and the bar represents the position of the first and third quartiles for the same distribution.

applied force F , the indentation δ is found by subtracting the deformation of the cantilever from the total tip displacement. Figure $1(c)$ show the stiffness profile using both calculation methods. To superimpose $N = 58$ different indented capsid profiles, we aligned them according to the two lowest points of the profile. Then, the indentation positions *r* were normalized using the capsid radius $R_{cap,AFM}$ obtained from the capsid height profile [Fig. $1(a)$]. Finally, we determined a unique stiffness profile where the diamond and square represent the median of all the k_{cap} values for position zero, for example. The upper and lower error bars are 75% and 25% of the data [see Fig. $1(d)$].

III. MODELING AFM NANOINDENTATION ON VIRAL CAPSIDS

A. Thin shell model for intrinsic stiffness heterogeneity

Thin shell elasticity is very well documented both in standard textbooks [\[19–22\]](#page-10-0) and outstanding research publications [\[23–25\]](#page-10-0). We will mainly recall in this subsection the main results of interest for our study. In particular, we present scaling derivations for the stiffness, which have been shown to be appropriate in the regime of linear elasticity. Within standard three-dimensional (3D) elasticity, the deformation of a material from an unconstrained configuration is characterized mainly by two tensors: the strain tensor, which quantifies the relative material deformation, and the stress tensor, which quantifies the strength and orientation of forces arising from this deformation. In the simplest case of unidimensional material, the linear relationship between longitudinal strain ϵ and stress σ is called Hooke's law and it defines the Young modulus as $Y = \sigma/\epsilon$, which is an elementary measurement of the stiffness of the material.

The concept of stiffness can be extended to different material geometry. In the case of thin shells like viral capsids, it is defined as the ratio between the applied mechanical force and the indentation length. In the case of a homogeneous spherical thin shell of width t and radius R , the stiffness results from the balance of in-plane stretching/compression and out-of-plane bending. Introducing the 2D stretching modulus as $\kappa_s = Yt$ and the 2D bending modulus (also known as the flexural rigidity) as $\kappa_b = Yt^3/[12(1-\sigma^2)]$, where σ is the Poisson ratio of the material, we define the dimensionless number associated with stretching/bending balance, which is also known as the Föppl–von Karman number (FvK):

$$
\gamma = \frac{\kappa_s R^2}{\kappa_b}.\tag{1}
$$

The scaling of stiffness is obtained by estimating the energetic budget associated with small indentation Δ of the spherical shell [\[19,26–28\]](#page-10-0). This indentation modifies the curvature of the shell locally over a distance *d* from a value *R*−¹ to $\rho^{-1} \sim R^{-1} - 2\Delta/d^2$. The bending cost is of order $\Delta E_b \sim$ $\kappa_b(\rho^{-1} - R^{-1})^2 d^2 \sim \kappa_b \Delta^2 / d^2$. The meridians are also compressed by the indentation resulting in a strain $\epsilon \sim \Delta/R$. The compression cost is of order $\Delta E_s \sim \kappa_s \epsilon^2 d^2 \sim \kappa_s d^2 \Delta^2 / R^2$. The actual deformation spread *d* is obtained by minimizing the sum of bending and compression energies leading

to $d_e \sim R^{1/2} (\kappa_b/\kappa_s)^{1/4} \sim R \gamma^{-1/4}$. The net energy associated with the indentation is therefore written as $\Delta E_{\text{tot}} \sim k\Delta^2$ with the stiffness

$$
k \sim \left(\frac{\kappa_s \kappa_b}{R^2}\right)^{1/2} \sim \kappa_s \gamma^{-1/2}.
$$
 (2)

This result is strictly valid for small enough indentation, and more importantly for small FvK numbers. Indeed, it has been shown by Lidmar *et al.* that closed shells with large FvK numbers are strongly *faceted* with icosahedral symmetry [\[29\]](#page-10-0). This phenomenon is related to the presence of topological defects, not taken into account in the scaling derivation presented so far. From a structural point of view, these defects are associated with a few numbers of capsomers (local group of proteins) having five proteins, while all other capsomers are composed of six proteins. This can be understood first from a geometrical point of view: the virus shape and in particular its icosahedral symmetry can be reproduced by a simple triangulation network model in which each triangle represents three proteins, according to the classical Caspar and Klug approach [\[30\]](#page-10-0). Now equilateral triangles in a hexagonal phase are known to produce a perfect tiling of a flat plane or any surface having zero curvature in at least one direction. In contrast, pentagons made of equilateral triangles form a 3D conical-like structure naturally and therefore curve a surface locally in two distinct directions. This observation can be made more precise by considering the distribution of stress within thin shells. The equations describing the mechanical equilibrium of thin shells are known as the two Föppl–von Karman equations. These highly nonlinear coupled equations involve the out-of-plane deflection $w(\mathbf{r})$ and the in-plane stress $\sigma_{ij}(\mathbf{r})$. More precisely, with the introduction of Airy stress function χ , which is related to the stress tensor σ_{ij} by $\sigma_{ij} = \epsilon_{ik} \epsilon_{jl} \partial_k \partial_l \chi$, where ϵ_{ij} is the antisymmetric unit tensor, the second FvK equation reads

$$
\frac{\nabla^4 \chi(\mathbf{r})}{\kappa_s} = s(\mathbf{r}) - K_G(\mathbf{r}),
$$
\n(3)

where $s(\mathbf{r})$ is the defect density and $K_G(\mathbf{r})$ is the Gaussian curvature $[23]$. With this equation, it is straightforward to realize that the morphology of a surface with nonzero Gaussian curvature is necessarily coupled to the presence of defects, in order to minimize the elastic stress within the shell. In the simplest case these defects, known as *disclinations*, are the pentamers. Moreover, the number of these defects is fixed by the topology of the surface, as can be seen by the Euler relation [\[31\]](#page-10-0). For closed shells, this constraint imposes having exactly 12 pentamers. It has been shown by several authors that most structures obtained using self-assembly will have defects that are regularly spaced, giving rise to the icosahedral symmetry previously mentioned [\[32,33\]](#page-10-0).

As the presence of these defects is obviously related to the mechanical properties of the shell, it is expected that the stiffness is different at these locations. Several authors have analyzed both numerically and experimentally the influence of these defects [\[26,28,34–36\]](#page-10-0). In the limit of small FvK numbers, for which the shell shape is spherical, it is possible to extend the previous scaling for the stiffness estimate by modifying the stretching contribution to the energy. Indeed,

for a hemispherical cap of curvature *R*−¹ and lateral extension *d*, subjected to vanishing boundary force $\sigma_{rr}(r = d) = 0$, the in-plane stretching energy yields $\Delta E_s = (\pi/384)\kappa_s d^6/R^4$ [\[37–39\]](#page-10-0). (This approximation has been made for the sake of simplicity. We checked that the scaling results are not altered when nonzero boundary force σ_b is taken into account.) Note that this formula allows us to recover the previous scaling result for stretching energy as a function of indentation Δ by considering the hemisphere's curvature change $R^{-1} \rightarrow \rho^{-1} \sim$ $R^{-1} - 2\Delta/d^2$. In the presence of a central disclination of charge *s*, the stretching energy becomes

$$
\Delta E_{s5} = \Delta E_s + \frac{\kappa_s d^2}{32\pi} \left(s^2 - \frac{\pi s d^2}{R^2} \right). \tag{4}
$$

Expanding this energy in Δ upon curvature change leads to correcting the stiffness associated with indentation on pentamers $(s = \pi/3)$ to

$$
k_5 \sim k_6 - \frac{\pi \kappa_s}{48} \tag{5}
$$

with $k_6 \sim \kappa_s \gamma^{-1/2}$. This estimate shows that there is a slight reduction in stiffness associated with pentamers, but its amplitude is not expected to be significant.

In the opposite limit of large FvK number, the scaling behavior has been derived by Widom *et al.* [\[28\]](#page-10-0). In this limit, the viral capsid becomes faceted with icosahedral symmetry, and indentation experiments are expected to give different results, whether a vertex (i.e., a pentamer) or a face (i.e., a hexamer) of the icosahedron is probed. In the latter case, indentation is performed on a flat triangular face. The resulting stiffness in this idealized geometry can be found in the classical textbook by Timoshenko [\[22\]](#page-10-0) by solving the first FvK equation for the out-of-plane deformation profile with appropriate boundary conditions. The result scales like

$$
k_{\text{face}} \sim \frac{\kappa_b}{R^2} \sim \kappa_s \gamma^{-1}.
$$
 (6)

Finally, upon indentation of an icosahedron's vertex, the force is mostly transmitted within the plane of each face. This problem has also been considered in the classical textbook by Landau [\[19\]](#page-10-0). In this case, the in-plane deformation induced by a force localized at the edge of a face has been shown to depend in a logarithmic way on the distance to the edge. The stiffness scales therefore as

$$
k_{\text{vertex}} \sim \frac{\kappa_s}{\ln R/a} \sim \kappa_s (\ln \gamma)^{-1},\tag{7}
$$

where *a* is a small cutoff distance associated with the presence of the core of the pentamer. The comparison of these two stiffnesses k_{face} and k_{vertex} is expected to be more favorable to experimental discrimination. In particular, the stiffness of pentamers is expected to be larger in this regime. These results have been derived by Widom *et al.* using the expected elastic behavior of an icosahedron, so they should be valid for whatever shell structure fulfills this symmetry. This is expected to be the case for large FvK numbers, where the dominant stretching cost over bending induces such a shape, but it might also be valid for very small structures which are intrinsically faceted like $T = 1$ viral capsid. These structures are so small that they are composed of pentamers only, giving rise to a strongly faceted shape.

B. Numerical simulations of nanoindentation

To complement our analytical and experimental approaches, we developed numerical simulations of the indentation of thin shells. The elastic properties are reproduced by using the classical model of a triangulated surface. Within such a numerical model, in-plane deformations are taken into account with bond deformations, varying their lengths, whereas out-of-plane deformations are taken into account varying dihedral angles between adjacent faces or triangles [\[23\]](#page-10-0). The elastic energy is written as

$$
E_{\text{elas}} = \sum_{\langle i,j\rangle} \frac{1}{2} k_s (\ell_{\langle i,j\rangle} - \ell_0)^2 + \sum_{\alpha,\beta} k_b (1 - \cos \theta_{\alpha,\beta}), \quad (8)
$$

where $\ell_{\langle i,j \rangle}$ is the length of the edge $\langle i, j \rangle$. When the latter length is different from the preferred one ℓ_0 , the bond is stretched or compressed by a spring force proportional to the spring stiffness k_s . Furthermore, k_b sets the bending energetic cost for nonzero angle $\theta_{\alpha,\beta}$ between the two given adjacent faces α , β . Following the works of Seung *et al.*, for example, the relation between stretching and bending modulus in the analytical and numerical model is given by [\[23\]](#page-10-0)

$$
k_s = \frac{\sqrt{3}}{2} \kappa_s, \quad k_b = \frac{2}{\sqrt{3}} \kappa_b. \tag{9}
$$

To mimic as much as possible real viral capsids, we analyze the structure of AAV8, which has a *T* number of 1. Since this small structure might not be large enough in terms of a triangular subunit, we divide each capsomer with a 5×5 subtriangular lattice preserving the icosahedral symmetry and the nonskew capsid shape. For AAV8, we used a radius of $R = 12.5$ nm.

The thin shell interacts both with the flat substrate on which it is lying, and with the approaching spherical or conical tip. This is realized by adding an adhesion energy to all vertices, $E_{\text{ads}} = \sum_{i} V(\mathbf{r}_i)$. For both interactions, we chose a Morse potential $\overline{V}(r) = V_0(1 - e^{-r/a})^2$, which is naturally repulsive at a very short distance and attractive at a larger distance. In the limit of very large distance, the adhesion force vanishes exponentially. We added a steric cost to each face to prevent the tip from entering the triangular network, i.e., we also consider the repulsion of the barycenter of each face. Contact points with the substrate modeled as an infinite rigid flat plate are not allowed to slide tangentially, and they are fixed at their initial position. In our simulation, we set $V_0 = 2k_B T \approx 1.2$ kcal/mol and $a = 2$ Å. For some simulations, we forbade the motion of bottom vertices in order to prevent the shell from rolling.

The relaxation is done using Langevin dynamics:

$$
\mathbf{r}_i(t+dt) = \mathbf{r}_i(t) - \frac{dt}{\zeta} \nabla (E_{\text{elas}} + E_{\text{ads}}) + \left(2\frac{k_B T}{\zeta} dt\right)^{1/2} \eta,
$$
\n(10)

where \mathbf{r}_i represents the position of the vertex, η is a random force, and ζ is the damping coefficient. We chose $\zeta =$ 1 kcal/mol/nm to get good agreement with the experimental diffusion time of a capsomer, and the integration time step $dt = 400$ fs. We assumed quasistatic nanoindentation, so that the tip is lowered by 9 pm between each relaxation iteration that lasts $2 \mu s$. We ensured that an equilibrium state is reached between each tip descent. Finally, since we restricted our

investigations to spherical viruses or weakly faceted viruses for which the FvK number $\gamma \simeq O(1)$ or 10, we set $k_s =$ $200k_BT \simeq 120$ kcal/mol and $k_b = 80 k_BT \simeq 50$ kcal mol⁻¹ at ambient temperature $T = 298$ K.

From the simulations, the elastic energy of relaxed configurations is obtained, and the position of the center of the AFM tip is saved and subtracted from its initial position to give the indentation depth. The maximal indentation depth is set to $\delta_{\text{max}} = 0.3R$, provided the tip has not reached the substrate in the off-centered configurations. When coordinates of the tip in the *x*-*y* plane are not zeros, the position of the tip is carefully computed so that indentations begin at the contact of the numerical cap. Data are taken between each relaxation step, and each nanoindentation experiment is repeated several times. A quadratic fit of the form $E_{\text{elas}} = \frac{k}{2} \Delta^2$ using the NONLINEARFIT method of MATHEMATICA enables us to extract the effective stiffness *k* of the virus for each experiment, and error bars are the corresponding root-mean-square deviations resulting from the repetitions.

C. Geometrical model for position-dependent stiffness

The scaling results from the previous section assume that the indentation is performed thanks to a pointlike force applied precisely on the top of the virus. A real AFM tip has, of course, a finite radius of curvature of order a few tens of nanometers, and its value is often similar to the radius of curvature of the virus itself. This observation suggests that great care should be the rule while considering the geometry of indentation experiments. Anticipating any misalignment of the tip with the highest point of the virus, we model in this section the variation of stiffness while several lateral positions relative to the top point of the shell are vertically indented with a finite-size tip. We consider two types of tip: a spherical tip and a conical tip ending with a hemisphere.

The precise geometry considered, and the definition of the model parameters, are shown in Fig. [2.](#page-5-0) In the case of a spherical tip, these cartoons illustrate an obvious effect that leads to strong variation of measured stiffness as the tip goes from the top to the edge of the spherical shell: upon performing identical *vertical* indentations at different lateral positions from the top to the edge, the corresponding shell deformation decreases while approaching the edge. In the figure, this is highlighted by simply observing the intersection of the two spheres. However, real indentation is expected to induce deformation at a larger scale, as was already discussed in the previous section. Accordingly, the spreading of the deformation is estimated by $d_e \sim R\gamma^{-1/4}$. Nevertheless, the estimation of the overlap region provides a simple way of quantitatively evaluating the influence of the lateral tip position. In particular, the quantity of interest is the *radial* indentation length Δ_r , which represents the maximal radial deformation along the new rotated axis in direction θ . After a lengthy but straightforward geometrical calculation, one gets for a thin shell of radius R and a spherical tip of radius R_t

$$
\Delta_r = \frac{1 + \frac{\tilde{\Delta}}{R - \tilde{\Delta}} - \sqrt{1 + \left(\frac{\tilde{\Delta}}{R - \tilde{\Delta}}\right)^2 \left(\frac{1 - \cos \theta^2}{\cos \theta^2}\right)}}{1 + \frac{\tilde{\Delta}}{R - \tilde{\Delta}}} \tag{11}
$$

FIG. 2. Geometrical model for indentation. (a) The shell to be indented has a radius *R*. Upon vertically indenting on a length Δ , the shell surface deforms horizontally over a typical length 2*d*. The deformed area is assumed to have a constant curvature radius ρ . (b) Ideal force-indentation curve in the linear regime. The vertical downward arrow represents the change of force needed to perform similar vertical indentation at a different lateral position. The horizontal rightward arrow represents the change in indentation reached at a similar force at a different lateral position. (c) and (d) True deformations (yellow areas) achieved by identical vertical indentation for two different lateral positions, for a spherical tip (*C*) or a conical tip (*D*).

together with the relation between lateral tip position *r* and angle $\cos \theta = \sqrt{1 - (\frac{r}{R + R_t})^2}$ and the new variable $\tilde{\Delta} = \Delta \frac{\cos \theta}{1 + \frac{R_t}{R}}$. In the limit of small indentation, this result simplifies to $\Delta_r \simeq \Delta \sqrt{1 - (\frac{r}{R + R_t})^2}$. A weaker deformation of the shell in the radial direction is therefore associated with a weaker elastic response [see Fig. $2(b)$]. Under the assumption of linear elasticity, the effective stiffness which is position-dependent is written, therefore, as

$$
k_{\text{eff}}^{(\text{sphere})}(r) = k(0)\sqrt{1 - \left(\frac{r}{R + R_t}\right)^2}.
$$
 (12)

In the case of a conical tip of opening angle α , and terminated with a hemisphere of radius R_t , the *radial* indentation also depends on the position, at least when the spherical part of the tip is in contact with the shell. However, when the conical part is in contact with the shell, the radial indentation associated with a vertical indentation Δ is identical regardless of the lateral position. This is because one of two principal curvatures of the cone is zero, and therefore the overlap with a spherical shell along this direction is the same whenever there is a contact. More precisely, the indentation is independent of the position for $r > (R + R_t) \cos \alpha$, i.e., when $\theta = \pi/2 - \alpha$, and its constant value is simply $\Delta_r = \Delta \sin \alpha$. The stiffness is therefore

$$
k_{\text{eff}}^{(\text{cone})}[r > (R + R_t)\cos\alpha] = k(0)\sin\alpha. \tag{13}
$$

Note that although the radial indentation is constant, it is expected that transverse deformation slightly increases with the position as the conical imprint on the shell becomes larger. The two estimations Eqs. (12) and (13) are simply based on the geometry of shapes overlapping with the shell. It can be complemented, for example, by the estimation of the boundary length of the overlap region. This is demonstrated in the Supplemental Material [\[18\]](#page-10-0). Nevertheless, the major positiondependent effect is found with the true indentation length.

The previous estimation assumes that the shell does not move laterally upon vertical indentation due to strong adhesion with the substrate. In the limit of weak adhesion, the shell is indeed expected to escape the imposed constraint of indentation by rolling or sliding sideways. In the intermediate adhesion regime, the shell will be both deformed and slightly shifted sideways. This additional shift in the position is expected to lower the true indentation further, and therefore this effect might reduce further the effective stiffness. The relevance of this effect can be simply estimated in the following way. We assume that the vertical indentation produces, in addition to the true indentation along the symmetry axis of the deformation, a horizontal shift Δ_h . The relation between vertical indentation Δ , true indentation Δ_r , and horizontal shift is written as $\Delta_r = \Delta \sqrt{1 - (\frac{r + \Delta_h}{R + R_t})^2}$ to the lowest order in Δ . The equilibrium is maintained by assuming the presence of a horizontal restoring force $|F_{\text{adh}}| = k_{\text{ads}} \Delta_h$, which balances the horizontal contribution of the deformation force $|F_{\text{deform}}| = k(0) \Delta_r \frac{r + \Delta_h}{R + R_t}$. To the lowest order, the horizontal shift is finally estimated as

$$
\Delta_h \simeq \Delta \frac{k(0)}{k_{\text{ads}}} \frac{r}{R + R_t}.\tag{14}
$$

Therefore the horizontal shift increases with lateral position, and so does the horizontal force associated with adhesion bonds. This estimate shows that adhesion bond rupture might become more likely as the edge of the particle is approached and that the shell might slip or roll away from the tip. Note that the horizontal shift decreases for larger adhesion strength, as it is inversely proportional to k_{ads} .

IV. COMPARING THEORY AND EXPERIMENTS OF NANOINDENTATION OF VIRAL CAPSIDS

A. Geometrical stiffness heterogeneity

According to the protocol described in the previous experimental sections, we performed nanoindentation experiments with AFM on two different viral capsids: AAV8 and HBV $(T = 4$ capsids) (Fig. [1\)](#page-2-0). For each viral particle to be investigated, we recorded force-displacement curves at different lateral positions across the particle [Fig. $1(a)$]. The resulting curves for one particular but representative capsid is shown in Fig. $1(b)$, in which all the curves have been horizontally aligned such that force increases at the same vanishing displacement.

These curves show clearly that the elastic response is strongly dependent on the lateral position of the tip with respect to the top of the particle. Using the best forcedisplacement curve recorded for the substrate, one can obtain a similar picture on the force-indentation curves [Fig. $1(c)$]. We recall here that the indentation δ is the net deformation of the shell with respect to its initial configuration prior to tip-shell contact. Both types of curve exhibit some oscillations when vertical indentation is performed far from the top of the shell [Fig. $1(b)$]. This is likely to be associated with a partial loss of substrate adhesion: as was discussed in the analytical model, vertical indentation performed at a large

FIG. 3. Simulation of nanoindentation experiments. Snapshots of the indentation process of the triangulated surface by a spherical tip (a) or a conical tip (b). Orange double-arrows shows the area in which the deformation is not negligible. (c) Elastic energy as a function of the relative indentation Δ/R for two positions. Quadratic fit and parameters are also shown. (d) Effective stiffness as a function of the lateral position of the tip for sphere and cone. Simple geometrical model (thin and dashed lines) for effective stiffness. The stiffness is renormalized with classical prediction Eq. [\(2\)](#page-3-0).

distance from the top of the shell induces a weak deformation of the shell, but it also exerts an increasing horizontal force on the adhesion bonds between the capsid and the substrate.

Assuming a linear elastic response, the force curves can be used to estimate the stiffness, as is shown in Fig. $1(c)$, where this information is represented as a function of lateral position for a single particle. This demonstrates very clearly that the effective stiffness decreases as the indentation experiment is performed on even not so distant lateral positions from the capsid's top. This observation survives when the results on several particles are aggregated [Fig. $1(d)$]. Notice, however, that due to the oscillations in the force-displacement curves for large lateral positions, the effective stiffness in these regions has a larger uncertainty. We also performed similar indentation measurements on HBV ($T = 4$ viral capsids). In this case, we observed position-dependent stiffness, although with a weaker amplitude, as will be discussed later.

To investigate more quantitatively this position-dependent stiffness and its dependence on various parameters of the system, we used the molecular-dynamics simulations described in the previous modeling sections. The indentation process is reproduced by computing the change in elastic energy of a triangulated surface constrained by the presence of a spherical or conical tip [Figs. $3(a)$ and $3(b)$] and a flat substrate.

Fitting the expected quadratic dependence of elastic energy under the assumption of linear elasticity, we extract the stiffness for different lateral positions [Fig. $3(c)$]. The resulting stiffness variation, shown in Fig. $3(d)$, is similar to the observed experimental stiffness change. Using numerical simulations also allows us to investigate the influence of various parameters on the position-dependent stiffness, such as tip geometry (spherical or conical) and elastic parameters.

Comparing both experimental and numerical indentations, we observed first a strong variation of the stiffness. This effect can be qualitatively understood by a simple geometrical model previously mentioned: a vertical indentation performed at different lateral positions will correspond to weaker effective deformation of the shell, resulting in a weaker restoring force. From a quantitative point of view, this geometrical model provides a simple prediction that can be compared to both experiments and simulations. In the case of a spherical tip, the expected stiffness decreases toward vanishing value, and the simulations confirm this prediction. On the contrary for a conical tip, the stiffness should first decrease similarly to the spherical case, and then reach a nonzero constant value. Again, this is also observed both in the experiments and in the simulations. However, the latter curves show a stronger decrease in the stiffness compared to the simple geometrical model, suggesting the presence of other effects at play in the real system.

The first major correction that can be brought to the geometrical model is to consider horizontal displacement which is concomitant with the vertical indentation. Within the simulations, we can indeed compute for each lateral position of the tip the net horizontal shift of the shell center of mass as

FIG. 4. Lateral displacement upon off-centered vertical indentation in the simulation. The lateral displacement of the center of mass is shown as a function of the lateral position of the tip for (a) a spherical or (b) a conical tip. The dotted line shows the initial proportionality with the lateral position of the tip. (c) Stiffness profiles for various geometrical parameters, including the radius of the shell *R*, the radius of the tip R_t , and the angle of the cone α . The corrected geometrical model taking into account the leading order for horizontal displacement is shown for various configurations with dashed lines.

compared to the original position prior to indentation. This is shown in Figs. $4(a)$ and $4(b)$, respectively, for a spherical and a conical tip. It is observed within these plots that the horizontal shift increases first linearly with the lateral position, in qualitative agreement with scaling model prediction Eq. [\(14\)](#page-5-0). This horizontal shift could be associated with partial loss of adhesion sites between the shell and substrate or with a more global deformation of the shell. Plugging a linear horizontal displacement $\Delta_h = Ar$ in the original geometrical model introduces a rescaling correction to the geometrical model Eq. [\(12\)](#page-5-0) in the form $k_{\text{eff}}^{(\text{sp})} = k(0)\sqrt{1 - [r/(R + Rt)]^2(1 + A)^2}$. This predicts indeed a stronger decrease of stiffness as compared to the original geometrical model. This argument can also be applied to the conical model. Using *A* as an adjustable parameter, the corrected geometrical model shows a better agreement with experimental (Supplemental Material Fig. 2 [\[18\]](#page-10-0)) and simulated data [Fig. $4(c)$].

Simulations were also used to address the influence of various parameters on the effective stiffness measured. This is shown in Fig. $4(c)$, for example. The geometrical model predicts that the value of constant stiffness for the conical tip depends only on the opening angle of the cone [see Eq. (13)] but not on the value of sphere radius terminating the cone. This behavior is observed in the figure with two identical plateaus of stiffness for two different sphere radii. Moreover, the value of the plateau is expected to increase with α . Again, this prediction is quantitatively reproduced in the simulations [Fig. 4(c)].

B. Intrinsic stiffness heterogeneity

Up to this point, we demonstrated and discussed stiffness heterogeneity depending strongly on the lateral position of the tip. This raises the question of monitoring intrinsic stiffness heterogeneity due to the shell's structure itself. To address this question, we changed the value of elastic parameters and designed specific numerical experiments. The most promising candidates for intrinsic stiffness heterogeneity are the pentamers and hexamers of the shell. We imposed the orientation of the shell on the substrate such that either a pentamer or a hexamer is facing up the tip. For this last case, there are several possible choices. However, the simplest case is to choose a hexamer equidistant from three neighboring pentamers, i.e., this corresponds to any threefold symmetry axis of the icosahedron. In other words, this hexamer is supposed to lie at the center of the faces in the faceted shell representation. Next we induced the same vertical indentation for different lateral positions, and we deduced the effective position-dependent stiffness. For the sake of simplicity, we used $T = 9$ shells to performed nanoindentation experiments along different lateral profiles, and we achieved a different degree of faceting by using different elastic parameters. Using this protocol, we were able to compare the stiffness for different *tip trajectories*: from central pentamer to peripheric pentamer, central pentamer to peripheric hexamer, central hexamer to peripheric hexamer, and central hexamer to peripheric pentamer. Note that this procedure is different from most numerical works in the literature, where indentation is performed on top of the shell for the various shell orientations encountered by the adsorption process on the substrate. At low FvK number or equivalently for roughly spherical shells, we observed that the stiffness decrease as a function of the position is similar, regardless of the trajectory [Fig. $5(b)$]. This agrees with the scaling prediction of thin shell theory as presented earlier. In this regime, the geometrical stiffness heterogeneity is dominant, and intrinsic heterogeneity is likely to be very hard to measure in real AFM data, as suggested by the observation in the simulations.

On the contrary, we anticipate different behavior for conditions in which faceted shells are expected, namely at higher FvK number or for $T = 1$ shells. In this case, the stiffness variation upon different tip trajectories is distinct [Fig. $5(c)$]. First, the pentamers on top of the shell are stiffer than hexamers. This is also confirmed by the value of the stiffness plateau: for trajectories starting from a central pentamer, the

FIG. 5. Heterogeneity of stiffness profiles in the simulation. The profiles corresponding to particular paths along the shell's surface are shown in (a) as colored dashed lines on faceted or spherical shells. The stiffness values associated with the profiles are shown for faceted shells with $\gamma \simeq 576.5$ (b) or spherical shell with $\gamma \simeq 57.65$ (c). The underlying connectivity of the shell was kept constant and corresponds formally to the *T* = 9 icosahedral shell. Four trajectories have been considered in both cases: pentamer to pentamer (bright maroon), pentamer to hexamer (dark maroon), hexamer to hexamer (blue), and hexamer to pentamer (red). Profiles are similar for spherical shells, while they show some heterogeneity for faceted shells.

plateau is higher if the trajectory ends on a peripheric pentamer than on a peripheric hexamer. The analysis of the last two trajectories is more involved, as it results from the balance of both intrinsic and position-dependent stiffness. Going from one central hexamer toward a distal pentamer, the stiffness first increases, reflecting the intrinsic or structural variation, before decreasing at larger distances, showing the geometrical effect. This could be due to the presence of a ridge joining two pentamers on the trajectory. Finally, going from one central hexamer to a peripheric hexamer might end up in the deformation of aligned pentamers, for a large lateral position. The dispersion of trajectories shows nevertheless that the effective position-dependent stiffness is strongly dependent on the trajectory itself. At this stage, we are led to the following partial conclusion: a closed shell that has a roughly spherical shape does not exhibit significant intrinsic stiffness heterogeneity, as all stiffness profiles superimpose. Therefore, the presence of pentamers and hexamers does not necessarily lead to stiffness heterogeneity. On the contrary, when the shell exhibits significant faceting, stiffness heterogeneity is observed, and different trajectories lead to different stiffness profiles.

Next, we tried to observe similar heterogeneity signatures in the experiments on both viral capsids AAV8 ($T = 1$) and HBV $(T = 4)$ using several indentations performed at

various lateral positions. First, we compared the average trajectory *stiffness versus position* along AAV8 and HBV capsids measured for several viral particles [Fig. $6(b)$]. For both capsids, the general trend is a decrease of stiffness from the center to the edge of the particle, in agreement with the analytical and numerical model. It is possible to rescale the stiffness with the top value and the position with the appropriate size $R + R_t$, so that both profiles collapse onto a single curve, up to the experimental error [Fig. $6(c)$]. This is consistent with the geometrical modeling proposed so far. The median stiffness value on the top, 0.8 N/m for AAV8 and 0.3 N/m for HBV, is also consistent with other values from the literature [\[8,40,](#page-10-0)[41\]](#page-11-0). Interestingly, we observed that the trajectories are relatively spread around the average trajectory, the effect being larger for AAV8 than HBV capsids, as can be noticed from the values of the first and third quartile of the local stiffness distribution. To relate these results with the ones obtained in the simulation [Fig. 5(b)], we extracted first the stiffness distribution on the top of the viruses for AAV8 and HBV [Figs. $7(a)$ and $7(b)$]. These distributions show multimodal features. It is possible to use these modes of top stiffness distribution to select subpopulations of profile [Figs. $7(c)$ and $7(d)$]. In practice, this is performed by inspecting the cumulative top stiffness distribution,

FIG. 6. Comparison between AAV8 and HBV average stiffness profiles. (a) Expected geometrical shape for AAV8 and HBV, based on their T-numbers. (b) Ensemble-averaged stiffness profiles as a function of lateral positions for AAV8 *(blue curve)* and HBV *(gray curve)*. For each position, the empty circle and extremities of the bar are, respectively, the median and the first and third quartile of the local stiffness distribution. (c) Rescaling of the two average profiles by their respective maximal amplitude, showing similar profiles.

FIG. 7. Heterogeneity of stiffness profiles in the experiments. Stiffness distribution on the top of AAV8 (a) and HBV (b) viral capsids. Different subpopulations (identified by different colors) have been identified using slope ruptures in the cumulative top stiffness distributions for each curve. (c) Subpopulations of AAV8 stiffness profile with similar colors as in (a). (d) Subpopulations of HBV stiffness profile with similar colors as in (b).

which shows several significant slope ruptures (see Supplemental Material Fig. 4 [\[18\]](#page-10-0)). These subpopulation profiles are clearly distinguishable, and they strongly suggest the presence of stiffness inhomogeneity. Now, within the simulations, the exact orientation of the adsorbed particles is known, and therefore it is likely that the observed subpopulations of profiles in the experiment are representative of the very same feature: the difference in orientation of adsorbed viral particles generates different stiffness profiles. Using subpopulation profiles, we recover the observations made in Fig. [6\(b\):](#page-8-0) relative inhomogeneity is stronger for AAV8 than for HBV capsids. Moreover, since the simulation suggests that heterogeneous stiffness profiles correlate to a faceted shape, we confirm from our mechanical experiment that stiffness heterogeneity of AAV8 viral capsids is associated with a more faceted shape than HBV viral capsid. This conclusion is in agreement with the structural information on these viral capsids, as AAV8 is a $T = 1$ shell (strongly faceted), and HBV is a $T = 4$ shell (weakly faceted, more spherical) [\[40](#page-10-0)[,42,43\]](#page-11-0).

V. CONCLUSION

In the present work, we investigated the sources of stiffness heterogeneities that can arise in indentation experiments of nanosized shells like viral capsids. We first identified and rationalized a purely geometrical effect that reduces the stiffness of the shell as the vertical indentation axis is shifted towards the border of the shell. This is due to the decrease of the net deformation for identical vertical indentation. When compared to simulations, the prediction of a geometrical model that we proposed shows additionally that the adhesion properties of the shell to the substrate are rather important to account quantitatively for the observations. To our knowledge, this geometric effect was already observed on the influenza virus [\[44\]](#page-11-0), but this time it has been quantitatively investigated. It has some important consequences in the interpretation of nanoindentation experiments, as the stiffness value might be

underestimated in the case of measurements with poor positioning precision.

The recognition of this geometrical effect allowed us to define a new procedure to highlight intrinsic stiffness heterogeneity coming from the structural and elastic properties of the shell, and not from the relative geometry between the tip and the shell. Within this procedure, heterogeneity is evidenced by comparing stiffness trajectories obtained by performing several indentation experiments while scanning the investigated shell. Using simulations, we showed that if the shell is rather spherical, all trajectories collapse onto the expected geometrical profile. On the contrary, for a faceted shell resembling an icosahedron, stiffness trajectories are rather dispersed, reflecting intrinsic stiffness heterogeneity. We measured this effect experimentally for two examples of small viruses, namely AAV8 and HBV $(T = 4)$. The different degrees of intrinsic stiffness heterogeneity found for these viruses are consistent with their structure: AAV8 is a $T = 1$ shell, with strong faceting, while HBV capsid is a $T = 4$ shell with weaker faceting (more spherical shape); accordingly, the dispersion of stiffness profiles is larger for AAV8 than for HBV.

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- [1] R. Zandi, B. Dragnea, A. Travesset, and R. Podgornik, On virus growth and form, [Phys. Rep.](https://doi.org/10.1016/j.physrep.2019.12.005) **847**, 1 (2020).
- [2] I. L. Ivanovska, P. J. de Pablo, B. Ibarra, G. Sgalari, F. C. MacKintosh, J. L. Carrascosa, C. F. Schmidt, and G. J. Wuite, Bacteriophage capsids: tough nanoshells with complex elastic properties, [Proc. Natl. Acad. Sci. USA](https://doi.org/10.1073/pnas.0308198101) **101**, 7600 (2004).
- [3] J. P. Michel, I. L. Ivanovska, M. M. Gibbons, W. S. Klug, C. M. Knobler, G. J. Wuite, and C. F. Schmidt, Nanoindentation studies of full length and empty viral capsids and the effects of [capsid protein mutations on elasticity and strength,](https://doi.org/10.1073/pnas.0601744103) Proc. Natl. Acad. Sci. USA **103**, 6184 (2006).
- [4] C. Carrasco, A. Acrreira, I. A. Schaap, P. A. Serena, J. Gomez-Herrero, M. G. Mateu, and P. J. de Pablo, Dna-mediated [anisotropic mechanical reinforcement of a virus,](https://doi.org/10.1073/pnas.0601881103) Proc. Natl. Acad. Sci. USA **103**, 13706 (2006).
- [5] M. G. Mateu, Mechanical properties of viruses analyzed by [atomic force microscopy: A virological perspective,](https://doi.org/10.1016/j.virusres.2012.06.008) Vir. Res. **168**, 1 (2012).
- [6] M. G. Mateu, Assembly, stability and dynamics of virus capsids, [Arch. Biochem. Biophys.](https://doi.org/10.1016/j.abb.2012.10.015) **531**, 65 (2013).
- [7] C. Zeng, M. Hernando-Perez, B. Dragnea, X. Ma, P. van der Schoot, and R. Zandi, Contact Mechanics of a Small Icosahedral Virus, Phys. Rev. Lett. **119**[, 038102 \(2017\).](https://doi.org/10.1103/PhysRevLett.119.038102)
- [8] C. Zeng, S. Moller-Tank, A. Asokan, and B. Dragnea, Probing the link among genomic cargo, contact mechanics, and nanoin[dentation in recombinant adeno-associated virus 2,](https://doi.org/10.1021/acs.jpcb.6b10131) J. Phys. Chem B **121**, 1843 (2017).
- [9] N. Martin-Gonzalez, P. Ibanez-Freire, A. Ortega-Esteban, M. Laguna-Castro, C. San Martin, A. Valbuena, R. Delgado-Buscalioni, and P. J. de Pablo, Long-Range Cooperative Disassembly and Aging During Adenovirus Uncoating, Phys. Rev. X **11**[, 021025 \(2021\).](https://doi.org/10.1103/PhysRevX.11.021025)
- [10] F. Eghiaian, I. A. Schaap, A. des Georges, J. J. Skehel, and C. Veigel, The influenza virus mechanical properties are dominated by its lipid envelope, Biophys. J. **96**[, 15 \(2009\).](https://doi.org/10.1016/j.bpj.2008.12.981)
- [11] N. Kol, Y. Shi, M. Tsvitov, D. Barlam, R. Z. Shneck, M. S. Kay, and I. Rousso, A stiffness switch in human immunodeficiency virus, Biophys. J. **92**[, 1777 \(2007\).](https://doi.org/10.1529/biophysj.106.093914)
- [12] M. Castelnovo, Viral self-assembly pathway and [mechanical stress relaxation,](https://doi.org/10.1103/PhysRevE.95.052405) Phys. Rev. E **95**, 052405 (2017).
- [13] M. Aznar, A. Luque, and D. Reguera, Relevance of capsid structure in the buckling and maturation of spherical viruses, Phys. Biol. **9**[, 036003 \(2012\).](https://doi.org/10.1088/1478-3975/9/3/036003)
- [14] A. Salvetti, S. Oreve, G. Chadeuf, D. Favre, Y. Cherel, P. Champion-Arnaud, J. David-Ameline, and P. Moullier, Factors influencing recombinant adeno-associated virus production, [Hum. Gene Ther.](https://doi.org/10.1089/hum.1998.9.5-695) **9**, 695 (1998).
- [15] L. Lecoq, S. Wang, T. Wiegand, S. Bressanelli, M. Nassal, B. Meier, and A. Bockmann, Solid statement [13c-15n] nmr res[onance assignment of hepatitis b virus core protein,](https://doi.org/10.1007/s12104-018-9810-y) Biomolec. NMR Assign. **12**, 205 (2018).
- [16] J. Z. Porterfield, M. S. Dhason, D. D. Loeb, M. Nassal, S. J. Stray, and A. Zlotnick, Full-length hepatitis b virus core protein packages viral and heterologous rna with similarly high levels of cooperativity, J. Virol. **84**[, 7174 \(2010\).](https://doi.org/10.1128/JVI.00586-10)
- [17] J. Heger-Stevic, P. Zimmermann, L. Lecoq, B. Bottcher, and M. Nassal, Hepatitis b virus core protein phosphorylation: identification of the srpk1 target sites and impact of their occupancy on

[rna binding and capsid structure,](https://doi.org/10.1371/journal.ppat.1007488) Plos. Pathogens **14**, 1007488 (2018).

- [18] See Supplemental Material at [http://link.aps.org/supplemental/](http://link.aps.org/supplemental/10.1103/PhysRevE.104.064408) 10.1103/PhysRevE.104.064408 for additional figures and calculations.
- [19] L. D. Landau and E. M. Lifshitz, *Theory of Elasticity* (Pergamon Press, Oxford, 1970).
- [20] E. Ventsel and T. Krauthammer, *Thin Plates and Shells: Theory, Analysis and Applications* (CRC Press, Boca Raton, FL, 2001).
- [21] C. R. Calladine, *Theory of Shell Structures* (Cambridge University Press, Cambridge, 2010).
- [22] S. Timoshenko and S. Woinowsky-Krieger, *Theory of Plates and Shells*, 2nd ed. (McGraw-Hill, India, 2010).
- [23] H. S. Seung and D. R. Nelson, Defects in flexible membranes with crystalline order, Phys. Rev. A **38**[, 1005 \(1988\).](https://doi.org/10.1103/PhysRevA.38.1005)
- [24] M. Bowick, D. R. Nelson, and A. Travesset, Interacting topo[logical defects on frozen topographies,](https://doi.org/10.1103/PhysRevB.62.8738) Phys. Rev. B **62**, 8738 (2000).
- [25] A. Travesset, Ground state of a large number of particles on a frozen topography, Phys. Rev. E **72**[, 036110 \(2005\).](https://doi.org/10.1103/PhysRevE.72.036110)
- [26] M. Buenemann and P. Lenz, Elastic properties and mechanical [stability of chiral and filled capsids,](https://doi.org/10.1103/PhysRevE.78.051924) Phys. Rev. E **78**, 051924 (2008).
- [27] S. Komura, K. Tamura, and T. Kato, Buckling of spherical shells adhering onto a rigid substrate, [Eur. Phys. J. E](https://doi.org/10.1140/epje/e2005-00038-5) **18**, 343 (2005).
- [28] M. Widom, J. Lidmar, and D. R. Nelson, Soft modes near [the buckling transition of icosahedral shells,](https://doi.org/10.1103/PhysRevE.76.031911) Phys. Rev. E **76**, 031911 (2007).
- [29] J. Lidmar, L. Mirny, and D. R. Nelson, Virus shapes and buck[ling transitions in spherical shells,](https://doi.org/10.1103/PhysRevE.68.051910) Phys. Rev. E **68**, 051910 (2003).
- [30] D. L. Caspar and A. Klug, Physical principles in the construction of regular viruses, [Cold Spring Harbor Symp. Quant. Biol.](https://doi.org/10.1101/SQB.1962.027.001.005) **27**, 1 (1962).
- [31] L. Giomi and M. Bowick, Crystalline order on Riemannian manifolds with variable Gaussian curvature and boundary, Phys. Rev. B **76**[, 054106 \(2007\).](https://doi.org/10.1103/PhysRevB.76.054106)
- [32] S. Li, P. Roy, A. Travesset, and R. Zandi, Why large icoshedral [viruses need scaffolding proteins,](https://doi.org/10.1073/pnas.1807706115) Proc. Natl. Acad. Sci. USA **115**, 10971 (2018).
- [33] L. Menou and M. Castelnovo, Mechanical stress relaxation in molecular self-assembly, Soft Matter **15**[, 6180 \(2019\).](https://doi.org/10.1039/C9SM00761J)
- [34] M. Gibbons and W. Klug, Influence of non uniform geometry on nanoindentation of viral capsids, Biophys. J. **95**[, 3640 \(2008\).](https://doi.org/10.1529/biophysj.108.136176)
- [35] W. S. Klug, W. H. Roos, and G. J. L. Wuite, Unlocking Internal [Prestress from Protein Nanoshells,](https://doi.org/10.1103/PhysRevLett.109.168104) Phys. Rev. Lett. **109**, 168104 (2012).
- [36] A. Llauro, B. Schwartz, R. Koliyatt, P. de Pablo, and T. Douglas, Tuning viral capsid nanoparticle stability with symmetrical morphogenesis, ACS Nano **10**[, 8465 \(2016\).](https://doi.org/10.1021/acsnano.6b03441)
- [37] S. Schneider and G. Gompper, Shapes of crystalline domains on spherical fluid vesicles, [Europhys. Lett.](https://doi.org/10.1209/epl/i2004-10464-2) **70**, 136 (2005).
- [38] G. M. Grason, Defects in crystalline packings of twisted fila[ment bundles. i. continuum theory of disclinations,](https://doi.org/10.1103/PhysRevE.85.031603) Phys. Rev. E **85**, 031603 (2012).
- [39] A. Azadi and G. M. Grason, Defects in crystalline packings of twisted filament bundles. ii. dislocations and grain boundaries, Phys. Rev. E **85**[, 031604 \(2012\).](https://doi.org/10.1103/PhysRevE.85.031604)
- [40] W. H. Roos, M. M. Gibbons, A. Arkhipov, C. Uetrecht, N. R. Watts, P. T. Wingfield, A. C. Steven, A. J. R. Heck, K. Schulten,

W. S. Klug, and G. J. L. Wuite, Squeezing protein shells: how continuum elastic models, molecular dynamics simulations, and [experiments coalesce at the nanoscale,](https://doi.org/10.1016/j.bpj.2010.05.033) Biophys. J. **99**, 1175 (2010).

- [41] C. Uetrecht, C. Verluis, N. R. Watts, W. H. Roos, G. J. L. Wuite, P. T. Wingfield, A. C. Steven, and A. J. R. Heck, High resolution spectroscopy of viral assemblies: molecular composition [and stability of dimorphic hepatitis b virus capsids,](https://doi.org/10.1073/pnas.0800406105) Proc. Natl. Acad. Sci. USA **105**, 9216 (2008).
- [42] S. A. Wynne, R. A. Crowther, and A. G. W. Leslie, The crystal [structure of the human hepatitis b virus capsid,](https://doi.org/10.1016/S1097-2765(01)80009-5) Mol. Cell **3**, 771 (1999).
- [43] X. Yu, L. Jin, J. Jih, C. Shih, and Z. H. Zhou, 3.5 angstroms cryoem structure of hepatitis b virus core assembled from fulllength core protein, PLoS ONE **8**[, 69729 \(2013\).](https://doi.org/10.1371/journal.pone.0069729)
- [44] S. Li, F. Eghiaian, C. Sieben, A. Herrmann, and I. A. T. Schaap, [Bending and puncturing the influenza lipid envelope,](https://doi.org/10.1016/j.bpj.2010.12.3701) Biophys. J. **100**, 637 (2011).