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Quantifying the Contribution of Dispersion Interaction and Hydrogen Bonding to the Anisotropic Elastic Properties of Chitin and Chitosan

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Abstract

² The elastic tensors of chitin and chitosan allomorphs were calculated using density ³ functional theory (DFT) with and without dispersion correction, and compared with ⁴ experimental values. The longitudinal Young's moduli were 114.9 GPa or 126.9 GPa for ⁵ α -chitin depending on hydrogen bond pattern, 129.0 GPa for β -chitin, and 191.5 GPa for ⁶ chitosan. Furthermore, the moduli were found to vary between 17.0 to 52.8 GPa in the

transverse directions, and between 2.2 to 15.2 GPa in shear. Switching off the dispersion
 correction lead to a decrease in modulus by up to 63%, depending on the direction. The
 transverse Young's moduli of α-chitin strongly depended on the hydroxylmethyl group
 conformation coupled with the dispersion correction, suggesting a synergy between
 hydrogen bonding and dispersion interactions. The calculated longitudinal Young's
 moduli were in general higher than experimental values obtained in static conditions.

⁸ Introduction

Chitin is a linear natural polymer consisting of β -1,4 linked N-acetyl-glucosamine, which can 9 be seen as an analogue to cellulose with the N-acetylamine group substituting the hydroxyl 10 group in the C2 position. It is biosynthesized by various living organisms and is found 11 as structural component in, e.g., the exoskeleton of arthropods, and in the cell wall of 12 fungi and yeast.¹ By deacetylation, anhydro-N-acetyl-glucosamine residues can be converted 13 to anhydro-glucosamine and its polymer β -1,4-linked glucosamine, or chitosan. Industrial 14 chitosan is usually a randomly deacetylated product and is not necessarilly deacetylated to 15 100%. However, structural studies are usually done on 100% deacetylated samples which is 16 a homopolymer of glycosamine residue. 17

One particular aspect of chitin and chitosan compared to cellulose is their bioresorbability. While cellulose cannot be metabolized by mammalians, including humans, both chitin and chitosan can be degraded inside the mammalian body giving them an important advantage over cellulose for biomedical applications. The intrinsic mechanical properties, e.g. stiffness, are another attractive aspect² of chitin and chitosan, which becomes important in the context of bio-based functional materials.

²⁴ Depending on its biological origin, the chitin polymer chain is usually assembled into ²⁵ either of the two crystalline allomorphs, namely α -chitin³ and β -chitin.⁴ The main difference ²⁶ between them is the chain polarity, which is antiparallel in α -chitin and parallel-up in β -



Figure 1: Structural details of α -chitin-A, α -chitin-B, β -chitin, and chitosan. The upper four snapshots present the cross-section morphology. The middle one shows the hydroxymethyl isomerization in α -chitin-A and α -chitin-B. The bottom ones illustrate the HB network.

chitin.⁵ In both cases, they form crystalline microfibrils of nanometer-sized lateral dimension, 1 similar to native cellulose. Fully deacetylated chitosan can be prepared from chitin in solid 2 state keeping the macroscopic arrangement.⁶ Structural studies are carried out on model 3 systems of high crystalline form that are thus either fully N-acetylated or fully deacetylated. 4 The longitudinal modulus is of interest since the function of these fibrils primarily is to 5 sustain tensile loads in structural and functional materials. Furthermore, the longitudinal 6 modulus is the least difficult one to approach experimentally. There are two reports of 7 the crystal modulus of α -chitin, one for chitosan and none for the β -chitin. Using X-ray 8 diffraction, Nishino et al.⁷ reported a longitudinal modulus of 41 and 65 GPa of the crystalline 9 regions in α -chitin and in chitosan, respectively. They also concluded the modulus of α -chitin 10 to be constant in the temperature range -190 to 150 °C. Using a similar approach, Ogawa et 11 al.⁸ reported a slightly higher value of 59.3 \pm 11.3 GPa for α -chitin. 12

¹³ Transverse mechanical properties and shear modulus are also important as the crystals

are subject to complex stress environments, both in their native setting (in the case of
chitin) and during processing. In addition to the experimental studies mentioned above, the
mechanical properties of chitin have been investigated using atomistic and coarse-grained
force-field based molecular dynamics simulation, ⁹⁻¹² but determination of the full elastic
tensors has not yet been reported.

Poisson's ratio is another interesting parameter to look at, since the measurement is
less prone to artifact and uncertain assumptions. Even if stress transfer in the material or
accurate measurement of the cross-sectional area is challenging, the Poisson's ratio does not
depend on the absolute value of stress on the sample and can be robust.

In our recent study, we analyzed the effect of dispersion interactions on cellulose crystals,¹³ by switching the dispersion correction¹⁴¹⁵ on and off. Here we apply the same approach to chitin and chitosan. For α -chitin the naturally occurring isomerization of the hydroxymethyl group¹⁶ (here denoted α -chitin-A and α -chitin-B, see computational details) further allowed us to analyze the contribution of hydrogen bonds (HBs) on the elastic modulus was quantified.

¹⁶ Computational method and experimental section

Periodic DFT calculations were performed using the quantum espresso package (version 17 6.6)^{17,18} either with or without applying pairwise DFT-D2 correction.¹⁹ The ElaStic package 18 was used for the estimation of the elastic tensors.^{20,21} Initial crystal coordinates were derived 19 from X-ray and neutron data.^{3,4,6,22} Three crystals, namely, α -chitin, β -chitin, and chitosan, 20 in their anhydrous form, were considered in this work. Due to the hydroxymethyl group 21 induced structural complexity of α -chitin, two different structures of α -chitin taken from the 22 DFT optimization of Deringer et al.¹⁶ were considered, denoted α -chitin-A (qt for O6 in four 23 anhydrous N-acetyl-glucosamine residues, shown in Figure 1), and α -chitin-B (qq for four 24 O6) respectively. A third one, with mixed qq and qt for O6, departed from the orthorhombic 25

unit cell during geometry optimization and was thus abandoned. A k-grid of $2 \times 2 \times 2$ was 1 applied to α -chitin-A, α -chitin-B, β -chitin, and chitosan. The kinetic energy cutoff was 160 2 Ry. The convergence threshold of total energy and forces for ionic minimization are 1.0e-6 3 Ry and 1.0e-5 Ry/bohr, respectively. Periodic calculations were carried out using Quantum 4 Espresso (version 6.6)^{17,18} in combination with ElaStic,^{20,21} a universal tool for calculating 5 elastic constants from first principles. Both α -chitin and chitosan have orthorhombic unit 6 cells,^{3,6} while β -chitin⁴ has a monoclinic one. The elastic tensor was determined using the 7 energy-strain approach. Here, 9 (orthorhombic) or 13 (monoclinic) combinations of strain 8 deformations were applied to deduce the corresponding 9 or 13 tensor elements. For each 9 strain deformation, 11 frames with the strain amplitude varying linearly from -0.01 to 0.0110 were optimized with fixed unit cell and the energy fitted with a parabolic function. To 11 visualize the feature of calculated elastic tensors, the tensors were further decomposed into 6 12 eigenvalues λ_i and 6 pairs of eigentensors of stress and strain equivalent where $\sigma_i = \lambda_i \epsilon_i$ The 13 stress/strain tensors have 6 independent components and can be visualized by 3D surface 14 contour plots as proposed in PAScal software.²³ The plots were generated using the online 15 Anisotropic calculator from Zuluaga et al.²⁴ and further processed using the PARAVIEW 16 software.²⁵ 17

¹⁸ The experimental Poisson's ratio was obtained from the displacement of the diffraction ¹⁹ spots in previous studies of crystal moduli measurements.⁸ The X-ray diffraction strain ²⁰ measurement of β -chitin was carried out by using extracellular crystalline filaments from ²¹ *Thalassiosira weissflogii* embedded in PVA, following the same protocol and analyss as recent ²² study on cellulose.²⁶ The strain of d-spacings of equatorial spots with respect to the strain ²³ meridian spots are also shown in Fig S1 and S2.

Stiffness tensor (GPa)												
	C11	C12	C13	C14	C15	C16						
	C12	C22	C23	C24	C25	C26						
	C13	C23	C33	C34	C35	C36						
	C14	C24	C34	C44	C45	C46						
	C15	C25	C35	C45	C55	C56						
	C16	C26	C36	C46	C56	C66						
		Ι	Disp cor	rectior	1			No	o disp co	orrectio	on	
	27.8	4.5	1.3	0	0	0	21.7	1.2	1.6	0	0	0
		30.6	6.1	0	0	0		12.6	2.9	0	0	0
			126.9	0	0	0			110.4	0	0	0
α -chitin-A				10.1	0	0				5.7	0	0
					3.9	0					3.4	0
						4.6						3.3
	26.4	8.4	2.3	0	0	0	16.4	-1.8	1.1	0	0	0
		52.8	4.1	0	0	0		26.2	4.6	0	0	0
α -chitin-B			114.9	0	0	0			115.3	0	0	0
				14.8	0	0				11.3	0	0
					2.2	0					2.6	0
						5.3						4.3
	21.9	5.7	5.9	0	0	-1.8	17.8	3.6	4.1	0	0	-1.3
		17.0	7.6	0	0	4.1		9.9	5.6	0	0	0.7
β -chitin			129.0	0	0	3.6			106.4	0	0	0.2
				15.2	1.2	0				6.9	-0.0	0
					4.1	0					3.9	0
						4.0						3.4
	24.4	17.1	12.7	0	0	0	18.8	4.3	8.0	0	0	0
chitosan		27.5	8.7	0	0	0		10.1	3.0	0	0	0
			191.5	0	0	0			142.6	0	0	0
				3.5	0	0				2.4	0	0
					10.1	0					7.6	0
						4.3						3.2

Table 1: Elastic tensor elements of chitin and chitosan crystals with or without of dispersion correction. The unit is GPa.

¹ Result and discussion

² Estimation of stiffness tensor from DFT calculation

Table 1 shows calculated elastic tensor elements of the four crystals. The one corresponding 3 to longitudinal deformation C33 was 126.9 and 114.9 GPa for α -chitin in the two different 4 hydroxymethyl group conformations (A and B, respectively). It was 129.0 GPa for β -chitin, 5 and 191.5 GPa for chitosan. The corresponding longitudinal Young's moduli, calculated 6 as $1/S_{33}$ from the compliance matrix S_{ij} (see Table S1 and S2) were 125.6, 114.4, 122.7, 7 and 184.8 GPa, respectively. Simulated moduli are indeed much higher than experimentally 8 determined values (41 and 59.3 \pm 11.3 GPa⁷ and 65 GPa,⁸ using two different X-ray tech-9 niques). This difference is probably related to the low off-diagonal elements calculated by 10 DFT.²⁶ In reality, when the fibril is stretched in the axial direction, the lateral dimension 11 decreases to keep the volume unchanged, and thus the energy increase is attenuated. This 12 process is probably slow and the molecular deformation to accommodate the available space 13 might be incompatible with strict periodicity in all directions. The experimental estimates 14 for chitin and chitosan by Nishino et al.⁷ look reasonable especially as the f-value, i.e. the 15 modulus multiplied by the chain cross-sectional area, are similar. Also in previous DFT 16 calculations on crystalline cellulose, moduli were high compared to the static deformation 17 experiments,^{26,27} and the simulated Poisson's ratio was low even in classical MD calcula-18 tion.²⁸ Thus, there are definitely limitations for both DFT and classical MD approaches. 19

Figure 1 shows the molecular structures of the four crystals. Chain polarity is antiparallel in both α -chitin and chitosan, and parallel-up in β -chitin. Both β -chitin and chitosan have their hydrated crystalline forms, but not α -chitin. The interchain HB network only extended along the *a*-axis within a single molecular sheet, but not within the sheets parallel to the *b*-axis, in both α -chitin-A and β -chitin. In both α -chitin-B and chitosan, however, the interchain HB network extended along both the *a* and *b* axes simultaneously.

Transverse elastic constants varied from 17.0 to 52.8 GPa and the shear moduli were in



Figure 2: 3D representation of the elastic modulus isosurfaces of crystalline chitin and chitosan with (grid contour) or without dispersion correction (solid contour) from two different perspectives, the isosurfaces in B is enlarged for α -chitin-A, α -chitin-B, β -chitin, chitosan, for better visualization. The views of two crystalline chitin allomorphs and chitosan labeled with unit cell and deformation vectors as well as the corresponding characteristic crystalline planes. The cross-section snaphots of each crystal are shown in the same viewing angle as the corresponding iso-surface in B. The unit cell parameter *a* is parallel to *x*, *c* is parallel to *z*, and *y* lies in the *ab* plane.

- ¹ the range 2.2 GPa to 15.2 GPa (Table 1), both of which are much smaller than that of the
- ² longitudinal ones, indicating strong anisotropy in all crystals.

Table 2: The unit cell parameters of crystalline α -chitin,³ β -chitin,⁴ and chitosan,²² from experiment and DFT calculation.⁶ Lengths in Å, angles in degrees and volumes in Å³. The difference in percentage is calculated through (No disp.-Expt.)/Expt.×100%.

		a	b	с	α	β	γ	Volume
α -chitin-A	Expt. ³	4.749	18.890	10.33	90.0	90.0	90.0	926.7
	Disp.	4.587	18.495	10.32	90.0	90.0	90.0	875.5
	No disp.	4.908	19.591	10.43	90.0	90.0	90.0	1002.9
	Rel. incr.	3.3%	3.7%	1.0%				8.2%
α -chitin-B	Expt. ³	4.749	18.890	10.33	90.0	90.0	90.0	926.7
	Disp.	4.656	18.166	10.33	90.0	90.0	90.0	873.7
	No disp.	4.971	18.953	10.47	90.0	90.0	90.0	986.4
	Rel. incr.	4.7%	0.3%	1.4%				6.4%
β -chitin	Expt. ⁴	4.819	9.239	10.38	90.0	90.0	97.2	458.5
	Disp.	4.725	8.952	10.36	90.0	90.0	95.3	436.3
	No disp.	4.969	9.791	10.44	90.0	90.0	98.7	502.1
	Rel. incr.	3.1%	6.0%	0.6%				9.5%
chitosan	Expt. ²²	8.129	8.347	10.31	90.0	90.0	90.0	699.6
	Disp.	7.942	8.173	10.37	90.0	90.0	90.0	673.1
	No disp.	8.217	8.855	10.44	90.0	90.0	90.0	759.6
	Rel. incr.	1.1%	6.1%	1.3%				8.6%

³ Influence of Dispersion Energy

With dispersion correction switched off, the estimated volume of the unit cell increased by 4 5-9%, and most of the unit cell parameters expanded (Table 2), indicating a relative looser 5 assembly of macromolecular chains than with dispersion correction switched on. At the same 6 time, estimated elastic constants decreased between -18% to -63% depending on the unit cell 7 direction (see Figure 1 and Table 1). The largest decrease occurred in the direction of the b-8 axis for all crystals (except for α -chitin-B), which is normal to the glucosamine ring plane for 9 chitosan, but parallel to the acetylated glucosamine ring plane in both α -chitin and β -chitin 10 (Figure 1). It is interesting to note that the E33 of α -chitin-B is not affected by switching 11 off dispersion correction. The relative decrease was smaller along the unit cell a-axis due to 12

the resistance from the interchain HB network, which strengthens chain association in that
direction, as discussed in detail below.

³ Contribution from HB

From a structural perspective, both the O6-HO6 group and the nearby N2-H2 group in 4 α -chitin-A orient themselves toward the oxygen in the C=O groups, thus forming a dual 5 HB between the (100) planes and within (010) sheets, while the gt conformation of the 6 hydroxymethyl group prohibits HBs between (010) planes. In the other form, α -chitin-B, 7 in which O6 is found in the gg conformation, the N2-H2···O=C HBs dominate within the 8 (010) planes, while O6HO6...O6 HBs are present between (010) planes (See Figure 1). Such 9 different OH isomerization leads to different density of interchain HBs between adjacent 10 (010) planes in A and B: one per anhydrous glucosamine residue in type B and zero in type 11 A respectively (see Table 3 and Figure 3). The HB density between (100) planes within 12 the same (010) sheet is also different between the A and B. It is two per residue in the A 13 form and one per residue in B. These differences in HB pattern originating from different 14 conformations lead to significant differences for the elastic tensor in the transverse direction. 15 As shown in Figure 1, and in both Table 1 and Table 3, the effects from HBs and dispersion 16 energy on the transverse modulus are coupled. A nearly two-fold difference in E22 (30.6 17 versus 52.8 GPa) is observed between α -chitin-A and α -chitin-B. By switching off dispersion 18 correction while retaining the HO6...O6 interchain HBs, these two values reduce to 12.6 and 19 26.2 GPa, respectively. On the other hand, E11 is 27.8 and 26.4 for A and B, respectively, 20 and deceases to 21.7 and 16.4 as the dispersion correction is cancelled, thus showing much 21 smaller reduction. Therefore, in the direction with more HBs, cancellation of dispersion leads 22 to larger reduction of the modulus. This is because the HBs lead to tighter chain assembly 23 and hence larger dispersion interaction. 24

²⁵ The effect of HBs on the moduli can be further estimated by comparison between cellulose ²⁶ I β and α -chitin. The number of HBs between two chains within the (200) plane of cellulose

		E22			E11	
	α -chitin-A	α -chitin-B	B-A	α -chitin-A	α -chitin-B	B-A
HB density	0	0.5	0.5	2	1	-1
E (Disp)	30.6	52.8	22.2	27.8	26.4	-1.4
E (No disp)	12.6	26.2	13.6	21.7	16.4	-5.3
ΔE	+18.0	+26.6	+8.6	+6.1	+10.0	+3.9

Table 3: Effect of HB density (number of HBs per anhydrous glucan residue) and dispersion correction on transverse elastic modulus E (GPa).

I β is one per glucose residue. For α -chitin-A and B within the (110) and (1-10) planes, this 1 number is zero and one respectively. Although the HB density is the same between cellulose 2 I β and α -chitin-B. The HB directions in I β is perfectly parallel to the (200) plane whereas 3 the (110) and (1-10) planes in α -chitin-B both showed a tilt angle of 15° with respect to 4 the (100) plane. Therefore the "effective" HB density that contributes to C22 is less than 5 one per residue. The Young's modulus decreased monotonously as the density of interchain 6 HBs decreased (Figure 3), and was calculated to 99.2, 52.8, and 30.6 GPa, respectively.¹³ 7 With dispersion switched off, these values decreased to 68.2, 26.2, and 12.6 GPa, thus being 8 reduced by 31.0, 26.6, and 18.0 GPa, respectively. Along the chain staggering direction (010 9 plane), the HB density is 0, 1, and 2 for I β , α -chitin-B, and α -chitin-A. Here, C11 increased 10 monotonously as 17.6, 26.4, and 27.8 GPa with dispersion on, and 5.3, 16.4, 21.7 GPa with 11 dispersion off. 12

Table 4: The elastic tensile and shear modulus of chitin and chitosan (GPa).

		Disp corre	ection		No disp correction				
	α -chitin-A	α -chitin-B	β -chitin	chitosan	α -chitin-A	α -chitin-B	β -chitin	chitosan	
E11	27.8	26.4	21.9	24.4	21.7	16.4	17.8	18.8	
E22	30.6	52.8	17.0	27.5	12.6	26.2	9.9	10.1	
E33	126.9	114.9	129.0	191.5	110.4	115.3	106.4	142.6	
G23	10.1	14.8	15.2	3.5	5.7	11.3	6.9	2.4	
G13	3.9	2.2	4.1	10.1	3.4	2.6	3.9	7.6	
G12	4.6	5.3	4.0	4.3	3.3	4.3	3.4	3.2	

Besides the non-covalent interactions, the modulus along one specific transverse direction is also partially governed by the covalent bonds. With dispersion correction applied, the es-



Figure 3: Illustration of interchain HB patterns and numbers in α -chitin-A, α -chitin-B, and cellulose I β .

timated transverse modulus is larger in the direction of high covalent bond density, as can be 1 seen by comparing E11 and E22 in Table 4. However, the contribution of HBs and dispersion 2 energy as well as the covalent bonds to elastic modulus is strongly correlated and cannot 3 be decomposed in a simple way. For instance, one cannot conclude that HBs contribute a 4 certain percentage to the stiffness by (artificially) removing the HBs and calculate the mod-5 ulus. This is because of (1) removing HBs also alters other parameters simultaneously (chain 6 packing etc.) that affects the stiffness too, and (2) separate contributions to the stiffness are 7 not necessarily additive. However, if one scales the absolute difference in modulus by the 8 number of HBs in that direction, one finds that the contribution from dispersion interaction 9 to the elastic mechanical properties is comparable to that from HBs, both along the b and a10 axis of α -chitin. 11

¹² The HB pattern in β -chitin is similar to that in α -chitin-A. The difference between the ¹³ two allomorphs is (1): the adjacent molecular chains, which are anti-parallel in the α , but parallel in β, and (2): the chain packing, which gives a zig-zag patterned cross-section in α,
and a more planar arrangement in β. The longitudinal modulus of the two forms are almost
the same (126.9 GPa versus 114.9 GPa) due to the same β-1,4-glycosidic linkages, the same
intrachain HO3... O5 HBs along the chain direction, as well as similar density. However, the
transverse modulus is slightly smaller for the β form. In the elastic tensor surface in Figure
2B, one recognizes that the maximum elastic constant around x and y axis (Figure 1) in the
cross-section is similar for α-chitin-A and β-chitin.

The higher longitudinal modulus E33 of chitosan compared to the other crystals (Table 9 4) is partly due to more efficient chain packing leading to higher number of chains per unit 10 area, but after normalization by volume it is still about 17% higher than the stiffest chitin 11 allomorph.

¹² Energy decomposition

Several numerical methods have been developed specifically for decomposing non-covalent 13 interactions in DFT calculations, such as reduced density gradient²⁹ and independent gra-14 dient method.³⁰ In the present work, decomposition was achieved using a simpler approach 15 where dispersion and electrostatic energy components were estimated by systematic unit cell 16 enlargement while simultaneously switching the dispersion correction on and off during DFT 17 geometry optimization. As shown in Figure 4, the original unit cell of β -chitin was enlarged 18 by a factor of two in either the a or b direction, or both. With periodic boundary condition 19 applied, this results in four different structures that represent the condensed phase, two dif-20 ferent cases of separated sheets, and the isolated chains, respectively. These are referred to 21 as 1x1, 2x1, 1x2, and 2x2, respectively (Figure 4). For α -chitin and chitosan, whose unit 22 cells contains two chains, a fifth structure was created by deleting one of the chains in the 23 fully separated case. This structure is here denoted $2x2^*$ (Figure S3 & S4 & S5). The energy 24 difference between these structures is the intermolecular energy needed for the separation of 25 molecular sheets or chains. 26



Figure 4: Graphical representation of the four enlarged cells of β -chitin for DFT energy optimization.

By switching off and on dispersion correction, eight (or ten) terms of total DFT energies 1 and four (or five) terms of dispersion correction energies were obtained after energy mini-2 mization. These are shown in Table 5. The total energy is split into intra- and inter-chains 3 energies and further into their respective electrostatic and dispersion contributions. The 4 total non-covalent interaction is the sum of four components: the inter-chain electrostatic 5 energy (E_{interE}) , the intra-chain electrostatic energy (E_{intraE}) , the inter-chain dispersion en-6 ergy (E_{interD}) , and the intra-chain dispersion energy (E_{intraD}) . Since the morphology and 7 the geometry of the individual polymer chains did not vary significantly, we assume both 8 E_{intraE} and E_{intraD} to be constant. The E_{intraD} (per unit cell) is -141.87 kJ/mol, calculated 9 from the 2x2 structure. Since the total dispersion energy, $E_{disp.corr.} = E_{intraD} + E_{interD}$, both 10 E_{interE} and E_{interD} can be obtained and are presented in Table 5. The results show that 11 E_{interD} is comparable to E_{interE} . The HB energy can be considered as one part of E_{interE} , 12 thus necessarily smaller than the total value. Therefore, we conclude that the intermolecular 13 dispersion energy is larger than the HB energy in β -chitin. Thus, the dispersion influences 14

- 1 the elastic muduli more than HB does. α -chitin and chitosan also follows this pattern as
- ² shown in Table 5.

Table 5: The energy decomposition of chitin and chitosan. \ast stands for one chain in the unit cell.

Unit ce	ell	Energy (kJ/mol per unit cell)								
		$E_{vdwcorr.}$	$E_{novdwcorr.}$	$E_{disp.corr.}$	E_{InterE}	E_{IntraD}	E_{InterD}	E_{IntraE}		
β -chitin	1x1	-370.93	-125.47	-260.77	-110.16	-141.87	-118.90			
	2x1	-313.45	-109.74	-210.77	-103.28	-141.87	-68.30			
	1x2	-180.83	-10.24	-176.81	-4.01	-141.87	-34.94	Very large		
	2x2	-146.96	0	-141.87	-5.09	-141.87	0			
α -chitin-A	1x1	-364.82	-118.81	-260.97	-103.84	-140.88	-120.09			
	2x1	-335.89	-111.88	-234.56	-101.34	-140.88	-93.68			
	1x2	-176.28	-7.56	-170.28	-5.99	-140.88	-29.40	Very large		
	2x2	-158.47	-3.02	-154.58	-3.88	-140.88	-13.70			
	2x2*	-142.70	0.00	-140.88	-1.82	-140.88	0.00			
α -chitin-B	1x2	-347.00	-96.45	-263.12	-83.88	-136.18	-126.93			
	2x1	-303.09	-82.54	-231.55	-71.54	-136.18	-95.37			
	1x2	-176.83	-20.45	-168.66	-8.17	-136.18	-32.47	Very large		
	2x2	-16.133	-6.37	-152.48	-8.85	-136.18	-16.30			
	2x2*	-135.99	0	-136.18	0.19	-136.18	0.00			
Chitosan	1x1	-296.24	-88.26	-224.20	-72.04	-105.26	-118.94			
	2x1	-240.69	-73.53	-172.93	-67.76	-105.26	-67.66			
	1x2	-170.32	-19.60	-156.41	-13.91	-105.26	-51.15	Very large		
	2x2	-116.57	-10.45	-128.41	11.85	-105.26	-23.15			
	2x2*	-104.89	0.00	-105.26	0.37	-105.26	0.00			

³ Eigentensor analysis

⁴ To facilitate the interpretation of the elastic tensor it was decomposed into six independent ⁵ second-order stress and strain eigentensors that fulfill the relation $\sigma = \lambda \epsilon$, where λ is a scalar. ⁶ If there are no off-diagonal elements in the elastic tensor, the material only responds to the ⁷ direction of deformation, and each eigentensor has only one non-zero element. However, the ⁸ presence of off-diagonal elements leads to coupling between different modes. The eigentensors ⁹ can be graphically represented by surfaces representing the axial stress/strain component in ¹⁰ the corresponding direction.

The eigentensors are shown in Figure 5 in the order of the magnitude of the correspond-1 ing eigenvalues. Simple normal stress/strain gives a single dumbbell shape and pure shear 2 results in two crossed dumbbells of different colors. α -chitin has relatively simple anisotropy 3 with eigenvalues highest for axial deformation along the chain axis followed by b-axis direc-4 tion which is roughly parallel to the pyranose plane and a-axis. The shear directions are ac, 5 ab, bc in the descending order of eigenvalues. Strong coupling between two normal directions 6 leads to one doughnut shape and one shear-like component, exemplified by, e.g., the second 7 and third eigenvalues in α -chitin-A with dispersion correction (top row) which correspond to 8 the a and b directions, respectively. Relatively strong coupling in the two directions perpen-9 dicular to the chain axis can also be seen for the B type without dispersion correction (3rd 10 eigentensor). Also β -chitin and chitosan exhibit strong coupling in the direction perpendic-11 ular to the chain direction especially with dispersion correction (2nd and 3rd eigentensor for 12 β -chitin and 2nd and 4th eigentensor for chitosan). 13

¹⁴ Poisson's ratio

¹⁵ The Poisson's ratios for orthorhombic unit cell when stretched along the fiber direction, can ¹⁶ be simply defined along the two orthogonal crystallographic axes. They are summarized in ¹⁷ table S3. For α -chitin, the experimental value was 0.46 along the *b*-direction (ν_{32}) whereas ¹⁸ the DFT gave 0.06-0.2 depending on the hydrogen bonding pattern, but significantly lower ¹⁹ than experimental value in any case. The experimental Poisson's ratio perpendicular to the ²⁰ pyranose plane was 0.045, comparable to DFT values of 0.01-0.08 depending on the hydrogen ²¹ bond pattern, but small in any case.

For monoclinic system, the eigenvectors of deformation orthogonal to fiber direction also have to be determined. For β -chitin, the major axis of Poisson's ratio makes about 70° to the *a*-axis and 27° to the *b*-axis, which is almost parallel to the pyranose plane. The corresponding Eigenvalue was 0.5. The minor axis orthogonal to this direction has an Eigenvalue of 0.16. On the other hand, the large absolute value of s_{36} of DFT with dispersion correction indicates



Figure 5: Graphical representation of the unit cells and eigenvalue tensors of α -chitin-A, α -chitin-B, β -chitin, and chitosan with and without dispersion correction.

that the axial stress is accompanied by large shear in the ab plane. From another point of view, the Eigenvectors are close to 45° with respect to our reference frame, with Eigenvalues of 0.8 and -0.4. When the dispersion is switched off, s_{36} is close to zero, and the calculated Poisson's ratios are much closer to the experimental value (Table S3).

5 Conclusion

Using DFT calculations the elastic tensors of α -chitin, β -chitin, and chitosan in their anhy-6 drous forms were studied. The longitudinal, transverse and shear moduli successively differ 7 by one order of magnitude, with longitudinal modulus > transverse modulus > shear modu-8 lus. For α -chitin-B, α -chitin-A, β -chitin, and chitosan, the outstanding longitudinal moduli 9 are estimated to be 114.9, 126.9, 129.0, and 191.5 GPa, respectively, indicating their intrinsic 10 potential as structural materials. The transverse moduli (C11 and C22, unit in GPa) are 11 estimated as 52.8 and 27.8, 30.6 and 27.8, 17.0 and 21.9, as well as 27.5 and 24.4 GPa, re-12 spectively. The shear moduli (C44, C55, C66, unit in GPa) are calculated as (10.1, 3.9, 4.6), 13 (14.8, 2.2, 5.3), (15.2, 4.1, 4.0), and (3.5, 10.1, 4.3), respectively. Switching off the dispersion14 energy correction reveals a significant contribution from dispersion interaction to mechanical 15 performance, which has often been ignored. The hydroxymethyl group isomerization allows 16 quantification of the contribution of HBs to mechanical properties, which is found to be as 17 important as the contribution from dispersion energy. These findings contribute towards our 18 fundamental understanding of the mechanical properties of crystalline polysaccharides, and 19 are important for explaining structure-property relationships of chitin- and chitosan-based 20 functional materials. 21

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Supporting Information Available 5

Tables of compliance matrices (Table S1 & S2); Possion's ratios (Table S3); Experimental 6

lattice strain ratio of α -chitin (Figure S1) and β -chitin (Figure S2); Crystal structures in PDB 7

format for α -chitin-A, α -chitin-B, β -chitin, and chitosan. 8

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¹ TOC Graphic

