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Anisotropic Circular Dichroism of Light-Harvesting Complex II in Oriented Lipid Bilayers: Theory Meets Experiment

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11 Complex II in Oriented Lipid Bilayers: Theory
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15 Meets Experiment
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3 KEYWORDS
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6 photosynthetic complexes, model membranes, optical spectroscopy, excitonic CD, exciton theory,
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8 oriented CD
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12 ABSTRACT
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16 Anisotropic circular dichroism (ACD) spectroscopy of macroscopically aligned molecules reveals
17 additional information about their excited states that is lost in the CD of randomly-oriented
18 solutions. ACD spectra of light-harvesting complex II (LHCII)—the main peripheral antenna of
19 photosystem II in plants—in oriented lipid bilayers, were recorded from the far-UV to the visible
20 wavelength region. The ACD spectra show a drastically enhanced magnitude and level of detail
21 compared to the isotropic CD spectra, resolving a greater number of bands and weak optical
22 transitions. Exciton calculations show that the spectral features in the chlorophyll Q_y region are
23 well reproduced by an existing Hamiltonian for LHCII, providing further evidence for the identity
24 of energy sinks at chlorophylls $a603$ and $a610$ in the stromal layer and chlorophylls $a604$ and $a613$
25 in the luminal layer. We propose ACD spectroscopy to be a valuable tool linking the three-
26 dimensional structure and the photophysical properties of pigment–protein complexes.
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Introduction

In photosynthetic pigment–protein complexes (PPCs), electrostatic (Coulomb) interactions among pigment molecules give rise to partially delocalized molecular exciton states, which largely determine the photophysical and spectroscopic properties of the system.^{1,2} Light-harvesting complex II (LHCII), the major trimeric peripheral antenna of photosystem II (PSII), binds 14 chlorophylls (Chls) and 4 carotenoid xanthophylls per monomer.³ The spectral characteristics of protein-bound Chls are modulated by their protein environment giving rise to Chl forms having different transition (site) energies. In principle, all pigments in LHCII participate in all exciton states, but depending on the strength of coupling and site energies, only 1–3 Chls have a predominant contribution to a given state⁴. Since the publication of the LHCII crystal structure, several groups have applied structure-based approaches to calculate the exciton states.^{4–11} The site transition and interaction energies have been calculated for all Chls in the LHCII trimer—using atomic coordinates from the crystal structure and applying various approximations and theoretical approaches to construct the electronic Hamiltonian of the complex and evaluate the complete set of eigenstates.

The exciton model is refined by reconstructing a set of experimental optical spectroscopic data and finding the best fit. Standard optical measurements are linear absorption, fluorescence, circular dichroism (CD) and linear dichroism (LD) spectra. Among these, CD is the most sensitive to the exciton coupling strength and the mutual geometry of the coupled chromophores, thus it is a useful structural probe for multi-chromophore aggregates such as photosynthetic PPCs. Excitonic CD, in simple cases, can be easily recognized by a split conservative band structure—positive and negative bands with zero integrated area. It has been shown that the excitonic CD of LHCII depends strongly on the folding,¹² pigment composition,^{13,14} oligomerization and aggregation

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3 state^{15,16} of the complex and is widely used for characterizing isolated native and reconstituted
4 complexes. The CD spectrum is also sensitive to the molecular environment.^{17,18}
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8 A few theoretical studies have been dedicated to modelling the visible-range CD spectrum of
9 LHCII based on exciton theory. Georgakopoulou *et al.*¹⁹ were able to reproduce the main spectral
10 features of trimeric and monomeric LHCII forms as well as alterations due to site-directed
11 mutations affecting pigment binding. However, the model required considerable structural
12 adjustments to reproduce the fine spectral details and showed that small changes in the degree of
13 rotation of the transition dipole moments significantly alter the CD signal.
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22 Untangling the underlying exciton structure from the CD spectra is evidently a challenge. In
23 principle, this can be facilitated by measuring anisotropic (or oriented) CD (ACD)—the CD of
24 molecules with fixed macroscopic spatial orientation. ACD has been developed and used with
25 chiral anisotropic phases, such as liquid crystals, and for small chiral molecules.^{20,21} It has been
26 shown to reveal additional independent structural information not contained in the isotropic CD.
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40 A clear example of using ACD to separately probe exciton transitions is the CsmA baseplate
41 protein found in the chlorosome antenna of photosynthetic green-sulfur bacteria, where
42 bacteriochlorophyll *a* effectively forms exciton dimers.²³ The ACD spectra of the CsmA protein
43 were theoretically calculated following the molecular exciton approach.^{24,25} More recently, a
44 simplified theoretical description of ACD has been presented²⁶ that can be of general use in the
45 verification and optimization of structural models of such light-harvesting systems.
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3 The aim of this work was to obtain high-fidelity reproducible ACD spectra in the visible, and
4 far- and near-UV regions of macroscopically aligned isolated LHCII complexes. ACD spectra will
5 help the understanding and interpretation of the excitonic CD features, providing additional
6 physical information directly related to the structural and photophysical properties of the LHCII
7 complex. This will reduce the number of free parameters in exciton calculations. To achieve this
8 scope, we embedded LHCII trimers in lipid membranes orienting them as compressed gel slabs or
9 surface-supported films. We show that the method greatly enhances the power and sensitivity of
10 CD spectroscopy to resolve excitonic transitions. The qualitative features of the ACD spectra could
11 be reproduced with an existing excitonic model and were proven more sensitive to changes in the
12 model parameters than the CD spectra.
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25 **Experimental methods**

26 *Preparation of LHCII proteoliposomes*

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29 LHCII trimers were isolated by solubilizing PSII-enriched membrane fragments with 0.7% n-
30 Dodecyl β -D-maltoside (β -DDM) followed by sucrose density gradient ultracentrifugation.²⁷
31 Reconstituted membranes of LHCII and plant thylakoid lipids were prepared as described
32 previously.²⁸
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40 *Absorption and circular dichroism spectroscopy*

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42 Absorption spectra were recorded using an Evolution 500 dual-beam spectrophotometer
43 (Thermo Scientific, USA). CD spectra in the visible region were recorded with a J-815 (Jasco,
44 Japan) and a Chirascan Plus (Applied Photophysics, UK) spectropolarimeters. Absorption and CD
45 spectra were recorded with spectral bandwidth of 1 nm and 3 nm, respectively. Measurements
46 were carried out at room temperature. The samples were diluted in buffer to an absorbance of ca.
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3 1 at the red maximum in the 600–700 nm region and placed in standard glass cuvettes of 1 cm
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5 optical path length.
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7 *Linear dichroism and anisotropic circular dichroism spectroscopy*

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10 Room-temperature LD and ACD spectra were recorded from macroscopically aligned samples.
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12 LHCII membranes were aligned by polyacrylamide gel compression as in ref. 22 or by drying the
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14 vesicles on a quartz surface under N₂ gas purging. The latter method promotes vesicle fusion into
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16 planar bilayer membrane patches. ACD was measured in face-aligned direction, with incident light
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18 parallel to the axis normal to the membrane plane (Supporting Figure S1). LD was measured in
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20 edge-aligned direction, with incident light perpendicular to the membrane normal.
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23 *Synchrotron-radiation circular dichroism spectroscopy*

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26 UV CD/ACD measurements were performed at the Diamond B23 Beamline for synchrotron
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28 radiation circular dichroism (SRCD). SRCD spectra were measured under purging N₂ at room
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30 temperature in the range 180–540 nm. Liquid suspensions of the samples were placed in quartz
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32 cells with optical path length of 0.2 mm. Dry films with LHCII membrane patches were mounted
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34 on a Linkam motorized XY stage held on the vertical sample chamber²⁹, where the incident beam
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36 was focused onto the membrane patches with a 10x microscope objective. In this manner, a 5x5
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38 mm grid was scanned at 1 mm spatial intervals resulting in 25 SRCD spectra.
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41 **Results**

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44 We incorporated LHCII into lipid vesicles to facilitate the macroscopic orientation of the
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46 complexes and measured CD spectra in the visible as well as the far- and near-UV regions. CD in
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48 the visible region originates exclusively from the pigment cofactors, Chls and carotenoids, and is
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50 mostly due to excitonic interactions between the pigments. In contrast, the UV region contains
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52 contributions from the pigments and the apoprotein (secondary structure in the far-UV and
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aromatic side chains and disulfide bonds in the near-UV). The trimeric complexes are embedded in the planar lipid bilayer such that their axis of C_3 symmetry is parallel to the axis normal to the membrane plane \mathbf{n}_\perp . The membranes were in turn oriented either in compressed gels or by forming flat surface-supported bilayers on fused silica window substrates. ACD measurements were conducted in the face-aligned direction, where the angle α between the light propagation direction \mathbf{k} and the axis normal to the membrane plane \mathbf{n}_\perp was 0° .

ACD in the visible region

The isotropic CD spectrum of lipid vesicles in buffer and the ACD spectra of dehydrated membrane patches in the visible region are shown in Figure 1. In addition to the measured face-aligned ACD spectrum (ACD_{face}), the figure shows the complementary edge-aligned spectrum at $\alpha = 90^\circ$ (ACD_{edge}), calculated considering the relationship²²

$$CD_{\text{iso}} = \frac{1}{3}ACD_{\text{face}} + \frac{2}{3}ACD_{\text{edge}} \quad \#(1)$$

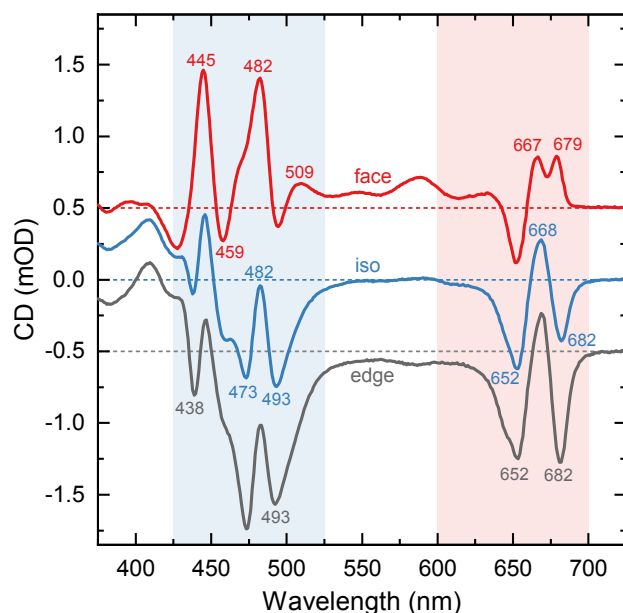


Figure 1. CD and ACD spectra of reconstituted LHCII membranes. Red curve – face-aligned ACD recorded from dehydrated membrane patches; blue curve – isotropic CD measured in buffer

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3 medium; gray curve – edge-aligned ACD calculated as $(3CD_{\text{iso}} - ACD_{\text{face}})/2$.^{22,26} The spectra are
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5 plotted in units of $\Delta A \times 10^{-3}$ (mOD) and normalized to unity isotropic absorbance at 675 nm. For
6
7 clarity, the ACD spectra are shifted vertically by 0.5. For each spectrum the horizontal dotted line
8
9 is the corresponding zero line. Numbers indicate peak wavelengths.

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13 Compared with the well documented isotropic CD spectrum of membrane-inserted LHCII^{17,30,31},
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15 the face-aligned ACD spectrum (ACD_{face}) was strikingly different. Qualitatively, it exhibits the
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17 following characteristic features:

- 18 - a positive peak at 679 nm instead of the negative 682 nm peak in the isotropic spectrum;
- 19
- 20 - intense positive 445 nm and 482 nm bands;
- 21
- 22 - clearly pronounced bands originating from weak optical transitions—at 510, 540 and 590
- 23 nm;
- 24
- 25 - the negative peaks at 438 and 473 nm found in the isotropic spectrum vanish and the
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27 shoulders at 428 and 459 nm become negative peaks.
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34 Bands observed in the isotropic CD spectrum of LHCII membranes in solution but not in the
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36 face-aligned ACD spectrum, can evidently be found in the complementary edge-aligned ($\alpha = 90^\circ$)
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38 ACD spectrum. To this group belong the negative 438, 473 and 493 nm bands, and the negative
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40 shoulder around 645 nm, that is characteristic for LHCII trimers as opposed to monomers¹⁵.

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43 The interpretation of the spectra can be rather challenging as ACD can be even more prone to
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45 perturbations and optical artefacts than isotropic CD. To rule out the presence of any unwanted
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47 distortions, the spectra of reconstituted LHCII membranes oriented by two different methods were
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49 recorded and additionally compared with those of solubilized complexes as controls. After
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51 rehydration of the dry LHCII membrane patches, the isotropic CD spectrum was completely
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53 restored (Supporting Figure S2). Furthermore, drying of LHCII trimers solubilized in detergent
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micelles had no effect on their CD spectra (Supporting Figure S3). Therefore, it is highly unlikely that the ACD spectra contain optical artefacts or reflect structural alterations in the complexes that might have occurred upon drying.

As a further test, we compared the ACD spectra obtained by gel compression, which was applied earlier²² to isolated thylakoid membranes and PSII-enriched membrane fragments. Uniaxial gel compression deforms the spherical vesicles into flattened ellipsoids with the membrane normal n_{\perp} parallel to the compression direction.³² The face-aligned ACD spectrum and the CD spectrum of LHCII membranes in uncompressed gel are shown in Figure 2a.

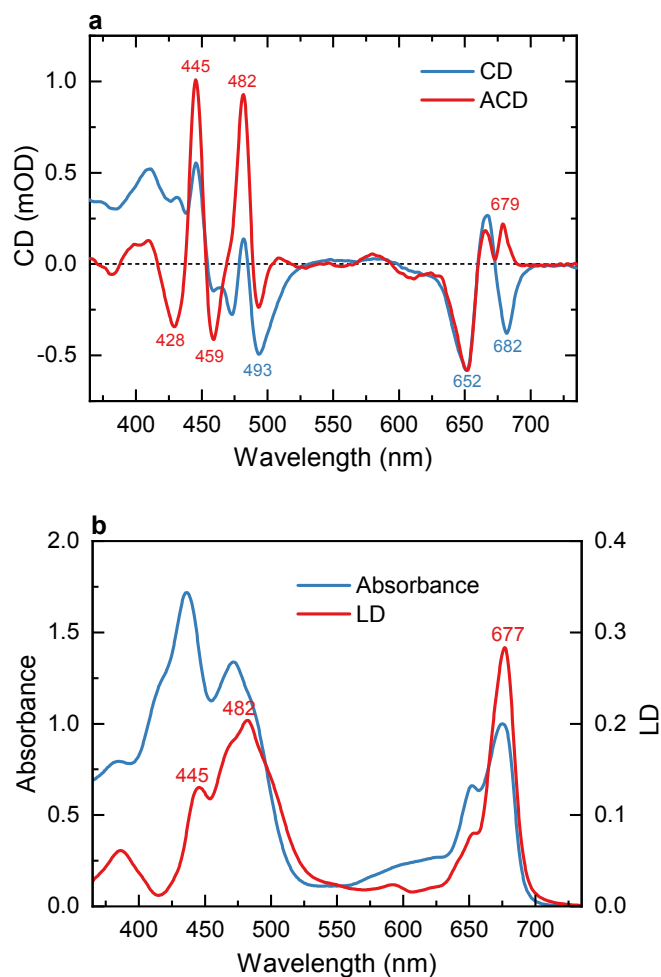


Figure 2. Optical spectra of reconstituted LHCII membranes in buffer and polyacrylamide gel. a). CD and ACD spectra measured on uncompressed or face-aligned uniaxially compressed gel,

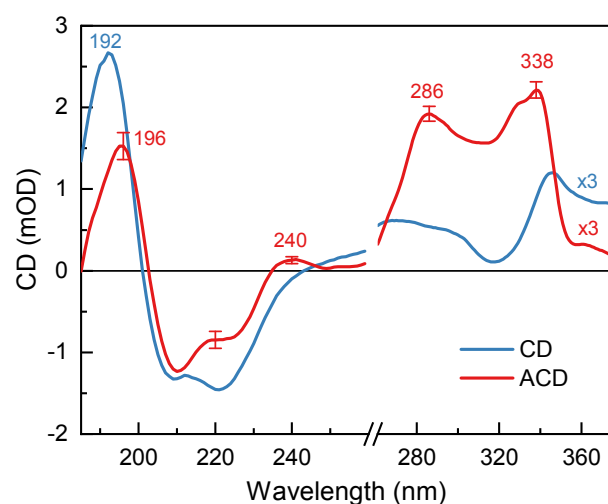
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3 respectively. b). Absorption spectrum (in buffer) and LD spectrum (edge-aligned compressed gel).
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5 The spectra are normalized to unity absorbance at 675 nm, 1 cm optical path length.
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9 Some minor changes can be detected in the CD spectrum of LHCII in gel compared to buffer
10 medium—the intensity of the negative 473 nm band is diminished. More importantly, the ACD
11 spectra in compressed gels and in dry membrane patches are virtually identical. The similarity
12 strongly suggests that (1) both methods effectively orient the membranes in the expected manner,
13 and (2) the spectra are free from artefacts such as inadvertent structural changes or optical
14 (flattening, scattering) effects. The LD measured in edge-aligned orientation, at an angle $\alpha = 90^\circ$,
15 is plotted in Figure 2b. Note that in randomly oriented samples, LD is zero. Due to rotational
16 symmetry, LD was also zero in the face-aligned orientation (data not shown) and the ACD
17 spectrum showed no change when rotating the sample (about axis of compression). Therefore,
18 artifacts in the ACD spectra arising from linear anisotropy or birefringence interference can be
19 completely ruled out. The LD spectrum of LHCII has peaks at 445, 482 and 677 nm that matched
20 the main positive peaks in the ACD spectrum. Positive LD means that the underlying transition
21 dipole moments are oriented preferentially in the membrane plane.³³ As a first approximation, it
22 can be expected that CD bands originating from the same transitions will be enhanced in the face-
23 aligned ACD spectrum due to photo-selection of those transitions. It is important to note that the
24 sign of the CD bands do not need necessarily to be positive.
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45 This agreement provides strong support that the features of the ACD spectrum are solely a
46 consequence of the alignment of the LHCII membranes with respect to the direction of the incident
47 beamlight. Additional support for this assumption is obtained from exciton simulations of the ACD
48 spectra reported below.
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54 *UV region*
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3 Isotropic and anisotropic UV CD spectra of reconstituted LHCII membranes (Figure 3) were
4 measured using Diamond B23 beamline for SRCD. The advantages of SRCD measurements are
5 not only in the increased sensitivity and signal-to-noise ratio, especially in the far-UV region, but
6 also in the ability to conduct microscopic measurements with the highly collimated microbeam.
7 This allowed us to probe the chiral heterogeneity of the dehydrated membrane patches. Note that
8 compressed gels are not suitable for UV measurements because of the high absorption from
9 polyacrylamide media.
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36 **Figure 3.** UV SRCD and ACD spectra of reconstituted LHCII membranes. The SRCD spectrum
37 was recorded from liquid suspension in a quartz cell of 0.2 mm path length. The ACD spectrum is
38 an average of 25 measurements at different spots of the 5x5 mm grid of a dried membrane patch,
39 and error bars show standard deviations. The spectra are normalized to unity absorbance at 675
40 nm and multiplied by 3 in the near-UV region (above 260 nm).
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48 The UV CD spectrum of LHCII in lipid membranes is virtually identical to that in detergent-
49 solubilized LHCII (Supporting Figure S4) retaining the characteristic profile of α -helical
50 secondary structures of the proteins—two negative bands of approximately equal amplitude at 210
51 nm and at 224 nm, and a strong positive band at 192 nm with nearly double the intensity of the
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3 negative bands.³⁴ The ACD spectrum recorded from dry membrane patches contains several
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5 notable differences with respect to the isotropic CD spectrum in the far-UV (below 260 nm):
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8 - lower intensity of the 192 nm band (up to one-half);
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10 - lower negative amplitude at 224 nm;
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12 - a characteristic positive peak at 240 nm, followed by a trough at 250–260 nm.
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15 Remarkable features were also observed in the near-UV region (260–350 nm): positive peaks at
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17 286 and 338 nm and a pronounced shoulder at 324 nm. The strong ACD signal in this region is
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19 surprising because of the small absorption and almost negligible isotropic CD.
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22 It must be noted that, whereas the visible and near-UV SRCD varied very little between
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24 measurements and sample batches, the shape of the ACD spectrum in the far-UV region,
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26 particularly the intensity of the 210 nm band displayed pronounced variability. To avoid artefacts
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28 from uneven drying and inhomogeneities in the membrane patches, SRCD spectra were scanned
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30 on 25 different spots using a highly focused microbeam. These spectra show qualitatively similar
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32 features despite variability in the relative intensities of the bands reflecting variable thickness.
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34 Importantly, the UV SRCD spectra of dry films of detergent-solubilized LHCII (not shown) were
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36 indistinguishable from the spectra of LHCII trimers in buffer solution.
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39 40 **Discussion**

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42 The landscape of excited states of an aggregate of coupled pigment molecules encompasses, in
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44 principle, couplings between all pigment excited states. For a system of N pigments, each with m
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46 energy levels, the total number of exciton levels to consider is $N^{(m-1)}$. The CD spectrum contains
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48 signals from all exciton states, which can be positive or negative, overlapping in energy and hence
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50 cancelling each other. It is generally not possible to extract unambiguously the energy levels and
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52 pigment couplings from CD/SRCD spectra alone, because different configurations can result in
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3 practically identical spectra. This is true even for the simplest possible scenario of a dimer of two-
4 level chromophores.^{2,26} ACD can provide, in principle, additional information inaccessible from
5 CD and LD, that imposes constraints on the possible geometries and electronic couplings between
6 pigment molecules.^{20,23,26}
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12 The applicability and usefulness of ACD for extracting or validating the excitonic Hamiltonian
13 of molecular aggregates can be demonstrated considering the simplest of exciton systems—the
14 dimer of identical molecules. The electronic eigenstates of the dimer can be approximated as
15 orthogonal linear combinations of the monomer wave functions. Taking into account only the
16 lowest-lying excited state of the monomers, the dimer will have two excitonic states that are split
17 in energy depending on the coupling strength, and whose transition dipole moments have
18 magnitude depending on the dimer geometry but are invariably perpendicular to each other. The
19 excitonic coupling gives rise to a pair (couplet) of CD bands of equal magnitude and opposite sign,
20 centered at the respective excitonic transition frequencies. Different geometries of the dimer and
21 different combinations of spectral bandwidths and exciton coupling strengths can result in
22 indistinguishable CD spectra. In principle, ACD measurements enable the observation of the
23 exciton transitions separately even if the coupling strength is so small that the bands largely overlap
24 and cancel each other in the isotropic phase. This was shown both theoretically and experimentally
25 for the chlorosome baseplate protein CsmA in green-sulfur photosynthetic bacteria, which
26 effectively contains dimers of bacteriochlorophyll *a*.^{23,26} When the baseplate is attached to the
27 membrane, one of the exciton transitions is polarized parallel to the axis normal to the membrane
28 plane and the other is in the membrane plane. The face-aligned ACD spectrum arises only from
29 the transition in the membrane plane revealing its exact wavelength position, bandwidth and the
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3 sign of the rotational strength. The extra information uniquely defines the exciton Hamiltonian and
4 the dimer geometry.²⁶
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7 *ACD of LHCII in the visible region*

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10 The simplest qualitative interpretation of the ACD spectra is based on the expectation that the
11 excitonic states with transition dipole moments oriented preferentially in the membrane plane are
12 selectively probed in the face-aligned direction ($\alpha = 0^\circ$), whereas transitions which tend to be
13 aligned perpendicularly to the membrane plane are suppressed. Accordingly, the CD bands at 445
14 nm, 482 nm and 679 nm must originate from excitonic transition dipoles lying parallel to the
15 membrane plane. To lend further support to this interpretation, positive peaks are observed in the
16 LD spectrum at these wavelengths (Figure 2b). Conversely, the 438 nm, 473 nm and 493 nm bands,
17 which are missing in the face-aligned ACD spectra, must reflect transitions perpendicular to the
18 membrane plane. This information is directly related to the relative geometry of the respective
19 pigment entities, but a more detailed analysis requires exciton calculations of optical spectra
20 presented in the following.
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35 *Exciton calculations of optical spectra*

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38 The information potential of the ACD measurements was further analyzed by exciton
39 calculations. The theory of optical spectra, based on the exciton Hamiltonian of the PPC, is
40 described in the Supporting Information. The parameters of the Hamiltonian that includes the
41 ground and the lowest-lying singlet excited states of the Chls in the LHCII have been previously
42 obtained¹⁰ from a combination of quantum-chemical and electrostatic computations, and then
43 refined by comparison with experimental absorption, CD and isotropic CD spectra. To account for
44 dynamic localization effects in the excitonic wavefunction, exciton domains are introduced^{10,35},
45 splitting the exciton Hamiltonian into blocks encompassing strongly coupled pigments. Exciton
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delocalization is only permitted within each exciton domain, and the optical spectra are obtained as a sum over the domain spectra. The expressions for linear absorption (OD), LD and CD of an exciton domain read (see e.g. ref. 1):

$$\text{OD}(\omega) \propto \omega \sum_M |\boldsymbol{\mu}_{M0}|^2 D_M(\omega - \omega_{M0}) \#(2)$$

$$\text{LD}(\omega) \propto \omega \sum_M |\boldsymbol{\mu}_{M0}|^2 (1 - 3\cos^2(\theta_M)) D_M(\omega - \omega_{M0}) \#(3)$$

$$\text{CD}(\omega) \propto \omega^2 \sum_M \sum_{m,n} c_m^{(M)} c_n^{(M)} (\mathbf{R}_{mn} \cdot (\boldsymbol{\mu}_{m0} \times \boldsymbol{\mu}_{n0})) D_M(\omega - \omega_{M0}) \#(4)$$

with the exciton transition dipole moment $\boldsymbol{\mu}_{M0} = \sum_m c_m^{(M)} \boldsymbol{\mu}_{m0}$ and where $\boldsymbol{\mu}_{m0}$ is the local transition dipole moment of molecule m , $c_m^{(M)}$ is the contribution of molecule m to the delocalized exciton state M (see Supporting Information), θ_M is the angle between the exciton transition dipole moment $\boldsymbol{\mu}_{M0}$ and the membrane normal \mathbf{n}_\perp , which is represented by the C_3 -symmetry axis of the LHCII trimer, and $\mathbf{R}_{mn} = \mathbf{R}_m - \mathbf{R}_n$ is the vector that connects the centers of pigments m and n . The $D_M(\omega - \omega_{M0})$ denotes a lineshape function, which contains vibrational sidebands and lifetime broadening induced by exciton relaxation.³⁶ Recently, an exciton-theory-derived formalism of ACD has been developed.²⁶ Here we tested the predictive power of the exciton Hamiltonian proposed earlier¹⁰ by calculating ACD spectra without further refinement of the model parameters. For light propagation in face direction, that is parallel to the membrane normal \mathbf{n}_\perp , the expression for the ACD signal reads²⁶

$$\text{ACD}_{\text{face}}(\omega) \propto \omega^2 \sum_M \sum_{m,n} c_m^{(M)} c_n^{(M)} \left\{ (\mathbf{R}_{mn} \cdot \mathbf{n}_\perp) (\boldsymbol{\mu}_{m0} \times \boldsymbol{\mu}_{n0}) \cdot \mathbf{n}_\perp + ((Q'_{m0} \cdot \mathbf{n}_\perp) \times \boldsymbol{\mu}_{n0}) \cdot \mathbf{n}_\perp \right\} D_M(\omega - \omega_{M0}) \#(5)$$

where Q'_{m0} is the electric quadrupole transitions moment of pigment m . The component $Q'_{m0}^{(\alpha\beta)}$ is defined as

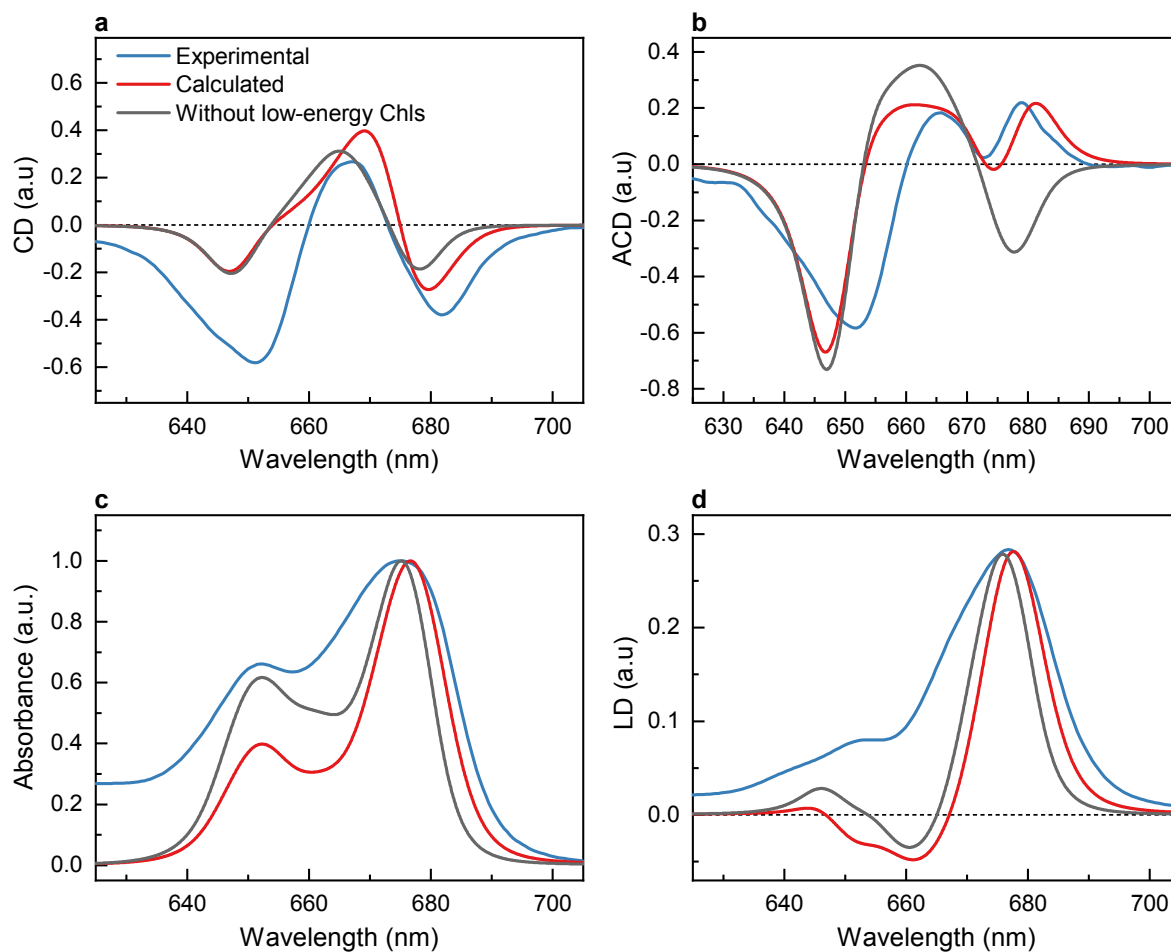
$$Q_{m0}^{(\alpha\beta)} = \sum_I q_I (\mathbf{R}_{\alpha,I}^{(m)} - \mathbf{R}_{\alpha}^{(m)}) (\mathbf{R}_{\beta,I}^{(m)} - \mathbf{R}_{\beta}^{(m)}) \# (6)$$

Here, $\mathbf{R}^{(m)}$ denotes the molecular center of pigment m , and q_I is the atomic transition charge, placed at the position $\mathbf{R}_I^{(m)}$ of the I^{th} atom of pigment m , obtained from a fit of the three-dimensional electrostatic potential of the ab-initio transition density of the Q_y transition of the pigments³⁷. The present transition charges were obtained with time-dependent density functional theory, using the B3LYP exchange-correlation functional and a 6-31G* basis set, in ref. ³⁷ for Chl *a* and in ref. ³⁸ for Chl *b*. The numerical values of these transition charges are given in the respective supporting informations.^{37,38}

The expressions for LD and CD in eqs. 3 and 4 assume ideal alignment of the PPC, whereas the measured ensemble spectra necessarily reflect a degree of randomness in the orientation. Accordingly, the calculated anisotropic spectra were represented as weighted sums of ACD of perfectly aligned complexes and isotropic CD. An orientation factor $\phi = 0.41$ was estimated from the relative ratio of the measured LD magnitude (Figure 2b) and the calculated ideal-orientation LD magnitude and used as a weighting coefficient for the ACD spectra.

The calculated CD, ACD, OD and LD spectra in the Chl Q_y region are compared with the experimental data in Figure 4. The calculated spectra in general contain somewhat narrower peaks than the experimental ones. The inhomogeneous width Δ_{inh} of the distribution function of site energies as well as the Huang–Rhys factor S , characterizing the dynamic modulation of site energies, might be larger at 300 K than at cryogenic temperatures, for which the present $\Delta_{inh} = 120$ cm^{-1} and $S = 0.5$ have been estimated.¹⁰ In our calculations the temperature dependence of the spectra is solely determined by the Bose–Einstein distribution function of vibrational quanta entering the homogeneous lineshape function (eqs. S7, S13–S14). From quasi-elastic neutron scattering experiments on LHCI evidence about a strong increase of protein flexibility above the

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3 glass transition temperature of 240 K, leading to an increase of the dynamic and/or static disorder,
4 was reported,³⁹ which could be responsible for the larger broadening of the experimental spectra
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44 **Figure 4.** Calculated CD (a), ACD in face configuration (b), absorption (c) and LD (d) spectra of
45 LHCII trimers based on the model including (red curves) and excluding (gray curves) the four
46 low-energy pigments, compared to the respective experimental spectra (blue curves, data from
47 Figure 2). The LD and ACD spectra are calculated assuming orientation factor $\phi = 0.41$. All spectra
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53 were measured and calculated at room temperature (300 K).
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3 The calculated CD spectrum in Figure 4a shows a negative band at 679 nm and a positive peak
4 at 670 nm, which are consistent with the experimental spectral features. The negative peak in the
5 calculated spectrum at 648 nm is weaker than in the experimental spectrum. This deviation is likely
6 due to the non-conservative nature of the spectrum, that is, $\int_0^\infty \text{CD}(\omega) d\omega \neq 0$. Similar deviations
7 were found in CD spectra of the CP29 light-harvesting complex of PSII and has been recently
8 explained by excitonic coupling between higher Chl transitions as well as carotenoid transitions to
9 the Q_y Chl transitions.³⁵

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20 The exciton model readily reproduces the characteristic qualitative shape of the measured ACD
21 spectrum (Figure 4b)—in particular, the apparent sign inversion of the low-energy band around
22 680 nm between the experimental isotropic CD and ACD spectra. Because the ACD spectrum is
23 extremely sensitive to the pigment geometry, this agreement between theory and experiment
24 provides evidence that the pigments contributing to the low-energy exciton states are correctly
25 assigned in the model. Removing the four lowest-energy Chls from the calculation results in
26 qualitatively similar CD, absorption and LD spectra, but the ACD spectrum, particularly in the
27 680 nm region, is severely misrepresented (the gray curves in Figure 4), highlighting the
28 importance of ACD in the assignment of the lowest-energy states.

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41 The energy sinks were identified earlier by Müh and Renger¹⁰ as Chl *a*603 and *a*610 in the
42 stromal layer and Chl *a*604 and *a*613 in the luminal layer. Each of those four sites contributes to
43 the sign inversion of the ACD spectrum, with the largest effects by Chls *a*604 and *a*610, as
44 observed by removing these two pigments in the calculations of the spectra (Supporting Figure
45 S5). There is general consensus in the literature about the energy sink at Chl *a*610 in the stromal
46 layer^{5,10,11}, whereas our second energy sink at Chl *a*603 in the luminal layer was not found in the
47 other studies^{5,11} (Supporting Table S1). Concerning the luminal layer there is also partial
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3 agreement. The present assignment of Chl *a*604 agrees with the model of Zucchelli et al.⁵ and that
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5 of Chl *a*613 agrees with Novoderezhkin et al.¹¹ Calculations with all three site energy sets^{5,10,11}
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7 result in qualitatively similar optical spectra, but the strong positive 680 nm ACD band is best
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9 reproduced with the site energies of ref. 10 (Supporting Figure S6). In our comparison we only
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11 replaced the site energies of ref. 10 by those of ref. ⁵ and 11 and did not take into account other
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13 differences, as, e.g., in the excitonic couplings and the optical lineshape theories, because the
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15 present couplings go beyond the point-dipole approximation and the present lineshape theory
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17 avoids artefacts from dynamic localization effects. These improvements could be part of the reason
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19 why some of the low energy sites were assigned differently before, in particular in the work of
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21 Novoderezhkin et al. ¹¹, where this assignment was solely based on a fit of optical spectra. The
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23 importance of Chl *a*604 for reproducing the specific CD and ACD spectral shape around 680 nm
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25 points to the correct assignment of this pigment as belonging to the group of low-energy Chls, as
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27 opposed to having an intermediate energy as in Novoderezhkin's model. A more quantitative
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29 investigation of the ACD spectrum seems to be suitable to remove the remaining ambiguity in the
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31 assignment of low-energy sites.
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38 From experiments on LHCII aggregates¹⁶, single molecule experiments⁴⁰ and molecular
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40 dynamics simulations⁴¹ it is known that the LHCII complex can adopt different conformations
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42 affecting the low-energy excited states, thought to be related to photoprotection^{42,43}. Indeed, the
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44 site energies had to be adjusted¹⁰ in order to describe the spectra of small lamellar aggregates of
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46 LHCII³⁵. This adjustment—model B by Müh and Renger¹⁰—leads to qualitatively similar ACD
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48 spectra as for the site energies inferred for solubilized LHCII trimers—model A (Supporting
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50 Figure S6). This is in line with the observed very similar ACD spectra of oriented lamellar
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52 aggregates of LHCII (data not shown).
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3 The calculated ACD spectrum (Figure 4b) deviates stronger below 670 nm from the
4 experimental data. These deviations could be due to the excitonic couplings involving higher Chl
5 transitions, which appear to affect the CD in the same spectral range as the ACD spectrum. A
6 further refinement of the present exciton theory by including these higher excited states of the Chls
7 and also of the carotenoids will help to improve the agreement between calculated and measured
8 spectra over the whole Q_y spectral range.
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11 The OD and LD spectra are in good qualitative agreement with the present experiments above
12 665 nm (Figure 4c and d). Please note that the deviations in the OD and LD spectra between the
13 present calculations and the experimental data occurring at short wavelengths below 665 nm are
14 due to the contributions of intramolecular vibronic Chl transitions, which do not affect the CD and
15 ACD spectra and were not taken into account in the present work to simplify the calculations. A
16 comparison of OD and LD spectra calculated including these intramolecular vibrations with
17 experimental data is given in ref. 10.
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21 Another subtlety in the modeling of optical spectra of LHCII, also related to conformational
22 transitions, concerns the evidence¹⁰ that the transition dipole moments of some of the low-energy
23 pigments must have a different orientation in solubilized trimers than in the crystal structure. In a
24 first attempt, to explain the reduced LD spectrum of ideally oriented complexes, Chl *a*604 has been
25 rotated by 45° such that its transition dipole moment is in the membrane plane.¹⁰ Such a rotation
26 would still be possible, since the low-energy part of the ACD spectrum is not significantly affected
27 by it (Supporting Figure S7). A systematic investigation of this question, including possible
28 changes of other low-energy pigments, has to await a refinement of the theory, as discussed above.
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3 In summary, the ACD data presented herein and the calculations ratified the exciton Hamiltonian
4 of LHCII proposed earlier¹⁰, and a refined theory of ACD, including higher excited electronic
5 states, will allow us to further refine this Hamiltonian.
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8 9 10 *UV ACD*

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12 The UV CD of peptides and proteins is widely used to study their structure, folding and
13 conformational dynamics, membrane insertion, and effects of ligand binding. CD spectra
14 characteristic to the polypeptide secondary structure content—e.g. α -helix, β -strand, left handed
15 extended helix of polyproline II (PPII), β -turn, and unordered structure—occur in the far-UV
16 region and arise mainly by $n\text{-}\pi^*$ and $\pi\text{-}\pi^*$ transitions of the peptide bond.^{44,45} The $n\text{-}\pi^*$ transition
17 gives rise to a negative peak in the CD spectrum at about 224 nm. According to Moffitt's theory⁴⁶,
18 excitonic interactions in α -helices result in three $\pi\text{-}\pi^*$ transitions with split energies: one gives rise
19 to the negative peak at around 210 nm and the other two are degenerate, at around 190 nm, with
20 amplitudes strongly depending on the probing direction. The CD spectrum of LHCII membranes
21 represented the expected signatures of an α -helix (Figure 3). Horn and Paulsen³⁴ found a value of
22 42% α -helices and 8% β -strands in detergent-solubilized native LHCII, the former is reasonably
23 close to the 41% seen in the LHCII crystal structure.
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33 Additional information regarding the angle of orientation of the secondary structures with
34 respect to the axis normal to the membrane plane can be extracted from the ACD spectra.⁴⁷⁻⁵⁰
35 Oriented helical peptides bound to a lipid bilayer exhibit a specific ACD spectrum, from which
36 the orientation of the helix can be extrapolated. A distinguishing feature of the ACD of α -helices
37 is the presence or absence of the negative band at around 210 nm, being indicative of surface or
38 transmembrane helix alignment, respectively.⁵¹
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3 Wu et al.⁴⁷ employed Moffitt's theory to successfully reproduce the expected CD and ACD
4 features of oriented transmembrane peptides. Following this approach, we calculated CD and ACD
5 spectra in the far-UV region for each of the five α -helices in LHCII taking into account their
6 respective inclination angles and lengths from the crystal structure³ (see Supporting Information).
7 While the calculated CD spectrum (Supporting Figure S8) is very similar to the experimental one,
8 not all features of the ACD spectrum described above could be reproduced. In the calculated ACD
9 spectrum, the intensity of the 208 nm band was smaller than that of the calculated CD spectrum,
10 whereas the experimental ACD (Figure 3) did not show such a change. Also, in the near-UV range,
11 the enhanced peak around 240 nm in the LHCII ACD spectrum was not observed in the UV CD
12 of oriented helical peptides. This result leads us to conclude that the UV CD spectrum of LHCII
13 includes sizeable contributions not only from the backbone but also from the aromatic side chain
14 residues as well as from interactions of the protein and the pigment cofactors (Chls and
15 carotenoids). Since the ACD spectrum appears to be sensitive to these details it will represent a
16 critical check for future semi-empirical calculations of UV spectra of pigment–protein complexes.
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38 **Conclusions**

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40 A new spectroscopic technique for the characterization of photosynthetic pigment–protein
41 complexes, ACD, was presented and applied to LHCII. The ACD spectra revealed a great deal of
42 information otherwise lost in the rotational averaging of the isotropic CD spectra. The CD and
43 ACD spectra in the UV region reflect the large content of α -helical secondary structure but also
44 reveal that pigments and possibly pigment–protein couplings also contribute to the UV CD. The
45 ACD spectra in the visible spectral region are determined by the optical properties of the protein-
46 bound pigments and their excitonic interactions. We demonstrated that ACD spectroscopy,
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3 combined with structure-based model calculations, is a valuable approach to identify the pigments
4 contributing to the observable excitonic transitions with higher degree of confidence. Further
5 evidence for the energy sinks in LHCII formed by Chls *a603* and *a610* in the stromal layer and
6 Chls *a604* and *a613* in the luminal layer is obtained from the present ACD calculations that
7 reproduce the experimental sign switch of the low-energy band of LHCII with respect to the CD
8 spectrum.

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10 The non-conservative nature of the experimental isotropic CD spectrum suggests that there is an
11 influence of higher excited Chl and carotenoid states that could explain the remaining deviations
12 between calculated and measured ACD spectra, in particularly on the high-energy side of the
13 visible spectrum. Inclusion of carotenoid states will help to further understand the higher-energy
14 spectral regions, including the blue and the near-UV, where the ACD spectra possess the most
15 characteristic and striking features. ACD will serve as an important complement to CD for the
16 characterization of carotenoid-Chl interactions and for the verification and refinement of exciton
17 Hamiltonians of light-harvesting complexes in general.

38 ASSOCIATED CONTENT

39
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41 The following files are available free of charge.

42
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44 **Supporting Information.** Supporting figures, theory for calculating optical spectra. (PDF)

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16 17 **Author Contributions**

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19 P.H.L., G.G. and P.A. designed the experiments. P.A. isolated LHCII and prepared reconstituted
20
21 membranes. P.A., M.L., K.P., M.D. and P.H.L. performed CD measurements in the visible
22
23 region. P.A., M.L., P.H.L., O.Z., T.J., G.S., B.U. and G.G. performed SRCD measurements. D.L.
24
25 and T.R. did theoretical calculations. The manuscript was written through contributions of all
26
27 authors. All authors have given approval to the final version of the manuscript.
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47
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51 “CALIPSOplus”).
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ABBREVIATIONS

ACD, anisotropic circular dichroism; CD, circular dichroism; Chl, chlorophyll; LD, linear dichroism; LHCII, light-harvesting complex II; PPC, pigment–protein complex; PSII, photosystem II; SRCD, synchrotron radiation circular dichroism.

REFERENCES

- (1) van Amerongen, H.; Valkunas, L.; van Grondelle, R. *Photosynthetic Excitons*; World Scientific: Singapore, 2000.
- (2) Cantor, C. R.; Schimmel, P. R. *Biophysical Chemistry: Part II: Techniques for the Study of Biological Structure and Function*; W. H. Freeman and Company: Oxford, 1980.
- (3) Liu, Z. F.; Yan, H. C.; Wang, K. B.; Kuang, T. Y.; Zhang, J. P.; Gui, L. L.; An, X. M.; Chang, W. R. Crystal Structure of Spinach Major Light-Harvesting Complex at 2.72 Å Resolution. *Nature* **2004**, *428*, 287–292.
- (4) Müh, F.; Madjet, M. E.-A.; Renger, T. Structure-Based Identification of Energy Sinks in Plant Light-Harvesting Complex II. *J. Phys. Chem. B* **2010**, *114*, 13517–13535.
- (5) Zucchelli, G.; Santabarbara, S.; Jennings, R. C. The Q_y Absorption Spectrum of the Light-Harvesting Complex II as Determined by Structure-Based Analysis of Chlorophyll Macrocycle Deformations. *Biochemistry* **2012**, *51*, 2717–2736.
- (6) Novoderezhkin, V. I.; Palacios, M. A.; van Amerongen, H.; van Grondelle, R. Energy-Transfer Dynamics in the LHCII Complex of Higher Plants: Modified Redfield Approach. *J. Phys. Chem. B* **2004**, *108*, 10363–10375.
- (7) Novoderezhkin, V. I.; Palacios, M. A.; van Amerongen, H.; van Grondelle, R. Excitation Dynamics in the LHCII Complex of Higher Plants: Modeling Based on the 2.72 Å Crystal Structure. *J. Phys. Chem. B* **2005**, *109*, 10493–10504.
- (8) Renger, T.; Madjet, M. E.-A.; Knorr, A.; Müh, F. How the Molecular Structure Determines the Flow of Excitation Energy in Plant Light-Harvesting Complex II. *J. Plant Physiol.* **2011**, *168*, 1497–1509.
- (9) Linnanto, J.; Martiskainen, J.; Lehtovuori, V.; Ihalainen, J.; Kananavicius, R.; Barbato, R.; Korppi-Tommola, J. Excitation Energy Transfer in the LHC-II Trimer: A Model Based on the New 2.72 Å Structure. *Photosynth. Res.* **2006**, *87*, 267–279.
- (10) Müh, F.; Renger, T. Refined Structure-Based Simulation of Plant Light-Harvesting Complex II: Linear Optical Spectra of Trimers and Aggregates. *Biochim. Biophys. Acta* **2012**, *1817*, 1446–1460.

- 1
2
3
4 (11) Novoderezhkin, V.; Marin, A.; van Grondelle, R. Intra- and Inter-Monomeric Transfers in
5 the Light Harvesting LHCII Complex: The Redfield–Förster Picture. *Phys. Chem. Chem.*
6 *Phys.* **2011**, *13*, 17093–17103.
- 7 (12) Hobe, S.; Prytulla, S.; Kuhlbrandt, W.; Paulsen, H. Trimerization and Crystallization of
8 Reconstituted Light-Harvesting Chlorophyll *a/b* Complex. *EMBO J.* **1994**, *13*, 3423–3429.
- 9 (13) Lichtenthaler, H. K. Chlorophylls and Carotenoids: Pigments of Photosynthetic
10 Biomembranes. *Methods Enzymol.* **1987**, *148*, 350–382.
- 11 (14) Morosinotto, T.; Breton, J.; Bassi, R.; Croce, R. The Nature of a Chlorophyll Ligand in Lhca
12 Proteins Determines the Far Red Fluorescence Emission Typical of Photosystem I. *J. Biol.*
13 *Chem.* **2003**, *278*, 49223–49229.
- 14 (15) Nussberger, S.; Dekker, J. P.; Kuhlbrandt, W.; van Bolhuis, B. M.; van Grondelle, R.; van
15 Amerongen, H. Spectroscopic Characterization of Three Different Monomeric Forms of the
16 Main Chlorophyll *a/b* Binding Protein from Chloroplast Membranes. *Biochemistry* **1994**, *33*,
17 14775–14783.
- 18 (16) Ruban, A. V.; Calkoen, F.; Kwa, S. L. S.; van Grondelle, R.; Horton, P.; Dekker, J. P.
19 Characterisation of Lhc II in the Aggregated State by Linear and Circular Dichroism
20 Spectroscopy. *Biochim. Biophys. Acta* **1997**, *1321*, 61–70.
- 21 (17) Akhtar, P.; Dorogi, M.; Pawlak, K.; Kovács, L.; Bóta, A.; Kiss, T.; Garab, G.; Lambrev, P.
22 H. Pigment Interactions in Light-Harvesting Complex II in Different Molecular
23 Environments. *J. Biol. Chem.* **2015**, *290*, 4877–4886.
- 24 (18) Lambrev, P.; Várkonyi, Z.; Krumova, S.; Kovács, L.; Miloslavina, Y.; Holzwarth, A. R.;
25 Garab, G. Importance of Trimer–Trimer Interactions for the Native State of the Plant Light-
26 Harvesting Complex II. *Biochim. Biophys. Acta* **2007**, *1767*, 847–853.
- 27 (19) Georgakopoulou, S.; van der Zwan, G.; Bassi, R.; van Grondelle, R.; van Amerongen, H.;
28 Croce, R. Understanding the Changes in the Circular Dichroism of Light Harvesting
29 Complex II Upon Varying Its Pigment Composition and Organization. *Biochemistry* **2007**,
30 *46*, 4745–4754.
- 31 (20) Kuball, H.-G. CD and Acd Spectroscopy on Anisotropic Samples: Chirality of Oriented
32 Molecules and Anisotropic Phases—a Critical Analysis. *Enantiomer* **2002**, *7*, 197–205.
- 33 (21) Kuball, H.-G.; Dorr, E.; Höfer, T.; Türk, O. Exciton Chirality Method. Oriented Molecules –
34 Anisotropic Phases. *Monatsh. Chem./Chem. Mon.* **2005**, *136*, 289–324.
- 35 (22) Miloslavina, Y.; Lambrev, P. H.; Jávorfí, T.; Várkonyi, Z.; Karlický, V.; Wall, J. S.; Hind,
36 G.; Garab, G. Anisotropic Circular Dichroism Signatures of Oriented Thylakoid Membranes
37 and Lamellar Aggregates of LHCII. *Photosynth. Res.* **2012**, *111*, 29–39.
- 38 (23) Nielsen, J. T.; Kulminskaya, N. V.; Bjerring, M.; Linnanto, J. M.; Ratsep, M.; Pedersen, M.
39 O.; Lambrev, P. H.; Dorogi, M.; Garab, G.; Thomsen, K. et al. *In Situ* High-Resolution
40 Structure of the Baseplate Antenna Complex in *Chlorobaculum Tepidum*. *Nat. Commun.*
41 **2016**, *7*, 12454.
- 42 (24) Hansen, A. E. Molecular Exciton Approach to Anisotropic Absorption and Circular
43 Dichroism I. General Formulation. *Monatsh. Chem./Chem. Mon.* **2005**, *136*, 253–273.
- 44 (25) Hansen, A. E. Molecular Exciton Approach to Anisotropic Absorption and Circular
45 Dichroism II. The Partial Optic Axis and Its Application in Molecular Exciton Theory.
46 *Monatsh. Chem./Chem. Mon.* **2005**, *136*, 275–287.
- 47 (26) Lindorfer, D.; Renger, T. Theory of Anisotropic Circular Dichroism of Excitonically
48 Coupled Systems: Application to the Baseplate of Green Sulfur Bacteria. *J. Phys. Chem. B*
49 **2018**, *122*, 2747–2756.
- 50
51
52
53
54
55
56
57
58
59
60

- 1
2
3
4 (27) Caffarri, S.; Croce, R.; Breton, J.; Bassi, R. The Major Antenna Complex of Photosystem II
5 Has a Xanthophyll Binding Site Not Involved in Light Harvesting. *J. Biol. Chem.* **2001**, *276*,
6 35924–35933.
- 7 (28) Akhtar, P.; Lingvay, M.; Kiss, T.; Deák, R.; Bóta, A.; Ughy, B.; Garab, G.; Lambrev, P. H.
8 Excitation Energy Transfer between Light-Harvesting Complex II and Photosystem I in
9 Reconstituted Membranes. *Biochim. Biophys. Acta* **2016**, *1857*, 462–472.
- 10 (29) Siligardi, G.; Hussain, R. CD Spectroscopy: An Essential Tool for Quality Control of
11 Protein Folding. In *Structural Proteomics*; Springer, 2015; pp 255-276.
- 12 (30) Yang, C.; Boggasch, S.; Haase, W.; Paulsen, H. Thermal Stability of Trimeric Light-
13 Harvesting Chlorophyll *a/b* Complex (Lhciib) in Liposomes of Thylakoid Lipids. *Biochim.*
14 *Biophys. Acta* **2006**, *1757*, 1642–1648.
- 15 (31) Moya, I.; Silvestri, M.; Vallon, O.; Cinque, G.; Bassi, R. Time-Resolved Fluorescence
16 Analysis of the Photosystem II Antenna Proteins in Detergent Micelles and Liposomes.
17 *Biochemistry* **2001**, *40*, 12552–12561.
- 18 (32) Kiss, L. I.; Ganago, A. O.; Garab, G. I. Quantitative Method for Studying Orientation of
19 Transition Dipoles in Membrane Vesicles of Spherical Symmetry. *J. Biochem. Biophys.*
20 *Methods* **1985**, *11*, 213–225.
- 21 (33) Garab, G.; van Amerongen, H. Linear Dichroism and Circular Dichroism in Photosynthesis
22 Research. *Photosynth. Res.* **2009**, *101*, 135–146.
- 23 (34) Horn, R.; Paulsen, H. Folding *in vitro* of Light-Harvesting Chlorophyll *a/b* Protein Is
24 Coupled with Pigment Binding. *J. Mol. Biol.* **2002**, *318*, 547–556.
- 25 (35) Lindorfer, D.; Müh, F.; Renger, T. Origin of Non-Conservative Circular Dichroism of the
26 CP29 Antenna Complex of Photosystem II. *Phys. Chem. Chem. Phys.* **2017**, *19*, 7524–7536.
- 27 (36) Renger, T.; Marcus, R. On the Relation of Protein Dynamics and Exciton Relaxation in
28 Pigment–Protein Complexes: An Estimation of the Spectral Density and a Theory for the
29 Calculation of Optical Spectra. *J. Chem. Phys.* **2002**, *116*, 9997–10019.
- 30 (37) Madjet, M. E.; Abdurahman, A.; Renger, T. Intermolecular Coulomb Couplings from Ab
31 Initio Electrostatic Potentials: Application to Optical Transitions of Strongly Coupled
32 Pigments in Photosynthetic Antennae and Reaction Centers. *J. Phys. Chem. B* **2006**, *110*,
33 17268–17281.
- 34 (38) Renger, T.; Trostmann, I.; Theiss, C.; Madjet, M. E.; Richter, M.; Paulsen, H.; Eichler, H.;
35 Knorr, A.; Renger, G. Refinement of a Structural Model of a Pigment–Protein Complex by
36 Accurate Optical Line Shape Theory and Experiments. *J. Phys. Chem. B* **2007**, *111*, 10487–
37 10501.
- 38 (39) Vrandečić, K.; Rätsep, M.; Wilk, L.; Rusevich, L.; Golub, M.; Reppert, M.; Irrgang, K.-D.;
39 Kühlbrandt, W.; Pieper, J. r. Protein Dynamics Tunes Excited State Positions in Light-
40 Harvesting Complex II. *J. Phys. Chem. B* **2015**, *119*, 3920–3930.
- 41 (40) Krüger, T. P.; Iliaia, C.; Johnson, M. P.; Ruban, A. V.; Van Grondelle, R. Disentangling
42 the Low-Energy States of the Major Light-Harvesting Complex of Plants and Their Role in
43 Photoprotection. *Biochim. Biophys. Acta* **2014**, *1837*, 1027–1038.
- 44 (41) Liguori, N.; Periole, X.; Marrink, S. J.; Croce, R. From Light-Harvesting to Photoprotection:
45 Structural Basis of the Dynamic Switch of the Major Antenna Complex of Plants (LHCII).
46 *Sci. Rep.* **2015**, *5*, 15661.
- 47 (42) Horton, P.; Wentworth, M.; Ruban, A. Control of the Light Harvesting Function of
48 Chloroplast Membranes: The LHCII-Aggregation Model for Non-Photochemical
49 Quenching. *FEBS Lett.* **2005**, *579*, 4201–4206.
- 50
51
52
53
54
55
56
57
58
59
60

- 1
2
3 (43) Johnson, M. P.; Goral, T. K.; Duffy, C. D.; Brain, A. P.; Mullineaux, C. W.; Ruban, A. V.
4 Photoprotective Energy Dissipation Involves the Reorganization of Photosystem II Light-
5 Harvesting Complexes in the Grana Membranes of Spinach Chloroplasts. *Plant Cell* **2011**,
6 tpc. 110.081646.
7
8 (44) Woody, R. W.; Tinoco Jr., I. Optical Rotation of Oriented Helices. III. Calculation of the
9 Rotatory Dispersion and Circular Dichroism of the Alpha- and 3_{10} -Helix. *J. Chem. Phys.*
10 **1967**, *46*, 4927–4945.
11 (45) Woody, R. W. Improved Calculation of the $n\pi^*$ Rotational Strength in Polypeptides. *J.*
12 *Chem. Phys.* **1968**, *49*, 4797–4806.
13 (46) Moffitt, W. Optical Rotatory Dispersion of Helical Polymers. *J. Chem. Phys.* **1956**, *25*, 467–
14 478.
15 (47) Wu, Y.; Huang, H. W.; Olah, G. A. Method of Oriented Circular Dichroism. *Biophys. J.*
16 **1990**, *57*, 797–806.
17 (48) Towell, J. F.; Manning, M. C. Analysis of Protein Structure by Circular Dichroism
18 Spectroscopy. In *Analytical Applications of Circular Dichroism*; Purdie, N., Brittain, H. G.,
19 Eds.; Elsevier Science B.V., 1994; pp 175–205.
20 (49) Dave, P. C.; Billington, E.; Pan, Y. L.; Straus, S. K. Interaction of Alamethicin with Ether-
21 Linked Phospholipid Bilayers: Oriented Circular Dichroism, ^{31}P Solid-State NMR, and
22 Differential Scanning Calorimetry Studies. *Biophys. J.* **2005**, *89*, 2434–2442.
23 (50) Bürck, J.; Wadhvani, P.; Fanghänel, S.; Ulrich, A. S. Oriented Circular Dichroism: A
24 Method to Characterize Membrane-Active Peptides in Oriented Lipid Bilayers. *Acc. Chem.*
25 *Res.* **2016**, *49*, 184–192.
26 (51) Bürck, J.; Roth, S.; Wadhvani, P.; Afonin, S.; Kanithasen, N.; Strandberg, E.; Ulrich, A. S.
27 Conformation and Membrane Orientation of Amphiphilic Helical Peptides by Oriented
28 Circular Dichroism. *Biophys. J.* **2008**, *95*, 3872–3881.
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31
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