

*Overview*

## **Comparative structural and catalytic properties of oxygen-evolving photosystem II preparations\***

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**Abstract.** Biochemical techniques now exist to produce the oxygen-evolving complex of photosystem II (PSII) and its associated photochemical redox reactions in various states of purity. These preparations permit one to assess the structural roles of polypeptides in promoting activity by using selective extraction techniques which remove certain polypeptides, to carry out reconstitution studies which re-establish activity, and, in the case of more recently developed, highly purified preparations discussed in this overview, to identify the minimal polypeptide complement necessary for photosynthetic oxygen evolution activity. These comparative investigations also suggest a tentative structure for an oxygen-evolving PSII core complex whose primary constituents are a hydrophobic complex of polypeptides, manganese, calcium and chloride, and the 33 kDa extrinsic polypeptide.

**Abbreviations:** DCBQ–2,6 dichloro-p-benzoquinone, Chl–chlorophyll, LHCP–light-harvesting chlorophyll proteins, PS–photosystem

### **Introduction**

The use of non-ionic detergents to fractionate thylakoid membranes (Berthold et al. 1981, Kuwabara and Murata 1982b) yields PSII-enriched membrane sheets (Goodman-Dunahay et al. 1984) and has produced new material for the study of the structure and function of the oxygen-evolving reaction. Briefly, these PSII preparations have provided material for a closer examination of the role(s) of extrinsic 33, 23, and 17 kDa proteins (see Ghanotakis and Yocum (1985) Govindjee et al. (1985) and Babcock (1987)

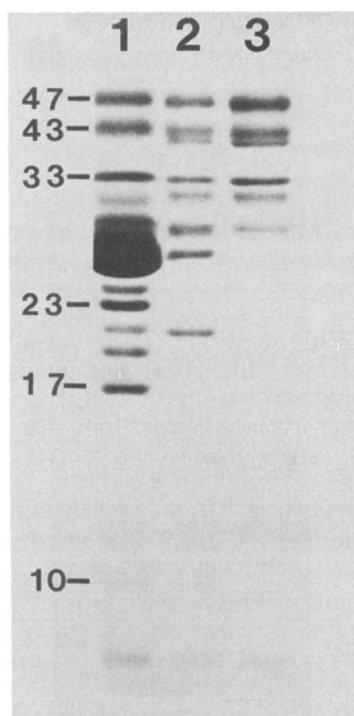
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for reviews), as well as a vehicle for preparing samples in which spectroscopic studies of PSII-associated electron carriers has been facilitated owing to the absence of PSI. More recently, attempts to use the PSII membranes as the starting material to produce more highly purified PSII preparations have succeeded, opening the way to further investigations on the polypeptide structure of PSII, as well as on the function of these polypeptides in facilitating photo-induced electron transfer from the oxygen-evolving complex to the reducing side of the photo-act. At the same time, these new preparations possess properties markedly different from those associated with the starting material, the more complex preparations of Kuwabara and Murata (1982b) and Berthold et al. (1981). In this overview, we discuss the comparative properties of the various types of PSII preparations available to researchers at the present time.

### **Polypeptide composition and comparative properties of electron transfer in PSII preparations**

Figure 1 presents a SDS-polyacrylamide gel of three PSII preparations isolated in the author's laboratory. Lane 1 presents the polypeptide composition of PSII membrane sheets, Lane 2 the composition of a reaction center complex, and Lane 3 the composition of a reaction center "core" preparation. The PSII membrane sheet (Lane 1) preparations contain the hydrophobic polypeptides (47, 43, 34, 32, 9 and 5 kDa) of the photochemically active PSII "core" (as originally defined by Satoh (1982) previous to the more recent isolation of the reaction center preparation (Nanba and Satoh 1987) which consists of 34 kDa (D2), 32 kDa (D1, the  $Q_B$  or herbicide binding protein) and cytochrome  $b_{559}$ ). Polypeptides in the 29–24 kDa region of the gel are associated with the binding of light-harvesting Chl a and b (collectively called LHCP). The extrinsic 33, 23, and 17 kDa polypeptides found in this preparation are also noted on the gel; following their discovery by Kuwabara and Murata (1982a) and Åkerlund et al. (1982), these extrinsic species have been studied in both everted thylakoid vesicles and in the PSII membranes sheets to generate an extensive (and still expanding) literature; the current status of research on the extrinsic proteins may be briefly, but by no means comprehensively, summarized as follows:

1. All three proteins are essential for long-term stability and function of the oxygen-evolving reaction in higher plants.
2. The 17 and 23 kDa proteins are necessary to maintain the structural integrity of the oxygen-evolving apparatus in a state where the reaction can function in an environment of low, physiological concentrations of



*Fig. 1.* SDS-polyacrylamide gel electrophoresis patterns of three oxygen-evolving PSII preparations. Lane 1: PSII membrane sheets; Lane 2: reaction center complex; Lane 3: oxygen-evolving reaction center core.

calcium and chloride; in addition the 23 kDa protein, presumably in concert with the 33 kDa species, is necessary to form a shield around the functional manganese cluster which protects it from exogenously added reductants (Waggoner and Yocum 1987).

3. The 33 kDa protein is essential in maintaining the native catalytic activity of the oxygen-evolving reaction. Although the 33 kDa protein can be removed from membrane sheets with full retention of functional manganese, these extracted systems do not retain the control rates of oxygen evolution activity. Recent thermoluminescence work in the laboratory of Inoue (1987) on samples from which the 33 kDa protein has been removed provides evidence that the 33 kDa protein may also have an effect on reactions at the reducing side of PSII.

The ability to utilize elevated concentrations of calcium and chloride to supplant in part the function of the 23 and 17 kDa polypeptides has permitted the application of treatments for the further purification of PSII, treatments which (at least at present) lead to the removal of the 23 and

17 kDa proteins along with other polypeptides associated with membrane sheets. This approach is exemplified by the work of Ikeuchi et al. (1985), who combined octylglucoside treatment with sucrose density gradient centrifugation of dissociated PSII membranes to produce a more highly refined PSII preparation. These results were followed by a report from Ghanotakis and Yocum (1986), who employed high concentrations of sodium chloride along with octylglucoside and differential centrifugation to isolate material (whose polypeptide composition is shown in Lane 2 of Fig. 1) which is similar, but not identical to the preparation of Ikeuchi et al. (1985). A major structural feature of such preparations is evident from their polypeptide composition, which shows that the major polypeptides of the light-harvesting apparatus have been removed by the purification procedures. This observation, based on gel electrophoresis, correlates with an approximate 4 to 5-fold enrichment in reaction centers on a chlorophyll basis, as summarized in Table 1. A principal difference between the two more highly purified reaction center complex preparations reported in the literature to date resides in the absence (in the Ikeuchi et al. (1985) preparation) of a polypeptide with an apparent molecular mass of 22 kDa. There are also differences in electron transfer activity on the reducing side of the two preparations (Table 2) which are reflected in a preference for ferricyanide as the acceptor in a DCMU-insensitive reaction when the 22 kDa protein has been removed. The Ikeuchi et al. preparation of the complex also requires digitonin (0.1%) in the assay reaction mixture to produce optimal rates of oxygen evolution. With regard to oxidizing side function, the two preparations appear quite similar, both in terms of rates of oxygen evolution as well as in terms of a requirement for elevated concentrations of calcium and chloride for optimal rates of oxygen evolution activity (owing to the absence of the 17 and 23 kDa polypeptides). A characterization of the calcium and chloride responses of the Ikeuchi et al. preparation has been reported by Ikeuchi and Inoue (1986).

Using the more highly refined preparations, Ghanotakis et al. (1987a, 1987b) have removed proteins of 28, 22, and 10 kDa from the oxygen-evolv-

Table 1. Oxygen-evolving preparations: enrichment of PSII components and activity

	Preparation		
	Berthold et al.	Ikeuchi et al.	Ghanotakis/Yocum
Mn/Chl	4/250	4/68	4/68
Z <sup>+</sup> /Chl	1/260	n.d.	1/73
V <sub>O<sub>2</sub></sub> <sup>a</sup>	550–700 <sup>b</sup>	850–1070 <sup>c</sup>	940 <sup>b</sup>

<sup>a</sup> expressed as  $\mu\text{moles O}_2/\text{hr/mg Chl}$

<sup>b</sup> with DCBQ as the electron acceptor

<sup>c</sup> with ferricyanide as the acceptor in the presence of 0.1% digitonin

ing reaction center complex by FPLC gel filtration to produce a 7-polypeptide reaction center core PSII preparation (Lane 3 of Fig. 1) which is capable of high rates of activity; ferricyanide now acts in this preparation as the preferred, but largely DCMU-insensitive acceptor. A similar type of preparation possessing substantially lower rates of oxygen evolution, isolated by the use of ion-exchange purification on DEAE Toyopearl, has been reported by Yamada et al. (1987), and Ikeuchi and Inoue (1986) have further purified their preparation. The reader should note that the purified oxygen-evolving "core" preparations possess reducing side electron transfer properties which are similar, if not identical to, those exhibited by the original reaction center preparation described by Ikeuchi et al. (1985), namely the predominance of DCMU-insensitive, ferricyanide-catalyzed activity as shown in Table 2, rather than DCMU-sensitive activity catalyzed by DCBQ.

Taken together, the results so far available from research on the oxygen-evolving reaction center "cores" implicate the hydrophobic polypeptides (47, 43, 34 (D2), 32 (D1), 9 and 5 kDa) which form the photochemically active core of PSII, along with manganese, calcium, chloride and the extrinsic 33 kDa protein as the minimal structural-cofactor unit which is capable of oxygen evolution activity. One of the common properties of these oxygen-evolving "core" preparations is the absence of an extrinsic 10 kDa protein, which was earlier proposed to be an essential component of PSII preparations capable of water oxidation (Ljungberg et al. 1984a). The relationship of polypeptides found in PSII membrane sheets to activity in the highly purified preparations described here remain to be elucidated. Other questions are raised by the properties of the reaction center complex and "core" preparations which have not yet been addressed, and a brief overview of the comparative properties of the various PSII preparations raises the following issues:

Table 2. Acceptor function in PSII reaction center complexes

Acceptor/Inhibitor Additions	O <sub>2</sub> evolution activity <sup>a</sup> assayed in preparations according to:	
	Ikeuchi et al.	Ghanotakis/Yocum
DCBQ	893	940
DCBQ + DCMU	765	530
Fe(CN) <sub>6</sub> <sup>3-</sup>	1481	230
Fe(CN) <sub>6</sub> <sup>3-</sup> + DCMU	1369	190

<sup>a</sup> expressed as  $\mu\text{moles O}_2/\text{hr}/\text{mg Chl}$

1. Are there new functional lesions created on the oxidizing side of PSII by the treatments required to produce more highly purified material, which are reflected in the loss of polypeptides found in membrane sheets? The rates of oxygen evolution activity reported for refined preparations is consistent with an enrichment in reaction centers on a chlorophyll basis, but the activities so far reported are lower than expected for activity in a system where a 4- to 5-fold purification has been obtained. Is this due to damaged centers in which manganese remains esr silent? Is it possible that removal of the LHCP complex has caused drastic alterations in quantum efficiency? In utilizing further detergent treatments, have important lipid constituents been stripped away from reaction center polypeptides? Further investigations are obviously necessary to clarify these questions.
2. To what extent can the structural properties of PSII membrane sheets be reconstituted in the more purified material? We (Merritt et al. 1987) have presented preliminary evidence for rebinding of 17 and 23 kDa proteins to the preparation lacking the LHCP complex, and it will be useful to know whether similar reconstitutions are possible with the oxygen-evolving reaction center core. At the same time, future research on isolation of complexes should determine whether purification procedures can be developed which permit the removal of LHCP complex without extraction of the extrinsic 23 and 17 kDa species.
3. What is the origin of the changes observed in reducing side function after removal of the 22 kDa protein? Is the appearance of DCMU-insensitive ferricyanide reduction a fortuitous consequence of the purification procedure itself? Is it possible that the 22 kDa protein is responsible for binding of extrinsic proteins, as implied by the data of Ljungberg et al. (1984b) and in addition responsible for regulation of electron transfer on the reducing side of PSII? Comparative investigations by several groups (Ghanotakis et al. 1987b, Henrysson et al. 1987, Ikeuchi and Inoue 1986, Yamada et al. 1987) correlate the loss of quinone-catalyzed electron transfer activity with the extraction of the 22 kDa protein and the appearance of DCMU-insensitive ferricyanide reduction activity. It will be interesting to see whether further research can establish the 22 kDa species as the PSII equivalent of the "H" subunit in photosynthetic bacteria.
4. Are small polypeptides required for PSII activity? Henrysson et al. (1987) have presented evidence for the presence of a number of tenaciously bound small polypeptides in a number of PSII preparations, including oxygen-evolving reaction center complexes. Of these species,

three (4, 5, 5.5 kDa) were present in all PSII preparations surveyed by the authors. In light of this finding, further investigations on low-molecular weight polypeptides are necessary to establish their role(s) in oxygen evolution and/or photochemical electron transfer reactions.

## Conclusions

Table 3 summarizes the properties of oxygen-evolving PSII preparations which are now available for use in biochemical and biophysical research on primary electron transfer and the oxygen-evolving reaction. As is apparent from the Table, new, more purified material offers the immediate advantage to the investigator of an enrichment in reaction center components on a chlorophyll basis. The disadvantages, at the present time is that these preparations represent structurally as well as functionally altered forms of the better-characterized membrane sheet preparations. Although high concentrations of calcium and chloride will activate the oxygen-evolving complex in the more purified preparations, one must at the same time realize that changes have also occurred to activity on the reducing side of the photoact. It is likely that experiments now in progress in several laboratories will better define the activities of PSII preparations containing fewer polypeptides than the PSII membrane sheets used as starting material, and that with this knowledge, these new preparations will prove to be useful adjuncts to the membrane sheet preparations, which are now widely used for PSII research.

Table 3. Comparative properties of oxygen-evolving PSII preparations

	Membrane sheets	Complex	"Core"
$V_{O_2}^+$	500-700	> 1000	> 1000
Ca <sup>2+</sup> /Cl <sup>-</sup> requirement	Low	High	High
TRAP/CHL ratio	1/250	1/68	1/40*
Acceptor preference	DCBQ	DCBQ** Fe(CN) <sub>6</sub> <sup>3-</sup> ***	Fe(CN) <sub>6</sub> <sup>3-</sup>
ESR multiline signal	Present	Present	Present****
Polypeptides released by purification procedures	PSI b <sub>6</sub> /f CF <sub>1</sub> /CF <sub>0</sub>	LHCP 23 kDa 17 kDa 22 kDa***	28 kDa (CP 29 ?) 22 kDa 10 kDa

+ expressed in  $\mu\text{moles O}_2/\text{hr}/\text{mg Chl}$

\* based on Mn content; see Ikeuchi and Inoue (1986)

\*\* for the reaction center complex prepared as in Ghanotakis and Yocum (1986)

\*\*\* for the reaction center complex prepared as in Ikeuchi et al. (1985)

\*\*\*\* reported in the preparation described in Yamada et al. (1987)

Certainly, the availability of oxygen-evolving PSII preparations in which the number of polypeptides can be varied either by purification procedures, or by the use of extraction and reconstitution methods, provides an opportunity to explore in detail the roles of these polypeptides as either structural or catalytic components in the oxygen-evolving reaction of PSII.

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