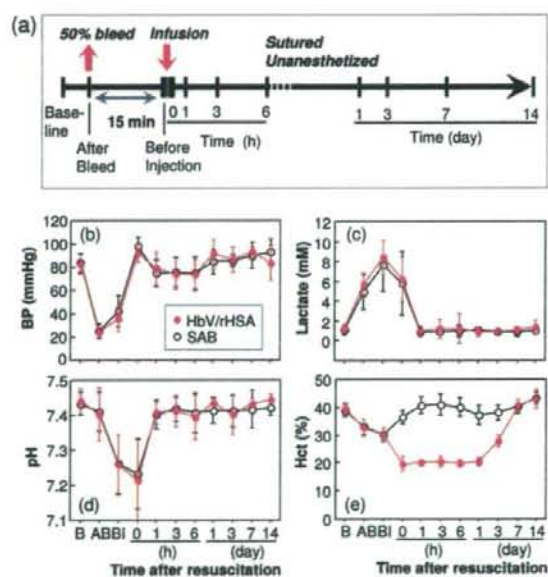


**Figure 9.** Regulation of physicochemical properties of HbV for versatile applications. (a) Oxygen dissociation curves of HbVs. Oxygen affinity ( $P_{50}$ , partial pressure of oxygen at which Hb is half-saturated with oxygen) is regulated by coencapsulation of PLP (105, 106). (b) Colloid osmotic pressure (COP) of chemically modified Hb solutions increase with the Hb concentration (15). In contrast, HbV particles have no oncotic effect. The figure shows 20 Torr when HbV is suspended in 5% rHSA. XLHb, intramolecularly cross-linked Hb. (c) Rheological properties of HbV suspended in various plasma substitute solutions (122). [Hb] = 10 g/dL, 25 °C.

with  $O_2$ ) of HbV can be regulated by coencapsulation of an allosteric effector (105, 106) (Figure 9a). The  $P_{50}(O_2)$  of purified Hb in a saline solution (in the presence of  $Cl^-$ ) is about 14 Torr; Hb strongly binds  $O_2$  and does not release  $O_2$  at 40 Torr (partial pressure of mixed venous blood). Historically, it has been regarded that the  $O_2$  affinity is expected to be regulated similarly to that of RBC, namely, about 25–30 Torr, using an allosteric effector or by a direct chemical modification of the Hb molecules. This enables sufficient  $O_2$  unloading during blood microcirculation, as evaluated by the arterio-venous difference in the levels of  $O_2$  saturation in accordance with an  $O_2$  equilibrium curve. Pyridoxal 5'-phosphate (PLP) is coencapsulated in HbV as an allosteric effector to regulate  $P_{50}(O_2)$  (105, 106). The main binding site of PLP is the N-terminal of the  $\alpha$ -chain and  $\beta$ -chain and  $\beta$ -82 Lysine within the  $\beta$ -cleft, which is part of the binding site of the natural allosteric effector, 2,3-diphosphoglyceric acid (2,3-DPG). The bound PLP retards the dissociation of the ionic linkage between the  $\beta$ -chains of Hb during conversion of deoxy to oxyHb in the same manner as 2,3-DPG does. Therefore, the  $O_2$  affinity of Hb decreases in the presence of PLP. The  $P_{50}(O_2)$  of HbV can be regulated to 8–150 Torr by coencapsulating the appropriate amount of PLP or inositol hexaphosphate as an allosteric effector. Equimolar PLP to Hb (PLP/Hb = 1/1 by mol) was coencapsulated, and  $P_{50}(O_2)$  was regulated to 18 Torr. Furthermore,  $P_{50}(O_2)$  was regulated to 32 Torr when the molar ratio PLP/Hb was 3/1. The  $O_2$  affinities of HbV can be regulated easily without changing other physical parameters, whereas in the case of the other modified Hb solutions, their chemical structures determine their  $O_2$  binding affinities. Consequently, regulation is difficult. The present HbV contains PLP at PLP/Hb = 2.5 by mol; the resulting  $P_{50}(O_2)$  is about 25–28 Torr, which shows sufficient  $O_2$  transporting capacity as a transfusion alternative. Actually, HbV has been shown to provide  $O_2$ -transport capacity that is both sufficient and comparable to that of RBCs in experiments related to extreme blood exchange (68, 69, 79, 105, 107, 108) and fluid resuscitation from hemorrhagic shock (102, 109–112) (Figure 10). A recent experiment of HbV as a priming solution for cardiopulmonary bypass (CPB) in a rat model showed that HbV protects neurocognitive function by transporting  $O_2$  to brain tissue even when the hematocrit is reduced markedly (113).

The appropriate  $O_2$  binding affinities for  $O_2$  carriers have not yet been decided completely. However, the easy regulation of the  $O_2$  binding affinity might be useful to meet the requirement of clinical indications such as oxygenation of ischemic tissues. The  $P_{50}(O_2)$  of HbV without PLP and  $Cl^-$  is 8–9 Torr. This formulation is effective for targeted  $O_2$  delivery to anoxic tissues caused by reduced blood flow (107, 114, 115).



**Figure 10.** (a) Scheme depicting the experimental protocol of hemorrhagic shock and resuscitation (102). Shock was induced by withdrawing 50% of circulating blood volume from Wistar rats. After 15 min, they were resuscitated with either HbV suspended in recombinant human serum albumin (HbV/rHSA) or shed autologous blood (SAB). (b) blood pressure. (c) lactate, (d) pH, and (e) hematocrit (Hct). Mean  $\pm$  SD. B, Baseline; AB, after bleeding; BI, before injection (B–6 h,  $n = 24$ ; 1–14 days,  $n = 5$ ).

**2.6. Rheological Properties and Their Physiological Implications for Tissue Oxygenation.** The extremely high concentration of the HbV suspension ([Hb] = 10 g/dL; [lipids] = 6 g/dL, volume fraction, ca. 40 vol %) imparts an  $O_2$  carrying capacity that is comparable to that of blood. The HbV suspension does not possess a colloid osmotic pressure (COP), because one HbV particle (ca. 250 nm diameter) contains about 30 000 Hb molecules. In fact, HbV acts as a particle, not as a solute. Therefore, HbV must be suspended in or cojoined with an aqueous solution of a plasma substitutes. This requirement is identical to that for emulsified perfluorocarbon, which does not possess COP (116, 117); it contrasts to characteristics of other Hb-based  $O_2$  carriers, intramolecular cross-linked Hbs, polymerized Hbs, and polymer-conjugated Hbs, which all possess very high COP as protein solutions (15, 118) (Figure 9b).

**Table 4. Publications of Preclinical Studies Aiming at Applications of Hb-Vesicles for a Transfusion Alternative and for Oxygen Therapeutics**

indication	ref
1. Resuscitative fluid for hemorrhagic shock	102, 109–112
2. Hemodilution	68, 69, 79, 101, 105, 107, 108
3. Priming fluid for extracorporeal membrane oxygenator (ECMO) for cardiopulmonary bypass	113
4. Perfusate for resected organs (transplantation)	24, 129
5. Oxygenation of ischemic brain (stroke)	130
6. Oxygenation of ischemic skin flap (plastic surgery)	115, 127, 128
7. Tumor oxygenation for sensitization to irradiation	131
8. CO carrier for cytoprotection at reperfusion	132

Animal tests of HbV suspended in plasma-derived HSA or rHSA showed an O<sub>2</sub> transporting capacity that is comparable to that of blood (110, 113). We reported previously that HbV suspended in plasma-derived HSA or rHSA was almost Newtonian: no aggregation was detected microscopically (68, 69). In Japan, rHSA was very recently approved for clinical use, in May 2008 (119), but various plasma substitutes are used worldwide, such as hydroxyethyl starch (HES), dextran (DEX), and modified fluid gelatin (MFG). The selection among these plasma substitutes is best determined not only according to their safety and efficacy, but also according to their associated price, experience of clinicians, and customs of respective countries. Water-soluble polymers generally interact with particles such as polystyrene beads, liposomes, and RBCs to induce aggregation or flocculation (120, 121). For that reason, it is important to determine the compatibility of HbV with these plasma substitutes. With that background, we studied rheological properties of HbV suspended in these plasma substitute solutions using a complex rheometer and a microchannel array (122). The rheological property of an Hb-based O<sub>2</sub> carrier is important because the infusion amount is expected to be considerably large, which might affect the blood viscosity and hemodynamics.

The HbV suspended in rHSA was nearly Newtonian (Figure 9c). Its viscosity was similar to that of blood, and the mixtures with RBCs at various mixing ratios showed viscosities of 3–4 cP. Other polymers, HES, DEX, and MFG, induced flocculation of HbV, possibly by depletion interaction, and rendered the suspensions as non-Newtonian with the *shear-thinning* profile (122). These HbV suspensions showed high viscosity and a high storage modulus ( $G'$ ) because of the presence of flocculated HbV. On the other hand, HbV suspended in rHSA exhibited a very low  $G'$ . The viscosities of HbV suspended in DEX, MFG, and high-molecular-weight HES solutions responded quickly to rapid step changes of shear rates of 0.1–100 s<sup>-1</sup> and a return to 0.1 s<sup>-1</sup>, indicating that flocculation formation is both rapid and reversible. Microscopically, the flow pattern of the flocculated HbV perfused through microchannels (4.5 μm deep, 7 μm wide, 20 cmH<sub>2</sub>O applied pressure) showed no plugging. Furthermore, the time required for passage was directly related to the viscosity.

It has been regarded that lower blood viscosity after hemodilution is effective for tissue perfusion. However, microcirculatory observation shows that, in some cases, lower “plasma viscosity” decreases shear stress on the vascular wall, causing vasoconstriction and reducing the functional capillary density (123). Therefore, an appropriate viscosity might exist, which maintains the normal tissue perfusion level. The large molecular dimension of HbV can result in a transfusion fluid with high viscosity. A large molecular dimension is also effective to reduce vascular permeability and to minimize the reaction with NO and CO as vasorelaxation factors (24, 25, 30, 31)

**Figure 11. Crystal structure of HSA with myristate (PDB ID: 1BJ5) from ref (136).**

(see Figure 3). These new concepts suggest reconsideration of the design of artificial O<sub>2</sub> carriers (124). Actually, new products are appearing, although they are in the preclinical stage, not only HbV but also zero-link polymerized Hb (125) and others with larger molecular dimensions and higher O<sub>2</sub> affinities (126). Erni et al. clarified that HbV with a high O<sub>2</sub> binding affinity (low  $P_{50}(O_2)$ , such as 8–15 Torr) and high viscosity (such as 11 cP) suspended in a high-molecular-weight HES solution was effective for oxygenation of an ischemic skin flap (115, 127, 128). That study showed that HbV retains O<sub>2</sub> in the upper arterioles, then perfuses through collateral arteries and delivers O<sub>2</sub> to the targeted ischemic tissues, a concept of targeted O<sub>2</sub> delivery by an Hb-based O<sub>2</sub> carrier (114). A high O<sub>2</sub> binding affinity (low  $P_{50}(O_2)$ ) would also be effective to improve the O<sub>2</sub> saturation of Hb in pulmonary capillaries when exposed to a hypoxic atmosphere or with an impaired lung function. Some plasma substitutes cause flocculation of HbV and hyperviscosity. However, reports show that hyperviscosity would not necessarily be deteriorative and might be, in some cases, advantageous in the body (32). HbV provides a unique opportunity to manipulate the suspension rheology,  $P_{50}(O_2)$ , and other physicochemical properties, not only as a transfusion alternative, but also for other clinical applications such as oxygenation of ischemic tissues and ex vivo perfusion systems (129–132) (Table 4).

### 3. ALBUMIN-HEMES AS O<sub>2</sub> CARRYING PLASMA PROTEINS

**3.1. HSA Incorporating Synthetic Fe<sup>2+</sup> Porphyrin (HSA-FeP).** HSA (Mw: 66 500) has a remarkable ability to bind a wide variety of exogenous compounds in the human circulatory system (133). This carrier protein comprises 3 homology helical domains (I–III) with 9 loops formed by 17 disulfide linkages; each domain contains 2 subdomains (A and B) (Figure 11) (134–137). It is known that many common drugs such as warfarin, diazepam, and ibuprofen bind to one of the two primary sites (site 1 in subdomain IIA, site 2 in subdomain IIIA) (138). In fact, HSA helps to solubilize these compounds to achieve high concentration in the bloodstream; otherwise, they would easily aggregate and be poorly distributed.

In 1995, we found that tetrakis(*o*-pivalamido)phenylporphyrinatoiron bearing a covalently linked proximal imidazole (FePI, Figure 12) was incorporated into HSA, yielding a red HSA-FePI hybrid (11). This synthetic hemoprotein can reversibly bind and release O<sub>2</sub> under physiological conditions (pH 7.4, 37 °C) in much the same way as Hb. The HSA host adsorbs a maximal eight FePI molecules. Their stepwise binding constants ( $K_1$ – $K_8$ ) range from 1.2 × 10<sup>6</sup> to 1.2 × 10<sup>4</sup> (M<sup>-1</sup>) (Table

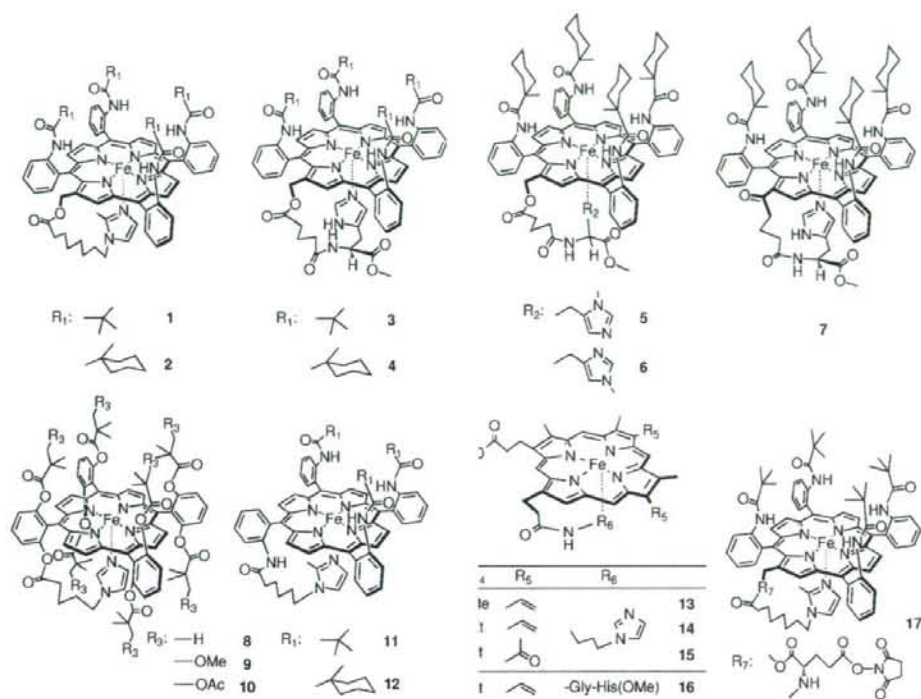


Figure 12. Structure of FePs in HSA-FePs.

3) (139, 140). Solution properties of the HSA-FeP1 solution ( $[r\text{HSA}] = 5 \text{ wt } \%$ ,  $\text{FeP}/\text{HSA}=1-8$ , mol/mol) are almost identical to those of HSA itself: the specific gravity, 1.013; viscosity, 1.1 cP; and COP, 20 Torr. Circular dichroism (CD) spectroscopy and isoelectric focusing measurement revealed that the second-order structure and surface charge distribution of HSA were unaltered after binding of FeP1. The obtained solution showed a long shelf-life of over two years at room temperature (141). Furthermore, HSA-FeP1 has no effect on the morphology of blood cell components (142) and does not engender immunological reaction and platelet activation (143). Upon addition of  $\text{O}_2$  gas through this solution, the visible absorption spectrum immediately changed to that of the  $\text{O}_2$  adduct complex. After exposure to CO gas, a stable carbonyl complex of HSA-FeP1 was formed (139, 140). The coordination structure of FeP1 and spin-state of the central ferrous ion was characterized by IR, resonance Raman, and magnetic circular dichroism (MCD) spectroscopy (139, 140, 144). The carbonyl HSA-FeP1 moved to the NO adduct complex after bubbling NO gas (145). Subsequent ESR spectroscopy revealed that FeP1 in albumin formed a six-coordinate nitrosyl complex. The proximal imidazole moiety does not dissociate from the central ferrous ion when NO binds to the trans side (146).

The  $P_{50}(\text{O}_2)$  value of HSA-FeP1 is always constant (33 Torr, 37 °C) independent of the binding number of FeP1 (eq 1, Table 5)



$$K(\text{L}) = [\text{P}_{50}(\text{L})]^{-1} = k_{\text{on}}(\text{L})/k_{\text{off}}(\text{L}) \quad (1)$$

The  $\text{O}_2$  binding equilibrium curve shows no cooperativity. However, the  $\text{O}_2$  transporting efficiency between the lungs [ $P(\text{O}_2)$ : ca. 110 Torr] and muscle tissue [ $P(\text{O}_2)$ : ca. 40 Torr] is 22%, which is identical to that for RBC.

Table 5. Solution Properties and Characteristics of HSA-FeP1

binding number of FeP1 ( <i>n</i> )	1-8
binding constant of FeP1	$1.2 \times 10^6 - 1.2 \times 10^4 \text{ M}^{-1}$
$M_w$	$(66.5 + 1.3n) \text{ kDa}$
<i>pI</i>	4.8
viscosity <sup>a,b</sup>	1.1 cP
COP <sup>a,b,c</sup>	20 Torr
shelf life <sup>a,d</sup>	> 2 years

<sup>a</sup> In phosphate buffered solution (pH 7.3), [HSA]: 5 g/dL. <sup>b</sup> At 37 °C. <sup>c</sup> A membrane filter with a cutoff ( $M_w, 30 \times 10^3$ ) was used. <sup>d</sup> At 25 °C.

The  $\text{O}_2$  association and dissociation rate constants [ $k_{\text{on}}(\text{O}_2)$  and  $k_{\text{off}}(\text{O}_2)$ ] can be measured using laser flash photolysis (147, 148). Interestingly, the rebinding process of  $\text{O}_2$  to HSA-FeP1 included two phases (fast and slow phase), perhaps because of the different environment around each FeP1 in the protein (149). The  $P_{50}(\text{O}_2)$  value can be controlled by tuning the chemical structure of FeP1. We have synthesized quantities of Fe(II)porphyrins (Figure 12) and evaluated the  $\text{O}_2$  binding parameters of their HSA-FeP hybrids (Table 6).

Actually, FeP2 has a bulky 1-methylcyclohexanamide group on the porphyrin ring plane (150). The  $\text{O}_2$  binding affinity of HSA-FeP2 [ $P_{50}(\text{O}_2)$ : 35 Torr] was almost identical to that of HSA-FeP1. However, the stability of the oxygenated complex increased to 4.5 times its usual value [half-life  $\tau_{1/2}(\text{O}_2)$ , 9 h; pH 7.3; 37 °C] (150).

In general, the basicity and structure of the proximal base greatly influences the  $\text{O}_2$  binding property of  $\text{Fe}^{2+}$ porphyrin. Both FeP3 and FeP4, similar analogues having an His ligand, showed high  $\text{O}_2$  binding affinities [ $P_{50}(\text{O}_2)$ : 3 Torr] (Figure 13, Table 6). Kinetically, substitution of the 2-methylimidazole to His reduces the  $\text{O}_2$  dissociation rate constant (150). Although one might think that high  $\text{O}_2$  binding affinity is not useful as a blood substitute, it can be efficient for oxygenation of hypoxic

Table 6. O<sub>2</sub> Binding Properties of HSA-FePs in Phosphate Buffered S

FeP	$k_{on}(O_2)$ ( $\mu M^{-1} s^{-1}$ )		$k_{off}$
	fast	slow	fast
1	34	9.5	0.75
2	46	7.3	0.98
3	36	6.1	0.059
4	54	8.8	0.089
5	54	6.8	0.02
6	54	8.1	0.62
7	34	4.5	0.045
8	11	1.5	0.50
9	11	2.0	0.41
10	8.9	2.3	0.34
12	29	4.4	1.10
13	—	—	—
14	—	—	—
15	—	—	—
16	—	—	—
17	28	—	0.33

<sup>a</sup> At 37 °C in parentheses.

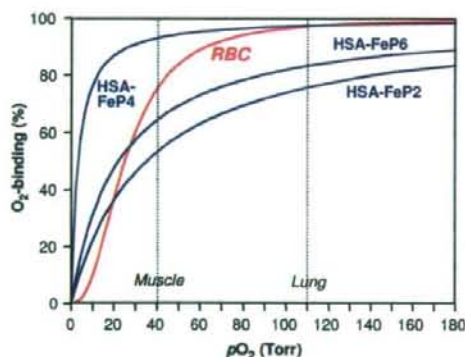


Figure 13. O<sub>2</sub> binding equilibrium curve of HSA-FePs under physiological conditions (pH 7.3, 37 °C).

regions in tumors. Furthermore, the HSA-FeP4 showed long  $\tau_{1/2}(O_2)$  of 25 h (37 °C), which is 13-fold longer than that of HSA-FeP1.

Another HSA-FeP5, in which the active porphyrin has 3-methyl-L-histidine as a proximal base, exhibits an extraordinarily high O<sub>2</sub> binding affinity [ $P_{50}(O_2)$ : 1 Torr] that approaches those of relaxed-state Hb and Mb (151). It is remarkable that replacement of the 3-methyl-L-histidine moiety by 1-methyl-L-histidine isomer (HSA-FeP6) reduced O<sub>2</sub> binding affinity to 1/35th of its former level. The low O<sub>2</sub> affinity of FeP6 is predominantly reflected by the high  $k_{off}(O_2)$  value. The axial Fe–N(1-methyl-L-histidine) coordination might be restrained by steric interaction between the 4-methylene group of the His ring and the porphyrin plane (151).

The proximal histidyl side chain can be introduced easily into the  $\beta$ -pyrrolic position of the porphyrin via an acyl bond in two steps, FeP7 (152). Although an electron-withdrawing acyl group is bound at the porphyrin periphery, the O<sub>2</sub> binding affinity of HSA-FeP7 is slightly higher than that of HSA-FeP4. The rigid His–Gly(carboxy)butanoyl spacer of FeP7 probably produces a favorable geometry to fix the imidazole onto the central Fe<sup>2+</sup> of the porphyrin.

Double-sided porphyrins (FeP8, FeP9, and FeP10) were also incorporated into HSA (153). We expected that steric encumbrances on both sides of the porphyrin enable the HSA-FePs to form a stable O<sub>2</sub> adduct complex. Actually, the  $\tau_{1/2}(O_2)$  of HSA-FeP8 was 5 h, which is 2.5-fold longer than that of HSA-FeP1. In addition, HSA-FeP8 showed high stability against hydrogen

ion (pH 7.3, 25 °C)

$k_{off}(O_2)$ ( $ms^{-1}$ )	$P_{50}(O_2)^a$ (Torr)	refs
0.20	13 (33)	139, 149, 150
0.16	13 (35)	150
0.010	1 (3)	150
0.014	1 (3)	150
0.0024	0.2 (1)	151
0.093	7 (22)	151
0.0059	0.8 (2)	152
0.069	28	153
0.076	23	153
0.088	23	153
0.16	22 (45)	154
—	0.1	155, 156
—	0.1	155, 156
—	0.4	156
—	0.1	155, 156
—	9 (27)	157

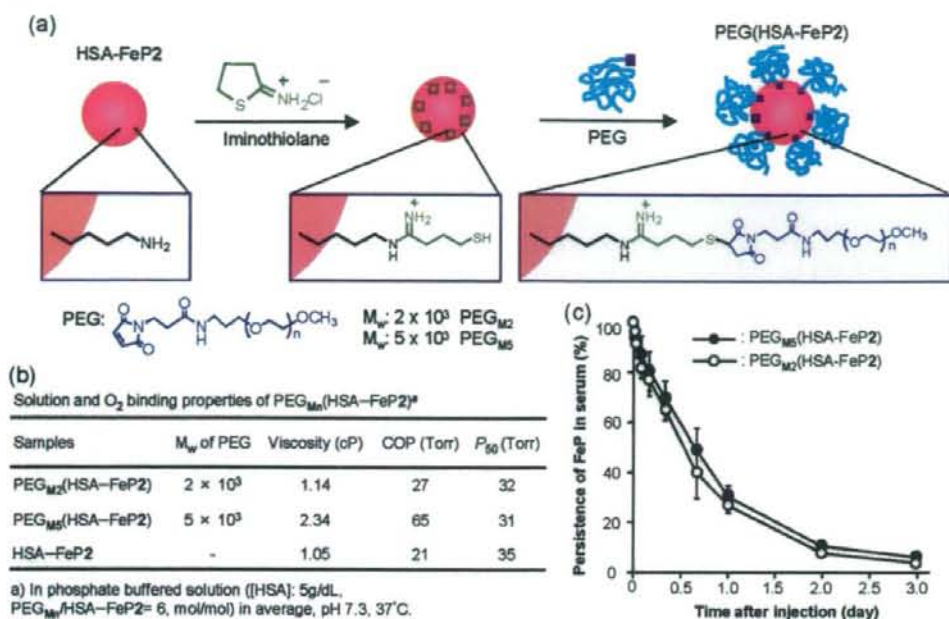
peroxide. The HSA incorporating double-sided porphyrins would be useful for the synthetic analogue of the oxidation enzyme.

Tailed porphyrins having an  $\alpha,\alpha,\alpha,\beta$ -conformer, FeP11 and FeP12, were synthesized easily via four steps from atropisomers of tetrakis(*o*-aminophenyl)porphyrin relative to eight steps of FeP1 (154). Although HSA-FeP12 binds O<sub>2</sub> reversibly, HSA-FeP11 was quickly oxidized by O<sub>2</sub>. We concluded that the 1-methylcyclohexanamide groups are necessary for the tailed porphyrin to form an O<sub>2</sub> adduct complex under physiological conditions.

Investigations have also revealed that heme [Fe<sup>2+</sup>-protoporphyrin IX; protoheme] derivatives having a proximal base at the propionate side chain (FeP13–FeP16) were incorporated into HSA (155, 156). The oxidation process of HSA-FeP13(O<sub>2</sub>) to the inactive ferric state obeyed first-order kinetics, suggesting that the  $\mu$ -oxo dimer formation was prevented by the immobilization of FeP13 into albumin. In fact, HSA-FeP15 showed lower O<sub>2</sub> binding affinity [high  $P_{50}(O_2)$ ] than the others did. The acetyl groups at the 3,8-positions of FeP15 decrease the electron density of the porphyrin macrocycle, thereby reducing the O<sub>2</sub> binding affinity. Actually, HSA-FeP16, in which the His–Gly tail coordinates to the Fe<sup>2+</sup> center, showed the most stable O<sub>2</sub> adduct complexes [ $\tau_{1/2}(O_2)$ , 90 min; pH 7.3; 25 °C] of any of these heme compounds.

**3.2. Surface-Modified HSA-FeP with PEG.** A remaining defect of HSA-FeP is that the active Fe<sup>2+</sup> porphyrin sites dissociate slowly from HSA when infused into animals because FeP is bound noncovalently to albumin. One possible solution is to bind the FeP molecule covalently to the protein. We have synthesized FeP17 having a succinimide side chain; it can react with the Lys amino group of HSA (157). The O<sub>2</sub> binding property of HSA-FeP17 is almost identical to that of HSA-FeP1.

Another approach is surface modification with PEG. Actually, PEG decollations of proteins and liposomes are well-known to enhance their plasma half-life, thermostability, nonimmunogenicity, and solubility in organic solvents (158–163). We surmised that surface modification of HSA-FeP2 by PEG might help to prolong the circulation lifetime of FeP2 and retain its O<sub>2</sub> transporting ability in vivo for a long period. Consequently, HSA-FeP2 (FeP2/HSA = 4/1, mol/mol) was modified with maleimide-PEG, and the solution properties, O<sub>2</sub> binding behavior, and circulatory persistence of the PEG-modified HSA-FeP2 [PEG(HSA-FeP2)] were examined (164). A thiolation reagent, iminothiolane, first reacted with the Lys amino groups of HSA to create active thiols that bind to  $\alpha$ -maleimide- $\omega$ -methoxy PEG (Figure 14a). Mass spectroscopy measurements and quantification of the mercapto group of PEG(HSA-FeP2)



**Figure 14.** Surface modification of HSA-FeP with poly(ethylene glycol). (a) Synthetic scheme of PEG<sub>Mn</sub>(HSA-FeP2). (b) Solution and O<sub>2</sub> binding properties. (c) Persistence of FeP2 in serum after administration of PEG<sub>Mn</sub>(HSA-FeP2) into Wistar rats. Each value represents the mean ± SD of four rats.

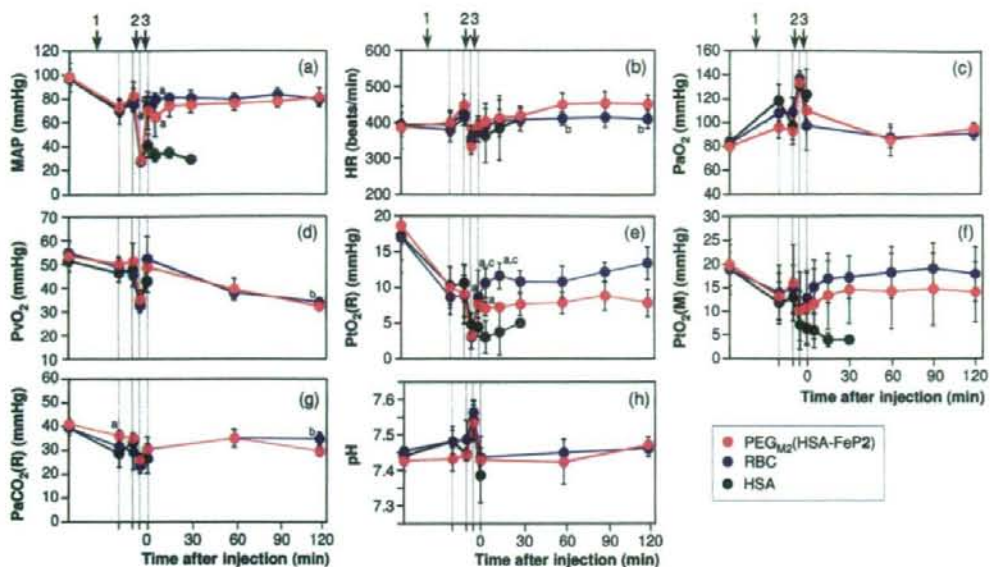
revealed the conjugation of six PEG chains on the HSA-FeP2 surface. The initial FeP2/HSA ratio 4/1 (mol/mol) was unchanged after the PEG binding. The adjustment of viscosity is important to design an artificial O<sub>2</sub> carrier. Maintenance of viscosity is necessary to preserve shear stress on the vascular wall that prevents loss of the functional capillary density (123, 165). The viscosity and COP of PEG(HSA-FeP2) were modulated to some degree by changing the molecular weight of PEG [M<sub>w</sub>: 2 × 10<sup>3</sup> (PEG<sub>M2</sub>) and 5 × 10<sup>3</sup> (PEG<sub>M5</sub>)] (Figure 14b). In fact, PEG<sub>M2</sub>(HSA-FeP2) showed almost identical values of viscosity and COP to those of the nonmodified HSA-FeP2. In contrast, PEG<sub>M5</sub>(HSA-FeP2) showed a higher viscosity and more pronounced hyperoncotic property relative to those of HSA-FeP2. Nevertheless, PEG<sub>M5</sub> conjugate may be useful as an efficient plasma expander (118, 166).

Under physiological conditions, PEG<sub>Mn</sub>(HSA-FeP2) binds and releases O<sub>2</sub>. The P<sub>50</sub>(O<sub>2</sub>) values were almost identical to those of the original HSA-FeP2, indicating that the O<sub>2</sub> binding equilibrium was not influenced by the presence of the PEG chains (Figure 14b). Surface modification by PEG delays proton-driven oxidation of the O<sub>2</sub> adduct complex, giving HSA-FeP2 the τ<sub>1/2</sub>(O<sub>2</sub>) of 12 h, which is almost equal to that of a natural hemoprotein, Mb [τ<sub>1/2</sub>(O<sub>2</sub>), 12 h; pH 7, 35 °C] (167). The conjugated PEG might change the local proton concentration of the HSA interior compared to the outer aqueous solution.

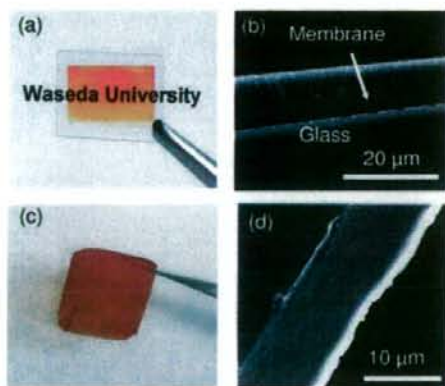
The circulation persistence of FeP2 in the bloodstream was measured after administration of PEG<sub>Mn</sub>(HSA-FeP2) to anesthetized rats (164). The PEG<sub>Mn</sub>(HSA-FeP2) solution (20% volume of the circulatory blood) was injected intravenously into rats from the tail vein. The concentration decays of PEG<sub>Mn</sub>(HSA-FeP2) in the blood showed single exponentials with half-life [τ<sub>1/2</sub>(FeP2)] of 13–16 h (Figure 14c). These values are considerably longer than those of the corresponding nonmodified HSA-FeP1 (168). Surface modification of HSA-FeP2 by PEG prevented the rapid clearance of the incorporated FeP2. On the basis of these findings, we can conclude that surface modification of HSA-FeP2 by PEG comprehensively improved its O<sub>2</sub> transporting ability.

We then proceeded to evaluate physiological responses to an exchange transfusion with PEG<sub>M2</sub>(HSA-FeP2) in an acute anemia rat model (169) (Figure 15). The animals were first placed in a 65 vol % hemodilution with 5 g/dL HSA. They subsequently underwent a 30 vol % blood replacement with the PEG<sub>M2</sub>(HSA-FeP2) solution. As negative and positive control groups, a 5 g/dL HSA solution (HSA group) and washed RBC suspension (RBC group) were infused, respectively, to similarly operated rats in hemorrhage. The isovolemic 65% hemodilution with HSA reduced the Hb concentration, thereby decreasing the O<sub>2</sub> supply to the tissue. Consequently, the mean arterial pressure (MAP), renal cortical O<sub>2</sub> partial pressure [PtO<sub>2</sub>(R)], and O<sub>2</sub> partial pressure of muscle tissue [PtO<sub>2</sub>(M)] were decreased. During hemorrhagic shock by 30% bleeding, significant decreases in the MAP, venous O<sub>2</sub> pressure (PvO<sub>2</sub>), PtO<sub>2</sub>(R), and PtO<sub>2</sub>(M) were observed by the loss of the circulation blood volume. The heart rate (HR) and respiration rate were also decreased. In contrast, arterial O<sub>2</sub> pressure (PaO<sub>2</sub>) increased to about 160% of the basal value (b.v.). The arterial CO<sub>2</sub> pressure (PaCO<sub>2</sub>) decreased to about 62% of the b.v.; the pH increased to 7.55.

The injection of the sample solutions increased the blood volume and improved the circulatory flow. Lactate was washed out from the tissues and into the circulatory system, which decreased the pH to the initial level of 7.43 in all groups. The administration of HSA restored no parameters: death occurred within 41 min. In contrast, the infusion of PEG<sub>M2</sub>(HSA-FeP2) or RBC kept all the rats alive until the end of measurements. After injection of PEG<sub>M2</sub>(HSA-FeP2), the animals showed marked and rapid recovery in MAP, HR, PaO<sub>2</sub>, PvO<sub>2</sub>, PaCO<sub>2</sub>, and pH, resembling that shown in the RBC group. These results demonstrate the O<sub>2</sub> transporting capability of the PEG<sub>M2</sub>(HSA-FeP2) solution as a resuscitative fluid. We observed that albumin-based oxygen carrier does not induce hypertensive action, because of its low permeability through the vascular endothelium in comparison with that of Hb molecules. The heart rate responses after the injection were also negligibly small. Visualization of the intestinal microcirculatory changes clearly



**Figure 15.** Effect of PEG<sub>M2</sub>(HSA-FeP2) solutions on (a) MAP, (b) HR, (c) PaO<sub>2</sub>, (d) PvO<sub>2</sub>, (e) PtO<sub>2</sub>(R), (f) PtO<sub>2</sub>(M), (g) PaCO<sub>2</sub>, and (h) pH in anesthetized rats subjected to hemodilution and hemorrhage. Each value represents the mean  $\pm$  SD of five rats [red, PEG<sub>M2</sub>(HSA-FeP2) group; blue, washed RBC group; and black, HSA group]. Arrows (1), (2), and (3), respectively, indicate the periods of 65% hemodilution, 30% bleeding, and sample infusion. \* $p < 0.05$  versus HSA group (Tukey-Kramer test). <sup>b</sup> $p < 0.05$  versus PEG<sub>M1</sub>(HSA-FeP2) group (unpaired *t*-test), and <sup>c</sup> $p < 0.05$  versus PEG<sub>M2</sub>(HSA-FeP2) group (Tukey-Kramer test).



**Figure 16.** The solid membrane of PEG<sub>M2</sub>(HSA-FeP). (a) Photograph of the membrane on the glass, (b) SEM of the membrane section, (c) photograph of the flexible film peeled from the poly(styrene) dish, and (d) SEM of the isolated film.

revealed the widths of the venule and arteriole to be fairly constant (170).

Reversible oxygenation of PEG<sub>M2</sub>(HSA-FeP2) was observed even in the solid state (171). The aqueous solution of PEG<sub>M2</sub>(HSA-FeP2)(CO) complex was spread on the glass plate and dried overnight at room temperature, producing a red transparent solid membrane (Figure 16a). In contrast, HSA-FeP2 without PEG decollation yielded a brittle membrane with many cracks. Scanning electron microscopy (SEM) observations of the PEG<sub>M2</sub>(HSA-FeP2) membrane showed a uniform thickness of 15  $\mu$ m and a smooth surface (Figure 16b). The  $\tau_{1/2}(\text{O}_2)$  was 40 h, which is three times longer than the value in water. The O<sub>2</sub> binding affinity was about a half that of the monomeric PEG<sub>M2</sub>(HSA-FeP2).

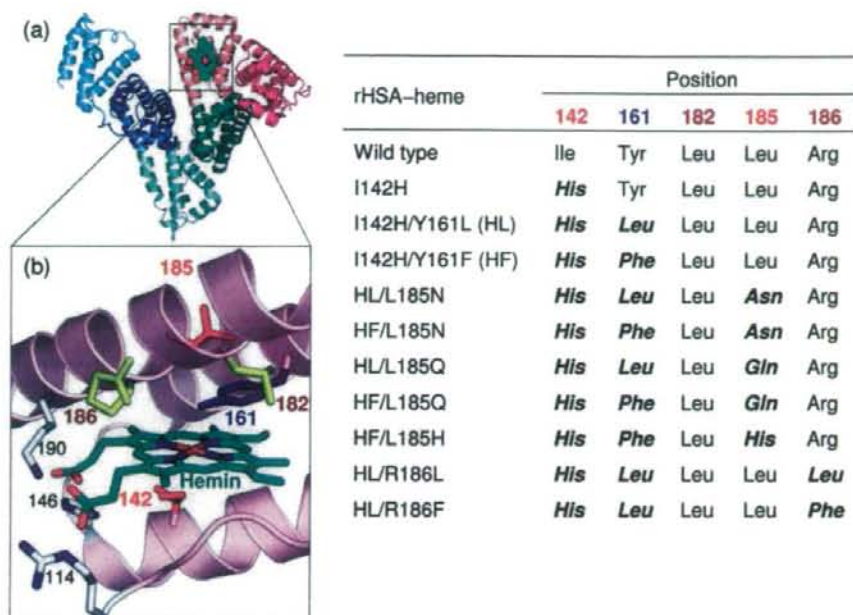
We subsequently added hyaluronic acid (HA) as a supporting polymer to the protein solution and prepared the solid membrane

on a poly(styrene) dish. Actually, HA is known as a glycosaminoglycan component of connective tissues, hyaline bodies, and extracellular matrix (172). Water evaporation of the PEG<sub>M2</sub>(HSA-FeP2)/HA mixture ([HSA]: 2.5 wt % and [HA]: 0.2 wt %) produced a uniform red solid membrane that was easily peeled from the dish, yielding a free-standing homogeneous thin film of the PEG(HSA-FeP2)/HA hybrid (Figure 16c,d).

The PEG<sub>M2</sub>(HSA-FeP2) solution is useful as a valuable O<sub>2</sub>-carrying plasma. Membranes of PEG<sub>M2</sub>(HSA-FeP2) with micrometer thickness can serve as a RBC substitute that can be preserved anywhere and reproduced as a saline solution at any time.

**3.3. Recombinant HSA-Heme (rHSA-Heme) Prepared Using Site-Directed Mutagenesis.** Hemin [Fe<sup>3+</sup>protoporphyrin IX] released from metHb during enucleation of RBC or through hemolysis is captured by HSA with a high binding constant ( $K \approx 10^8 \text{ M}^{-1}$ ) (173). Crystallographic studies have revealed that hemin is bound within a narrow D-shaped hydrophobic cavity in subdomain IB with axial coordination of Tyr-161 to the central ferric ion and electrostatic interactions between the porphyrin propionates and a triad of basic amino acid residues (Arg-114, His-146, and Lys-190) (Figure 17) (174, 175). In terms of the general hydrophobicity of this  $\alpha$ -helical heme pocket, the subdomain IB of HSA potentially has similar features to the heme binding site of Hb or Mb. However, when one reduces HSA-hemin to obtain the ferrous complex, it is autoxidized rapidly by O<sub>2</sub>, even at low temperature ( $\sim 0^\circ \text{C}$ ), because HSA lacks the proximal His, which, in Hb and Mb, enables the prosthetic heme group to bind O<sub>2</sub>. Knowledge of the detailed architecture of the heme binding site in HSA enables us to design mutagenesis experiments to construct a tailor-made heme pocket for stable O<sub>2</sub> binding. Therefore, we used site-directed mutagenesis to introduce an His into the heme binding site that was expected to provide axial coordination to the central Fe<sup>2+</sup> atom of the heme and thereby promote O<sub>2</sub> binding.

Results of our modeling experiments suggested that a favorable position for the axial imidazole insertion would be Ile-142 (Figure 17). The N<sub>i</sub>(His)-Fe distances were estimated

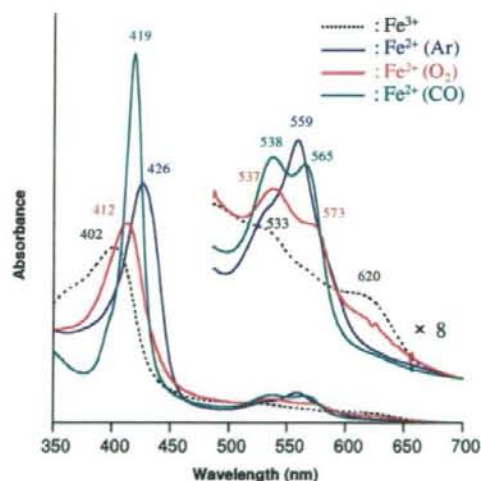


**Figure 17.** (a) Crystal structure of HSA-hemin complex (1O9X) from ref 174. Hemin is shown in a space-filling representation. (b) Heme pocket structure in subdomain IB and positions of amino acids where site-specific mutations were introduced. Abbreviations of rHSA(mutant)s are shown in the table.

as 2.31 Å for H142 (compared to 2.18 Å for Mb). We therefore designed and produced two single mutants I142H and a double mutant I142H/Y161L (HL) (176).

In the UV-vis absorption spectrum of rHSA(HL)-hemin, the ligand-to-metal charge transfer band at 625 nm was weakened because of the Y161L mutation. The MCD spectrum of rHSA(HL)-hemin showed a similar S-shaped pattern in the Soret band region resembling that of ferric Mb (177, 178). These results suggest that rHSA(HL)-hemin is in a predominantly ferric high-spin complex having a water molecule as the sixth ligand. The rHSA-hemin was easily reduced to the ferrous complex by adding a small molar excess of aqueous sodium dithionite under an Ar atmosphere (Figure 18). A single broad absorption band ( $\lambda_{\max}$ : 559 nm) in the visible absorption spectrum and the MCD spectrum of rHSA(HL)-heme indicated the formation of a five-N-coordinate high-spin complex (176, 177, 179). The heme therefore appears to be accommodated in the mutated heme pocket with an axial coordination involving His-142. Upon exposure of rHSA(HL)-heme solution to O<sub>2</sub>, the UV-vis absorption changed immediately to that of the O<sub>2</sub> adduct complex (Figure 18). It formed a carbonyl complex under a CO atmosphere. The single mutant rHSA(I142H)-heme, which retains Y161, was unable to bind O<sub>2</sub>. The polar phenolate residue at the top of the porphyrin plane is likely to accelerate the proton-driven oxidation of the Fe<sup>2+</sup> center. The replacement of Tyr-161 in rHSA(I142H)-heme by Leu enhanced stabilization of the O<sub>2</sub> adduct complex.

To evaluate the kinetics of O<sub>2</sub> and CO bindings to rHSA-hemes, laser flash photolysis experiments were carried out (Tables 7 and 8). It is noteworthy that the absorbance decay accompanying the CO recombination to rHSA(HL)-heme was composed of double-exponential profiles, which is normally not observed in Mb (the faster phase is defined as species I; the slower phase is defined as species II). The ratio of the amplitude of the species I and the species II was approximately 3:2. On the other hand, the rebinding of O<sub>2</sub> to rHSA(HL)-heme followed a simple monophasic decay. Numerous investigations of syn-



**Figure 18.** UV-vis absorption spectral changes of rHSA(HL)-heme in potassium phosphate buffered solution (pH 7.0).

thetic model hemes have helped to reveal the relation between the structure around the hemes and their O<sub>2</sub> and CO binding abilities (4, 147, 148). A bending strain in the proximal base coordination to the central Fe<sup>2+</sup> atom, the "proximal-side steric (proximal pull) effect", is known to be capable of both increasing the dissociation rate for CO and decreasing the association rate. Simultaneously, it increases the O<sub>2</sub> dissociation rate without greatly altering the O<sub>2</sub> association kinetics. Consequently, one possible explanation for the existence of the two phases is that two different geometries of the axial His (His-142) coordination to the central ferrous ion of the heme might exist, each one accounting for a component of the biphasic kinetics of CO rebinding.

**Table 7.** O<sub>2</sub> Binding Parameters of rHSA(Mutant)-Heme Complexes in Phosphate Buffered Solution (pH 7.0) at 22 °C

hemoproteins	$k_{on}(O_2)$ ( $\mu M^{-1}s^{-1}$ )	$k_{off}(O_2)$ (m s <sup>-1</sup> )		$P_{50}(O_2)$ (Torr)	
		I	II	I	II
rHSA(HL)-heme	7.5	0.22	1.70	18	134
rHSA(HF)-heme	20	0.10	0.99	3	31
rHSA(HL/L185N)-heme	14	0.02	0.29	1	14
rHSA(HF/L185N)-heme	26	0.10	1.03	2	24
rHSA(HL/R186L)-heme	25	0.41	8.59	10	209
rHSA(HL/R186F)-heme	21	0.29	7.01	9	203
Mb <sup>a</sup>	14	0.012		0.51	
RBC <sup>b</sup>				8	

<sup>a</sup> Sperm whale myoglobin in 0.1 M potassium phosphate buffer (pH 7.0, 20 °C); ref 180. <sup>b</sup> Human red cell suspension in isotonic buffer (pH 7.4, 20 °C); ref 181.

**Table 8.** CO Binding Parameters of rHSA(Mutant)-Heme Complexes in Phosphate Buffered Solution (pH 7.0) at 22 °C

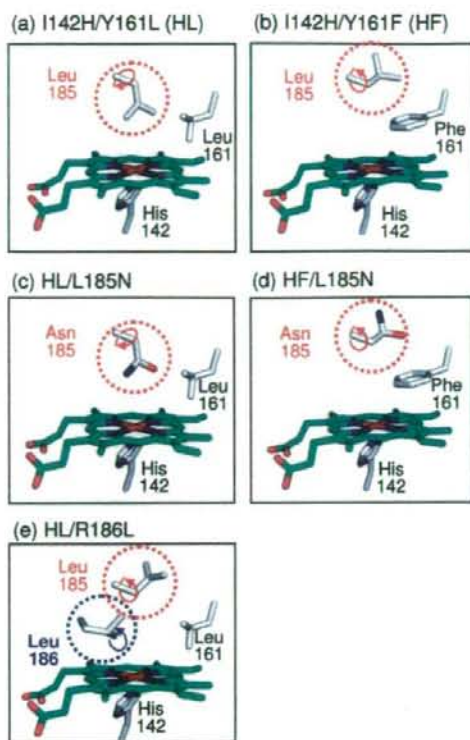
hemoproteins	$k_{on}(CO)$ ( $\mu M^{-1}s^{-1}$ )		$k_{off}(CO)$ (s <sup>-1</sup> )		$P_{50}(CO)$ (Torr)	
	I	II	I	II	I	II
rHSA(HL)-heme	2.0	0.27	0.013	0.079	0.0053	0.240
rHSA(HF)-heme	6.8	0.72	0.009	0.061	0.0011	0.068
rHSA(HL/L185N)-heme	6.8	1.60	0.008	0.039	0.0010	0.020
rHSA(HF/L185N)-heme	7.7	1.09	0.008	0.043	0.0008	0.032
rHSA(HL/R186L)-heme	5.0	0.57	0.011	0.165	0.0018	0.234
rHSA(HL/R186F)-heme	7.9	1.12	0.010	0.148	0.0010	0.107
Mb <sup>a</sup>	0.51		0.019		0.03	

<sup>a</sup> Sperm whale myoglobin in 0.1 M potassium phosphate buffer (pH 7.0, 20 °C); ref 180.

**3.4. Modulation of O<sub>2</sub> Binding Property of rHSA-(mutant)-Heme.** To control the O<sub>2</sub> binding affinity of rHSA-heme, we designed and produced diverse rHSA(mutant)-hemes in which bulky hydrophobic or hydrophilic amino acids were introduced around the O<sub>2</sub> binding site (Tyr-161, Leu-182, Leu-185, and Arg-186) (Figure 17). More recently, the beneficial effect of low-dose CO on the microcirculation by a hemoglobin-based artificial oxygen carrier has been discussed (132, 182). Control of the CO binding affinity of rHSA-heme is also tempting.

**A. Substitution of Tyr-161 with Leu or Phe.** The first, Tyr-161, was substituted to noncoordinating and hydrophobic amino acids (Leu or Phe). The O<sub>2</sub> and CO binding properties of rHSA(HL)-heme and rHSA(I142H/Y161F)-heme [rHSA(HF)-heme] showed that the presence of a Phe rather than a Leu at position 161 results in 6-fold and 4-fold increases in the O<sub>2</sub> binding affinity for species I and II, respectively (Table 7). This enhancement is mainly attributable to an increase in the O<sub>2</sub> association rate constant. The same trend was observed for CO binding [3-fold increase in  $k_{on}(CO)$ ] (Table 8). The substitution of Leu-161 (102 Å<sup>3</sup>) by Phe-161 (137 Å<sup>3</sup>) (183) replaces an isopropyl group with a rigid benzyl group within the heme pocket. In rHSA(HL), the small side chain of Leu-161 might enable free rotation of the side chain of neighboring Leu-185, thereby reducing the volume on the distal side of the porphyrin plane (Figure 19a,b). On the other hand, the bulkier aromatic side chain of Phe-161 might prevent rotation of the isopropyl group of Leu-185 and thereby provide greater room of the distal pocket; this effect might provide easier access to the heme Fe<sup>2+</sup> atom and account for the increased association rates for O<sub>2</sub> and CO.

**B. Substitution of Leu-185 with Polar Amino Acid.** Leu-185 was substituted with a more hydrophilic amino acid (Asn, Gln, or His), which was expected to interact with the coordinated O<sub>2</sub> by hydrogen bond and to stabilize the O<sub>2</sub> adduct complex similarly to Hb and Mb. In rHSA(mutant)-hemes in which Gln

**Figure 19.** The proposed configuration of Leu-185 in (a) rHSA(HL)-heme and (b) rHSA(HF)-heme, Asn-185 in (c) rHSA(HL/L185N)-heme and (d) rHSA(HF/L185N)-heme, and Leu-186 in (e) rHSA(HL/R186L)-heme.

or His was introduced into Leu-185, they formed ferrous six-coordinated low-spin complexes under an Ar atmosphere. That result suggests that the introduced amino acid coordinates to the heme iron as a sixth ligand under an Ar atmosphere. Upon exposure of the solutions to O<sub>2</sub>, they were oxidized. Bis-histidyl hemochromes are known to be oxidized by O<sub>2</sub> rapidly via an outer sphere mechanism (184–186). On the other hand, rHSA(HL/L185N)-heme and rHSA(HF/L185N)-heme in which Asn was introduced at Leu-185 formed ferrous five-coordinated high-spin complexes under an Ar atmosphere. They formed O<sub>2</sub> adduct complexes under O<sub>2</sub> atmosphere. The introduced Asn is estimated to be too far to coordinate to the heme.

Marked differences were apparent in a comparison of the O<sub>2</sub> and CO binding parameters for rHSA(HL)-heme and rHSA(HL/L185N)-heme. First, the presence of Asn rather than Leu at position 185 caused 2-fold and 3–6-fold increases, respectively, in the  $k_{on}(O_2)$  and  $k_{on}(CO)$  values. The Asn might partly rotate upward, which provides somewhat greater space of the distal pocket. Second, Asn-185 induced 18-fold and 10-fold increases in the O<sub>2</sub> binding affinity for species I and II, because the  $k_{off}(O_2)$  values were 1/6–1/11 of their former values. This corresponds to a free energy difference of  $-1.8$  kcal mol<sup>-1</sup> at 22 °C. The magnitude of the effect seems to be reasonable considering that, in HbO<sub>2</sub> and MbO<sub>2</sub>, the distal His-64 stabilizes the coordinated O<sub>2</sub> by  $-0.6$  to  $-1.4$  kcal mol<sup>-1</sup> because of the hydrogen bond (187). In contrast, The O<sub>2</sub> and CO binding parameters for rHSA(HF)-heme and rHSA(HF/L185N)-heme showed no significant differences. The bulky benzyl side chain of Phe-161 can prevent rotation of the polar amide group of Asn-185 and thereby decrease the effect of polarity and size on O<sub>2</sub> and CO binding parameters (Figure 19c,d) (188).



**C. Substitution of Arg-186 with Leu or Phe.** For administration into the human circulatory system, it would be better if the affinity were similar to the human RBC [ $P_{50}(O_2)$ : 8 Torr, 25 °C]. It is expected that providing a certain degree of hydrophobicity into the distal side of the heme by insertion of a nonpolar residue would reduce the  $O_2$  binding affinity of the rHSA-heme complex. The most suitable position for that introduction might be at Arg-186, which is the entrance of the heme pocket and which is rather close to the central Fe(II) ion. Therefore, rHSA(HL/R186L)-hemin and rHSA(HL/R186F)-hemin were prepared. The  $O_2$  dissociation rate constants of rHSA(HL/R186L)-heme and rHSA(HL/R186F)-heme were 3–4-fold higher than that of rHSA(HF)-heme, which reduced the  $O_2$  binding affinities [larger  $P_{50}(O_2)$ ]. This reduction might be attributable to the increased hydrophobicity in the distal pocket. The  $O_2$  binding affinities of rHSA(HL/R186L)-heme [ $P_{50}(O_2)$ : 10 Torr] and rHSA(HL/R186F)-heme [ $P_{50}(O_2)$ : 9 Torr] have become equivalent to those of human RBC. The important structural factor in these mutants is Y161L, which enables the rotation of the isopropyl group of Leu-185 above the  $O_2$  coordination site. Unexpectedly, the  $k_{on}(O_2)$  and  $k_{on}(CO)$  values of rHSA(HL/R186L)-heme and rHSA(HL/R186F)-heme were 3-fold and 3–4-fold higher than those of rHSA(HL)-heme and in the same range as that of rHSA(HF)-heme. In fact, Leu-161 is small, but the hydrophobic Leu-186 or Phe-186 might push up the neighboring Leu-185 residue (Figure 19e) (188).

We have engineered mutant rHSA-heme complexes that can bind  $O_2$ . Principal modifications to the heme pocket that are necessary to confer reversible  $O_2$  binding are (i) replacement of Tyr-161 by hydrophobic amino acid (Leu or Phe), and (ii) introduction of His as a proximal base at position Ile-142. Furthermore, (iii) modification of the distal amino acid has a considerable effect on the modulation of  $O_2$  and CO binding affinities.

#### 4. CONCLUSIONS

The structures of our artificial  $O_2$  carriers differ greatly from those of sophisticated RBCs. However, clear advantages of simplified artificial  $O_2$  carriers are readily apparent: the absence of blood-type antigens and infectious viruses, stability for long-term storage at room temperature for any emergency, all of which overwhelm the functionality of RBCs. The shorter half-life of artificial  $O_2$  carriers in the bloodstream (ca. 3 days) limits their use, but they are applicable as a transfusion alternative for shorter periods of use. Easy manipulation of physicochemical properties such as  $P_{50}(O_2)$  and viscosity supports their possible development of tailor-made  $O_2$  carriers to suit various clinical indications. The achievements of ongoing research described above give us confidence in advancing the further development with the expectation of its eventual realization.

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原著

## ヘモグロビン小胞体を含む血液検体の臨床検査 -デキストラン添加による干渉作用の回避-

### Clinical Laboratory Test of Blood Specimens Containing Hemoglobin-vesicles - Interference Avoidance by Addition of Dextran-

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## 和文抄録

ヘモグロビン (Hb) 小胞体 (250 nm) は赤血球 (8 μm) に比べ1/30程度の大きさの人工酸素運搬体である。Hb小胞体を含む血液検体を遠心分離操作するとHb小胞体は血清に浮遊し、Hb小胞体による干渉作用が血液生化学検査を阻害することが解っているが、その他にも免疫学検査、凝固線溶検査、血糖検査などにおけるHb小胞体の干渉作用を検討する必要がある。本研究では、これらの検査におけるHb小胞体の干渉作用を明らかにすると共に、その回避法を明らかにすることを目的とした。遠心分離により血球と同時にHb小胞体を沈降分離させるため、各種分子量のデキストラン (Dex) を添加して血液中のHb小胞体を凝集させる条件を設定した。さらにヒト血液にHb小胞体を15vol%混合した試料についてこの分離条件を適用し、各検査項目について干渉の有無を検討した。Dexの分子量 (487 kDa) および濃度 (終濃度: 2.6g/dL) の設定により、通常の遠心分離 (3000-5000 rpm, 10 min) でHb小胞体を沈降分離できることを確認した。Hb小胞体は生化学および凝固線溶検査の多くの項目で干渉作用を示したものの、Dexを添加して遠心除去することにより大部分の項目で干渉を回避できた。ただし、Dex添加により生化学検査でリボプロテインの低下、および凝固線溶検査で von Willebrand factor (vWF) 活性の低下、トータル plasminogen activator inhibitor type-1 (PAI-1) の上昇を認め、これらについてはDex添加による干渉を受ける項目として注意を要する。血糖検査として検討したグルコースとグリコヘモグロビンはHb小胞体とDexの干渉なく測定できた。従って、Hb小胞体投与後の血液検体に本法を利用すれば、従来通り遠心分離で血清や血漿が得られ、生化学・免疫学検査、凝固線溶検査、血糖検査の多くの項目で干渉なく検査ができる。

## Abstract

Hemoglobin-vesicle (HbV) is an artificial oxygen carrier of which size (250 nm) is 30 times smaller than red blood cells (8 μm). HbVs remain in serum after centrifugation because of its small size and as a result, HbV interferes with the clinical laboratory test. Here we examine the interference of HbV in other clinical laboratory tests such as immunological test, coagulation fibrinolysis examination, and blood sugar test. To precipitate the HbV by conventional centrifugal separation of blood, we determined an appropriate condition to aggregate HbV by an addition of Dextran (Dex). The obtained plasma or serum was evaluated by common clinical laboratory test. HbV could be precipitated by addition of Dex (Mw. 489 kDa, final concentration: 2.6 g/dL in blood) with conventional centrifugation (3000-5000 rpm, 10 min). Though the presence of HbV interfered with the measurement of many analytes, the interference could be removed by the addition of Dex. However, it should be cautioned that this method underestimates lipoprotein concentration and von Willebrand factor (vWF) activity, and overestimates total plasminogen activator inhibitor type-1 (PAI-1). Blood sugar tests for glucose and glycated hemoglobin (Hb<sub>A1c</sub>) could be performed without interference effect of HbV and Dex. Taken together, the present method will be useful to separate serum

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and plasma from the blood specimens containing HbV for accurate clinical laboratory tests.

## Keywords

Hemoglobin-vesicles, oxygen carrier, blood specimen, clinical laboratory test, interference, dextran

### 1. はじめに

輸血代替物として人工酸素運搬体の使用が想定される救命救急や大手術では、各種の臨床検査やモニタリングにより患者の容態の管理が行われる。大量に投与される人工酸素運搬体の臨床応用を円滑に進めるには、人工酸素運搬体の投与により周辺の医療技術や機器にどのような影響を及ぼすか把握しておく必要がある。これまでに、血液検体検査のほか、Co-oximetry, pulse oximeter, あるいは magnetic resonance (MR) oximetry による酸素飽和度のモニタリングなどにおいて人工酸素運搬体の影響が調べられてきている<sup>1-10)</sup>。Hbベース、パーフルオロカーボン (PFC) ベースを含め、人工酸素運搬体は血球成分に比較して小粒径のため、遠心分離を経てなお血漿層に浮遊する。このため、人工酸素運搬体の特性吸収や濁度が血液生化学検査の多くの項目で測定値に影響を与える。血漿タンパク質と同程度の大きさの修飾Hbは実験的にも血漿から分離するのが極めて困難であり、干渉作用の少ない機器の選定や修飾Hb濃度の関数として干渉分の補正係数を算出することで対処せざるを得ない<sup>11-13)</sup>。ただし、干渉は濃度だけでなく、経時的に変化するMetHb含量などの影響を受けるため<sup>11)</sup>、実際にはこれらの因子を複合的に考慮した補正が必要となる。特定の測定機種について独自の補正を含めた解析手法は臨床試験データの詳細解析には有効と思われるが、補正係数が測定機種に依存するため一般に普及する方法としては適していない。

一方、リン脂質二分子膜でHb溶液を被覆したHb小胞体 (250 nm) は、赤血球 (8 μm) に比較すれば1/30程度の大きさであるが、血漿タンパク質や修飾Hb (数nm) に比較すれば数十倍の大きさがある。この大きさになると、超遠心分離機を使用すれば比較的短時間で沈殿させることができ、また、デキストランなど高分子凝集剤の添加によりHb小胞体を凝集させれば、遠心分離操作で血清からHb小胞体を分離できる。このような分離法の工夫により血液生化学検査における干渉を回避できることがわかっている<sup>14)</sup>。本研究では、デキストラン添加によるHb小胞体の凝集現象を利用して、血液から一段階の遠心分離操作により血漿や血清を分離する条件を検討した。更にこの条件で得られる血清や血漿について生化学・免疫検査、凝固線溶検査、血糖検査を実施し、Hb小胞体による干渉作用を回避できる項目、およびデキストラン添加による干渉作用のある項目を明らかにすることを目的とした。

### 2. 実験方法

#### 2.1. デキストラン (Dex) 添加によるHb小胞体の凝集

市販の分子量の異なるDex粉末 (分子量: 11, 19.6, 40.2, 72.2, 124, および487 kDa, SIGMA) を生理食塩水に溶解させて分

子凝集剤として使用した。Hb小胞体分散液 (Hb: 0.05 g/dL, 3 mL) とDex溶液 (20 g/dL, 0.3 mL) を混合し (Dex終濃度: 1.8 g/dL), 25℃に静置して溶液濁度の変化 ( $\lambda = 700$  nm) から凝集を検出し、その経時変化を観測した。この結果からHb小胞体の凝集生起に要するDex分子量および時間に関する知見を得た。

#### 2.2. 血清分離条件の検討

Hb小胞体が浮遊する血液から血清を分離する実験では、出血蘇生試験のためHb小胞体を投与したビーグル犬から採取した血液を使用した<sup>15)</sup>。循環血液量の50%を脱血した後に、同量のHb小胞体分散液 ([Hb]=8.6 g/dL, 5% リコンビナントアルブミンに浮遊) を投与し、4時間経過した時点での採血液を利用した。採血液 (5 mL) とDex溶液 (20 g/dL, 0.5 ~ 0.88 mL) を混合し、25℃で10分間静置した。この混合液を遠心分離 (5000 rpm, 10分) して上澄み液からのHb小胞体の除去を観測した。これらの結果から、Dexの分子量と濃度についてHb小胞体を完全に除去できる条件を決定した。

#### 2.3. ヒト血液検体検査

健康成人から採血した新鮮血8.5容に対しHb小胞体を1.5容の割合で混合し各種採血管 (ベノJECT II, テルモ) に分注した。さらに予めDex (分子量400~500 kDa, SIGMA) を20 g/dLになるよう生理食塩水に溶解した溶液を終濃度2.6 g/dLで添加混合した。10分間室温放置後3000 rpm, 10分遠心分離し血清または血漿を得た。得られた血清の生化学検査 [総タンパク, アルブミン, 総ビリルビン, アスパラギン酸アミノトランスフェラーゼ (AST), アラニンアミノトランスフェラーゼ (ALT),  $\gamma$  グルタミルトランスペプチダーゼ ( $\gamma$ -GTP), 乳酸脱水素酵素 (LDH), ロイシンアミノペプチダーゼ (LAP), クレアチンキナーゼ (CK), コリンエステラーゼ (ChE), 尿素窒素, 尿酸, 総コレステロール, エステル型コレステロール, 遊離型コレステロール, トロンボポエチン (A), トリグリセライド, リン脂質, 遊離脂肪酸, 高密度リポタンパク質-コレステロール (HDL-C) 定量, カリウム, カルシウム, 無機リン, クレアチニン, C反応性蛋白 (CRP) 定量, フェリチン, ハプトグロビン], 免疫学検査 [免疫グロブリンG (IgG), 免疫グロブリンM (IgM), B型肝炎ウイルス表面 (HBs) 抗原, B型肝炎ウイルス表面 (HBs) 抗体, B型肝炎ウイルス (HCV) 抗体], 血漿の凝固線溶検査 [活性化部分トロンボプラスチン時間 (APTT), プロトロンビン時間 (PT), アンチトロンビンIII (ATIII), トータルプラスミノゲンアクチベーターインヒビター-1 (PAI-1), フィブリノゲン, フォン・ヴィレブランド因子 (vWF) 抗原, vWF活性, D-ダイマー, フィブリン分



解産物 (FDP)], 糖尿病関連検査 [血中グルコース, グリコヘモグロビン (Hb<sub>A1c</sub>) は血液検体で測定] をそれぞれ実施した。これら全ての検査を株式会社エスアールエル (東京) に依頼した。比較はHb小胞体添加血液に生理食塩水を添加し得られた血清または血漿と, 生理食塩水添加血液にDexまたは生理食塩水を添加し得られた血清または血漿とした。また血清採取用真空採血管は凝固促進剤, 血清分離剤が収容されているものなど幾つか種類があるので (Fig. 3の採血管収容物を参照), 採血管の違いによる影響も調べた。干渉作用の有無の判定では, 米国FDAのClinical Laboratory Improvement Amendments (CLIA) の規定する臨床試験に関する測定誤差の許容範囲 (CLIA limit) を判断基準とし<sup>24)</sup>, 基準値との較差 (誤差) が許容範囲を正に超える項目を (↑), 負較差を与える項目を (↓) と表記した。また, 基準値との較差が許容範囲である場合は干渉作用無 (none) と判断した。CLIA limitが規定されていない項目に関しては, 較差の許容範囲を20%として判定した。今回はコントロールとの比較による干渉の有無の判定を目的としたため, 血液にHb小胞体, Dex溶液ないし生理食塩水を添加して希釈率を統一 (1.35倍希釈) して比較した。希釈率の補正は行っていない。

### 3. 結果および考察

#### 3.1. Dex添加によるHb小胞体の凝集

まず, Hb小胞体に各種分子量のDexを添加し, 溶液濁度変化 ( $\Delta O.D.$ ) を経時的にモニターした結果を Fig. 1に示す。分子量72.1 kDa以下のDexをHb小胞体に添加してもほとんど溶液濁度の変化を認めないが, 分子量124 kDaのDexでは溶液濁度の上昇が観測され, さらに分子量487 kDaのDex添加により著しい溶液濁度の増大が観測された。粒子による光散乱強度は粒子径の6乗に比例するため, 凝集による粒子径の増大は溶液濁度の増大 (光散乱強度の増大) として検出される。高分子添加により懸濁粒子が凝集する現象はよく知られ, リン脂質小胞体は共存する高分子との相互作用が比較的詳しく調べられている

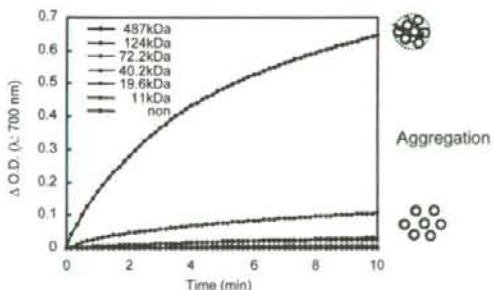


Fig. 1. Kinetics of aggregation of HbV in the presence of 1.8% dextran at 25°C. Increase of the  $\Delta O.D.$  indicates the formation of larger aggregate of HbV by the addition of Dex.

<sup>21,26)</sup>。高分子による小胞体凝集の主要な作用として, 小胞体表面の電荷 (ゼータ電位) の中和あるいは遮蔽による粒子の不安定化, 小胞体間の架橋, 高分子による粒子の排除 (枯乾効果) などが知られ, 一般に同一の繰り返し単位を有する高分子であれば高分子量ほど凝集能が高く, 凝集生起の臨界分子量の存在が認められる<sup>21)</sup>。今回の結果より, 分子量487 kDa程度のDexがHb小胞体の凝集に適していると考えられる。Hb小胞体の凝集は数分間で進行するため, Dexを添加してHb小胞体を凝集させる条件を室温 (25°C) で10分間静置に設定した。

#### 3.2. 血清分離条件の検討

Hb小胞体の浮遊する採血液に対し分子量の異なるDex溶液を添加し, 10分間静置した後に遠心分離した採血管を Fig. 2aに示す (Dex終濃度: 1.8 g/dl)。Dex分子量の効果は明確で, 487 kDaのDexを添加した系のみ透明な血漿が得られた。124 kDaではHb小胞体の沈殿を認めるものの不十分であり, それ以下の分子量では沈殿を認めなかった。遠心加速度 ( $r\omega^2$ ) による粒子の沈降速度 ( $v$ ) は次の式 (1) で表すことができる。

$$v = \frac{d^2}{18} \times \frac{(\sigma - \rho)}{\eta} \times r\omega^2 \quad (1)$$

ここで,  $d$  (cm): 粒子の直径,  $\sigma$  ( $g/cm^3$ ): 粒子の密度,  $\rho$  ( $g/cm^3$ ): 溶液の密度,  $\eta$  ( $g \cdot cm^{-1} \cdot s^{-1}$ ): 溶液の粘度。沈降速度は粒子直径の2乗に比例するため, 凝集により見かけ上の粒子の直径を大きくすることで沈降速度が増大する。Fig. 1の結果との対応から, Dex 487 kDaの添加によるHb小胞体の著しい凝集により遠心で沈降できる大凝集体が生起するものと考えられる。この結果より, Dexの分子量を487 kDaに設定した。

次に, Dex 487 kDaの濃度条件を検討した。7 ml採血管 (ブレン) に高分子凝集剤としてDex 487kDaの0.3–0.75 mLを添加した採血管を作成し, 採血液 (5 ml) を各採血管に採取して, 25°Cで10分間静置した。この混合液を遠心分離 (5000 rpm, 10分) して血漿層からのHb小胞体の除去の有無を観測した。結果として, 透明な血漿層が得られるのはDex 487kDaを終濃度で1.8 g/dl以上添加した場合であり, 更に血漿層を超遠心分離して分析すると, 2.6 g/dl以上では完全にHb小胞体が除去されることを確認した (Fig. 2b)。血漿層にHb由来の吸収はなくDex 487kDa添加による赤血球やHb小胞体の溶血もない。沈殿物は下層の血球層と上層のHb小胞体で明らかな界面を認め, 容易に各々の沈降占有容積率 (クリット値) を計測できる。またHb小胞体の沈殿と血漿の界面も明確であるため, 血漿採取は比較的容易である。以上より, Hb小胞体投与後には, Dex 487kDaが終濃度2.6 g/dlとなるように封入された採血管を使用すれば, 従来通りの遠心分離にて濁度やHb吸収の干渉作用のない血漿ないし血清を採取できることが示された。

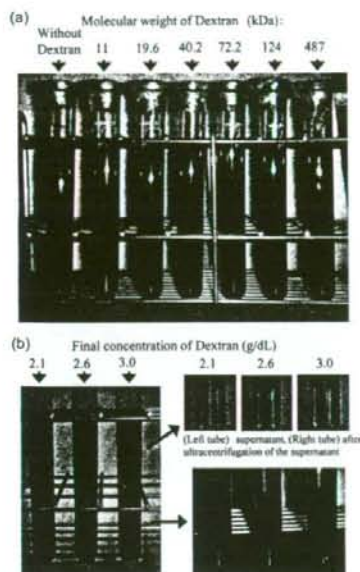


Fig. 2. Precipitation of the aggregated HbV by centrifugation. (a) Effect of the molecular weight of the Dex. (b) Effect of the concentration of the Dextran. HbV could be precipitated in centrifugation of blood sample in presence of 2.6 g/dL Dex (Mw. 487 kDa). Aggregated HbV forms pellet on a red blood cell pellet after centrifugation.

### 3.3. ヒト血液検体検査

高分子量Dex添加によりHb小胞体を除去する方法について、血液検体検査への適合性をヒト血清、血漿の外観と検査結果で評価した。Hb小胞体添加血液に生理食塩水を加えて遠心分離した血清、血漿はHb小胞体が浮遊しているため赤色であった。しかしDex添加によりHb小胞体を除去した血清と血漿 (Fig. 3) は一部の血清採取用採血管 (Fig. 3c, d) で少し赤みが帯びていたもののほぼ黄色であった。赤みが帯びていた血清はDexと血液の混和が不十分であったためHb小胞体を除去しきれなかったことが考えられる。Hb小胞体の除去効果は血清、血漿の外観から肉眼で確認可能であった。各検査においてHb小胞体浮遊血清、血漿は以下のような干渉作用を示した。生化学検査 (Table 1~3) においては総タンパク、アルブミン、LDH、CK、クレアチニン、CRP、ハプトグロビンの上昇が見られた。また、総コレステロール、エステル型コレステロール、遊離型コレステロール、リン脂質ではHb小胞体の脂質膜が影響し上昇した。尿素窒素とIgMは低下し、ChEは低下傾向を示した。総ビリルビン、AST、ALT、 $\gamma$ GTPにおいては検査不能と判断された。免疫学検査項目であるHBs抗原・抗体、HCV抗体いずれもHb小胞体による影響は認められなかった。これらの結果は4種すべての血清採取用採血管で同様であった。生化学・免疫学検査の全32項目中、Hb小胞体の干渉作用により適切な測定値が得られない項目は17ないし18項目に上った (Table 1~3)。一方、Dexの添加によりHb小胞体を除去することで、3ないし4項目 (遊離脂肪酸、リポプロテイン、遊離コレステロール、フェリチン、ALT、AST) を除き適切



Fig. 3. Blood collecting tubes and analytes of clinical laboratory tests. (a)-(g) Blood samples after centrifugation. The blood collecting tube contain blood+saline+saline (control), blood+saline+Dex (interference effect of Dex), blood+HbV+saline (interference effect of HbV), or blood+HbV+Dex (this method).

Table 1. Clinical chemistry and immunological tests (Blood collection tube without contents)

Analytes	Units	CLIA limits	Blood+Saline			Blood+HbV			
			+Saline	+Dex	IF*	+Saline	IF*	+Dex	IF*
			(Control)						
Total protein	g/dL	± 10%	5.1	4.8	none	14	↑	4.8	none
Albumin	g/dL	± 10%	3.3	3.2	none	4.3	↑	3.2	none
Total bilirubin	mg/dL	± 0.4mg/dL or ± 20%	0.2	0.1	none	impossible	×	0.1	none
AST	IU/L	± 20%	11	10	none	impossible	×	10	none
ALT	IU/L	± 20%	7	7	none	impossible	×	7	none
γ-GTP	IU/L	± 20%	8	8	none	impossible	×	7	none
LDH	IU/L	± 20%	134	118	none	270	↑	127	none
LAP	IU/L	ND, ± 20%	63	62	none	58	none	64	none
CK	IU/L	± 30%	46	46	none	89	↑	47	none
ChE	IU/L	ND, ± 20%	243	242	none	215	none	247	none
Urea nitrogen	mg/dL	± 2 mg/dL or ± 9%	6.1	6.0	none	1.2	↓	6.3	none
Creatinine	mg/dL	± 0.3 mg/dL or ± 15%	0.39	0.43	none	2.03	↑	0.47	none
Uric acid	mg/dL	± 17%	2.6	2.5	none	2.5	none	2.7	none
Total cholesterol	mg/dL	± 10%	129	119	none	334	↑	113	none
Cholesterol ester	mg/dL	ND, ± 20%	99	92	none	238	↑	92	none
Free cholesterol	mg/dL	ND, ± 20%	30	27	none	96	↑	21	↓
Triglyceride	mg/dL	± 25%	90	73	none	86	none	69	none
phospholipid	mg/dL	ND, ± 20%	163	156	none	235	↑	153	none
Free fatty acid	mEQ/L	ND, ± 20%	0.08	0.08	none	0.09	none	0.14	↑
HDL-C	mg/dL	ND, ± 20%	43	43	none	39	none	45	none
Lipoproteins	mg/dL	± 30%	12	5	↓	15	none	6	↓
K <sup>+</sup>	mEQ/L	± 0.5 mmol/L	2.4	2.5	none	2.5	none	2.6	none
Ca <sup>2+</sup>	mg/dL	± 0.25 mmol/L	6.2	6.2	none	5.9	none	6.2	none
Inorganic phosphate	mg/dL	ND, ± 20%	2.1	2.0	none	2.2	none	2.2	none
CRP	mg/dL	ND, ± 20%	≤ 0.02	≤ 0.02	none	0.1	↑	≤ 0.02	none
Ferritin	ng/mL	ND, ± 20%	3.5	3.0	none	3.4	none	2.8	↓
Haptoglobin	mg/dL	ND, ± 20%	80	74	none	117	↑	80	none
IgG	mg/dL	± 25%	747	734	none	655	none	744	none
IgM	mg/dL	ND, ± 20%	93	86	none	60	↓	88	none
HBs antigen	IU/mL	positive or negative	< 0.05	< 0.05	none	< 0.05	none	< 0.05	none
HBs antibody	mIU/mL	positive or negative	< 10.0	< 10.0	none	< 10.0	none	< 10.0	none
HCV antibody		positive or negative	0.0	0.0	none	0.0	none	0.0	none

AST: aspartate aminotransferase. ALT: alanine aminotransferase. γ-GTP: γ-glutamyltranspeptidase. LDH: lactate dehydrogenase. LAP:

leucine aminopeptidase. CK: creatine kinase. ChE: cholinesterase. HDL-C: high density lipoprotein cholesterol. CRP: C-reactive protein.

HBs: hepatitis B surface. HCV: hepatitis C virus. \* IF means interference. (↑) overestimation, (↓) underestimation, (none) no interference.



Table 2 Clinical chemistry and immunological tests (Blood collection tube containing clot activator)

Analytes	Units	CLIA limits	Blood+Saline			Blood+HbV			
			+Saline	+Dex	IF*	+Saline	IF*	+Dex	IF*
			(Control)						
Total protein	g/dL	± 10%	5.2	4.8	none	14.3	↑	4.9	none
Albumin	g/dL	± 10%	3.3	3.2	none	4.2	↑	3.2	none
Total bilirubin	mg/dL	± 0.4mg/dL or ± 20%	0.1	0.1	none	Impossible	×	0.1	none
AST	IU/L	± 20%	11	11	none	Impossible	×	12	none
ALT	IU/L	± 20%	5	8	↑	Impossible	×	8	↑
γ-GTP	IU/L	± 20%	7	7	none	Impossible	×	7	none
LDH	IU/L	± 20%	131	114	none	265	↑	115	none
LAP	IU/L	ND, ± 20%	61	62	none	58	none	62	none
CK	IU/L	± 30%	48	47	none	73	↑	48	none
ChE	IU/L	ND, ± 20%	239	236	none	220	none	241	none
Urea nitrogen	mg/dL	± 2 mg/dL or ± 9%	6.1	6.3	none	Impossible	×	5.9	none
Creatinine	mg/dL	± 0.3 mg/dL or ± 15%	0.45	0.41	none	1.80	↑	0.42	none
Uric acid	mg/dL	± 17%	2.5	2.5	none	2.3	none	2.7	none
Total cholesterol	mg/dL	± 10%	126	126	none	333	↑	119	none
Cholesterol ester	mg/dL	ND, ± 20%	95	96	none	239	↑	91	none
Free cholesterol	mg/dL	ND, ± 20%	31	30	none	94	↑	28	none
Triglyceride	mg/dL	± 25%	88	80	none	82	none	69	none
phospholipid	mg/dL	ND, ± 20%	162	157	none	229	↑	157	none
Free fatty acid	mEq/L	ND, ± 20%	0.08	0.08	none	0.09	none	0.15	↑
HDL-C	mg/dL	ND, ± 20%	41	42	none	39	none	43	none
Lipoproteins	mg/dL	± 30%	14	11	none	14	none	4	↓
K <sup>+</sup>	mEq/L	± 0.5 mmol/L	2.4	2.4	none	2.5	none	2.6	none
Ca <sup>2+</sup>	mg/dL	± 0.25 mmol/L	6.0	6.1	none	5.7	none	6.0	none
Inorganic phosphate	mg/dL	ND, ± 20%	2.1	2.0	none	2.8	none	2.1	none
CRP	mg/dL	ND, ± 20%	≤ 0.02	≤ 0.02	none	0.08	↑	≤ 0.02	none
Ferritin	ng/mL	ND, ± 20%	2.8	2.5	none	2.8	none	2.6	none
Haptoglobin	mg/dL	ND, ± 20%	78	78	none	118	↑	82	none
IgG	mg/dL	± 25%	736	721	none	640	none	742	none
IgM	mg/dL	ND, ± 20%	90	84	none	55	↓	85	none
HBs antigen	IU/mL	positive or negative	< 0.05	< 0.05	none	< 0.05	none	< 0.05	none
HBs antibody	mIU/mL	positive or negative	< 10.0	< 10.0	none	< 10.0	none	< 10.0	none
HCV antibody		positive or negative	0.0	0.0	none	0.0	none	0.0	none

AST: aspartate aminotransferase, ALT: alanine aminotransferase, γ-GTP: γ-glutamyltranspeptidase, LDH: lactate dehydrogenase, LAP: leucine amino-peptidase, CK: creatine kinase, ChE: cholinesterase, HDL-C: high density lipoprotein cholesterol, CRP: C-reactive protein, HBs: hepatitis B surface, HCV: hepatitis C virus. \* IF means interference. (↑) overestimation, (↓) underestimation, (none) no interference.