

## Antibodies to Core Lipopolysaccharide Determinants: Absence of Cross-reactivity with Heterologous Lipopolysaccharides

D. Heumann, J. D. Baumgartner,  
H. Jacot-Guillarmod, and M. P. Glauser

Division of Infectious Diseases, Department of Internal Medicine,  
Centre Hospitalier Universitaire Vaudois, Lausanne; Hoffmann-La Roche,  
Basel, Switzerland

Using monoclonal antibodies directed against defined epitopes of endotoxin core, this study demonstrated that the presentation of lipopolysaccharide (LPS) to antibodies is critical for measuring the specific binding of antibodies to LPS structures. False cross-reactive reactions apparently were observed when free core LPS or lipid A were used as antigens in ELISA, whereas coating with complexes of high-density lipoproteins with core LPS increased both the sensitivity and the specificity of the test compared with coating with free core LPS, so that nonspecific binding of antibodies was largely avoided. Using this technique, it was not possible to find broadly cross-reactive core LPS antibodies after immunization of rabbits and humans with rough mutants of gram-negative bacteria. These observations underscore the need for careful evaluation of the potential for cross-reactivity of antisera and of monoclonal antibodies directed against endotoxin core.

The core region of lipopolysaccharides (LPS) is relatively similar in structure among various gram-negative bacteria, whereas the outermost O-side chains are responsible for the marked antigenic diversity of gram-negative bacteria. O-side chain-specific antibodies afford a strain-specific protection but do not protect against heterologous strains in animal models. The core region of LPS is exposed at the surface of the O-side chains lacking rough mutants, among which *Escherichia coli* J5 and *Salmonella minnesota* R595 have been the most studied. After immunization with such mutants, antisera of rabbits or humans contain antibodies directed against core LPS. These antisera have been reported to protect against challenge with heterologous gram-negative bacteria or smooth LPS [1–5]. However, the demonstration that protection is due to core LPS antibodies has not been convincingly put forward so far, especially because of the difficulty in unequivocally demonstrating cross-reactions between core LPS antibodies and purified LPS extracted from heterologous stationary-phase bacteria [1, 6–13].

Investigations of the role of cross-reactive antibodies in immunity to endotoxins require an accurate and specific assay for antibodies directed against the various epitopes of core LPS and lipid A. A number of serologic tests have been proposed to measure these antibodies, including immunofluorescence [14, 15], passive hemagglutination [3, 16–18], and passive hemolysis [19–23]. In recent years, core LPS anti-

bodies have usually been measured by ELISA. Polyclonal rabbit antisera raised against core LPS and lipid A [9, 10, 17, 24–26], murine monoclonal hybridomas selected against core LPS or lipid A [12, 27–30], and human sera from normal volunteers and patients [31–34] were analyzed by this technique. These tests were done with LPS diluted in aqueous buffers as coating antigen. However, due to the limited solubility and self-aggregating properties of amphiphilic core LPS or lipid A, this procedure might induce a poorly reliable coating of the antigen and an increase in the nonspecific binding of immunoglobulins [23]. To immobilize antigen at the bottom of the wells, some techniques have relied on a chemical treatment before coating, which may modify or delete epitopes. Another proposed method has been to use core LPS and lipid A in association with bovine serum albumin to improve the coating [35–37].

In the present study, we investigated various ELISA methods for demonstrating cross-reactive antibodies against endotoxin core. To obviate the potential nonspecific binding of antibodies to the hydrophobic structures of LPS, we complexed LPS with high-density lipoproteins (HDL), a natural carrier in vivo of LPS both in animals and humans [38, 39]. The use of complexes of LPS or lipid A with HDL has three theoretical advantages: (1) a better coating of the plastic due to the protein carrier effect, (2) a removal of LPS contaminants (e.g., proteins and nucleic acids) during the ultracentrifugation step used to purify LPS-HDL complexes, and (3) a more physiologic presentation of the antigen since LPS is rapidly complexed with HDL in the circulation [38, 39].

We compared the binding of antibodies to LPS-HDL complexes with the binding to free LPS or lipid A. We used monoclonal antibodies directed against precisely defined epitopes of core LPS to determine the accuracy of the binding observed with the various antigen preparations. Finally, using the various ELISA techniques, we investigated the development of

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Reprints or correspondence: Dr. J. D. Baumgartner, Division of Infectious Diseases, CHUV, CH-1011 Lausanne, Switzerland.

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cross-reactive antibodies after immunization of animals and humans with *E. coli* J5.

## Material and Methods

**LPS and bacteria.** *E. coli* O111:B4, *E. coli* J5, *S. minnesota* S128 and its rough mutants were gifts from E. J. Ziegler (San Diego). LPS from *E. coli* O111:B4 (O111 LPS) was obtained from Sigma Chemical (St. Louis). LPS from *E. coli* J5 mutant (J5 LPS), LPS from *S. minnesota* R595 mutant (Re LPS), and lipid A from *S. minnesota* R595 mutant (lipid A) were obtained from Ribl Immunochem Research (Hamilton, MT). Lyophilized powders were reconstituted at a concentration of 1 mg/ml in pyrogen-free water, with 0.5% (vol/vol) triethylamine for Re LPS and lipid A. For preparation of radiolabeled LPS, *E. coli* O111:B4 and *E. coli* J5 were grown in trypticase soy broth containing 10 mCi [<sup>3</sup>H]acetate/l, and LPS was extracted by the water-phenol method [40]. [<sup>3</sup>H]LPS from *S. minnesota* R595 was a gift from R. J. Ulevitch (San Diego). Lipid A (0.5 mg; Ribl) was labeled with 100  $\mu$ Ci <sup>51</sup>Cr [41].

**Immunizations.** Boiled *E. coli* O111:B4 and *E. coli* J5 bacterial cells were suspended in 0.15 M NaCl and adjusted spectrophotometrically to a concentration of  $5 \times 10^9$  cells/ml (22% light transmission at 610 nm). New Zealand rabbits were immunized with six intravenous injections of 1 ml of boiled cells administered during a 2-week period. Samples of serum were collected 7 days after the last injection. Lipid A and Re LPS vaccines were prepared by reacting 10 ml of sheep red blood cells with 5 ml of lipid A or Re LPS at 1 mg/ml. After extensive washing in NaCl (0.9%), 0.5 ml of sensitized sheep red blood cells were given intravenously three times per week for 2 weeks and the postimmunization serum was obtained 7 days after the last injection. Healthy human volunteers were immunized with J5 vaccine (provided by E. J. Ziegler) as described [4, 5].

**Monoclonal antibodies.** Mouse monoclonal antibody C117 (IgG2b), a gift from B. J. Appelmek (Amsterdam) was obtained by immunizing mice with *S. minnesota* R595 [42]. It recognizes an epitope consisting of parts of the 3-deoxy-D-manno-2-octulosonic acid (KDO) disaccharide and lipid A; it is specific for Re LPS but specificity for lipid A alone is absent [43]. Clones 8A1 (IgG1) and IC3 (IgG3, an isotype switch variant of 8A1 with the same specificity), developed by Centocor (Malvern, PA), were provided by Hoffmann-La Roche (Basel, Switzerland); they were from immunizations of mice with heat-killed *E. coli* J5 boosted with a mixture of J5 LPS and Re LPS. Both of the latter monoclonal antibodies are directed against a yet undefined epitope within the lipid A structure [29, 30]. The two human IgM hybridomas F117 and F136 were obtained by one of us (H. J.-G.) from fusions of a human heteromyeloma cell line with peripheral blood lymphocytes of human volunteers vaccinated with heat-killed *E. coli* J5 bacteria. F117 is directed against a yet undefined epitope of the lipid A, and F136 exclusively recognizes the J5 LPS core, as revealed by immunoblotting techniques.

**Affinity purification of human J5 LPS antibodies.** Plasma obtained from human volunteers immunized with heat-killed *E. coli* J5 bacteria [5] were screened by ELISA, selected for high titer of antibody against J5 LPS, and pooled. A high-titered pool was used for purification of immunoglobulins. After enrichment of immunoglobulins by precipitation of the pool with saturated (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> at a final concentration of 50%, crude fractions enriched in IgM or IgG were

obtained by gel filtration through Bio-gel A (1.5 m; Bio-Rad Laboratories, Richmond, CA) equilibrated in 20 mM phosphate buffer, 150 mM NaCl, pH 7.3 (PBS). Absence of the other immunoglobulin was checked in both fractions, using Ouchterlony plates developed with precipitating commercial anti-human IgG or IgM antisera (Cappel Laboratories, Malvern, PA). The IgG-enriched fraction was further purified by affinity adsorption on protein-A sepharose (Pharmacia, Uppsala, Sweden), resulting in a homogenous population of IgG devoid of IgM, as revealed both by Ouchterlony analysis and SDS-PAGE. The IgM-enriched fraction was further passed through a second gel filtration column, using Sepharose Cl-2B (Pharmacia) equilibrated in PBS to remove high-molecular-weight contaminants, and finally purified by ion-exchange chromatography on DEAE-sepharose [44] to eliminate  $\alpha_2$ -macroglobulin. When developed in Ouchterlony plates with a polyclonal anti-human IgM antiserum or with a polyclonal anti-whole plasma antiserum, this IgM preparation gave one single line of identity, confirmed by the presence of a single band in SDS-PAGE run in nonreduced conditions.

The final step consisted of affinity purification using J5 LPS-agarose equilibrated in PBS. The affinity resin containing J5 LPS (J5 LPS-agarose) was prepared as described [38] after activation of diamino dipropylamine agarose (Pierce Chemical, Rockford, IL) with 6% glutaraldehyde and coupling with purified core glycolipid in 50 mM sodium borate buffer (pH 8.2). Both purified IgG and IgM fractions were passed through the affinity column; after loading, the column was extensively washed with PBS before specific anti-LPS antibodies were eluted with 0.1 M glycine-HCl buffer (pH 2.5). Eluted antibodies were immediately neutralized with 1 M TRIS and dialyzed against PBS. Specific antibodies were measured ( $\mu$ g/ml) using radial immunodiffusion plates prepared for detection of human IgG or IgM (Nor Partigen, Behring, Germany).

**Preparation of LPS-HDL complexes.** LPS-HDL complexes were prepared as described [45] (except that 100  $\mu$ g of LPS/ml of plasma was added instead of 10  $\mu$ g). Briefly, a pool of fresh human plasma was made 20 mM in EDTA; 100  $\mu$ g aliquots of LPS or lipid A were added for each milliliter of plasma. Incubation was done for 1 h at 37°C. LPS-HDL complexes were then isolated by ultracentrifugation with increasing concentrations of potassium bromide (KBr), as described for purification of HDL [46]. In this manner we prepared J5 LPS-HDL, Re LPS-HDL, lipid A-HDL, and O111 LPS-HDL complexes and isolated normal HDL as a control for ELISA. After ultracentrifugation in KBr, complexes were dialyzed against 75 mM TRIS/150 mM NaCl (pH 7.3) containing 0.1% EDTA and kept at 4°C. The final volume of the solution of purified HDL or HDL-LPS was one-fifth the initial volume of plasma used for the extraction. The amount of LPS within HDL was quantified using radiolabeled LPS.

**ELISA.** The assay was done in 96-well Immulon ELISA plates (Dynatech Laboratories, Alexandria, VA). For coating with PBS buffer, antigen was dissolved at 10  $\mu$ g/ml and 100  $\mu$ l aliquots were distributed per well, incubated for 2 h at 37°C, and let stand overnight at 4°C. With LPS-HDL complexes, 100  $\mu$ l aliquots of complexes dissolved 1:200 in PBS were coated in the same way as for native antigen in PBS. For coating in ethanol, 100  $\mu$ l aliquots of antigen dissolved at 10  $\mu$ g/ml in ethanol were left overnight at room temperature, allowing complete evaporation of the solvent. After antigen coating, plates were shaken empty, extensively washed with PBS, and remaining antigen-attachment sites were blocked by add-

ing 10% fetal calf serum in PBS (PBS-FCS) and incubating for 90 min at 37°C. After additional PBS washings, 100  $\mu$ l of test sera diluted in PBS-FCS were added in triplicate and incubation proceeded for 2 h at 37°C. The plates were again washed thoroughly with PBS and incubated for 75 min with a 1:1000 dilution in PBS-FCS of horseradish-peroxidase conjugates of various specificity: These included goat antibodies specific for human  $\gamma$  or  $\mu$  chain (Sigma), goat antibodies specific for rabbit  $\gamma$  chain (Sigma) or rabbit  $\mu$  chain (Cappel), and goat antibodies specific for mouse  $\gamma$  chain (Sigma). After extensive washing with PBS, substrate (50 mg *o*-phenylenediamine, 20  $\mu$ l H<sub>2</sub>O<sub>2</sub> 30% (vol/vol) in 60 ml of 50 mM citrate buffer, pH 5.5) was added. Enzymatic reaction at room temperature was stopped with 40  $\mu$ l H<sub>2</sub>SO<sub>4</sub> 4 N after various periods, depending on the intensity of reaction. The A<sub>490</sub> was measured using a Dynatech MR600 photometer (Dynatech, Chantilly, VA).

After initial studies to determine the optimal conditions of coating, a concentration of 10  $\mu$ g/ml of free antigens in PBS or ethanol or a dilution of 1:200 of the solution of LPS-HDL in PBS (i.e., 1–2  $\mu$ g/ml LPS) was adopted. Controls were included using HDL alone as a solid-phase coating when ELISA was run with LPS-HDL complexes or using buffer or ethanol for the other two techniques. Usually background values ranged between 0.02 to 0.2 A<sub>490</sub>, with slightly higher background values with HDL than with the other two techniques. Control values were subtracted from total readings.

## Results

**Demonstration of LPS within LPS-HDL complexes.** Plasma (3 ml) containing various concentrations of radiolabeled LPS were processed for isolating LPS-HDL complexes. Table 1 shows that 50%–85% of the radioactivity was recovered within the HDL fraction. The final volume of LPS-HDL complexes was one-fifth the initial plasma volume. Since we used an initial concentration of 100  $\mu$ g/ml LPS in plasma for preparing the LPS-HDL complexes for ELISA, the concentration of LPS within the complexes was between 250  $\mu$ g/ml for O111 LPS-HDL and 425  $\mu$ g/ml for lipid A-HDL. When preparing the plates for ELISA, the complexes were diluted 1:200, corresponding to a final concentration of  $\sim$ 1–2  $\mu$ g/ml. HDL alone had a pattern identical to that of LPS-HDL complexes after protein staining of samples run in SDS-PAGE. The major protein was apoprotein A1. There was a small band of albumin (data not shown).

**Studies with monoclonal antibodies (table 2).** Cross-reactions outside the expected specificity were obtained with monoclonal antibodies when ELISAs were done using free LPS either in PBS or in ethanol solvent for coating. The apparent cross-reactivity observed in these conditions was probably an artifact, since C117 was found to bind to lipid A, whereas it has no specificity for lipid A alone. A similar, probably misleading, cross-reactivity was seen with monoclonal antibody F136, an antibody specific for J5 LPS core sugars; it bound to lipid A, a molecule lacking core sugars. That these findings may be associated with the physicochemical state of LPS is suggested by the observation that the roughest LPS had the strongest apparent cross-reaction; in contrast, non-

**Table 1.** Amount of radiolabeled lipopolysaccharide (LPS) binding to the high-density lipoprotein (HDL) fraction during the preparation of LPS-HDL complexes.

Complex	LPS added ( $\mu$ g/ml)	Total counts per minute (cpm)		
		Plasma	LPS-HDL fraction	% recovered
J5 [ <sup>3</sup> H]LPS	10	7425	4070	55
	100	77,230	43,200	56
Re [ <sup>3</sup> H]LPS	100	75,000	59,220	79
O111 [ <sup>3</sup> H]LPS	100	32,500	15,980	50
[ <sup>51</sup> Cr] lipid A	20	15,680	13,330	85
	60	39,060	26,400	65

NOTE. A known amount of radiolabeled LPS was added to EDTA plasma, and the total cpm was measured in plasma. The LPS-HDL complexes were then extracted [45]. Total cpm in LPS-HDL fractions was measured and the percentage recovered was calculated as (total cpm in LPS-HDL)/(total cpm in plasma)  $\times$  100.

**Table 2.** Comparison of coating with LPS-HDL complexes or with free LPS for measuring monoclonal antibodies against lipid A or core LPS.

Antigen	Mouse monoclonal IgG			Human monoclonal IgM	
	Re LPS C117	Lipid A 8A1	Lipid A 1C3	J5 LPS F136	Lipid A F117
HDL-J5 LPS	0	0	0	>2	0
HDL-Re LPS	1.98	0	0	0	0.11
HDL-lipid A	0	1.67	1.95	0.11	1.16
HDL-O111 LPS	0	0	0	0	0
J5 LPS*	0.15	0.21	0.21	>2	0
Re LPS*	1.98	0.12	0.30	0.09	0
Lipid A*	0.57	>2	>2	0.20	1.57
O111 LPS*	0	0	0	0	0
J5 LPS <sup>†</sup>	0.55	0.53	0.52	>2	0.74
Re LPS <sup>†</sup>	1.84	0.86	0.82	1.04	1.74
Lipid A <sup>†</sup>	1.60	>2	>2	1.16	>2
O111 LPS <sup>†</sup>	0	0	0	0	0.33

NOTE. Values are shown as optical densities (OD) obtained under these conditions: monoclonal antibody concentration,  $\sim$ 0.5  $\mu$ g/ml; dilution of conjugated antibody, 1:1000; incubation of peroxidase substrate, 6 min. Results are expressed after subtraction of background values. OD values <0.05 are shown as 0. LPS, lipopolysaccharide; HDL, high-density lipoprotein.

\* Free LPS or lipid A diluted in PBS.

<sup>†</sup> Free LPS or lipid A diluted in ethanol.

specific binding was not observed when using a hydrophilic antigen like the LPS from the smooth strain *E. coli* O111.

In contrast, when ELISAs were done using LPS within HDL complexes for coating, antibodies appeared highly specific for the corresponding antigens. This was the case for monoclonal C117 directed against Re LPS, monoclonal antibodies 8A1, 1C3, and F117 directed against lipid A, and the monoclonal antibody F136 recognizing J5 LPS. Cross-reactions were nonexistent for IgG antibodies and minimal for IgM antibodies.

**Studies with polyclonal rabbit antisera (table 3).** Using the various antigens for coating, we evaluated the ELISA pat-

**Table 3.** Comparison of coating with LPS-HDL complexes or with free LPS for measuring antibodies against lipid A or core LPS in rabbit sera.

Antibodies, antigen	Antibodies measured in antisera from rabbits immunized with				Non-immune rabbit sera
	<i>E. coli</i> J5	LPS Re	Lipid A	<i>E. coli</i> O111	
<b>IgG</b>					
HDL-J5 LPS	>2	0	0	0	0
HDL-Re LPS	0	>2	0	0	0
HDL-lipid A	0	0	>2	0	0
HDL-O111 LPS	0	0	0	>2	0
J5 LPS*	0.82	0.18	0	0	0
Re LPS*	0.17	0.94	0	0	0
Lipid A*	0.08	0.26	1.07	0	0
O111 LPS*	0	0.22	0	1.88	0
J5 LPS†	1.44	0.10	0.18	0	0
Re LPS†	0.11	1.42	0.26	0	0
Lipid A†	0.33	0.55	>2	0	0
O111 LPS†	0	0	0.15	0.62	0
<b>IgM</b>					
HDL-J5 LPS	>2	0	0.22	0	0
HDL-Re LPS	0	1.65	0.16	0	0
HDL-lipid A	0	0.85	>2	0	0
HDL-O111 LPS	0	0	0	>2	0
J5 LPS*	1.84	0.18	0.14	0	0
Re LPS*	0.12	0.51	0.27	0	0
Lipid A*	0	0.18	0.81	0	0
O111 LPS*	0	0.53	0.11	>2	0
J5 LPS†	>2	0.24	0.44	0.10	0.15
Re LPS†	0.21	1.11	0.44	0	0
Lipid A†	0.21	0.70	1.22	0	0.13
O111 LPS†	0	0.34	0.42	1.89	0

NOTE. Optical densities (OD) were measured as follows: dilution of rabbit antisera, 1:500; dilution of conjugated antibody, 1:1000; incubation of peroxidase substrate, 30 min. OD values < 0.05 are shown as 0. LPS, lipopolysaccharide; HDL, high-density lipoproteins.

\* Free LPS or lipid A diluted in PBS buffer.

† Free LPS or lipid A diluted in ethanol.

tern of reactivity of polyclonal antisera from rabbits immunized with *E. coli* J5 or *E. coli* O111 whole cells, Re LPS, or lipid A extracted from Re LPS. When free LPS in PBS were used as antigens for coating, cross-reactions were observed both with polyclonal rabbit IgG and IgM, including cross-reactions with O111 LPS. This apparent cross-reactivity was increased when ethanol was used as solvent. As for monoclonal antibodies, when LPS-HDL were used as antigens for coating, polyclonal rabbit IgG showed antibody specificities exclusively directed against the corresponding immunizing antigens. Some cross-reactivity against core LPS antigens was observed when polyclonal rabbit IgM were tested, but no antiserum to core LPS cross-reacted with the smooth O111 LPS.

*Studies with human polyclonal antibodies (table 4.)* Similar comparisons using various coating conditions for ELISA were done to evaluate the pattern of reactivity of polyclonal human IgG and IgM antibodies. We used IgG and IgM fractions purified from a pool of plasma from human volunteers

**Table 4.** Comparison of coating with LPS-HDL complexes or with free LPS for measuring antibodies against lipid A or core LPS in humans.

Antigen	Fractionated pooled*		Immunopurified anti-J5 LPS†	
	IgG	IgM	IgG	IgM
HDL-J5 LPS	1.608	1.345	0.735	>2
HDL-Re LPS	0.095	0.090	0	0
HDL-lipid A	0.109	0	0	0
J5 LPS‡	0.620	0.851	0.836	>2
Re LPS‡	0.258	0.312	0	0.130
Lipid A‡	0.321	0.175	0.118	0.166
J5 LPS§	1.029	1.110	0.622	>2
Re LPS§	0.507	0.847	0.067	0.326
Lipid A§	0.423	1.215	0.142	0.851

NOTE. Optical densities (OD) are shown after subtraction of background values under these conditions: primary antibody diluted 1:50, conjugates diluted 1:1000, incubation with peroxidase substrate 20 min. OD values < 0.05 are shown as 0. LPS, lipopolysaccharide; HDL, high-density lipoprotein.

\* Total IgG and IgM fractionated from a pool of plasma with a high titer of antibodies against J5 LPS.

† Anti-J5 LPS IgG and IgM antibodies were immunopurified from total IgG and IgM fractions using a column of J5-LPS agarose.

‡ Free LPS or lipid A diluted in PBS buffer.

§ Free LPS or lipid A diluted in ethanol.

vaccinated with *E. coli* J5. In addition, each immunoglobulin fraction was affinity-purified through J5 LPS-agarose and tested in the various ELISA conditions. The contents of anti-Re LPS or anti-lipid A antibodies in both total IgG or IgM pools appeared considerably higher when free LPS in PBS or ethanol were used for coating than when LPS-HDL complexes were used. Similarly, immunopurified anti-J5 LPS human IgG or IgM had apparent cross-reactivity when free antigens in ethanol or PBS were used for coating. Here again, this cross-reactivity was more important for IgM than for IgG. In contrast, when using LPS-HDL complexes for coating, immunopurified anti-J5 LPS IgG or IgM were exclusively directed against J5 LPS and did not cross-react with Re LPS or lipid A.

### Discussion

The existence of a broad cross-reactivity of core LPS antibodies has not been unequivocally shown so far because of the difficulty in demonstrating in vitro that polyclonal or monoclonal antibodies to core LPS can bind to a variety of LPS extracted from smooth strains [1, 6-13]. In addition, although the structure of the core LPS or of the lipid A is more conserved than that of the O-side chains, there is nevertheless significant variability among strains. Recently, in a study of 29 murine monoclonal core LPS antibodies, Pollack et al. [12] found that cross-reactivity was restricted by inter- and intraspecies differences in covalent core structures and by epitope concealment by overlying O-side chains and core sugars. However, it has been argued that the LPS usually used for in vitro studies of cross-reactivity were extracted from

stationary-phase bacteria and might therefore differ from the LPS harbored by bacteria during infectious processes. Indeed, core LPS determinants may be more exposed on LPS from growing bacteria [26] or from bacteria submitted to antibiotic drugs [47] than from stationary-phase bacteria.

Although the controversies about *in vitro* cross-reactivity of core LPS antibodies are not resolved, recent studies with monoclonal antibodies [29, 48] seem to support the concept that core LPS antibodies can be cross-protective *in vivo*. However, the results of animal experiments in this field are subject to caution because the published data have not always been replicated [8, 20, 49–54]. Thus, the adequate characterization of the binding specificities of antibodies directed against core LPS or lipid A constitutes an important prerequisite for the understanding of their biologic role. To detect cross-reactive antibodies of the various immunoglobulin classes, ELISA appears more appropriate than hemagglutination assays, which measure predominantly IgM antibodies, or than passive hemolysis assays, which cannot discriminate between immunoglobulin classes.

We investigated various ELISA conditions, considering a factor that has recently been emphasized, that LPS from rough strains are hydrophobic structures that can have significant variability of their physical properties (e.g., their degree of hydration and state of aggregation) depending on how they are prepared and diluted. These physical properties have a profound impact on their biologic activities [55–57] and, at least for lipid A, on the nonspecific binding of antibodies to it [23].

There are at least three reasons why LPS-HDL complexes may improve the presentation of LPS to antibodies in ELISA. (1) Due to the physicochemical properties of the complexes, exposure of hydrophilic determinants would favor the binding of specific antibodies while masking the hydrophobic structures that promote nonspecific binding [23]. In addition, the protein content of the LPS-HDL complexes could improve the reliability of the coating of core LPS to plastic surfaces. (2) The preparation of LPS-HDL complexes includes an ultracentrifugation step on KBr gradient that might remove proteins and nucleic acids that are coextracted with LPS during the standard extraction procedures. For instance, phenol-extracted LPS (Sigma) has a bacterial protein content of 0.5%–3% and a bacterial RNA content of 0.5%–3%. These amounts of contaminants may be sufficient to cause significant reactions with antibodies in ELISA when 10–100 µg of LPS/ml are used for coating, since proteins are better coating agents than LPS. (3) HDL is an important physiologic carrier of LPS in serum [38, 39]: If cross-reactive antibodies can act as neutralizing antibodies in protection against endotoxin shock, they would recognize exposed LPS epitopes within LPS-HDL complexes.

In the present experiments, when free lipid A or free core LPS diluted in PBS buffer or in ethanol were used for coating ELISA plates, cross-reactive bindings against core LPS or

lipid A were detected in various core LPS antisera, immunoglobulin preparations, or monoclonal antibodies. In contrast, when using LPS-HDL complexes as antigens, the same antibody preparations showed a very narrow specificity that was restricted to the antigens used for immunization. One possible explanation for the cross-reactivity detected when using free LPS for coating was that nonspecific binding of antibodies occurred due to the exposure of the hydrophobic deepest parts of LPS on the ELISA plates. In support of that explanation was the pattern of reactivity of the monoclonal antibodies, particularly of C117 whose precise specificity is known in detail. The epitope of this anti-Re LPS antibody is part of lipid A and part of the KDO disaccharide. This clone does not recognize lipid A alone or J5 LPS [43].

The methodology used to define the specific epitope of C117 involved inhibition studies and gel precipitation using chemically defined KDO and lipid A-derived synthetic molecules [58]. Thus the reactivity of C117 to free lipid A in the present experiments showed the apparent artifactual nature of this binding. Of particular relevance is our finding that when the binding of C117 was measured using ethanol-coated LPS or lipid A ELISA plates, the nonspecific interaction observed between C117 and lipid A was nearly the same magnitude as its specific interaction with Re LPS. Therefore, our data demonstrate that apparently false cross-reactivities of core LPS antibodies may occur with standard ELISA, a phenomenon that was absent or minimal when LPS-HDL complexes were used for coating.

A possible explanation for the lack of broad cross-reactivity of anti-core LPS antibodies in the assay with LPS-HDL complexes may be the hiding of LPS epitopes within the HDL molecule. For C117, this explanation is unlikely because it was shown by other methods that this antibody does not specifically recognize lipid A. The observed lack of binding of this clone to HDL-lipid A was therefore most likely due to the abolishment of nonspecific interactions. With the other antibodies or antisera, the possibility that epitopes were concealed within HDL complexes seemed unlikely since the specifically defined antibodies reacted fully against their corresponding antigens within HDL complexes. In addition, the specific reactions were more pronounced as revealed by higher optical density values when using LPS-HDL complexes as antigens than when using free LPS, perhaps because the exposure of hydrophilic antigenic determinants was favored in LPS-HDL complexes. Therefore, by analogy with the findings with C117, the apparent cross-reactivity observed when the other antibodies were tested with free core LPS probably resulted from nonspecific interactions.

Although we have tested only one LPS extracted from a smooth strain, *E. coli* O111, the absence of cross-reactivity of core LPS antibodies with this LPS is of particular relevance because the strain is the parent of the rough mutant J5 and has been extensively studied in protection experiments in animals, including in experiments with purified O111 LPS prepared by the same method as the LPS used in our ELISA

[2, 48, 59–61]. Since the core LPS antibodies or antisera, which seemed protective in vivo against O111 LPS, did not recognize it in vitro, the mechanisms responsible for the in vivo protection demonstrated by others remain enigmatic.

Our data do not rule out the possibility that cross-reactivity might be demonstrated in other systems nor do they exclude the possibility that HDL may prevent some cross-reactive sites from interacting with antibody. However, the present study demonstrates that apparently artefactual cross-reactivities of core LPS antibodies may be shown by standard ELISA and that the use of LPS-HDL complexes as coating antigens significantly decreases this phenomenon. Therefore, the results of experiments of cross-reactivities performed with ELISA using free core LPS or lipid A as coating antigens must be interpreted with caution. Indeed, definite proof for true cross-reactive antibodies requires measurements of affinity constants for clearly defined epitopes within the LPS molecule.

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