



RAPID COMMUNICATION

Detection of *H pylori* antibody profile in serum by protein array

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Supported by National 863 Research Project of China, No. 2002AA232031

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Received: 2005-10-21

Accepted: 2005-12-22

serum samples. It is especially useful for large scale epidemiological investigation of the infection of *H pylori*.

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Key words: *Helicobacter pylori*; Protein array; Antibody; Immunogold

Han FC, Li XJ, Jiang H, Qin LP, Li D, Guo YH, Liu ZG, Zhang L, Yan XJ. Detection of *H pylori* antibody profile in serum by protein array. *World J Gastroenterol* 2006; 12(25): 4044-4048

<http://www.wjgnet.com/1007-9327/12/4044.asp>

Abstract

AIM: To detect multiple *H pylori* antibodies in serum samples of individuals who carry *H pylori* by protein array.

METHODS: Recombinant *H pylori* antigens, urease B subunit (UreB), vacuolating toxin A (VacA) and cytotoxin associated gene A protein (CagA), were prepared and immobilized in matrixes on nitrocellulose membrane by robotics to bind the specific immunoglobulin G (IgG) antibodies in serum. Staphylococcus protein A (SPA) labeled by colloid gold was used to integrate the immuno-complex and gave red color signal. The scanner based on charge-coupled device (CCD) could collect the image signal and convert it into digital signal.

RESULTS: When human IgG was printed on the membrane in increasing concentrations and incubated with immunogold, a linear dose response curve was obtained and the detection limit for IgG was about 0.025 ng. The cutoff values, which were defined as the mean grey level plus 3 times of standard deviation, were 27.183, 28.546 and 27.402, for anti-UreB IgG, anti-CagA IgG and anti-VacA IgG, respectively, as 400 human serum samples with negative *H pylori* antibodies were detected by the protein array. When 180 serum samples from patients in hospital were employed for detection of IgG against UreB, CagA and VacA, the sensitivity of the protein array was 93.4%, 95.4%, 96.0%, and the specificity was 94.8%, 94.4% and 97.5%, respectively, as compared with the results obtained by ELISA. The assay also showed high reproducibility, uniformity and stability, and the results were available within 30 min.

CONCLUSION: The protein array is a very practical method for rapid detection of multiple antibodies in

INTRODUCTION

H pylori chronically infect more than half of the world population and are associated with chronic gastritis, peptic ulcer and even gastric cancer^[1-7]. Several virulence factors of *H pylori*, such as urease, vacuolating toxin A (VacA) and cytotoxin associated gene A protein (CagA) have been characterized^[8-11]. Urease is produced by all *H pylori* and functions to hydrolyze urea into CO₂ and NH₃ which can buffer the acid environment and permit the bacterium survival in stomach. All *H pylori* strains carry *vacA* gene and only 50% of them express VacA protein, which assembles into a flower-shaped oligomer, alters intracellular vesicular trafficking and induces vacuole formation in eukaryotic cells. The most important feature that distinguishes *H pylori* strains is the presence or absence of the *cag* pathogenicity island (*cag* PAI). It contains about 30 genes and codes for a type IV secretion machinery system (TFSS), through which CagA is introduced into host cell^[12,13]. Phosphorylated CagA in cytoplasm dephosphorylates host cell protein (cortactin), altering cytoskeletal structure and forming a hummingbird phenotype^[14,15]. CagA is implicated in host cell signal transduction system^[16,17], and CagA positive *H pylori* are much more closely related with peptic ulcer and gastric cancer in western country^[4,6]. When *H pylori* infect human being, multiple antibodies are generated against various antigenic compounds and an antibody library may form in serum. Therefore, screening the antibodies against these virulence factors in serum of *H pylori* infected individual is useful for detection and classification of the pathogen.

Serological assays for diagnosis of *H pylori* infection are included among the noninvasive methods recommended

by the European *H. pylori* study group^[18,19]. At present, the common method to detect *H. pylori* antibodies in serum is enzyme-linked immunosorbent assay (ELISA). The procedure is time-consuming especially when multiple antibodies are detected at the same time. Although there are reports regarding the detection of multiple proteins in an antibody-based protein microarray system^[20-21], the labor-intensive procedures and the expensive instrumentations have limited its application^[22]. Based on the previous work, we developed a low-cost protein array system for rapid detection of multiple *H. pylori* antibodies in serum samples.

MATERIALS AND METHODS

Preparation of antigens

Recombinant urease B subunit (UreB, 40 Ka), N-terminal fragment (amino acid: 1-284, 38 Ka) of CagA and middle region fragment (amino acid: 579-907, 30 Ka) of VacA were produced in our institute previously^[23]. The purity of these proteins was 95%, 96% and 96%, respectively, as identified by coomassie stained gel and evaluated by dual-wave length flying-spot scanner CS-9000 (Shimadzu). The antigenicity was defined as the dilution titer of the antigen to give 1.0 OD in ELISA when the standard positive serum samples (Xinkang Company, Shenzhen) were used. If the original concentration was 1 mg/mL, the dilution titer of UreB, CagA, and VacA was 1:800, 1:1000, and 1:600, respectively.

Preparation of protein array

Nitrocellulose membrane (Pharmacia) with a pore diameter of 0.45 μm was immersed in 0.05 mol/L carbonate buffer (pH 9.0) and dried at room temperature. The antigens (0.1%) in 0.01 mol/L phosphate-buffered saline (PBS, pH 7.0) were transferred from the micro-well plate to the membrane by use of the stainless steel solid pin (0.7 mm in diameter) and dotted on the nitrocellulose membrane by using computer-controlled high speed robotics, MicroGrid II (BioRobotics). The printing was performed in a cabinet at 4°C, with 60% humidity. Each pin was estimated to transfer 10 nL of the solution. The protein array consisted of 4 matrixes, in three of which the antigens were printed in 9 replicates. Another one was the control matrix printed of rabbit myosin (R.M, Sigma) and human IgG (H. IgG, Sigma) (Figure 1A). The membrane was blocked by 5% bovine serum albumin (BSA, Sigma) at 4°C for 2 h and then washed twice by weakly shaking in the wash buffer for 10 min. After drying at room temperature, the membrane was cut into small pieces (1.2 cm \times 1.2 cm) according to the positions properly marked and assembled into the *H. pylori* protein array apparatus. The longitudinal section of the protein array apparatus was illustrated in Figure 2.

Preparation of immunogold

Colloid gold was made according to the sodium citrate reduction method^[23]. In brief, 200 mL of 0.01% chloroauric acid (analytical grade) was heated to the boiling point and 8 mL of the 1% sodium citrate (analytical grade) was then added under vigorously stirring condition. When the solution became dark red in color, the boiling was

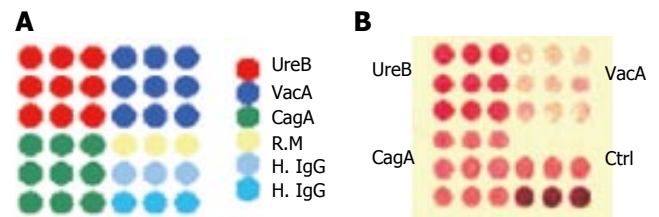


Figure 1 A: Schematic map of protein distribution on the protein array. Four squares were involved and different proteins were printed in different areas: UreB in upper left area; VacA in upper right area; CagA in left bottom area; R.M (10 ng) in the upper row of right bottom area for negative control (Ctrl), H. IgG in the middle row for positive control and at the bottom row for strong positive control; B: positive results of anti-UreB IgG, anti-CagA IgG and anti-VacA IgG detected by the protein array.

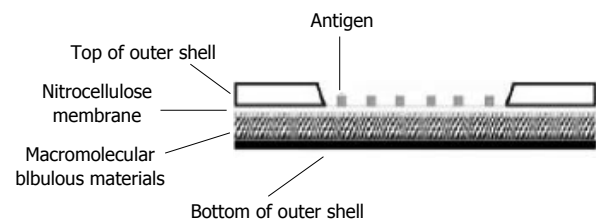


Figure 2 Longitudinal section of the protein array apparatus. The shell of the apparatus was made of plastic materials. A window was opened on the top of the outer shell and the margin of the nitrocellulose membrane printed with antigens was sealed to the inner surface of the window. Under the nitrocellulose membrane was the macromolecular bilubous material.

lasted further for 5 min. The colloid gold should have the highest light absorbance at 525 nm. The shape and size were observed and the diameter (15 nm) was measured under transmission electron microscope. One hundred microlitres of colloidal gold was taken and the pH value was adjusted to 6.2 by 0.1 mol/L potassium carbonate. Staphylococcus protein A (SPA) was added to the final concentration of 1.5 mg% under stirring condition. Ten minutes later, 10 mL of 5% BSA was added and the colloidal gold was stirred for 5 min, making sure that there was no bubble. The immunogold was centrifuged at 4000 g, 4°C for 5 min. The supernatant was then loaded to the Sephacryl S-400 column (Pharmacia) and eluted with 0.02 mol/L tris-buffered saline (pH 7.4). The main red portion was collected and centrifuged at 12 000 g, 4°C for 10 min. The pellets were then suspended in 100 mL solution (0.01 mol/L PBS, 1% BSA) and stored at -20°C for further use.

Procedures to detect antibodies in serum samples

The procedures are as follows: Firstly, 4 drops of the washing solution were added to the center of the nitrocellulose membrane to immerse it evenly. Secondly, 100 μL of the five times diluted serum was added onto the membrane, and 6 drops of the washing solution were dripped onto it to wash the residual serum 5 min later. Thirdly, 100 μL of the immunogold was added onto the membrane, and again, 6 drops of the washing solution were dripped onto it 5 min later. Finally, the protein array was placed into the scanner, which adopts charge-coupled device (CCD) camera to collect the grey level of the central part (0.5 mm in diameter) of every spot and converts the image signal into digital signal. The results

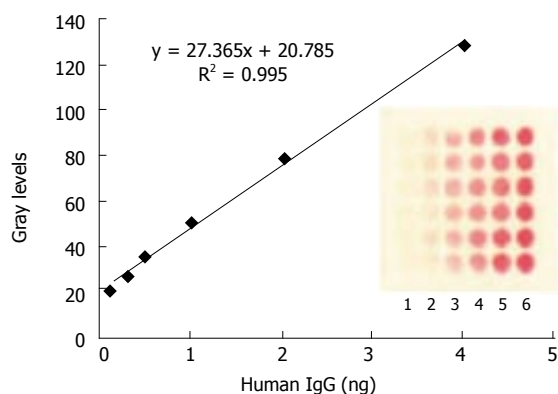


Figure 3 The mean grey level of the spot against various amount of H.IgG immobilized on the protein array. The concentrations of human IgG spotted from lane 1 to lane 6 are as follows: 1.25, 2.5, 5, 10, 20, 40 μ g/mL. The optimized amount of SPA binding to colloid gold was 12 μ g/mL. The data in the plot were obtained from the image of CCD by averaging the mean grey levels of the 6 replicate spots.

were given according to the positions of the matrixes and the mean grey levels of the spots in the matrixes.

Detection of the antibodies in serum samples

Eight hundred and sixty-five serum samples were collected from blood centers in Xi'an and Shenzhen. *H. pylori* antibodies (anti-UreB IgG, anti-CagA IgG and anti-VacA IgG) negative serum samples were screened by using the ELISA kits (kit to detect IgG against *H. pylori* CagA, Jingying Company, Shanghai; kits to detect IgG against *H. pylori* CagA and VacA, Xingkang Company, Shenzhen). The cutoff values of anti-UreB IgG, anti-CagA IgG and anti-VacA IgG were then determined. One hundred and eighty serum samples, collected from patients with different background in Xi'an Center Hospital, were employed to evaluate the sensitivity and specificity of the protein array in detection of *H. pylori* antibodies. Rapid urease test (RUT) and ELISA were also used to detect the infection of *H. pylori*.

Statistical analysis

All experiments were performed in replicate. χ^2 and Student's *t* test were used for statistic analysis.

RESULTS

Limit of detection

Human IgG was printed in 6 replicates in increasing concentrations and incubated with 100 μ L of immunogold on nitrocellulose membrane. A linear dose response curve was obtained (Figure 3) and the detection limit for IgG was about 0.025 ng bound on membrane in this system.

Amount of antigen immobilized

Each antigen with different concentrations ranging from 0.1 to 2.0 ng/nL was printed on the membrane to make the protein array. Five serum samples with *H. pylori* antibody titer of 1:32 in ELISA were used as references to detect the relevant antibody by protein array. The least amount

Table 1 Cutoff values determined in *H. pylori* antibody negative sera

Items	Number	Mean grey values	SD	Cutoff values
Anti-UreB IgG	400	18.060	3.041	27.183
Anti-CagA IgG	400	17.992	3.518	28.546
Anti-VacA IgG	400	17.730	3.224	27.402

of antigen that defined all the samples as positive was designated as the optimized one. In this way, the amount of antigen chosen to immobilize on membrane was 12, 10, and 16 ng for UreB, CagA and VacA, respectively.

Determination of the cutoff values

The cutoff values, which were defined as the mean grey level plus 3 SD, were 27.183, 28.546 and 27.402, for anti-UreB IgG, anti-CagA IgG and anti-VacA IgG, respectively, as 400 serum samples with negative *H. pylori* antibodies were detected by the protein array (Table 1).

Uniformity and reproducibility

When a protein array was randomly chosen from the same batch and subjected to detection of the antibodies in a serum sample, the coefficient variation (CV) of the grey values of the 9 spots in every matrix was less than 8%. When 10 protein arrays were randomly chosen to detect the antibodies in a serum sample, the mean grey value of every matrix in every protein array was obtained and the CV of the mean grey values from the ten relevant matrixes was less than 10%. Ten quality control serum samples, 5 positive and 5 negative for the three antibodies detected by ELISA, were used to screen the antibody profiles by protein array. The results were the same when the test was repeated 5 times.

Sensitivity and specificity

For the 180 serum samples collected from clinical patients, the antibodies were screened by both ELISA and protein array. The results showed that, of these samples, 117, 108 and 99 were positive for anti-UreB IgG, anti-CagA IgG and anti-VacA IgG respectively by protein array (positive results of the three antibodies are shown in Figure 1B), and 122, 109 and 101 were positive respectively by ELISA. The sensitivity of protein array in detection of anti-UreB IgG, anti-CagA IgG and anti-VacA IgG was 93.4%, 95.4%, and 96.0%, and the specificity was 94.8%, 94.4% and 97.5% respectively, compared with the results detected by ELISA (Table 2).

Relation of *H. pylori* antibody profiles to diseases

The positive rates of anti-UreB IgG in groups of gastric cancer (GC, 26/32) and peptic ulcer (PUD, 39/45) were significantly higher than that in group of non-ulcer dyspepsia (NUD, 52/103) ($P < 0.005$). There was no significant difference in the positive rates of anti-CagA IgG or anti-VacA IgG between different patient groups (Table 3).

Table 2 Results detected by ELISA and protein array

Protein array	ELISA					
	Anti-UreB IgG		Anti-CagA IgG		Anti-VacA IgG	
	P	N	P	N	P	N
P	114	3	104	4	97	2
N	8	55	5	67	4	77

P : positive; N: negative.

DISCUSSION

Biochip technique is an advanced tool that enables the binding of multiple molecules in a small area at one time^[24,25]. The molecules (protein, nucleic acid or amino acid) can be immobilized on a piece of support substance such as glass, silica, plastic, nitrocellulose and even metal. The fixation of the molecules to the support substance is by means of non-specific absorption, affinity absorption or covalent attachment, depending on the purpose to design the biochip. Methods based on protein array by non-specific absorption are successfully used for serodiagnosis of infectious diseases in some laboratories^[26-28]. The protein array in the current investigation was designed in accordance with the recent trends including mini-turization of assays and the simultaneous measurement of a panel of antibodies in a single assay.

Actually, this protein array is an immunoassay based on dot immunogold filtration assay. Nitrocellulose membrane was chosen as the chip and the recombinant antigens were immobilized on it by non-specific adsorption. The membrane can bind more amount of protein compared with the glass surface, as each pore of the membrane can be considered as a column and the antibodies can be accumulated when they pass through the membrane. Moreover, the proteins in the membrane are in three-dimensional structure and the binding of antigen and antibody is highly efficient^[29]. Colloid gold was used to label SPA as it was red in color and did not need any substance to produce signal for detection. The reader, which was CCD based video collection system to convert image signal into digital signal, made the results more objective. The entire system offers the simultaneous determination of several antibodies in one test with low cost and faster kinetic. It takes less than 30 min to finish the whole procedures. The stability was quite good and the specificity and sensitivity were close to those of ELISA when *H pylori* antibody profiles of 180 serum samples were detected. Therefore, it is the most practical means of obtaining accurate and precise results. All these make the method a promising future for application, especially for large scales investigation of *H pylori* infection.

In western countries, about 50% to 60% *H pylori* isolates are cagA positive, and the rates are much higher in Asia-Pacific areas^[30-32]. In this study, the positive rates of anti-UreB IgG in groups of GC and PUD were significantly higher than that in group of NUD, indicating that *H pylori* infection was more closely related with GC and PUD, whereas the role of anti-CagA IgG or anti-VacA IgG in prediction of specific diseases was limited.

Table 3 Relation of diseases to *H pylori* antibody (IgG) profiles

Diseases	Numbers				
	Total	RUT	Anti-UreB	Anti-CagA	Anti-VacA
GC	32	30	26 ^a	25	22
PUD	45	43	39 ^c	37	36
NUD	103	58	52	46	41

^aP < 0.005 vs NUD; ^cP < 0.005 vs NUD.

In the pathogenesis study of the *H pylori*, we know very well that the living conditions of this bacterium *in vivo* and *in vitro* are quite different. Some proteins contributing to virulence of the bacterium may not be expressed under *in vitro* culture conditions, and their expression may depend on certain *in vivo* stimuli, such as, cell to cell contact^[33]. So antibody profile may be a reliable indication to reflect the host immune response to the bacterium. Since two genomes of this bacterium have already been sequenced, functional study of these genes is becoming the most important target and many studies have been carried out in the disease related gene screening and identification^[34-37]. With more *H pylori* pathogenic genes being known, more proteins may be added onto the protein array. Obviously, to optimize the binding condition of antigen and antibody and to select one that is suitable for a panel of reactions are the key to success. By accumulating more data about the clinical background of the patients and the antibody profiles in sera, we can draw a clear picture of the relationship between the two aspects.

In summary, detection of multiple *H pylori* antibodies in serum by protein array has the prospect in clinical application, especially in developing countries or small hospitals. Further studies are still going on to increase the density of the protein array.

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