Title	Identification of antigenic peptides derived from B-cell epitopes of nucleocapsid protein of mouse hepatitis virus for serological diagnosis
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Identification of antigenic peptides derived from B-cell epitopes of nucleocapsid protein

## Abstract

Mouse hepatitis virus (MHV) infection is found commonly in laboratory mice and
this virus has been known to cause various diseases such as subclinical infection, enteritis,
hepatitis, and encephalitis. Serological tests are used commonly to diagnose MHV infection.
Complete MHV virions have been used primarily as antigens for serological diagnosis to date.
To develop an antigen that is more specific, more sensitive, and easier to prepare for
serological diagnosis, the antigenic sites in the MHV nucleocapsid (N) protein were screened
in this study. Sixteen antigenic linear sequences in the N protein were found using antisera
obtained from mice infected naturally with MHV and a peptide array containing overlapping
10-mer peptides covering the entire N protein. From these antigenic sequences, two
synthesized peptides, ILKKTTWADQTERGL and RFDSTLPGFETIMKVL, which were
consistent with positions 24-38 and 357-372 of the N protein respectively, were used as
antigens in ELISA. Evaluation of ELISA with these peptides revealed that both peptides were
specific to anti-MHV antisera. Furthermore, ELISA performed using these peptides was more
sensitive than commercial ELISA used for a screening sera from mice infected accidentally to
MHV maintained in cages, suggesting that these peptides are useful for serological diagnosis
of MHV infection.

Keywords: diagnosis, ELISA, epitope mapping, Mouse hepatitis virus, nucleocapsid protein

#### 1. Introduction

Mouse hepatitis virus (MHV) is an enveloped positive-stranded RNA virus, which belongs to the *Coronaviridae* (de Vries et al., 1997). MHV infection causes various diseases such as subclinical infection, enteritis, hepatitis, and encephalitis (Compton et al., 2004; Haring and Perlman, 2001; Matthews et al., 2002). MHV strains are classified as enterotropic or polytropic. Polytropic strains disseminate from the initial site of replication in the respiratory epithelium to neural, hepatic, vascular, and lymphatic tissues. Replication of enterotropic strains is limited generally to replication within the intestinal tract and these strains cause subclinical infection in adult mice. In contrast, polytropic strains are more likely to cause more severe diseases with clinical symptoms (Baker, 1998). Since MHV is one of the most prevalent viruses in laboratory mice (Jacoby and Lindsey, 1997), it is important to detect this virus in mice colonies by appropriate and sensitive screening tests during routine health surveillance programs.

Serological assays detecting antibodies against pathogens in sera are the most popular screening tests for diagnosis of infection (Compton and Riley, 2001). ELISA and the indirect immunofluorescent assay are the most common serological assays for diagnosis of MHV infection (La Regina et al., 1988; Peters et al., 1979). To date, infected cells or purified viruses are used usually as antigens in serological tests for detecting anti-MHV antibodies. Conversely, cross-reactivity between antibodies present in sera and nonspecific antigens used in the test may yield a false positive result. Therefore, the use of purified antigens such as recombinant proteins and synthesized antigenic peptides derived from pathogens will make serological assays more specific and sensitive. MHV comprises at least three proteins: the

envelope glycoprotein designated as the spike (S) protein, the envelope membrane (M) protein, and the nucleocapsid (N) protein (de Vries et al., 1997). The S protein is involved in cellular receptor binding and membrane fusion (Gallagher and Buchmeier, 2001). The N protein is associated with genomic RNA and plays a role in virus assembly though interactions with the viral RNA and the M protein (Hurst et al., 2005; Verma et al., 2006). The N protein consists of three domains conserved domains among the strains, which are separated by two variable regions (Parker and Masters, 1990). Several monoclonal antibodies against the S, N, and M proteins were reported previously, indicating that these proteins are immunogenic (Fleming et al., 1983; Kubo et al., 1994; Nelson et al., 2000; Stohlman et al., 1994; Stühler et al., 1991). Therefore, these proteins are potential antigens for serological diagnosis.

In this study, the antigenicity of the peptides derived from the MHV N protein was

demonstrated by using anti-MHV antisera and peptide array membrane, on which a set of overlapping peptides covering the entire N protein was synthesized. The results showed that 16 antigenic linear sequences existed in the N protein. In these antigenic sequences, two sites coding amino acids 24–38 (ILKKTTWADQTERGL) and 357–372 (RFDSTLPGFETIMKVL) of the N protein are strongly antigenic to anti-MHV antisera derived from mice infected naturally. ELISA performed using these peptides was specific and could detect anti-MHV antisera with higher sensitivity, than a commercial ELISA. These results indicate that these two peptides are useful for serological diagnosis of MHV infection in laboratory mice colonies.

#### 2. Materials and Methods

2.1. Mouse sera.

Sera from C57BL/6 mice infected accidentally with MHV were provided by Prof. Ichiro Miyoshi from the Center for Experimental Animal Science, Graduate School of Medical Sciences, Nagoya City University. As per the guidelines of the Institutional Animal Care and Use Committee (IACUC), Nagoya City University, for screening and eradication of MHV infection, the mice were euthanized and sera were obtained. Normal mice sera were acquired from BALB/c, C57BL/6N, and DBA/2 mice obtained from SLC (Shizuoka, Japan). These sera were derived from the samples obtained for routine screening during a health surveillance program in the animal facility as per the guidelines of IACUC of Graduate School of Veterinary Medicine, Hokkaido University.

#### 2.2. MHV-N peptide array

A set of overlapping 10-mer peptides, obtained by shifting one amino acid from the amino terminus to the carboxy terminus of the N protein that was derived from MHV-JHM strain (GenBank ID: X00990), were synthesized on derivatized cellulose membranes (INTAVIS Bioanalytical Instruments AG, Köln, Germany) using the Auto Spot Peptide Synthesizer ASP-222 (INTAVIS Bioanalytical Instruments AG) according to the manufacturer's protocol (Frank, 2002). Four hundred forty-six peptides were synthesized on the membrane, which are consistent with the peptides coding amino acids from 1–10 to 446–455 of the N protein.

2 2.3. Detection of antibodies in mouse sera binding to peptides on the peptide array membrane

Sera were diluted to 1:1,000 with 5% skim milk in PBS-Tween 20 (137 mM NaCl, 81 mM Na<sub>2</sub>HPO<sub>4</sub>, 15 mM KH<sub>2</sub>PO<sub>4</sub>, 27 mM KCl, and 0.1% Tween 20) and incubated with the peptide array membrane for 2 h at room temperature. The peptide array membrane was washed with PBS-Tween 20 three times and incubated with the HRP-conjugated anti-mouse IgG (GE Healthcare BioSciences, Uppsala, Sweden) diluted to 1:10,000 with 5% skim milk in PBS-Tween 20 for 2 h at room temperature. After reaction with the HRP-conjugated anti-mouse IgG, the membrane was washed with PBS-Tween 20 three times. The antibodies bound specifically to the peptides on the membrane were detected by the ECL Advance Western Blotting Detection Kit (GE Healthcare Biosciences). Visualization and calculation of chemiluminescence of each spot were performed using LAS-3000 imaging system (Fuji film, Tokyo, Japan).

2.4. ELISA

A commercial ELISA, MONILISA MHV (Wakamoto Pharmaceutical Co., Ltd., Tokyo, Japan), was used in this study. Sera used in this ELISA were diluted to 1:40. Evaluation of whether sera were MHV positive or negative was confirmed by comparison with immunoreactivity of positive control sera provided by the manufacture. The peptides coding amino acids 24–38 (ILKKTTWADQTERGL) and 357–372 (RFDSTLPGFETIMKVL) of the MHV-N protein were synthesized by Invitrogen (Carlsbad,

1	CA, USA). In a 96-well plate, 200 µl of 0.5 µM peptide in 0.1 M sodium
2	carbonate-bicarbonate buffer (pH 9.6) was seeded, and the plate was incubated overnight at 4
3	$^{\circ}$ C. The wells were washed three times with PBS-Tween 20 and incubated with 200 $\mu$ l of 1%
4	BSA in PBS-Tween 20 (blocking buffer) for 1 h at 37 °C. The wells were washed three times
5	with PBS-Tween 20, and 200 µl of sera diluted to 1:200 with blocking buffer was added and
6	the plate was incubated for 1 h at 37 °C. After washing the well three times with PBS-Tween
7	20, 200 μl of the HRP-conjugated anti-mouse IgG diluted to 1:10,000 was added to the wells
8	and incubated for 1 h at 37 °C. After washing the well three times with PBS-Tween 20, 200 µl
9	of 1.5 mg/ml $o$ -phenylenediamine with 0.01% $H_2O_2$ was added to each well and incubated for
10	10 min at 37 °C. Following this, 50 μl of 6 N H <sub>2</sub> SO <sub>4</sub> was added and OD was measured at 450
11	nm using a plate reader.

#### 2.5. Statistics

For statistical analyses, Student's t-test, Fisher's exact test, and chi square test were performed using Statview 5.0 software (SAS Institute Inc., Cary, NC, USA).

#### 3. Results

## 3.1. Epitope mapping of the MHV N protein

To determine antigenic peptides for detecting anti-MHV antibodies, B-cell epitope mapping of the MHV N protein was performed using the peptide array membrane. This array

membrane comprised 446 spots, which consisted of 10-mer peptides obtained by a single amino acid shift, to produce a complete overlapping set of the N protein. Sera from mice that were infected accidentally with MHV were incubated with the peptide array membrane. These mice were confirmed to be positive using commercial ELISA for detecting anti-MHV antibodies. As shown in Fig. 1, the IgG antibodies in the sera, bound specifically to the peptides on the spots, were detected. The serum used in Fig. 1 contained antibodies that recognized at least four rows of spots, positioned at 27–29, 84–86, 183–187, and 362–366. This result indicates the presence of four antigenic sites in the MHV N protein that are B-cell linear epitopes. In contrast, no significant intensity was observed when sera from control mice were used (data not shown). Sera from seven mice infected with MHV were then analyzed with the same method. The chemiluminescence intensity of each spot was measured, and the mean values are shown in the graph (Fig. 2). Highest intensity was observed around the 28th spot, on which a peptide consistent with amino acids 28-37 of the MHV N protein was synthesized. A similar intensity was observed at the 362nd spot consistent with the intensity observed at the sequence position 362–371. The peptides on the spots with an intensity higher than 2.0 in Fig. 2 were defined as antigenic, and therefore, as B-cell linear epitopes. Sixteen B-cell epitopes in the MHV N protein were found, and their position and sequence are shown in Table 1.

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- 3.2. Development of ELISA to detect anti-MHV antibodies using antigenic peptides derived
- 21 from the MHV-N protein

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The B-cell epitopes of the MHV N protein from MHV seen in Fig. 2 and Table 1

were expected to be antigens for serological diagnosis of MHV infection. To examine whether these epitopes were appropriate for diagnosis, we developed an ELISA using synthesized peptides corresponding to the B-cell epitopes of the MHV N protein. Two 16-mer peptides corresponding to the amino acids 24–39 (MHV-N 24–39) and amino acids 357–372 (MHV-N 357–372) of the N protein were employed as antigens for this ELISA. The sensitivity and specificity of ELISA using the peptide-antigens were examined by normal or MHV-positive mouse sera. When using the MHV-N 24–39 peptide, the OD values of 1:50-diluted MHV-positive sera were higher than those of the diluted MHV-positive sera, and the values decreased as the dilution increased (Fig. 3, upper graph). In contrast, the OD values of normal sera were almost the same as those without sera. In the case of the MHV N 357–372 peptide, a similar result was seen (Fig. 3, lower graph). These results indicate that both the MHV N peptides could be employed as specific antigens for detecting anti-MHV antibodies.

3.3. Evaluation of ELISA using the MHV-N peptides for serological diagnosis of MHV infection

To evaluate whether ELISA using the MHV-N peptides could be employed for diagnosis, its immunoreactivity was examined using sera from the MHV-contaminated colony that comprised mice infected accidentally with MHV and their cage mates. When sera from eight control mice were used, the mean and the SD were as follows: 0.113 and 0.025 for the MHV-N 24–39 peptides, and 0.181 and 0.023 for the MHV–N 357-372 peptides (Table 2). The means and SD of OD values from the 15 mice maintained in the MHV-contaminated

1 colony are also given in Table 2. The mean OD values of sera from mice in 2 MHV-contaminated colony were significantly higher than that from control mice. 3 Furthermore, these sera were evaluated by both commercial ELISA and ELISA using the 4 MHV-N peptides. As shown in Table 2, the cutoff OD values of ELISA using MHV-N 5 peptides for MHV-positive sera were determined to be the mean + 2 SD from the control mice. 6 All control samples were negative in each ELISA (Table 3). When sera from mice in the 7 MHV-contaminated colony were used, eight of the 15 samples were positive in commercial 8 ELISA. In contrast, 11 samples were positive in the MHV-N 24-39 and 12 samples were 9 positive in the MHV-N 357-372 ELISA. All samples that were positive in commercial 10 ELISA were also positive in ELISA performed using the two peptides (data not shown). 11 Furthermore, the positive/negative frequency was significantly different between the mice in 12 the control and MHV-contaminated colonies when using ELISA using each MHV-N peptides. 13 In contrast, according to the chi-square test results, the positive/negative frequency of sera 14 from mice in the MHV-contaminated colony was not significantly different between 15 commercial ELISA and ELISA using the MHV-N peptides. These results indicate that the 16 ELISA performed using peptides from the N protein is as specific as the commercial ELISA for the detection of MHV infection. 17

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#### 4. Discussion

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MHV infection is found commonly in laboratory mice and causes various diseases such as subclinical infection, enteritis, hepatitis, and encephalitis (Compton et al., 2004; Haring and Perlman, 2001; Matthews et al., 2002). Therefore, it is important to monitor

laboratory mice colonies for MHV infection. Serological assays are used generally for the diagnosis of MHV infection (Compton and Riley, 2001; La Regina et al., 1988; Peters et al., 1979). ELISA is used commonly for diagnosis of MHV infection. In the present study, several B-cell epitopes in the MHV N protein were identified using sera from MHV-positive mice and the MHV-N peptide array membrane. In addition, it was demonstrated that ELISA using the antigenic peptides derived from B-cell epitopes of the MHV N protein is useful for detection of anti-MHV antibodies in mouse sera.

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The N protein is one of the structural proteins of Coronaviridae (deVries et al., 1997). Previous studies indicated that the N protein of *Coronaviridae* is conserved relatively, immunogenic, and expressed abundantly during infection (Daginakatte et al., 1999; Fleming et al., 1983; King and Brian, 1982). Furthermore, a recent study demonstrates that the whole N protein is suitable as an antigen for serological diagnosis using ELISA (Kunita et al., 2011). Therefore, in this study, it was examined whether the peptides derived from the MHV N protein were suitable antigens for diagnosis. SPOT peptide array was used for screening of antigenic epitopes (Frank, 2002; Reineke et al., 2001; Reineke et al., 2002). In this study, the peptide array membrane was prepared that contained a set of nine amino acids overlapping the 10-mer peptides covering the entire N protein. This array was expected to clarify accurately the comprehensive antigenic linear B-cell epitopes in the N protein. In addition, sera from mice infected naturally with MHV were used for identification of antigenic sites in the N protein in live MHV. As expected, several antigenic sites were detected in the N protein (Figs. 1 and 2 and Table 1). Two sites positioned from amino acids 24 to 39 and from amino acids 357 to 373 were more antigenic than the other sites in the N protein. Previous reports demonstrated the location of epitopes of monoclonal antibodies against the N protein (Stohlman et al., 1994). The epitopes were classified into five regions: amino acids 1–133, 171–196, 231–249, 249–277, and 374–465. These antigenic regions corresponded, in part, to those in the present results in Table 1. Because sera used for epitope mapping in this study were derived from MHV-infected C57BL/6 mice, the same strain used in the study by Stohlman et al., this result is considered to be reasonable. In addition, some epitopes that have not been reported as yet were found in the present result, e.g., amino acids 150–159, 307–325, 335–346, and 357–373. In particular, the 357–373 amino acid segment was a strong antigenic site in this study. The present results indicate that the approach using peptide array was very efficient for screening the antigenic sites in the entire N protein to identify B-cell epitopes. In addition, these sites are not the same as the RNA binding site in the central domain and the interactive sites of the M protein in the highly conserved carboxy-terminal domain among the strains of MHV (Hurst et al., 2005; Nelson et al., 2000; Parker and Masters, 1990).

In this study, two sites corresponding to amino acids 24–39 and 357–372 of the N protein were selected as ELISA antigens. As shown in Fig. 3 and Table 2, ELISA using the MHV-N 24–39 or MHV-N 357–372 peptides showed no nonspecific cross-reaction to normal sera and was able to show specific reaction to anti-MHV antisera. The positive/negative frequency was significantly different between mice from the control and MHV-contaminated colonies when ELISA was performed with each MHV-N peptides (Table 3). Furthermore, from this ELISA, not only all MHV-positive sera but also some MHV-negative sera diagnosed by commercial MHV ELISA were detected to be positive (data not shown). In contrast, the antigen in commercial ELISA used in this study was the complete MHV virus (Kagiyama et al., 1990). These results indicate that the MHV-N 24–39 and MHV-N 357–372 peptides are as specific as the complete MHV virus for serological diagnosis of MHV

infection, and suggest that the ELISA using these peptides is more sensitive than commercial ELISA. In addition, the utilization of peptides as antigens, compared to the complete MHV as used in commercial ELISA, gives us the advantage in quality control, preparation, and economy of ELISA products. However, the positive/negative frequency of sera from mice in the MHV-contaminated colony was not significantly different between commercial ELISA and ELISA using the MHV-N peptides. In addition, the reactivity of sera from various mice strains was not examined by ELISA using the MHV-N peptides. Further studies are necessary for validation of sensitivity and specificity of the two peptides to sera from mice in the MHV-contaminated colony.

The sequence of the three carboxyl-terminal residues in the MHV-N 24–39 peptide (GLN) is polymorphic among the strains of MHV. For example, this sequence is substituted into GPN in polytropic MHV-1, MHV-3, and MHV-A59 strains and into AGN in polytropic MHV-S and enterotropic MHV-Y strains (Parker and Masters, 1990). Therefore, it is necessary to examine whether this peptide is reactive to antisera against various MHV strains. In contrast, the amino acids in the MHV-N 357–372 peptide are conserved completely among all MHV strains that protein sequence data were registered in GenBank database, suggesting that the MHV-N 357–372 peptide can detect infection caused by various strains of MHV.

In summary, 16 linear B-cell epitopes were identified in the MHV N protein in this study. The present results indicate that two peptides, ILKKTTWADQTERGLN and RFDSTLPGFETIMKVL, are useful for serological diagnosis of MHV infection using serological assays such as ELISA. Immunological assay with peptide array membrane and antisera is an effective method for identifying B-cell epitopes in the proteins derived from the pathogenic agents and for the development of specific antigens for serological diagnosis.

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**Table 1.** Antigenic sequences in the MHV N protein

Position of spots	Position of	Peptide sequences		
(Intensity $> 2.0$ in Fig. 2)	amino acids			
1, 2, 5	1–14	MSFVPGQENAGSRS		
24–30	24–39	ILKKTTWADQTERGLN		
49–55	49–64	PKQTATTQPNSGSVVP		
72, 73	72–82	ITQFQKGKEFQ		
81, 83	81–92	FQFAQGQGVPIA		
130	130–139	YLGTGPYAGA		
150	150-159	WVASQQAETR		
165	165–174	VERDPSSHEA		
184–186	184–195	VLPQGFYVEGSG		
193	193–202	GSGRSAPASR		
240–244	240–253	LVLAKLGKDAGQPK		
307–316	307–325	SDPQFPILAELAPTAGAFF		
335, 337	335–346	KNSGGADGPTKD		
347	347–356	VYELQYSGAV		
356–364	356–373	VRFDSTLPGFETIMKVLN		
384	384–393	GGADVVSPKP		

**Table 2.** Detection of anti-MHV antibody from mice sera by ELISA using the MHV-N peptides

	MHV-N 24–39 peptide			MHV-N 357–372 peptide		
	${ m OD}_{450}$			${ m OD}_{450}$		
	mean	SD	mean + 2 SD	mean	SD	mean + 2 SD
Control (n = 8)	0.113	0.025	0.164	0.181	0.023	0.226
MHV-contaminated	$0.326^{*}$	0.205		$0.388^{*}$	0.219	
colony $(n = 15)$						

<sup>\*</sup> p<0.05 vs control by Student's t-test.

Table 3. Evaluation of ELISA using the MHV-N peptides for detection of MHV infection

	Positive samples/total samples				
	Commercial	ELISA using	ELISA using		
	ELISA	MHV-N 24–39	MHV-N 357–372		
Control	0/8	0/8	0/8		
MHV-contaminated colony	8/15*	11/15*	12/15*		

<sup>\*</sup> p<0.05 vs control by Fisher's exact test.

### Figure Legends

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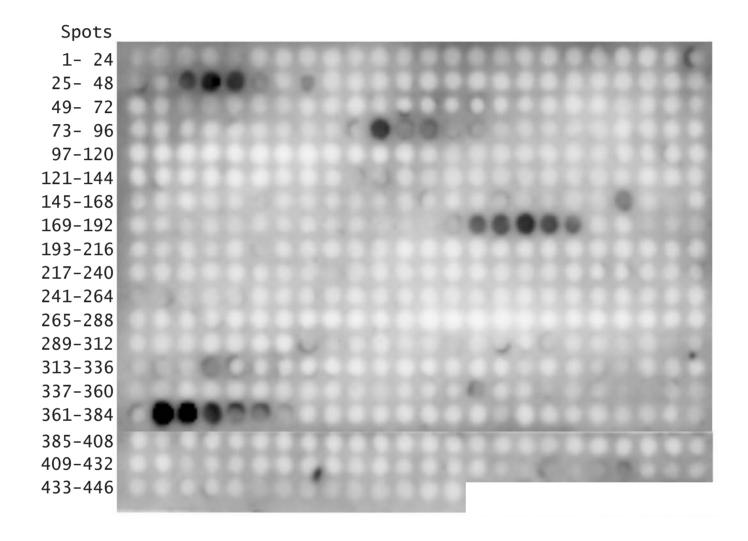
- 3 Fig. 1. Detection of antigenic sites in the MHV N protein using a peptide array membrane.
- 4 Antibodies in mouse sera bound to each peptide on a peptide array membrane were visualized
- 5 by chemiluminescence. A typical image from seven MHV-positive sera samples is shown.

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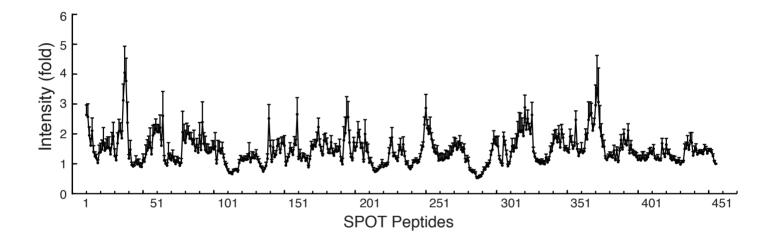
- 7 Fig. 2. Antigenicity of the MHV N protein. Peptide-binding antibodies in sera from mice
- 8 infected naturally were detected as shown in Fig. 1. The intensity of chemiluminescence at
- 9 each spot was quantified. The values are relative to the background and are presented as mean
- $\pm$  SEM of seven sera. The values on X-axis indicate the position of the spots.

- 12 **Fig. 3.** ELISA using the MHV-N peptides. MHV-positive or -negative sera were diluted and
- used for ELISA using the MHV-N peptides as described in Materials and Methods. The upper
- graph is a plot of OD<sub>450</sub> from ELISA using the MHV-N 24–39 peptide. The lower graph is a
- 15 plot of OD<sub>450</sub> from ELISA using the MHV-N 357–372 peptide. Open circles indicate
- MHV-negative sera, and closed squares indicate MHV-positive sera. (-) indicates the sample
- without serum. The plots are representatives from three independent experiments.

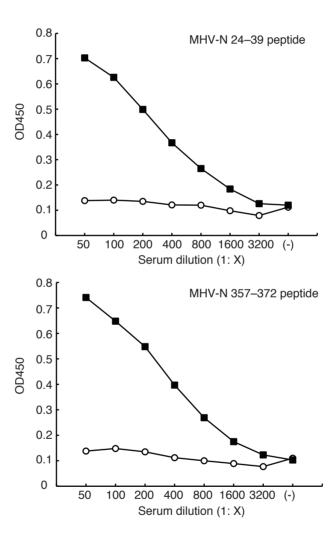
## FIGURE 1

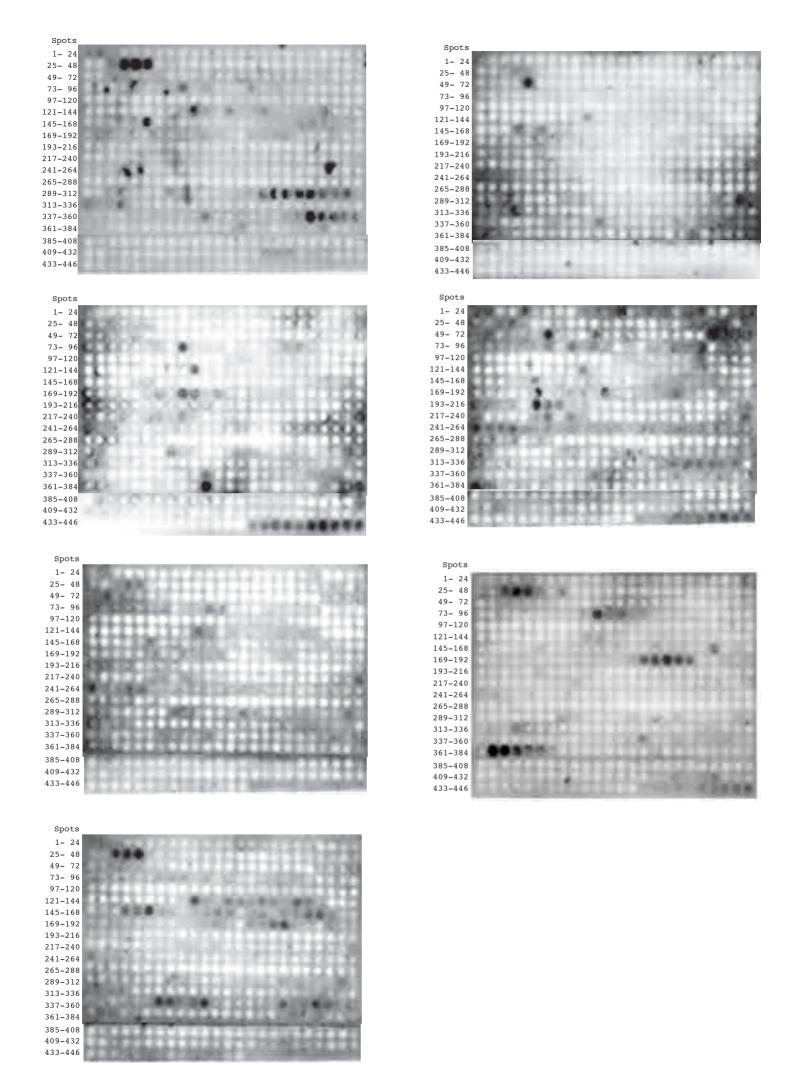


# FIGURE 2



# FIGURE 3





Supplemental Figure