Novel Immunogenic Peptides Elicit Systemic Anaphylaxis in Mice: Implications for Peptide Vaccines

Claire Mary Smith, Peter Bradding, Daniel Robert Neill, Helen Baxendale, Franco Felici and Peter William Andrew

_J Immunol_ 2011; 187:1201-1206; Prepublished online 27 June 2011; doi: 10.4049/jimmunol.1002152
http://www.jimmunol.org/content/187/3/1201

References

This article cites 19 articles, 6 of which you can access for free at:
http://www.jimmunol.org/content/187/3/1201.full#ref-list-1

Subscription

Information about subscribing to _The Journal of Immunology_ is online at:
http://jimmunol.org/subscription

Permissions

Submit copyright permission requests at:
http://www.aai.org/About/Publications/JI/copyright.html

Email Alerts

Receive free email-alerts when new articles cite this article. Sign up at:
http://jimmunol.org/alerts
Novel Immunogenic Peptides Elicit Systemic Anaphylaxis in Mice: Implications for Peptide Vaccines

Claire Mary Smith,* Peter Bradding,* Daniel Robert Neill,* Helen Baxendale,†
Franco Felici,‡ and Peter William Andrew*

Peptide-based therapies are showing increasing potential for the development of vaccines and in the treatment of many important diseases. We previously reported two peptide conjugate vaccines that protected mice against pneumococcal disease. During this study, we observed an unexpected phenomenon; several vaccine candidates induced a rapid, fatal anaphylaxis after booster injection of the peptide conjugate. Further investigation indicated the reaction was mediated by the production of peptide-specific IgE and the release of histamine. Notably, among seven peptides tested, all of which bound the same mAb that selected them from a phage library, only four elicited this severe reaction. Sequence alignment analysis of all peptides revealed unique clusters of acidic amino acid residues in the allergenic peptides. Substitution of the acidic amino acid residues, ED, of peptide MP2 with their amine equivalents, QN, eliminated the anaphylactic effects but did not affect the production of peptide-specific IgG. These results have important implications for both the study of allergens and the development of future peptide-based therapies. The Journal of Immunology, 2011, 187: 1201–1206.

Peptide therapies are showing increasing potential in the treatment of many important diseases ranging from influenza to allergic disease and, more recently, cancer (1, 2). The activity of these short peptides varies significantly from interfering with native protein activity to inducing protection against larger vaccine targets. A successful new approach to circumvent the problem of using poorly immunogenic bacterial polysaccharides as effective vaccine targets is to substitute the native polysaccharide Ag with peptides that mimic the protective epitope (3–7). We have recently described the ability of peptide mimics of pneumococcal polysaccharides to induce immunity and protect mice against Streptococcus pneumoniae (8). In this study, seven peptides were selected from a phage-display library using the antitype 6B polysaccharide mAb Db3G9. Surprisingly, immunization with four of seven 12-mer peptides induced symptoms of anaphylaxis in mice upon re-exposure to the Ag. This effect has not been described previously, and the results presented in this work could have important implications for both the study of allergens and the development of future peptide-based therapies.

Materials and Methods

Peptides

The peptides used in this study were selected from a random 12-aa phage-displayed peptide library using a human anti-6B pneumococcal polysaccharide mAb (Db3G9) as described by Smith et al. (8). Peptides of seven positive phage clones (MP2, MP10, MP13, MP14, MP15, MP17, and MP18, the sequences of which are shown in Table I) and the modified peptide MP2m were synthesized by standard F-moc solid phase peptide synthesis, and the peptide was conjugated to keyhole limpet hemocyanin (KLH) or BSA (at a molar ratio of 6:1) by Cambridge Research Biochemicals (Cambridge, U.K.). The purity of the peptides was >90% as assessed by HPLC. The molecular mass of purified peptides was confirmed by mass spectrometry. Each peptide conjugate was dissolved in PBS at a concentration of 5 mg/ml and stored at −20°C until use.

Mice and immunization protocol

Nine-week-old, outbred, female MF1 mice were purchased from Harlan-Olac (Bicester, U.K.). The immunization of the mice was performed as follows: a stock solution of the native peptide, peptide–KLH conjugate, or control suspension containing 0.5 mg/ml and emulsified with 0.5 volume of InjjectAlum adjuvant (Pierce, Rockford, IL). A control suspension containing 0.5 mg/ml KLH was prepared in the same way. The presence of endotoxins in the immunization solutions was excluded by means of a limulus amebocyte lysate test (Lonza, Slough, U.K.), which was negative (data not shown). Groups of 5–10 mice were sensitized by i.p. injection with 100 μl immunization solution. Sham-sensitized mice received KLH alone in PBS and adjuvant according to the same schedule. Three weeks after the initial injection, mice were challenged i.p. with a second dose of Ag as before. The health status of animals was monitored and their condition scored according to the scheme of Morton and Griffiths (9).

Assessment of hypersensitivity reactions

The health status and body temperature of all mice was recorded before and immediately after challenge for up to 2 h. Body temperature readings were performed using a 1319 K-type rectal thermometer (TES, Taipei, Taiwan) and recorded once the digital display displayed a value for more than 10 s. Anaphylactic signs were evaluated for up to 1 h after the second challenge dose. Mice were culled if significant morbidity was observed. Cardiac puncture under terminal anesthesia was performed to recover blood from mice that had reached the end stage of the experiment or at the equivalent time point from mice from the control group that did not develop disease signs.

T cell proliferation assay

To assess whether the peptides represent T cell epitopes, lymphocytes from two mice sensitized with MP2–KLH were obtained from homogenized spleens using standard techniques. Single-cell suspensions were labeled with 10 μM CFSE (Molecular Probes, Eugene, OR) and incubated for 10

---

The Journal of Immunology

*Department of Infection, Immunity and Inflammation, University of Leicester, Leicester LE1 9HN, United Kingdom; †Department of Immunology, University College Medical School, Royal Free Hospital Campus, London NW3 2PF, United Kingdom; Department of Science and Technology for the Environment and Territory, University of Molise, Contrada Fonte Lappone, 86090 Pesche (IS), Italy

Received for publication July 13, 2010. Accepted for publication April 19, 2011.

This work was supported by the European Union (Grant Code QLK2-CT-1999-2387) and partly by laboratories partially funded by European Regional Development Fund No. 05567.

Address correspondence and reprint requests to Prof. Peter William Andrew, Department of Infection, Immunity and Inflammation, University of Leicester, POB 138, University Road, Leicester LE1 9HN, United Kingdom. E-mail address: pwa@le.ac.uk

Abbreviation used in this article: KLH, keyhole limpet hemocyanin.

Copyright © 2011 by The American Association of Immunologists, Inc. 0022-1767/11/$16.00

www.jimmunol.org/cgi/doi/10.4049/jimmunol.1002152
min at 37°C. After 10 min, the labeling was halted by centrifugation and resuspension in RPMI 1640 containing 10% HI-FCS, 2 mM L-glutamine, and 0.1 mM 2-mercaptoethanol. Cells were cultured in round-bottom 96-well plates (Corning Costar) at a density of 1.25 × 10⁵ in the media above containing either 1:10 control solution (PBS), 10 μg/ml anti-mouse CD3 (plate-bound) with 2 μg/ml anti-mouse CD28 and 20 ng/ml IL-2, or 50 μg/ml MP2–KLH for up to 72 h. Proliferation was determined by FACS as previously described (10).

Quantification of peptide-specific serum IgE and IgG

To monitor serum IgE and IgG Ab responses, tail vein blood was obtained on days 0, 14, 36, and 56 after initial sensitization. Sera were prepared and stored at −80°C. Levels of peptide-specific IgE and IgG were measured by using ELISA. To separate the IgE fraction of the serum from the IgG, the sample was first passed through a Protein G column. The elimination of the IgG fraction from the IgE preparation was confirmed using ELISA. Microtiter plates (Maxisorp; Nunc, Roskilde, Denmark) were coated with 5 μg/ml peptide–BSA conjugate in coating buffer, pH 9.6 (50 mM NaHCO₃, pH 9.6, 0.02% [w/v] Na₂CO₃), 100 μl per well overnight at 4°C. The next day, plates were blocked with blocking buffer (5% [w/v] nonfat dry milk, 0.05% [v/v] Tween 20 in PBS) for 1 h at 37°C and washed three times with washing buffer (50 mM Tris HCl pH 7.5, 150 mM NaCl, 0.05% [v/v] Tween 20). Mouse sera were diluted 1:100 in blocking buffer and 100 μl added to the wells and incubated for 2 h at 37°C with shaking. The plates were washed three times as before, and binding Abs were detected using alkaline phosphatase-conjugated goat anti-mouse IgE or IgG secondary Ab (Fc specific; Sigma), diluted 1:5000, and 1 mg/ml p-nitrophenyl phosphate (N9389; Sigma) dissolved in 1 M diethanolamine pH 9.8 and 0.5 M MgCl₂. Absorbance was read at 405 nm after 1 h at 37°C, and titer was determined as the dilution factor required to obtain absorbance greater than double the background absorbance. All analyses were performed in triplicate.

Quantification of serum histamine levels

Whole-blood samples were collected by cardiac puncture at the end of the experiment and were analyzed for histamine content using a radioenzymic assay as described by Sanmugalingam et al. (11). Cleared serum was obtained by centrifugation of clotted blood for 10 min at 8000 × g.

Statistical analysis

Survival data of mice immunized with the peptide–KLH conjugates and those that received KLH alone were analyzed by the Kaplan–Meier test using GraphPad Prism 5 software. Condition scores were used in place of survival data for groups V1–V6 as control mice were culled alongside experimental groups to obtain matched control serum for use in further tests. A Student’s t test was used for group analyses of IgG and IgE concentrations, condition scores, body temperature, and histamine concentrations. Significant statistical differences were noted as p < 0.05.

Results

Induction of immediate anaphylactic reaction by peptide vaccines upon booster dosing

Mice that received a booster dose of MP2, MP14, MP17, and MP18 peptide–KLH conjugates developed signs of anaphylaxis and experienced a high mortality rate (Fig. 1A). In contrast, all mice in the control (KLH) group and the groups that received the other three peptide–KLH conjugates (MP10, MP13, and MP15) survived. When anaphylaxis developed, signs were seen within 10 min of administering a second dose, and mice became severely lethargic. Cardiovascular collapse occurred after 15 min as seen by increased difficulty obtaining a tail-bleed. Signs of abdominal

---

**FIGURE 1.** Peptide vaccines eliciting an immediate systemic reaction. One hundred microliters of the indicated peptides and control KLH alone were injected i.p. 36 d after the first injection. A, Percentage mortality of mice administered a booster dose of peptide vaccines, measured at 2 h after the injection. B, The IgG (black bars) and IgE (open bars) Ab response in mice immunized with one dose of MP2–KLH on day 0, n = 10. Serum was obtained by tail bleed on days 14, 36, and 56 postimmunization. The reactivity of no-sera control is also shown (shaded bars). The median serum dilution (1:X) performed was 500 ± 755 for IgG and 64 ± 227 for IgE. KLH-treated mice did not show any increase in serum IgG or IgE concentrations (data not shown). The graph shows a significant increase in serum Ig levels. C, The whole-blood histamine levels in mice immunized with MP2–KLH after the injection of a second dose of the peptide. Error bars represent the SEM. The graph shows a significantly higher whole-blood histamine concentration in mice that received two doses of peptide MP2–KLH compared with that in mice that received the immunization solution minus the Ag (n = 2–5). *p < 0.05.
FIGURE 2. The proliferative capacity of CFSE-labeled splenocytes obtained from mice sensitized with MP2–KLH. Cells were exposed to anti-CD3, anti-CD28 and IL-2, MP2–KLH, or control. With each division, the fluorescent intensity diminishes by half as the label is inherited by daughter cells. A. Representative samples from a flow cytometric assay for T cell proliferation after stimulation with allergenic peptide MP2–KLH. Division was apparent at 24 h postlabeling and pronounced by 72 h. B. T cell proliferation over time. Cells exposed to mouse anti-CD3, anti-CD28, and IL-2 showed the largest proportion of highly divided cells after 72 h (closed squares). Cells stimulated with MP2–KLH (closed circles) showed substantially higher levels of proliferation as early as 5 h compared with those of cells that received the control solution of PBS (open circles). The mean percentage (+SEM) of proliferated T cells is shown (n = 2).

SERUM IgE and histamine levels increase in mice immunized with peptide MP2

To investigate the nature of the anaphylaxis, naive mice were given one dose of MP2–KLH, and serum was collected 14, 36, and 56 d postimmunization. Analysis of these sera (by ELISA coating with MP2–BSA) showed that immunization with MP2–KLH caused a significant increase in anti-MP2 serum IgE after 36 d (p < 0.05) (Fig. 1B). Anti-MP2 serum IgG concentration increased 10-fold (in OD) over the same time frame. Directly after the administration of the second dose of MP2–KLH, the whole-blood histamine levels rose significantly (p < 0.05) compared with those of the control group (Fig. 1C). The highest histamine concentration recorded peaked at 11600 nmol/l 30 min after challenge compared with 3600 nmol/l in mice that received the control solution of PBS (open circles). The mean percentage (+SEM) of proliferated T cells is shown (n = 2).

The KLH carrier was not a factor in the development of anaphylaxis

To investigate whether the carrier molecule influenced the immunological response to MP2, mice were immunized with unconjugated MP2, followed by a booster dose of unconjugated MP2 by the same schedule as for MP2–KLH. Again, directly after the administration of the second dose of unconjugated peptide, mice experienced severe anaphylaxis and a high mortality rate (Fig. 1A).

Lymphocytes obtained from mice sensitized with MP2–KLH showed a marked increase in proliferation after secondary exposure to the peptide compared with that of the negative control. Flow cytometric analyses (Fig. 2A) indicated that 5 h after stimulation with the peptide, the rate of cell division was seven times higher (4.4 ± 1.1%) than that of those exposed to control solution (0.61 ± 0.2%). This difference was even more pronounced by 24 h and 72 h (Fig. 2B). These results, along with data supporting the development of anaphylaxis from immunization with unconjugated peptide (see earlier), indicate that MP2 is a T cell epitope.

TABLE I. Sequence alignment of peptides using ClustalW

<table>
<thead>
<tr>
<th>Peptide</th>
<th>Sequence Alignment</th>
<th>Overall Identity to MP2 (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>MP2</td>
<td>-NCPRUIPEDTY-</td>
<td>-</td>
</tr>
<tr>
<td>MP17</td>
<td>-DMTAVIDPR-</td>
<td>50</td>
</tr>
<tr>
<td>MP18</td>
<td>-MPOFWPPDEEV-</td>
<td>25</td>
</tr>
<tr>
<td>MP13</td>
<td>-DNPALFDDML-</td>
<td>25</td>
</tr>
<tr>
<td>MP14</td>
<td>-WEDNPRIELN-</td>
<td>16</td>
</tr>
<tr>
<td>MP10</td>
<td>-EIVPWPDKERS-</td>
<td>16</td>
</tr>
<tr>
<td>MP15</td>
<td>-Y-SLITPEPTDP-</td>
<td>8</td>
</tr>
</tbody>
</table>

Each peptide was placed based on its identity to the adjacent peptides by ClustalW.

Scheme: underlined font, small and hydrophobic (including aromatic) residues; boldface font, acidic residues; black, basic, hydroxyl, amine, and Q residues.

The hyphen (-) represents a space created by ClustalW alignment.
FIGURE 4. A, The human mAb Db3G9 (black bars) binds to peptide conjugates of both MP2 and MP2m equally \((n = 2)\). The reactivity of no-sera control is also shown (shaded bars). B, The total IgG Abs induced by immunization with peptides MP2 (open bars) or MP2m (black bars) bind to both peptides but, notably, more anti-MP2 Abs bind to MP2 than to MP2m \((n = 3)\). The reactivity of no-sera control is also shown (shaded bars).

Substituting the negatively charged acidic residues in MP2 eliminated the onset of anaphylaxis in mice

Sequence alignment analysis of all peptides revealed a high frequency of clustered regions of acidic amino acid residues (ED) that were not found in the nonallergenic peptides (see Fig. 3, Table I). To determine whether anaphylaxis was due to the presence of a high proportion of negatively charged acidic amino acid residues, a further peptide, MP2m, was synthesized. In this peptide, the glutamic acid (E) and aspartic acid (D) residues present in MP2 were replaced by glutamine (Q) and asparagine (N), respectively. This did not affect the binding of mAb Db3G9, which displayed an equal level of binding \((p > 0.05)\) to MP2 and MP2m in ELISA (Fig. 4A).

Six groups of five mice \((V1–V6)\) were exposed to different immunization schedules with MP2, MP2m, or serotype 6B pneumococcal polysaccharide (see Table II). Mice immunized with one dose of MP2m displayed significantly \((p < 0.05)\) lower serum IgE titer \((10 ± 0)\) than that of mice immunized with one dose of MP2 \((400 ± 100)\). Mice immunized with MP2 and later boosted with MP2m \((group V2)\) had greatly reduced morbidity (condition scores, Table II) compared with those of mice boosted with MP2m \((group V1)\) and showed a quicker recovery. To confirm the anaphylaxis, the serum histamine and body temperatures were recorded. Table II shows that those mice that received two doses of MP2 \((group V1)\) developed a significantly higher \((p < 0.05)\) histamine concentration \((664 ± 224 \text{ nmol/l})\) and a substantially greater drop in body temperature \((-2.8 ± 0.3°C)\) compared with those of mice immunized with either MP2 or MP2m and boosted with MP2m \((groups V2 and V3, respectively)\) and those of mice immunized with MP2m and boosted with MP2 \((group V5)\). The values for histamine concentration and body temperature mirror each other in their effects, providing confirmatory results.

Using ELISA, we compared the ability of the Abs raised from immunization with MP2–KLH and MP2m–KLH to bind MP2–BSA and MP2m–BSA (Fig. 4B). We found that the Abs bound to both peptides but, notably, less anti-MP2 Abs bound to MP2m than to MP2. However, anti-MP2m Abs do cross-react and are shown to bind MP2 and MP2m equally. This indicates that an immunogenic epitope of MP2 was not present in MP2m and that an important proportion of the Ab response to MP2 was generated against this epitope.

Discussion

In this study, we have shown that novel peptide vaccines can induce systemic anaphylaxis in mice. In a mouse model there are two pathways that can lead to systemic anaphylaxis: a classical pathway mediated by IgE, FcεRI, mast cells, histamine, and platelet-activating factor and an alternative pathway mediated by IgG, FcyRIII, macrophages, and platelet-activating factor (12). We have shown that immunization with a peptide conjugate vaccine induced a rapid, fatal anaphylaxis, provoked by the production of peptide-specific IgE and the release of histamine after booster injection. This suggests the response is primarily IgE-mast cell dependent, although we cannot exclude a role also for IgG.

Despite all seven peptides being recognized by the mAb Db3G9, only four provoked an allergic reaction in vivo upon booster injection. This indicated that it was not shape but the amino acid sequence that was critical in determining allergenicity. Using a sequence alignment tool, significant similarities between the allergenic peptides \((MP2, MP14, MP17, and MP18)\) were identified (Table I). Similarities included the pairing of tryptophan and proline residues \((PW, WP, or WGP)\), which are highly important to the structure of small peptides, although they also were found in the nonallergenic peptides MP10 and MP13. Proline residues are known to create important structural features such as an l-shaped bend that may serve a role in orienting important residues for Ab presentation (6). Another similarity was the occurrence of

<table>
<thead>
<tr>
<th>Group</th>
<th>First Dose</th>
<th>Median Ig Titer (1:(X) ± IQR^c)</th>
<th>Second Dose</th>
<th>Histamine ((\text{nmol/l})^b)</th>
<th>Condition Score(^d)</th>
<th>Change in Body Temperature (°C^f)</th>
</tr>
</thead>
<tbody>
<tr>
<td>V1</td>
<td>MP2–KLH</td>
<td>500 ± 100</td>
<td>250 ± 150</td>
<td>MP2–KLH</td>
<td>644.4 ± 224.5</td>
<td>23.4 ± 1.9</td>
</tr>
<tr>
<td>V2</td>
<td>MP2–KLH</td>
<td>400 ± 0</td>
<td>160 ± 60</td>
<td>MP2m–KLH</td>
<td>110.6 ± 10.9*</td>
<td>2.4 ± 1.7***</td>
</tr>
<tr>
<td>V3</td>
<td>MP2m–KLH</td>
<td>300 ± 100</td>
<td>10 ± 0*</td>
<td>MP2m–KLH</td>
<td>98.4 ± 24.0*</td>
<td>0.6 ± 0.6***</td>
</tr>
<tr>
<td>V4</td>
<td>MP2–KLH</td>
<td>400 ± 100</td>
<td>150 ± 60</td>
<td>6B PnCPS</td>
<td>64.4 ± 23.8*</td>
<td>0.6 ± 0.6***</td>
</tr>
<tr>
<td>V5</td>
<td>MP2m–KLH</td>
<td>200 ± 140*</td>
<td>10 ± 0*</td>
<td>MP2–KLH</td>
<td>111.6 ± 17.6*</td>
<td>1.2 ± 0.7***</td>
</tr>
<tr>
<td>V6</td>
<td>6B PnCPS</td>
<td>64 ± 36**</td>
<td>10 ± 0*</td>
<td>MP2–KLH</td>
<td>85.8 ± 32.5*</td>
<td>0.6 ± 0.6***</td>
</tr>
</tbody>
</table>

All values in last three columns represent the mean ± SEM \((n = 5)\). Significant differences to V1 are in boldface font.

\(^a\)Serum Ig response to primary Ag 20 d after first dose. This is displayed as the dilution factor required to obtain greater than double the background absorbance in ELISA.

\(^b\)The mean serum histamine concentration and change in body temperature in mice 30 min after challenge with a second dose of Ag \((n = 5)\).

\(^c\)The mean cumulative condition score as outlined by Morton et al. (9).

\(p < 0.05, **p < 0.01, ***p < 0.001\) (significant difference of groups V2–V6 compared with group V1, which received two doses of MP2–KLH).

IQR, interquartile range; PnCPS, pneumococcal polysaccharide.
glutamic acid (E) and aspartic acid (D) residues found as pairs or in close proximity to each other (i.e., ED, EWD, and DE). This feature was absent from all nonallergenic peptides, including peptide MP13, which otherwise showed a high level of similarity to peptide MP2. This feature has also been reported for other known allergens, with several researchers identifying IgE epitopes that contain a higher proportion of charged amino acids (D, E, and K) compared with IgG epitopes (13, 14). Furthermore, some allergens have been shown to lose their ability to bind IgE when these residues are mutated or removed (14, 15). The sequence of MP2 was chosen for further investigation because this peptide contains only one E and D residue and therefore would be the simplest to manipulate without markedly affecting the three-dimensional structure of the epitope. The removal or substitution of the motif that contained tryptophan or proline, however, would dramatically alter their conformation, and so removing these could confound the interpretation of the results. A modified MP2 peptide (designated MP2m) was created where the E and D residues were substituted by their uncharged derivatives glutamine (Q) and asparagine (N), respectively, which peptide contains a terminal amide group in place of a carboxylate group. These substitutions were expected to alter the hydrogen bonding potential of the peptide but maintain its overall size. In fact, the human mAb Db3G9 bound to MP2 and MP2m equally, and therefore, its overall shape and the number of peptides per KLH were not affected by this alteration.

Despite extensive efforts, the mechanism behind why some proteins provoke an allergic response remains unknown, but it is thought that allergens share particular molecular motifs. The predicted primary structures of some major allergens have indicated that most are relatively small (<70 kDa), negatively charged proteins with low hydrophobicity and high stability. Another feature is the high proportion of negatively charged (aspartic acid, glutamic acid), positively charged (lysine), and polar (serine, threonine, tyrosine, and cysteine) residues (14, 16).

Given that Ab Fab regions usually recognize only 8–15 aa, it is likely that the structure of small exposed epitopes and not the whole protein is important in provoking an allergenic immune response. Indeed, allergenic epitopes have been shown to be as short as five amino acids (14), and single amino acid substitutions have abolished Ab binding altogether (16). Such substitutions may markedly affect the three-dimensional structure of the epitope and thus reduce Ab reactivity. In addition, Ab binding sites can span several domains, with the amino acids involved being found in different parts of the linear sequence (15, 16). It is therefore difficult to predict exactly which epitopes provoke an allergic response, as eliminating or substituting particular amino acids could alter not only the epitope but also the overall structure of the protein.

This study has shown that immunizing mice with MP2m eliminated the allergic effects seen with peptide MP2. Mice immunized with MP2m displayed significantly reduced serum IgE levels compared with those of mice that received MP2. Furthermore, the raised serum histamine levels and decreased body temperature seen in anaphylaxis were significantly lower in mice immunized with two doses of MP2m. Notably, mice immunized with the allergenic MP2 did not suffer anaphylaxis upon receiving a subsequent dose of MP2m. Similarly, a primary dose of MP2m did not result in anaphylaxis upon boost with MP2. This suggested that the IgE-stimulating Abs produced in response to immunization with MP2 were raised to the region of the peptide containing the acidic amino acids. This effect is also commented on by another group who suggested that the mast cell activation and skin reaction-inducing capacity of three peptide cancer vaccines was correlated to their basic (electric) charge (17).

The mechanism behind these events is still unclear, but in our study, the elimination of anaphylaxis may be explained by the genetic restriction of the MHC class II, which plays an important role in the immune recognition of proteins and the development of Ab responses (18). Limited individual variation means that the peptides that bind MHC class II molecules are defined by anchor residues within the sequence (19). For example, peptides with sequences such as xWPxxxED, WxPxxxED, or xDExxxPWx may be recognized by the same MHC molecule, as the peptides share negatively charged residues located four positions away from a proline residue. Thus, it is not the whole sequence that is important to allergen determination but the positions of key residues (19). In this study, which is strengthened by the use of outbred mice, the substitution of ED may have led to altered binding to a particular MHC class II molecule, which in turn eliminated the allergic response. In addition, as MP2m did not induce anaphylaxis in MP2-immunized mice, it appears that the acidic residues are essential for IgE recognition. It seems unlikely that the response involves the cross-linking of IgE/FceRI receptors on mast cells by multiple peptides displayed on the carrier KLH molecule because anaphylaxis occurred with MP2 in the absence of KLH. Further work into why peptide MP2 provoked a Th2 response that was not seen with MP2m may help unravel important mechanisms in the induction of an allergic response.

In summary, this study has highlighted important new features that determine whether a peptide is allergenic and has shown certain amino acid motifs in a 12-aa sequence to be responsible for an Ab class switch to IgE. To date, there are no published examples of anaphylaxis due to peptide mimics of polysaccharide. The anaphylactic response in our study consisted of the characteristic clinical features described in mice (20) coupled with the detection of a significantly higher concentration of histamine in the serum of mice immunized with MP2m compared with that in the serum of control mice. With the current characterization of allergenic epitopes, this allergic response would not have been predicted. However, there was a robust IgE response after immunization with MP2, and this could be used as a surrogate for allergenic potential. We would therefore recommend that an assay for allergic potential be routine in the development of peptide vaccines, and that skin-prick testing in humans or RAST to detect circulating peptide-specific IgE be undertaken before administration of a second dose of vaccine. As a greater number of peptide Ags become available for use in experimental therapies, it is important to establish the safety of individual peptide mixtures. This knowledge may help prevent the induction of anaphylaxis in humans given peptide-based therapy.

Acknowledgments
We thank Dr. Santhosh Mukundan and Sarah Glenn for technical support.

Disclosures
The authors have no financial conflicts of interest.

References


