Peptide-based ELISAs are not sensitive and specific enough to detect muscarinic receptor type 3 autoantibodies in serum from patients with Sjogren's syndrome
Roescher, N.; Kingman, A.; Shirota, Y.; Chiorini, J.A.; Illei, G.G.

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ABSTRACT

Objectives The detection of autoantibodies to the muscarinic receptor type 3 (M3R) in the serum of patients with Sjögren’s syndrome (SS) by ELISA is controversial. A study was undertaken to test whether modification of M3R peptides could enhance the antigenicity and increase the detection of specific antibodies using an ELISA.

Methods A series of controlled ELISAs was performed with serum from 71 patients with SS and 37 healthy volunteers (HV) on linear, citrullinated and/or cyclised and multi-antigenic peptides (MAP) of the three extracellular M3R loops to detect specific binding.

Results Significant differences (p<0.05) in optical density (OD) between serum from patients and HV were detected for a cyclised loop 1-derived peptide and the negative control peptide. Furthermore, there were no statistically significant differences between the frequency of positive patients (defined as OD >2SDs above the mean of the HV) and HV on any of the peptides tested.

Conclusions Binding of serum from patients with SS to M3R-derived peptides does not differ from binding to a control peptide in an ELISA and no significant binding to M3R-derived peptides was found in the serum from individual patients compared with HV. These data suggest that peptide-based ELISAs are not sufficiently sensitive and/or specific to detect anti-MR3 autoantibodies.

The frequency of patients positive for muscarinic receptor type 3 (M3R) autoantibodies, thought to be a key pathogenic factor in the exocrine dysfunction in patients with Sjögren’s syndrome (SS), is reported to vary from 0% up to nearly 100% when detected by a linear peptide-based ELISA. Low anti-M3R titres, the inability of an antibody to recognise a linear peptide and/or the choice of epitope may all have a role in the variability observed. We tested sera from patients with SS and healthy volunteers (HV) on previously reported M3R peptides, as well as on cyclised, citrullinated and multi-antigenic peptide (MAP) versions. Serum from 79 patients (primary SS, mean age 49.5 years, 89% women) and 37 HV (mean age 45.5, 67% women) was obtained at our clinic and at the NIH blood bank and stored at −80°C until use. Several ELISA conditions were tested and optimised using a MAP-8 Ro peptide (LGEMPLTALLRLGKMT). Ten M3R-derived peptides (three linear loop 2 peptides, three cyclised loop 1, 2 or 3 peptides, two MAP-8 loop 2 peptides, one citrullinated loop 2 and one citrullinated and cyclised loop 2) and a SGSG control peptide were screened in a series of pilot ELISAs with 15 HV and 37 patients to determine the most antigenic peptides. Receiver operating characteristics (ROC) graphs were made and the area under the curve (AUC) was determined (for sequence see figure 1). These and the control SGSG peptide were re assayed in four tightly controlled ELISAs on four consecutive days with serum from all patients and HV (figure 1A). The analysis of variance results showed that the between-day variation in optical density (OD) scores was significant for all peptides (p≤0.002) except the SGSG peptide (p=0.207). Significant differences between HV and patients were found for the SGSG (p=0.037) and the loop 1 peptide (p=0.025). To determine sensitivity and specificity of each peptide, ROC graphs were made and the AUC was calculated (figure 1B). None of the AUC were statistically significant (p>0.100). Individual autoantibody positivity was defined as an average OD >2SDs of the mean of the HV, assuming a normal distribution in this group (table 1).

The present study demonstrates a poor reproducibility of M3R peptide-based ELISAs. The average patient OD of loop 1 was significantly different from the HV, but this was also observed with the SGSG peptide, and there were no differences in the number of positive individuals between the two groups.

There are some technical reasons that may account for these negative results, including microstructural changes in the tested peptides over time and inconsistency in binding of the peptide to the microtitre plate. Other reasons may be that M3R autoantibodies are reactive with other non-tested epitopes, only recognise an epitope in its native form or peptide:antibody interactions may be too weak/strong for detection.

<table>
<thead>
<tr>
<th>Peptide</th>
<th>HV, % of group (n)</th>
<th>P, % of group (n)</th>
<th>p Value</th>
</tr>
</thead>
<tbody>
<tr>
<td>SGSG</td>
<td>2.7 (1)</td>
<td>9.9 (7)</td>
<td>0.178</td>
</tr>
<tr>
<td>KRAI</td>
<td>5.4 (2)</td>
<td>7.0 (5)</td>
<td>0.743</td>
</tr>
<tr>
<td>MAP KRAI</td>
<td>2.7 (1)</td>
<td>7.0 (5)</td>
<td>0.350</td>
</tr>
<tr>
<td>Kc_c</td>
<td>2.7 (1)</td>
<td>5.6 (4)</td>
<td>0.492</td>
</tr>
<tr>
<td>Loop 1</td>
<td>2.7 (1)</td>
<td>4.2 (3)</td>
<td>0.691</td>
</tr>
</tbody>
</table>

The p values represent the significance of the corresponding two-independent sample t-test.
unstable. Moreover, the lack of a reliable positive control may have limited our efforts to optimise conditions for M3R antibodies.

Our data indicate that relatively simple peptide-based ELISAs are not sufficiently sensitive and specific to detect the putative anti-M3R autoantibodies. Functional assays using native receptors are more elaborate but may prove more reliable. Future studies should therefore use controls which have tested positive in functional assays. Establishing a reference set of such sera which could be shared among investigators would significantly aid in clarifying the controversies about the detection of anti-M3R antibodies.

N Roescher, A Kingman, Y Shiroti, J A Chiorini, G G Illei

Correspondence to: Nienke Roescher, NIH/NIDCR/MPTB, 10 Center Drive, Building 10, Room 1A21, Bethesda, MD 20892-1190, USA; roeschern@mail.nih.gov

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