Cut off selection in anti-acetylcholine receptor antibody determination

We have read the interesting paper by Somnier,1 and want to make an observation about cut off selection in anti-acetylcholine receptor antibody (AChR-Ab) determination. The value 0.5 nmol/l is chosen in an arbitrary way, and so assay specificity and sensitivity are also arbitrarily determined.2 Receiver operating characteristic (ROC) analysis, combined with information theory, is the most exact approach to cut off selection in a continuous variable.3 Because the author mentions that AChR-Ab titres of myasthenic patients have a log normal distribution (although no Kolmogorov-Smirnov goodness-of-fit test is reported) and, as control serum sample titres probably also follow a log normal distribution, a ROC analysis could be performed assuming the binormal hypothesis (two overlapping normal curves, one from control patients and other from myasthenic patients). In this way a maximum likelihood estimation could be obtained to plot a ROC curve4 (figure A). The point nearest (0,1) is the one where sensitivity and specificity are optimal, therefore the one for a proper cut off selection. A further and more accurate cut off selection could be achieved by taking into account the prevalence of myasthenia gravis in the sample as, according to Bayes' theorem, post-test probability is dependent on prior-probability (prevalence). Using ROC analysis and Bayes' theorem, the point that should be chosen as cut off is the one that offers more information at the prevalence of the sample6 (figure B).

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1 Somnier FE. Clinical implementation of anti-acetylcholine receptor antibodies. J Neurol Neurosurg Psychiatry 1993;56:496-504.

Somnier replies: The cut off selection (0.5 nmol/l) was based on a 10 year experience consisting of analysis of more than 3000 serum samples from 366 patients with myasthenia gravis (MG) and 540 control serum samples from healthy individuals and patients with neuromuscular diseases other than MG. The choice of cut off point was made on account of these experimental data and with the objective of obtaining a maximum of specificity without undue loss of sensitivity. The cut off point was certainly not chosen in an arbitrary way.

Anti-acetylcholine receptor (AChR) antibody titres of patients with MG appear to have a normal log distribution.1 The goodness-of-fit was 2p > 0.44 (r2 square test) and 2p > 0.38 (Kolmogorov-Smirnov test). In control serum samples, the distribution of titres (linear estimates and logarithmic transformed values) has a negative skewness. Provided that this is not a disqualifying feature, a receiver operating characteristic (ROC) analysis assuming the binormal hypothesis2 could be performed. Figure 1 shows the leftmost section of the normal deviate of the ROC curve, δM (delta mean of normalized curves) = −5.6189 and S (slope) = 3.518. The location of various cut off points from 0.1 to 1.0 nmol/l is marked on the graph, enabling the reader to estimate the corresponding Z scores. Figure 2 shows the cut off points for maximum information as a function of the prevalence of MG serum samples. As 0.10–0.20 is the most likely occurrence in relation to a centralised laboratory, the ROC analysis confirms the choice of 0.5 nmol/l as the most appropriate cut off point for a diagnostic application of radioimmunossay. The table shows the ratio of true and false positive samples as a function of cut off point. The theoretical (ROC) analysis and the observed true positive ratios are in agreement, but false positivity appears to be underrated by the ROC analysis. Consequently, ROC analysis does not add any new feature to the clinical implementation of the anti-AChR antibodies RIA.

1 Somnier FE. Clinical implementation of anti-acetylcholine receptor antibodies. J Neurol Neurosurg Psychiatry 1993;56:496-504.

Figure 1 Normal deviate of ROC curve. Anti-acetylcholine receptor antibodies radioimmunoassay.

Figure 2 Maximum information. Cut off point prevalence.

### Table 1: Anti-AChR antibodies radioimmunoassay

<table>
<thead>
<tr>
<th>Cut-off (nmol/l)</th>
<th>Theoretical true positive</th>
<th>Observed true positive</th>
<th>Theoretical false positive</th>
<th>Observed false positive</th>
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</thead>
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<tr>
<td>0.3</td>
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<td>0.5</td>
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<td>0.890</td>
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<td>1.0</td>
<td>0.842</td>
<td>0.871</td>
<td>0.003</td>
<td>0.000</td>
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