Chromosome studies in systemic sclerosis with consideration of antibodies to topoisomerase I

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Abstract
Chromosome studies were performed on 11 patients with systemic sclerosis and on 35 control subjects. Nine patients with systemic sclerosis were positive for antibodies to topoisomerase I and two were negative. Of the 1100 metaphases from these 11 patients, 171 (15-5%) had chromosome breaks, compared with 61 of 3500 (1-7%) metaphases from normal control subjects. There were no statistically significant differences in the numbers of chromosome breaks between men and women. The most common fragile site in patients with systemic sclerosis was at 3p14. The karyotype of all patients was normal. Chromosome breaks did not correlate with the presence of antibodies to topoisomerase I.

Patients and methods

PATIENTS WITH SYSTEMIC SCLEROSIS
Eleven patients with systemic sclerosis were identified as having diffuse scleroderma with the skin of the proximal portions of the extremities and trunk, or both, affected, as well as fingers, hands, and face (classic scleroderma), according to the preliminary criteria for the diagnosis of systemic sclerosis. The median age of the 11 patients with scleroderma was 55 years, ranging from 32 to 69 years and the group consisted of six men and five women. None of the patients with systemic sclerosis had received cytotoxic drugs or other drugs known to induce chromosomal changes before the study. Similar studies were performed on 35 control subjects recruited from the hospital and laboratory staff. The control subjects were matched for age and sex. The median age of the 35 control subjects was 57 years, ranging from 28 to 71 years, and there were 18 men and 17 women.

CHROMOSOME ANALYSIS
A 10 ml sample of blood was obtained from each patient and control subject and three blood cultures were established by injecting 0.5 ml of whole blood into 10 ml of culture medium consisting of M medium (modified F10 medium without thymidine, folic acid, and hypoxanthine, Gibco formula 785227) with 4.5% fetal calf serum, phytohaemagglutinin, and PSN (Gibco antibiotic mixture of penicillin, streptomycin, and neomycin). The cultures were incubated for four days, treated with colcemid (0.1 µg/ml) and 0.0075 M hypotonic potassium chloride for 12 minutes, and fixed in a minimum of three changes of methanol/glacial acetic acid (3:1) fixative. Cells in fresh fixative were dropped onto a clean slide. The slides were air dried and stained with Giemsa. All cultures were handled by the same technician. A minimum of 100 G banded metaphases from each subject was selected at random and photographed under 1000× magnification and scored for structural chromosomal abnormalities.

ANTIBODY TO TOPOISOMERASE I
All samples were screened for antibody to topoisomerase I using a previously described technique. Briefly, DNA topoisomerase I was purified from calf thymus as an antigen. Serum samples from patients and control subjects were diluted 1:200 and screened for the antibody to topoisomerase I by an enzyme linked immunosorbent assay (ELISA).
### Table 1. Comparison of chromosome breakage in patients with systemic sclerosis and normal control subjects

<table>
<thead>
<tr>
<th>Subject</th>
<th>No of metaphases</th>
<th>Chromosome breakage</th>
<th>p Value*</th>
<th>Breakage at 3p14</th>
<th>p Value†</th>
</tr>
</thead>
<tbody>
<tr>
<td>Systemic sclerosis (n=11)</td>
<td>1100</td>
<td>171</td>
<td>&lt;0·001</td>
<td>62</td>
<td>&lt;0·001</td>
</tr>
<tr>
<td>Normal (n=35)</td>
<td>3500</td>
<td>61</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

*p Value based on χ² test between all chromosome breakages in patients with systemic sclerosis and normal control subjects. †p Value based on χ² test breakage at 3p14 between patients with systemic sclerosis and normal control subjects.

### Results

Figure 1 gives an example of a metaphase showing chromosome breakage. The results of the cytogenetic observations in the 11 patients with systemic sclerosis compared with those in the 35 normal control subjects are given in Table 1. Of the 1100 metaphases from these 11 patients, 171 (15·5%) of the metaphases had chromosome breaks, compared with 61 (1·7%) of the 3500 metaphases from normal control subjects. The overall occurrence of structural chromosome breakages was significantly increased (p<0·001) in patients with systemic sclerosis. Chromosome breaks were found in 88 of 600 metaphases from six male patients and 83 of 500 metaphases from five female patients. There were no statistically significant differences between the numbers of chromosome breaks in male and female patients.

The distribution of chromosome breakage sites in all patients was analysed. The numbers of lesions at different chromosomal sites are shown in Table 2. Of a total 171 chromosome breaks from patients with systemic sclerosis, 62 breaks were at 3p14, compared with only seven (0·2%) of 3500 metaphases from normal control subjects. Chromosome breaks at 3p14 were 28 times more common in patients with systemic sclerosis than in normal control subjects. The chromosome region at 3p14 was the most common fragile site in patients with systemic sclerosis.

Figure 2 shows the relation of the antibody to topoisomerase I and chromosome breaks in patients with systemic sclerosis. The cut off points for the significantly increased levels of antibodies to topoisomerase I were determined on the basis of 21 healthy control subjects. The normal mean (SD) value of the absorbance was 0·406 (0·173). Values above 0·925 (normal value plus three SD) were regarded as increased levels of antibodies to topoisomerase I. The occurrence of the antibody to topoisomerase I in our patients with systemic sclerosis was 8/11 (73%) by ELISA. No close correlation between the presence of the antibody to topoisomerase I and chromosome breaks in patients with systemic sclerosis was found (r=0·77, p>0·05).

### Discussion

Our data provide additional confirmation that there is an increased occurrence of chromosome breaks in patients with systemic sclerosis. Although chromosome breaks are common in patients with systemic sclerosis, the common chromosome regions prone to breakage in these patients have not previously been reported. We found that the most common fragile site in patients with systemic sclerosis is 3p14. Previously Smeets et al. reported that 3p14 was the region most prone to breakage in normal human chromosomes. The chromosome region at 3p14 seems to be the fragile site in normal subjects and in patients with systemic sclerosis. Several workers have suggested that fragile sites play a preferential part in the chromosome translocations that are associated with certain diseases. The clinical significance of the fragile site in patients with systemic sclerosis is not clear.

It has been suggested that patients with systemic sclerosis may possess a serum clastogen and could induce chromosome breakage. Although the serum factor has not been isolated or identified, its presence in patients' serum indicates that scleroderma associated chromosome breakage has a molecular basis. Topoisomerase I has been associated with chromosomal decondensation and genetic recombination. Patients with diffuse scleroderma often have antibodies to topoisomerase I and high titres of the antibody to topoisomerase I are mostly present in patients with diffuse scleroderma. An evaluation of the relations between antibodies to topoisomerase I and chromosome breakage was included in this study. We found that there was no correlation...
between the presence of antibodies to topoisomerase I and chromosome breakage in patients with diffuse scleroderma. The occurrence of chromosome abnormalities among our patients who were positive for antibodies to topoisomerase I were similar to those in patients who were negative for such antibodies. Thus antibodies to topoisomerase I in patients with systemic sclerosis do not appear to be the chromosome breakage factor. Powell et al reported that the anticientromere antibody which is also present in patients with a variant form of systemic sclerosis does not appear to have clastogenic properties in vitro. The biochemical nature of the breakage factors remain unclear at present and need further investigation.

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