Clonal analysis of three morphologically distinct lymphomas occurring in the same patient

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Abstract

Aims—To determine whether three morphologically distinct lymphoid tissues (mucosa associated lymphoid tissue (MALT) lymphoma, monocytoid B cell lymphoma, and large cell anaplastic lymphoma), which occurred in the same patient, were in fact three morphological variants of the same lymphoproliferative process.

Methods—Previously described methods of clonal analysis using the polymerase chain reaction (PCR) were used to determine the pattern of rearrangements of the immunoglobulin heavy chain gene and the T cell receptor β and T cell receptor γ chain genes in the three lymphomas.

Results—All three morphological entities had identical patterns of gene rearrangements.

Conclusions—This finding confirms the association between MALT lymphoma and monocytoid B cell lymphoma, and also provides evidence that large cell anaplastic lymphoma may not only arise de novo but may also be an end stage morphological picture in lymphoma progression.

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The occurrence in one patient of more than one morphologically distinct lymphoid neoplasm, either concurrently or sequentially, is well recognised. Many cases represent blast cell transformation occurring in a low grade lymphoma and resulting in a more aggressive or high grade tumour. This phenomenon was noted in 40% of cases of centroblastic/centrocytic lymphomas in a necropsy study undertaken by Lennerström. Transformation is usually to centroblastic lymphoma, although less frequently the high grade lymphoma is composed predominantly of bizarre, anaplastic cells, and very occasionally transformation to a B immunoblastic lymphoma is seen. Another well recognised entity is diffuse large cell lymphoma arising in patients with a previous history of chronic lymphocytic leukaemia (CLL). This phenomenon is known as Richter’s syndrome. The reported incidence in CLL is 3–10%.

In 1988 Cogliatti et al described a series of patients with monocytoid B cell lymphoma (MBCL). A subset of their study group (33%) also had concomitant low grade B cell lymphoma of mucosa associated lymphoid tissue (MALT). Two of these cases later developed into unspecified high grade lymphomas. Investigations to determine clonality were not undertaken in this series, but recent work on both Richter’s syndrome and on blast cell transformation of centrocytic/centroblastic lymphomas, using restriction enzyme analysis (Southern blotting) to detect immunoglobulin gene rearrangements, has shown a clonal relation between the initial low grade neoplasm and the subsequent high grade lymphoma.

Here, we report a case of a 75 year old woman who developed three morphologically distinct malignant lymphomas over an 11 year period. In spite of the range of histological appearances shown by these tumours (a primary gastric lymphoma, a low grade rectal MBCL, and most recently a Ki-1 positive large cell anaplastic lymphoma (LCAL) in her cervical nodes), gene rearrangement studies using PCR were highly suggestive that all three lesions were the same clone.

Case report

A 63 year old woman, presented in 1979 with a 10 month history of epigastric pain and weight loss. Endoscopy revealed a 5 cm in diameter ulcer on the lesser curve. After biopsy, a radical gastrectomy was performed. Histological examination showed a diffuse high grade gastric lymphoma. She received adjuvant chemotherapy and subsequently abdominal radiotherapy.

She remained well until October 1990 when she presented again, this time with a short history of anorexia and general malaise. A rectal polyp was discovered, which on histology, was shown to be a monocytoid B cell lymphoma. Coincidentally she developed rapidly enlarging right upper cervical and left supraclavicular lymphadenopathy. A biopsy specimen of the supraclavicular node revealed a large cell anaplastic lymphoma. A computed tomography scan also revealed lymphadenopathy in the region of the coeliac axis, but a bone marrow examination was normal. From January 1991 to August 1991 she received seven courses of combination chemotherapy. She responded extremely well to treatment, but she was left with a small mass in the cervical chain that was cytologically negative for malignant cells. This node continued to regress in size during follow up.

Pathological findings

Gastric lesion (tumour 1); March 1979

A total gastrectomy specimen with attached greater omentum, spleen, and part of the pancreas was received fixed in formalin. Opening the stomach revealed a superficially ulcerated...
Figure 1  High grade gastric lymphoma displaying loosely cohesive sheets of large cells with a coarsely granular chromatin pattern and frequent mitoses.

Figure 2  Colonic polyp containing a mixture of cell types, including an important component of uniform larger neoplastic mononuclear cells with pale cytoplasm and indented nuclei.

Figure 3  Cervical lymph node replaced by large cell anaplastic lymphoma. Numerous bizarre cells, including multinuclear forms, are present.

thickened area 70 mm × 30 mm on the greater curve. Twelve lymph nodes were dissected off the specimen. Histological examination (fig 1) showed a diffuse high grade primary gastric lymphoma. The central part of the tumour was formed by solid masses of large lymphoid cells with coarsely granular nuclear chromatin and a high mitotic rate. Numerous reactive looking germinal follicles in the adjacent lymphoid tissue suggested transformation from a pre-existing low grade MALT lymphoma. The neoplastic cells infiltrated downwards into, but not through, the muscularis propria. One of the 12 lymph nodes was also focally affected, with tumour cells growing in a perifollicular, or mantle-zone pattern. No Helicobacter pylori organisms were identified in the sections.

Colonic lesion (tumour 2); October 1990
A 13 mm intact pedunculated polyp measuring 20 mm in maximal cross-sectional dimension and 13 mm in height was received in formalin. The polyp had a homogeneous, firm, cream-coloured cut face. Haematoxylin and eosin stained sections showed a lymphoproliferative mass covered by somewhat attenuated, but otherwise normal, colonic mucosa. The lymphoid tissue had a vaguely nodular pattern and was composed of mixed large and small lymphoid cells interspersed with prominent aggregates of granular macrophages and scattered polymorphs and eosinophils. High power examination of the lymphoid nodules revealed small reaction centres, numerous plasma cells and, in places, an infiltrate of uniform neoplastic mononuclear cells with abundant pale cytoplasm and large, slightly indented, or reniform nuclei (fig 2). This infiltrate focally impinged on the reaction centres. These morphological features are those of a monocytoid B cell lymphoma.

Cervical lymph node (tumour 3); December 1990
Part of a lymph node measuring 20 × 18 × 5 mm was received in formalin. The node had a slightly nodular cut face. Sections showed effacement of nodal architecture by a diffuse infiltrate of large lymphoid cells with vesicular nuclei, prominent nucleoli, and abundant cytoplasm (fig 3). A smaller population of multinucleate and bizarre forms was also present. Mitotic rate was high and focal necrosis was noted. The morphological appearances were consistent with large cell anaplastic lymphoma.

Immunohistochemical staining was undertaken on all three lesions using a panel of antibodies (table 1).

<table>
<thead>
<tr>
<th>Antibody</th>
<th>Source</th>
<th>Action</th>
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<tbody>
<tr>
<td>UCHL1 (CD45RO)</td>
<td>Dakopatts a/s</td>
<td>Pan-T cell marker; also marks a subset of B cells</td>
</tr>
<tr>
<td>L26 (CD20)</td>
<td>Dakopatts a/s</td>
<td>Pan B cell marker</td>
</tr>
<tr>
<td>BerH2 (CD30)</td>
<td>Dakopatts a/s</td>
<td>Detects the Ki-1 antigen in activated lymphoid cells</td>
</tr>
<tr>
<td>Mac387</td>
<td>Dakopatts a/s</td>
<td>Histiocytic cell marker</td>
</tr>
<tr>
<td>EMA</td>
<td>Dakopatts a/s</td>
<td>Marker of epithelial cells and some subsets of lymphoid cells</td>
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The results of immunostaining are shown in table 2.

### CLONAL ANALYSIS

Sections (25 μm) were cut from tissue blocks on which previous microscopy had shown considerable numbers of neoplastic cells. These were then dewaxed in xylene and the xylene removed by washing twice in 100% ethanol. DNA was then extracted by proteolysis and phenol-chloroform extraction as described before.9

PCR analysis of the immunoglobulin heavy chain (IgH), T cell receptor β (TCR β) and T cell receptor γ (TCR γ) genes was then undertaken using reaction conditions, dimers, and temperatures, as described before.10–12 The reactions were run out in 10% polyacrylamide gel and viewed under ultraviolet light.

Using this protocol, monoclonal proliferations generate discrete bands whereas polyclonal (reactive) proliferations and non-lymphoid proliferations generate a diffuse "smear" on the gel. The results showed that all three tumours showed identically sized partial rearrangements of TCR β (65 and 90 base pairs) and of TCR γ (90 base pairs). No rearrangements of the IgH gene were found (fig 4).

### Discussion

The occurrence of three morphologically different lymphomas in one patient over an extended period of time, as described in our case report, is unusual. There are, however, numerous cases in which two morphologically different non-Hodgkin's lymphomas (NHL) or even NHL and Hodgkin's disease occurred in the same patient.13 Although this phenomenon hinted at a pathogenetic association, it was seldom possible to demonstrate monoclonyality until the advent of molecular genetic techniques. We undertook clonal analysis in this case to investigate whether all three tumours represented progressively less well differentiated morphological manifestations of the same lesion.

The initial gastric lymphoma presenting in 1979 displayed features to suggest that its origin was in a pre-existing low grade MALT lymphoma. Low grade lymphomas of MALT are characterised by the presence of reactive B cell follicles surrounded by neoplastic centrocyte-like cells. These neoplastic cells infiltrate diffusely, and also selectively invade epithelial structures to produce characteristic lymphoepithelial lesions.14 The presence of numerous reactive-looking germinal centres at the periphery of the gastric lymphoma and the phenomenon of follicular colonisation15 identified in our case point towards the probability of a pre-existing low grade MALT lymphoma.16 The gastric lymphoma could therefore reasonably have been expected to behave in a fairly indolent clinical manner.

The MBCL occurring 10 years later is representative of a tumour type which, although related to MALT lymphoma,17 is not imbued with the same tendency to "home" into sites with MALT. A lymphoma composed of monotypic monocyted B cells was first described by McGinn et al in 198517 and the term MBCL was coined in 1986 by Sheikhani et al.18 The co-existence of an MBCL with a concomitant MALT lymphoma is a recognised association1 but the clinical course of the former is somewhat more aggressive.

The Ki-1 positive LCAL was the final and least well differentiated morphological manifestation of the neoplasm. Most Ki-1 positive LCALs in the series reported by Chott et al in 1990 exhibited either T cell lineage (68%) or displayed no markers at all (22%).19 However, the less common B cell variants (10%) displayed an equally aggressive clinical course with an overall median survival time of only 13 months from the time of diagnosis. Similar
Clonal analysis of three morphologically distinct lymphomas

patterns of immunoreactivity to those demonstrated by Chott have also been shown in LCAL by other workers.\(^8\)\(^9\) Negative immunohistochemical staining on paraffin wax sections for epithelial membrane antigen (EMA) (as found in our case) was reported in 42% of the cases considered by Chott et al.\(^8\) and highlights the point that EMA positivity is not a prerequisite for diagnosis of LCAL. Notably, only about half of all Ki-1 positive LCALs are positive for leucocyte common antigen (LCA) on paraffin wax sections; and the phenotype LCA negative/EMA positive was found in 30% of the cases reported by Chott et al. This phenotype may lead to diagnostic confusion between LCAL and anaplastic carcinoma in some circumstances.

Recent studies by Bitter et al.\(^10\) and by other workers\(^11\) have suggested that there are two morphological groups of Ki-1 positive lymphomas and that the presence of a unique chromosomal abnormality, t(2;5)(p23;q35) defines the subset of Ki-1 positive lymphomas displaying classic anaplastic morphology on haematoxylin and eosin staining, as opposed to the second group which displayed morphological features of high grade non-Hodgkin's lymphoma without specific unifying features. In 1989 Chan et al also recognised two morphologically distinct variants of anaplastic Ki-1 positive large cell lymphoma but did not relate their observations to the karyotype of the tumour cells.\(^12\)

Most cases of Ki-1 positive LCAL develop de novo (primary type),\(^13\)\(^-\)\(^21\) but a small number arise secondary to other types of lymphoma\(^21\)\(^24\)\(^25\) and may represent the end point of a dedifferentiating tumour sequence. Mycosis fungoides, lymphophtihelioid lymphoma, and angioimmunoblastic-type T cell lymphoma have all shown this transformation in their later stages.

It is interesting to speculate that Ki-1 positive LCAL arising secondarily may represent a subset of the group of LCALs without the t(2;5)(p23;q35) translocation. This suggestion would be supported by the dual observations: firstly, that the prognosis of secondary LCALs is worse than that of primary LCALs;\(^21\) and secondly, that Ki-1 positive lymphomas lacking the specific translocation have a worse prognosis than those with it.\(^22\)

To identify the rearrangements, we used the polymerase chain reaction technique (PCR). As only formalin fixed, paraffin wax embedded tissue was available for investigation, this study would not have been possible using the more conventional methodology of restriction enzyme analysis (such as Southern blotting). PCR is particularly suitable when analysing blocked archival material.\(^26\)

It may seem somewhat paradoxical, at first sight, that all three B cell tumours shared only T cell gene rearrangements. However, both partial TCR \(\beta\) gene and TCR \(\gamma\) gene rearrangements are often found in B cell neoplasms.\(^27\) The lack of a demonstrable clonal immunoglobulin heavy chain (IgH) gene rearrangement is also acceptable because of the significant false negative rates for PCR with IgH rearrangements.\(^28\) In this particular case all three tumours failed to show an IgH rearrangement, thereby providing additional supportive circumstantial evidence that they were in fact the same clone. The fact that TCR genes are rearranged in a B cell lesion highlights the importance of looking for multiple rearrangements in all lymphoid tumours.

In the case reported here we cannot conclusively prove clonality between the three tumours as to do so would require sequencing of the amplified rearrangements. This technique is unsatisfactory when undertaken on PCR products produced using degenerate primers. However, as the rearrangements demonstrated are identical in all three cases, the probability that they are the same tumour is extremely high.

We believe that PCR techniques, as applied here, demonstrate that PCR will continue to be a valuable tool in the further understanding of the pathogenesis of lymphoid neoplasms, without the inherent technical difficulties presented by restriction enzyme analyses. The case also demonstrates the importance of looking for multiple antigen receptor gene rearrangements to reduce the risk of false negative results.

We feel that the demonstration of probable monoclonality in this unusual case with three histologically distinct appearances is relevant to other cases of synchronous or metachronous lymphoid malignancies, and that it illuminates the complex relationship between the biology and morphology of lymphomas. The case also adds supporting evidence to the hypothesis that LCAL is a heterogeneous group of neoplasms, which may arise not only de novo, but also as the end stage in the progression of a lymphoproliferative disorder.

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A simplified method of detection of clonal rearrangements of the T-cell receptor-gamma chain gene. 


