Relationships between lymphomas linked to hepatitis C virus infection and their microenvironment

Antonino Carbone, Annunziata Gloghini

Antonino Carbone, Department of Pathology, Centro di Riferimento Oncologico Aviano, Istituto Nazionale Tumori, IRCCS, Aviano and Pordenone Hospital, 33081 Aviano, Italy
Antonino Carbone, Member of WHO IARC Monograph Working Group on Biological Agents, 2009 Lyon, France
Annunziata Gloghini, Department of Diagnostic Pathology and Laboratory Medicine, Fondazione IRCCS Istituto Nazionale dei Tumori, 20133 Milano, Italy

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Correspondence to: Antonino Carbone, MD, Chairman of the Department of Pathology, Centro di Riferimento Oncologico Aviano, Istituto Nazionale Tumori, IRCCS, Via F. Gallini 2, 33081 Aviano, Italy. acarbone@cro.it
Telephone: +39-0434-659085 Fax: +39-0434-659370
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Abstract
The relationships between lymphomas and their microenvironment appear to follow 3 major patterns: (1) an independent pattern; (2) a dependent pattern on deregulated interactions; and (3) a dependent pattern on regulated coexistence. Typical examples of the third pattern are hepatitis C virus (HCV)-associated marginal zone lymphomas (MZLs) and mucosa-associated lymphoid tissue lymphomas. In these lymphomas, a regulated coexistence of the malignant cells and the microenvironmental factors usually occurs. At least initially, however, tumor development and cell growth largely depend on external signals from the microenvironment, such as viral antigens, cytokines, and cell-cell interactions. The association between HCV infection and B-cell lymphomas is not completely defined, although this association has been demonstrated by epidemiological studies. MZL and diffuse large B-cell lymphoma are the histotypes most frequently associated with HCV infection. Many mechanisms have been proposed for explaining HCV-induced lymphomagenesis; antigenic stimulation by HCV seems to be fundamental in establishing B-cell expansion as observed in mixed cryoglobulinemia and in B-cell lymphomas. Recently, antiviral treatment has been proved to be effective in the treatment of HCV-associated indolent lymphomas. Importantly, clinically responses were linked to the eradication of the HCV-RNA, providing a strong argument in favor of a causative link between HCV and lymphoproliferation.

Key words: Hepatitis C virus-infection; B-cell lymphomas; Marginal zone lymphoma; Mucosa-associated lymphoid tissue lymphomas; Diffuse large B-cell lymphomas; Microenvironment

Core tip: The relationships between lymphomas and their microenvironment appear to follow 3 major patterns: (1) an independent pattern; (2) a dependent pattern on deregulated interactions; and (3) a dependent pattern on regulated coexistence. The association between hepatitis C virus infection and B-cell lymphomas is not completely defined, although this association has been demonstrated by epidemiological studies.

INTRODUCTION

Genetic alterations and abnormal microenvironmetal factors are involved in tumor development, cell growth and disease progression. Inflammatory cells and soluble mediators, i.e., cytokines and chemokines, are essential microenvironmetal factors that sustain cell growth and invasion, induce angiogenesis and suppress anti-tumor immune functions.

In multidimensional studies on hematolymphoid malignancies, a relevant clinical role of the tumor microenvironment has recently emerged, bringing new knowledge and suggesting new ideas and targets for treatment.

The relationships between lymphomas and their microenvironment appear to follow 3 major patterns: (1) an independent, largely autonomous pattern; (2) a dependent on deregulated interactions pattern; and (3) a dependent on regulated coexistence pattern. A typical example of the first pattern is Burkitt lymphoma where all tumor cells proliferate because of permanent MYC gene activation. A typical example of the second pattern is classic Hodgkin lymphoma, where Reed-Sternberg cells escape the regulated cell growth- and proliferation-control. Typical examples of the third pattern are Hepatitis C virus (HCV)-associated marginal zone lymphomas (MZLs) and mucosa-associated lymphoid tissue (MALT) lymphomas. In this pattern, a regulated coexistence of the malignant cells and the microenvironment is reminiscent of the pattern that the normal counterpart B cells engage in with their respective microenvironment. At least initially, tumor development and cell growth largely depend on external signals from the microenvironment, such as viral antigens, cytokines, and cell-cell interactions.

HCV infection is a worldwide problem. There are important regional differences in the prevalence of HCV infection: the lowest rates are reported in Northern Europe while prevalence estimates exceed 2% in Italy, Japan, Egypt and southern parts of United States. Among the carcinogenic viruses recognized by the recent International Agency for Research on Cancer (IARC) monograph Epstein-Barr virus (EBV), human papilloma virus (HPV), human T-lymphotropic virus type I (HTLV-I), and Kaposi sarcoma-associated herpesvirus (KSHV) play a direct role in carcinogenesis encoding oncoproteins which are able to promote cellular transformation. Conversely, HCV and Helicobacter pylori appear to have an indirect role, by inducing a chronic inflammation. Hepatitis B virus (HBV) has both a direct and indirect role in promoting hepatocellular carcinoma (HCC); as a matter of fact, chronic HBV carriers can develop HCC without developing cirrhosis.

PATHOGENETIC ASPECTS

HCV infection is the cause of chronic hepatitis, liver cirrhosis and hepatocellular carcinoma (HCC) (Table 1). HCV infection has been also associated to a spectrum of extra-hepatic lymphoproliferative disorders including mixed cryoglobulinemia (MC), the most well defined disorder associated with HCV infection, monoclonal gammopathies and B-cell lymphomas. HCV infection has been associated with B-cell low grade indolent lymphoma, especially of marginal zone origin, as well as with aggressive lymphomas, mainly diffuse large B-cell lymphomas (DLBCL) (Table 2). Authoritative studies have demonstrated that in HCV-infected patients with indolent lymphomas, eradication of HCV with antiviral treatment (AT) could directly induce lymphoma regression, providing a strong argument in favor of a causative link between HCV and lymphoproliferation.

INFLAMMATORY MICROENVIRONMENT

The liver is the main target of HCV infection and the major site of inflammatory events, including recruitment of inflammatory cells.

Occurrence of HCV enrichment in intrahepatic inflammatory infiltrates supports the notion that HCV is directly involved in the emergence and maintenance of these B-cell expansions. Intrahepatic B-cell clonalities are invariably associated with extrahepatic manifestations of HCV infection associated B-cell lymphomas.

HCV AND LYMPHOMAGENESIS

According to the recent IARC monograph on biological agents and carcinogenesis, the association between HCV infection and B-cell lymphomas is not completely defined. This association has been demonstrated by epidemiological studies in highly endemic geographical areas. The role of HCV infection in lymphomagenesis may be related to the chronic antigenic stimulation of B-cell response, similar to the well characterized induction of gastric MALT lymphoma development by Helicobacter pylori chronic infection. In fact, chronic HCV infection may sustain a multi-step evolution from MC to overt low grade lymphoma and eventually to high-grade lymphoma. During this process, additional genetic aberrations may induce independence from antigenic stimulation. The clonal component of MC is often an IgM with a rheumatoid factor activity that mirrors the expansion of a B-cell monoclonal population not only in bone marrow but also in liver. It has also been suggested that HCV antigens (NS3 and E7 or E2) can play a role in lymphomagenesis. Recently, it has been published that HCV-related cryoglobulins (either IgM and IgG) are mainly directed against core and NS3 proteins. Chromosomal alterations could also play a role in development of HCV-related lymphoproliferative disorders: for instance, MC with or without lymphoma is characterized by translocation t(14; 18) with the overexpression of the antiapoptotic bcl-2 gene leading to prolonged B-cell survival.

Importantly, cytokines and chemokines (IFNγ, TNFα, CXCL13 and BAFF in MC, as well as osteopontin) are involved in the mechanisms of HCV-
Splenic MZL

Variant

Specific Cholangiocarcinoma

psittaci have been involved in MALT lymphomas arising from Helicobacter pylori, Borrelia burgdorferi, and Chlamydia psittaci. For example, H. pylori infection has been well characterized in MZLs. Within indolent lymphoma subtypes, the association with HCV infection has been best characterized in MZLs. HCV infection has been associated with the development of various lymphoid proliferations, including lymphoid tissue hyperplasia, reactive lymphoid hyperplasia, and lymphoid neoplasms. The association of HCV infection and lymphomas has been of particular interest, as HCV infection is a well-known risk factor for hepatocellular carcinoma.

LYMPHOMAS

HCV infection and lymphomas

Table 1: Biological agents assessed by the International Agency for Research on Cancer Monographs Working Group

<table>
<thead>
<tr>
<th>Group</th>
<th>Agent</th>
<th>Cancers on which sufficient evidence in humans is based</th>
<th>Other sites with limited evidence in humans</th>
<th>Established mechanistic events</th>
</tr>
</thead>
<tbody>
<tr>
<td>Epstein-Barr virus</td>
<td>Nasopharyngeal carcinoma, Burkitt lymphoma, Immune-suppression-related non-Hodgkin lymphoma, Extranodal NK/T-cell lymphoma (nasal type), Hodgkin lymphoma</td>
<td>Gastric carcinoma, Lympho-epithelioma-like carcinoma</td>
<td>Cell proliferation, inhibition of apoptosis, genomic instability, cell migration</td>
<td></td>
</tr>
<tr>
<td>Hepatitis B virus</td>
<td>Hepatocellular carcinoma</td>
<td>Cholangiocarcinoma, Non-Hodgkin lymphoma</td>
<td>Inflammation, liver cirrhosis, chronic hepatitis</td>
<td></td>
</tr>
<tr>
<td>Hepatitis C virus</td>
<td>Hepatocellular carcinoma, Non-Hodgkin lymphoma</td>
<td>Cholangiocarcinoma</td>
<td>Inflammation, liver cirrhosis, liver fibrosis</td>
<td></td>
</tr>
<tr>
<td>Kaposi sarcoma herpes virus</td>
<td>Kaposi sarcoma, Primary effusion lymphoma</td>
<td>Multicentric Castleman’s disease</td>
<td>Cell proliferation, inhibition of apoptosis, genomic instability, cell migration</td>
<td></td>
</tr>
<tr>
<td>Human immunodeficiency virus, type 1</td>
<td>Kaposi sarcoma, Non-Hodgkin lymphoma, Hodgkin lymphoma</td>
<td>Cancer of the vulva, vagina, penile, Non-melanoma skin cancer,</td>
<td>Immortalization, genomic instability, inhibition of DNA damage response, anti-apoptotic activity</td>
<td></td>
</tr>
<tr>
<td>Human papillomavirus type 16 (For the other types see Table 2)</td>
<td>Carcinoma of the cervix, vulva, vagina, penis, anogenital, oral cavity, oropharynx and tonsil</td>
<td>Hepatocellular carcinoma</td>
<td>Inflammation, oxidative stress, altered cellular turnover, changes in gene expression, methylation, mutation</td>
<td></td>
</tr>
<tr>
<td>Helicobacter pylori</td>
<td>Non-cardia gastric carcinoma, Low-grade B-cell mucosa-associated lymphoid tissue</td>
<td>Gastric lymphoma</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

1 Newly identified link between virus and cancer. In red are highlighted the lymphoid proliferations; HBV has both a direct and indirect role in promoting hepatocellular carcinoma. Modified and adapted from Bouvard et al.

Table 2: Hepatitis C virus-associated indolent and aggressive lymphoid proliferations

<table>
<thead>
<tr>
<th>Subtype</th>
<th>Variant</th>
<th>Specific lymphoma sites</th>
</tr>
</thead>
<tbody>
<tr>
<td>Monoclonal B-cell lymphocytosis</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Tissue based monoclonal B cell and plasma cell proliferations of uncertain type</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Lymphoplasmocytic lymphoma/WM</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Chronic lymphocytic disorders (non CLL)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>MZL</td>
<td>Splenic MZL, Nodal MZL, MALT</td>
<td>Gastric, Extranodal non gastric, Salivary gland, Skin, Orbit, Liver</td>
</tr>
<tr>
<td>Diffuse large B-cell lymphoma</td>
<td></td>
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</tbody>
</table>

1WM: Waldenström’s macroglobulinemia; CLL: Chronic lymphocytic leukemia; MZL: Marginal zone lymphoma; MALT: Mucosa-associated lymphoid tissue.

HCV infection and specific lymphomas

Within indolent lymphoma subtypes, the association with HCV infection has been best characterized in MZLs. Other infectious agents have been involved in the pathogenesis of specific types of MZLs. For example, Helicobacter pylori, Borrelia burgdorferi, and Chlamydia psittaci have been involved in MALT lymphomas arising in stomach, skin and orbit, respectively. Conversely, chronic stimulation by HCV plays a role in development of splenic marginal zone lymphoma (SMZL) and primary nodal marginal zone lymphoma. Primary nodal marginal zone lymphoma is a distinct clinical-pathological subtype characterized by exclusive primary lymph node localization in the absence of extranodal site of involvement (Figure 1).

Splenomegaly is a common feature of SMZL.

Splenomegaly is a common feature of SMZL. Splenic and nodal MZL are indolent B-cell lymphomas corresponding to post-germinal center memory B cells that are supposed to derive from marginal zone B cells. These entities share some morphologic and pathogenic features, but have distinctive clinical presentation, immunophenotype, and molecular abnormalities. Histologically, when the marginal zone B cells surround normal follicles with benign mantle zones as a third outer layer, they produce a marginal zone pattern. As the marginal zone cells extend outwards into the interfollicular areas, they form confluent clusters resulting in an interfollicular pattern or a diffuse pattern in the absence of any follicles at later stages of the disease.

As the marginal zone cells extend outwards into the interfollicular areas, they form confluent clusters resulting in an interfollicular pattern or a diffuse pattern in the absence of any follicles at later stages of the disease. The marginal zone cells may also grow inwards into the follicles and produce either partial or complete follicular colonization (Figure 1). Gastric and non-gastric extranodal MZL are typically indolent diseases of middle and advanced age; disseminated disease is present in nearly one-third of cases. Interestingly, three specific MALT lymphoma sites showed an elevated prevalence of HCV infection: salivary glands, skin and orbit. The association of HCV infection and salivary glands lymphoma has been clearly demonstrated.

Moreover, a study on B-cell lymphoma in patients with Sjögren’s syndrome and HCV infection reported an
The lymphoma cells are most likely to be independent from first-line approach for HCV-associated DLBCL, because lymphoma, AT seems not to play a central role in the HCV-negative DLBCL.

In particular, residual signs of low-grade lymphoma and specific presentation with respect to HCV-negative DLBCL. Patients affected by HCV-associated DLBCL display specific characteristics that differentiate them from those with HCV-negative DLBCL. This pattern requires proper identification in order to avoid the misdiagnosis of the lymphoma. Three hypothetic models have emerged to understand the molecular mechanisms of HCV-associated lymphoma development: (1) continuous external stimulation of lymphocyte receptors by viral antigens and consecutive proliferation; (2) direct role of HCV replication and expression in infected B-cells; and (3) permanent B-cell damage, e.g., mutation of tumor suppressor genes, caused by a transiently intracellular virus (“hit and run” theory). Other non-exclusive hypotheses have been proposed over the past two decades. These hypotheses have variously emphasized the important role played by chromosomal aberrations, cytokines, or microRNA molecules.

CONCLUSION

Many epidemiological studies have provided evidence that HCV infection is associated with development of indolent and aggressive B-cell lymphoma. However, the causal association between HCV infection and B-cell lymphomas is not completely defined. The similarities shared by rearranged Ig genes present in B cells from patients with type II MC and malignant B-cells from HCV-positive patients affected by B-cell lymphoma support the possibility that the antigens that promote type II MC and B-cell lymphoma in HCV-positive patients are the same. These similarities also suggest that type II MC may be a precursor of B-cell lymphoma. Type II MC probably plays a central role in the development of B-cell lymphoma in HCV-positive patients with Sjögren’s syndrome.

Three hypothetic models have emerged to understand the molecular mechanisms of HCV-associated lymphoma development: (1) continuous external stimulation of lymphocyte receptors by viral antigens and consecutive proliferation; (2) direct role of HCV replication and expression in infected B-cells; and (3) permanent B-cell damage, e.g., mutation of tumor suppressor genes, caused by a transiently intracellular virus (“hit and run” theory).

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