

# Leukaemia Section

## Short Communication

### Intestinal T-cell lymphoma

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#### Identity

**Alias :** Enteropathy-type T-cell lymphoma

#### Clinics and pathology

##### **Phenotype/cell stem origin**

The disease originates from a CD3+, CD7+ T-lymphocyte lacking CD4 and CD8 expression.

##### **Epidemiology**

The disease affects mainly the adult, with male predominance, and it is frequently associated with gluten-sensitive enteropathy.

##### **Pathology**

The disease consists of ulcerated lesions involving the small intestine. Perforation may occur. Small and larger atypical lymphocytes with pale cytoplasm infiltrate the epithelial mucosa of the villi. The TCR-Beta and TCR-Gamma genes are clonally rearranged.

##### **Treatment**

Multiagent chemotherapy (CHOP or CHOP-like regimens) was used.

##### **Evolution**

The disease may derive from patients with coeliac disease not responding to gluten-free diet. The lymphoma may spread to regional lymph nodes.

##### **Prognosis**

Response to chemotherapy is suboptimal and patients are vulnerable to toxicity of treatment due to intestinal symptoms and malnutrition preceding the diagnosis of lymphoma. Survival at 2 years was 28% in a study (Daum et al., 2003).

#### Cytogenetics

##### **Cytogenetics molecular**

Extra copies of chromosome 9q centered around the 9q33-34 region was detected by CGH and FISH studies (Zettl et al., 2002). 9p deletion with p16 loss was found in 18% of the cases, and loss of heterozygosity at 9p21 with loss of p16 expression was documented in approximately half of those cases with a large cell component (Obermann et al., 2004). DNA gains may involve the 5q33-34 and 7q31 regions, with a 30% frequency. Loss of chromosome material was detected at 6p24; 7p21, 17q23-25 and 17p13 (Baumgartner et al., 2003).

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