

# Leukaemia Section

## Mini Review

### NK cell neoplasias

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#### Clinics and pathology

##### **Disease**

Neoplasms of natural killer (NK) cells are rare, and have not been well characterized until the past decade. In the new WHO classification of hematolymphoid tumors, three categories of NK cell neoplasms are recognized:

- Extranodal NK/T cell lymphoma,
- Aggressive NK cell leukemia, and
- Blastic NK cell lymphoma.

Blastic NK cell lymphoma is morphologically and immunologically different from the first two categories, lacks EBV association, and there is little compelling evidence that it truly represents an NK cell neoplasm. In fact, recent studies suggest that this may be a neoplasm of probable precursor dendritic cells related to plasmacytoid monocytes (plasmacytoid dendritic cells). Since its lineage is still uncertain, this entity will not be discussed.

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

NK cell represents a distinctive lineage of lymphocyte that is closely related to T cell. It shows many immunophenotypic and functional similarities with cytotoxic T lymphocyte, but differs in the lack of expression of surface CD3 molecule and T-cell receptor, and the absence of rearranged T-cell receptor genes. It characteristically expresses CD56 (neuronal cell adhesion molecule, N-CAM), which is also expressed in some cytotoxic T lymphocytes. NK cells can lyse target cells without prior sensitization (spontaneous antibody-independent MHC-unrestricted cytotoxicity) via the NK receptors.

##### **Etiology**

The exact etiology is unknown, but a very strong association with Epstein Barr virus (EBV) has been demonstrated.

##### **Epidemiology**

NK cell neoplasms show strong geographic differences in their prevalence. They are more common in Asia, Mexico, and South America, but are very rare in the Western populations.

##### **Clinics**

They occur predominantly in the nose/nasopharynx, but sometimes in extranasal sites (most commonly skin), in middle-aged to elderly patients. Systemic involvement is uncommon at diagnosis but rarely, they may present initially in a leukemic form. The most common presenting symptoms are nasal obstruction, nasal discharge and epistaxis. The full-blown midfacial destructive and ulcerative lesions (hence the name midline granuloma) are much less commonly seen nowadays. Patients with aggressive NK cell leukemia present with high swinging fever, systemic symptoms and hepatosplenomegaly; they are usually extremely ill, with deranged liver function and coagulation profile.

##### **Cytology**

The neoplastic NK cells are often heterogeneous in appearance but some (particularly the circulating leukemic cells) may resemble large-sized normal large granular lymphocytes with ample amount of pale or lightly blue cytoplasm that contains fine or coarse azurophilic granules.

## Pathology

The malignant infiltrate is diffuse, often with a prominent angiocentric and angiodestructive component. Coagulative necrosis and apoptosis are common. The cytological spectrum is variable, ranging from small, medium-sized, large or anaplastic cells, to a mixture of these cells. The cells often have irregularly folded nuclei and granular chromatin. In Giemsa-stained cytologic preparations, azurophilic granules are often detected in the cytoplasm. Reactive histiocytes with haemophagocytosis are sometimes found in the bone marrow, particularly for the leukemic form. NK cell neoplasms are characterised by an immunophenotype of CD2+, surface CD3-, cytoplasmic CD3e+, CD56+ and T cell receptor (TCR)-, lack of TCR gene rearrangement, and strong association with EBV.

## Treatment

The disease is often resistant to chemotherapy. For extranodal NK/T cell lymphoma, the best results are obtained by radiotherapy with or without aggressive chemotherapy/stem cell rescue. Plasma or serum EBV DNA and tissue p73 gene hypermethylation assay can be used for monitoring of disease status or detection of minimal residual disease. Aggressive NK cell leukemia is treated by chemotherapy, but response is typically poor.

## Evolution

Although extranodal NK/T cell lymphoma is usually localized at presentation, systemic progression often occurs, usually early in the course of disease. Common distant sites of involvement are the skin, liver, lung, gastrointestinal tract, testis, and rarely bone marrow. Patients with aggressive NK cell leukemia typically exhibit a rapidly progressive clinical course, with multi-organ failure and bleeding tendency.

## Prognosis

Clinical factors reported to have prognostic significance in extranodal NK/T cell lymphoma include stage and bulk of disease, B symptoms, age, performance status and International Prognostic Index. The overall survival for patients with extranodal NK/T cell lymphoma is 30-40%. Practically all patients with aggressive NK cell leukemia die from the disease within a few weeks or months of presentation.

## Genetics

### Note

In contrast to T cells, NK cells do not show rearrangements of the TCR genes. As expected from their proposed normal counterpart, NK cell neoplasms show a germline configuration of the TCR genes and do not express TCR proteins on the cell surface. The detection of single circularised episomal form of EBV

in the neoplasm by Southern blot analysis provides indirect evidence to the clonal nature. Molecular demonstration of X chromosome inactivation in female patients with NK cell neoplasms also provides evidence for clonality. However, the most direct evidence for clonality of this group of tumors has been provided by the detection of clonal chromosomal abnormalities (see section below).

It has been shown that in over 90% of NK cell neoplasms, a specific pattern of promoter CpG methylation occurs, with p73 being consistently involved. It has been further suggested that p73 may be an important target in the oncogenesis of NK cell neoplasms, and the demonstration of its methylation may serve as a useful molecular marker for disease monitoring.

## Cytogenetics

### Note

A variety of genetic abnormalities has been described, but so far no specific and consistent chromosomal translocation has been identified by conventional cytogenetics. In most instances, the genetic changes involve loss or gain of genetic materials such as del(6q), and i(1q). Frequent genetic losses in 6q and 13q have been confirmed by both comparative genomic hybridization (CGH) and loss of heterozygosity (LOH) analyses. Other non-random abnormalities include +X, i(1q), i(7q), +8, i(17q), and 11q23 rearrangement. Chromosomal deletion involving chromosome 6q at around q21-q25 is the commonest recurrent chromosomal abnormality, and fluorescence in situ hybridisation studies have shown that 6q22-q23 is the most frequently involved regions in the chromosome 6 deletions. A recent study using LOH and homozygosity mapping of deletion (HOMOD) analyses has, however, defined a distinct 3 Mb smallest region of overlapping on 6q25.

A possible involvement of 8p22-p23 in both NK cell neoplasms and NK cell line such as NK-92 has also been suggested. Translocation involving 8p23 has been reported in 3 cases of NK cell neoplasms, with the partner chromosomes being 8q13, 17q24 and 1q10. An add(8)(q23) abnormality has also been demonstrated in one case each of aggressive NK cell leukemia and extranodal NK/T cell lymphoma.

## References

- Kern WF, Spier CM, Hanneman EH, Miller TP, Matzner M, Grogan TM. Neural cell adhesion molecule-positive peripheral T-cell lymphoma: a rare variant with a propensity for unusual sites of involvement. *Blood*. 1992 May 1;79(9):2432-7
- Wong KF, Chan JK, Ng CS, Lee KC, Tsang WY, Cheung MM. CD56 (NKH1)-positive hematolymphoid malignancies: an aggressive neoplasm featuring frequent cutaneous/mucosal

involvement, cytoplasmic azurophilic granules, and angiocentricity. *Hum Pathol.* 1992 Jul;23(7):798-804

Chan JK, Sin VC, Wong KF, Ng CS, Tsang WY, Chan CH, Cheung MM, Lau WH. Nonnasal lymphoma expressing the natural killer cell marker CD56: a clinicopathologic study of 49 cases of an uncommon aggressive neoplasm. *Blood.* 1997 Jun 15;89(12):4501-13

Kwong YL, Chan AC, Liang R, Chiang AK, Chim CS, Chan TK, Todd D, Ho FC. CD56+ NK lymphomas: clinicopathological features and prognosis. *Br J Haematol.* 1997 Jun;97(4):821-9

Wong KF, Chan JK, Kwong YL. Identification of del(6)(q21q25) as a recurring chromosomal abnormality in putative NK cell lymphoma/leukaemia. *Br J Haematol.* 1997 Sep;98(4):922-6

Cheung MM, Chan JK, Lau WH, Foo W, Chan PT, Ng CS, Ngan RK. Primary non-Hodgkin's lymphoma of the nose and nasopharynx: clinical features, tumor immunophenotype, and treatment outcome in 113 patients. *J Clin Oncol.* 1998 Jan;16(1):70-7

Siu LL, Wong KF, Chan JK, Kwong YL. Comparative genomic hybridization analysis of natural killer cell lymphoma/leukemia. Recognition of consistent patterns of genetic alterations. *Am J Pathol.* 1999 Nov;155(5):1419-25

Wong KF, Zhang YM, Chan JK. Cytogenetic abnormalities in natural killer cell lymphoma/leukaemia--is there a consistent pattern? *Leuk Lymphoma.* 1999 Jul;34(3-4):241-50

Zhang Y, Wong KF, Siebert R, Matthiesen P, Harder S, Eimermacher H, Feller AC, Schlegelberger B. Chromosome aberrations are restricted to the CD56+, CD3- tumour cell population in natural killer cell lymphomas: a combined immunophenotyping and FISH study. *Br J Haematol.* 1999 Jun;105(3):737-42

Siu LL, Chan V, Chan JK, Wong KF, Liang R, Kwong YL. Consistent patterns of allelic loss in natural killer cell lymphoma. *Am J Pathol.* 2000 Dec;157(6):1803-9

Wong KF, Chan JK, Cheung MM, So JC. Bone marrow involvement by nasal NK cell lymphoma at diagnosis is uncommon. *Am J Clin Pathol.* 2001 Feb;115(2):266-70

Siu LL, Chan JK, Wong KF, Kwong YL. Specific patterns of gene methylation in natural killer cell lymphomas : p73 is consistently involved. *Am J Pathol.* 2002 Jan;160(1):59-66

Wong KF. Genetic changes in natural killer cell neoplasms. *Leuk Res.* 2002 Nov;26(11):977-8

Cheung MM, Chan JK, Wong KF. Natural killer cell neoplasms: a distinctive group of highly aggressive lymphomas/leukemias. *Semin Hematol.* 2003 Jul;40(3):221-32

Siu LL, Chan JK, Wong KF, Choy C, Kwong YL. Aberrant promoter CpG methylation as a molecular marker for disease monitoring in natural killer cell lymphomas. *Br J Haematol.* 2003 Jul;122(1):70-7

Sun HS, Su IJ, Lin YC, Chen JS, Fang SY. A 2.6 Mb interval on chromosome 6q25.2-q25.3 is commonly deleted in human nasal natural killer/T-cell lymphoma. *Br J Haematol.* 2003 Aug;122(4):590-9

Chim CS, Ma SY, Au WY, Choy C, Lie AK, Liang R, Yau CC, Kwong YL. Primary nasal natural killer cell lymphoma: long-term treatment outcome and relationship with the International Prognostic Index. *Blood.* 2004 Jan 1;103(1):216-21

Wong KF. A novel EBV-negative natural killer cell line. *Leuk Res.* 2004 Mar;28(3):225-7

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