

Leukaemia Section

Short Communication

t(4;14)(p16;q32)

Jean-Loup Huret, Jacky Bonaventure

Genetics, Dept Medical Information, University of Poitiers, CHU Poitiers Hospital, F-86021 Poitiers, France (JLH); Unité INSERM 393, Hopital Necker-Enfants Malades, 149 rue de Sèvres 75743, Paris Cedex 15, France (JB)

Published in Atlas Database: March 1998

Online updated version: <http://AtlasGeneticsOncology.org/Anomalies/t04142059.html>

DOI: 10.4267/2042/37437

This work is licensed under a Creative Commons Attribution-Non-commercial-No Derivative Works 2.0 France Licence.
© 1998 Atlas of Genetics and Cytogenetics in Oncology and Haematology

Clinics and pathology

Disease

Found in plasma cell leukaemia and multiple myeloma.

Phenotype / cell stem origin

Malignant plasma cells have the phenotype of mature terminally differentiated B-cells; their origin may be a pluripotent stem cell.

Epidemiology

Yet poorly described: found in 6 fresh tumours and 5 cell lines; might be frequent but karyotypically undetected.

Clinics

Data on clinics and cytogenetics are missing.

Cytogenetics

Cytogenetics, morphological

May be undetectable (telomere-telomere translocation).

Cytogenetics, molecular

Therefore molecular probes are indicated, and FISH is relevant.

Genes involved and Proteins

FGFR3

Location: 4p16.3

FGFR3 Ig-like trans-mb Tyr kinase

Protein

Contains an extracellular domain with Ig-like loops, a transmembrane domain, and intracellular tyrosine

kinase domains; localisation: plasma membrane; tyrosine kinase receptor; role in signal transduction.

IgH

Location: 14q32

Results of the chromosomal anomaly

Hybrid gene

Description

FGFR3 is translocated on der(14) which contains the 3' IgH enhancer.

Fusion protein

Description

No fusion protein, but promoter exchange between both partner genes; however, somatic mutations similar to what has been found in thanatophoric dwarfism have been identified in some cases; they may also contribute to abnormal FGFR3 activation.

Oncogenesis

Overexpression and activation of FGFR3 provides an oncogenic signal.

References

Bergsagel PL, Chesi M, Nardini E, Brents LA, Kirby SL, Kuehl WM. Promiscuous translocations into immunoglobulin heavy chain switch regions in multiple myeloma. Proc Natl Acad Sci USA 1996 Nov 26;93(24):13931-6.

Chesi M, Nardini E, Brents LA, Schröck E, Ried T, Kuehl WM, Bergsagel PL. Frequent translocation t(4;14)(p16.3;q32.3) in multiple myeloma is associated with increased expression and activating mutations of fibroblast growth factor receptor 3. Nat Genet 1997 Jul;16(3):260-4.

Richelda R, Ronchetti D, Baldini L, Cro L, Viggiano L, Marzella R, Rocchi M, Otsuki T, Lombardi L, Maiolo AT, Neri A. A novel chromosomal translocation t(4;14)(p16.3;q32) in multiple myeloma involves the fibroblast growth-factor receptor 3 gene. *Blood* 1997 Nov 15;90(10):4062-70.

This article should be referenced as such:

Huret JL, Bonaventure J. t(4;14)(p16;q32). *Atlas Genet Cytogenet Oncol Haematol.*1998;2(3):91-92.
