

# Leukaemia Section

## Short Communication

### t(3;5)(p21;q32)

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## Clinics and pathology

### Disease

MKPL-1 cell line, established from a 66-year-old male patient with an acute megakaryoblastic leukemia (M7-AML) and a karyotype apparently with -21,+3mar (Takeuchi et al., 1992), re-analysed for tyrosine kinase dysregulation (Gu et al., 2007).

### Epidemiology

Only one case to date.

## Genes involved and proteins

### RBM6

#### Location

3p21

#### Protein

From N-term to C-term, contains a BTB/POZ domain (mediates homomeric dimerization) and decamer repeat domains, responsible for multimerization/self-association of the protein, RRM1 and RRM2 (RNA recognition motif) domains, an octamer repeat, a C2H2 zinc finger, a nuclear localisation signal, and a G-patch (made of highly conserved glycines; may have RNA binding functions). RNA-binding protein. Binds poly(G). Splicing factor (Heath et al., 2010).

### CSF1R

#### Location

5q32

#### Protein

Contains Ig-like domains (extracellular), a transmembrane domain, and a split tyrosine kinase domain (intracellular), from N-term to C-term. Transmembrane glycoprotein, receptor for the ligand

colony stimulating factor-1 (CSF1). Upon binding of CSF1, CSF1R tyrosine phosphorylation is induced leading to RAS/RAF/MAPK, PI3K/AKT/mTOR and JAK/STAT (specifically STAT1, STAT3, and STAT5) pathways activation. CSF1R activation by CSF1 results in increased growth, proliferation and differentiation (Fischer et al., 2008).

## Result of the chromosomal anomaly

### Hybrid gene

#### Description

Fusion of RBM6 exon 2 to CSF1R exon 12; the reciprocal CSF1R-RBM6 was not detected.

### Fusion protein

#### Description

The RBM6-CSF1R fusion protein consists of the amino terminal 36 amino acids of RBM6, fused to the carboxy terminal 399 amino acids of CSF1R, including a polymerisation domain of RBM6, and the tyrosine kinase domain of CSF1R.

#### Oncogenesis

Constitutive tyrosine kinase activation.

## References

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