

# Leukaemia Section

## Mini Review

### t(3;4)(p21;q34)

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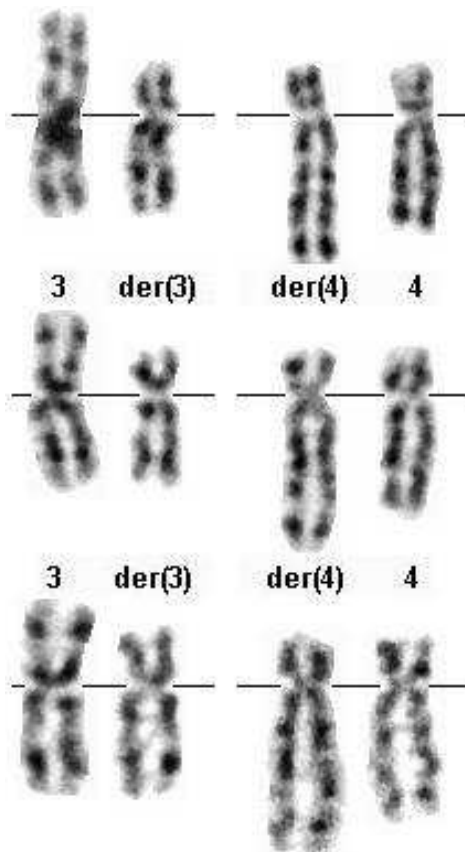
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## Identity



t(3;4)(p21;q34) G-banding

## Clinics and pathology

### Disease

Myeloid lineage, found in 1 myelodysplastic syndrome (MDS) and 1 Acute Myeloid Leukemia (AML).

### Phenotype / cell stem origin

MDS-RA and M1 AML by FAB criteria, a primitive myeloid progenitor is likely to be involved.

### Etiology

No known prior exposure.

### Epidemiology

Only 2 cases to date, a 69 year old female and a 31 year old male, sex ratio 1M/1F.

### Clinics

Elevated WBC ( $68 \times 10^9/l$ ), 93% blasts in blood, lymphadenopathy, hepatosplenomegaly, high LDH in AML patient.

### Cytology

Positive for CD 34, HLDR, CD33, CD68, MPO in AML.

### Treatment

Chemotherapy followed by bone marrow transplantation in AML.

### Evolution

After the first cycle of therapy, persistent bone marrow infiltration with 11% blasts.

### Prognosis

Survival 6 month in MDS, 15 month+ in AML.

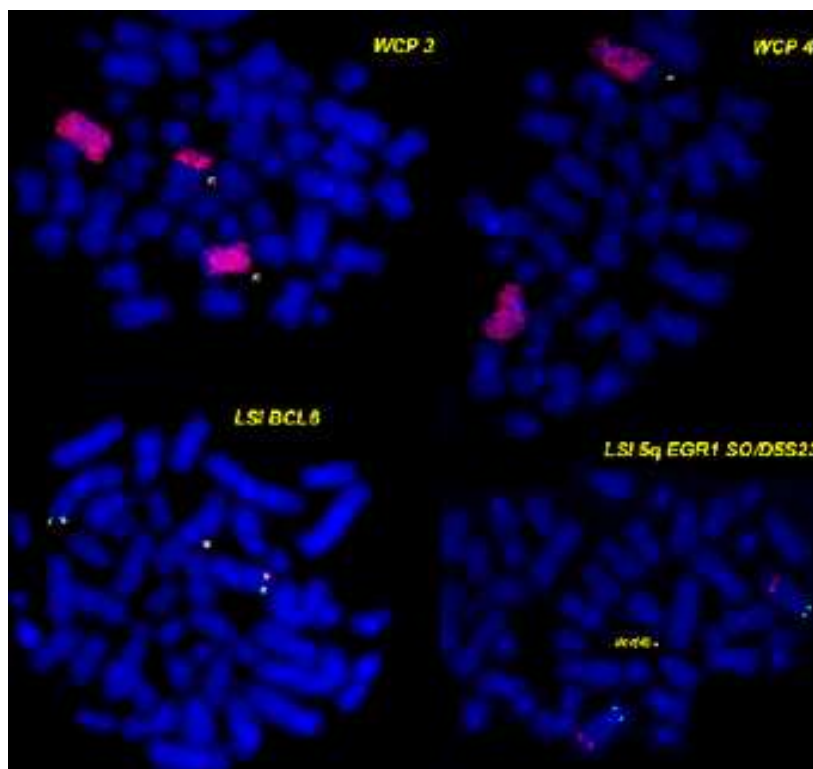
## Cytogenetics

### Cytogenetics morphological

May be misinterpreted as t(3;5) in suboptimal preparations.

### Cytogenetics molecular

FISH analysis is recommended to exclude the more frequent t(3;5).



FISH with WCP 3 and 4 and LSI BCL6 and 5q EGR1 probes.

### Probes

WCP 3 and 4 probes, locus specific BCL6 and 5q probes.

### Additional anomalies

t(3;4)(p21;q34) is part of a complex karyotype in MDS case associated with del(20q), sole abnormality in AML case.

### Genes involved and Proteins

**Note:** 3p21 is a recurrent breakpoint in MDS/AML and t-MDS/t-AML suggesting, 3p21 site is likely to contain a gene (genes) involved in the pathogenesis of t(3;4)(p21;q34). Frequent deletion or allelic loss of band 3p21 is common in solid tumors, indicating the presence of tumor suppressor genes on this chromosome arm. The association among structural chromosome 3 aberrations and fragile sites on 3p may indicate the importance of previous mutagen exposure in the etiology of these diseases.

Although several cancer-related genes have been located to 3p21, no gene has yet been identified to be related with hematological malignancies. One of the candidate genes may be the AF3p21 gene, a novel fusion partner of the MLL gene described in a patient who had developed therapy-related leukemia with t(3;11)(p21;q23). AF3p21 encodes a protein localized

exclusively in the cell nucleus, suggesting the possibility that AF3p21 protein plays a role in signal transduction in the nucleus.

### References

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