

# Cancer Prone Disease Section

## Mini Review

## Rothmund-Thomson syndrome (RTS)

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### Identity

**Alias:** Poikiloderma atrophicans and cataract

#### Note

Is a chromosomal instability syndrome with an increased risk of cancers.

#### Inheritance

Autosomal recessive; rare geno-dermatosis with increased frequency in females; 260 cases reported in the English medical literature.

### Clinics

#### Note

Clinical expression highly variable.

#### Phenotype and clinics

Main features include:

- Growth retardation.
- Skin defects appearing within the first year of life (90%) and persisting throughout life: atrophic dermatosis, poikiloderma, hyperpigmentation, teleangiectasia.
- Sparse hair which may progress to partial or total alopecia; dystrophic nails.
- Photosensitivity.
- Congenital skeletal defects - hypoplasia or absence of the radii and thumbs, osteopenia, cystic or sclerotic changes of the long bones - (>50%); bone age lower than chronological age.
- Juvenile cataract, corneal dystrophy (50%).
- Hypodontia.
- Hypogonadism (25%).
- Proportionate short stature.
- Premature aging.

Detailed definition of the clinical profile in a contemporary cohort of 41 RTS patients evidences some differences in the distribution of the clinical findings (figure 1), which should be kept into account to optimize diagnostic criteria.

Diagnosis: the diagnosis is difficult before the development of the erythema; differential diagnosis with:

- Werner syndrome,
- Dyskeratosis congenita,
- Cockayne syndrome,
- Bloom syndrome,
- Fanconi anaemia,
- Anhidrotic ectodermal dysplasia.

#### Neoplastic risk

Patients have an enhanced risk of bone cancer, specifically osteosarcoma (30 out of 300 - 10% - in the literature) and nonmelanoma skin cancers (squamous cell carcinoma, basal cell carcinoma) with an estimated prevalence around 5%.

Etiology: deficiency of RECQL4 helicase (see below), a protein involved in some aspect of replication and transcription, repair and recombination may suggest a mutational mechanism for cancer development, but the specificity for mesenchymal tumors is currently unknown.

#### Treatment

Only protection against sunlight is possible by means of sunscreens with both UVA and UVB protection; dermatologic therapies; surgical correction of skeletal malformations and cataracts;



regular and careful work-up of signs and symptoms of both cutaneous and internal malignancy; caution is warranted in administering chemotherapy to affected individuals due to their sensitivity to chemotherapeutic agents.

**Evolution**

The disease tends to progress during the first years of life, but becomes static so that patients may have a normal lifespan; the mortality from neoplastic disease

during the second or third decade is very significantly increased.

**Cytogenetics**

**Inborn conditions**

Spontaneous/induced chromatid breaks were found increased in only a very few studies; In contrast with (mainly negative) chromatid results, consistent clonal/non clonal structural chromosomal abnormalities

were evidenced in most studies, often involving chromosome 8, in cultured lymphocytes and in fibroblasts; low frequency trisomy 8 mosaicism has been reported in both lymphocyte and primary fibroblast cultures as well as in uncultured blood and buccal smears, indicating this characteristic chromosomal abnormality is present in vivo; a propensity to centromere misdivision with development of clones carrying isochromosomes, such as i(8q), i(8p), i(12p), i(12q) is peculiar of RTS.

### **Cytogenetics of cancer**

Marked chromosomal instability has been detected in mesenchymal tumours developed by RTS sibs.

## **Other findings**

### **Note**

Reduced unscheduled DNA synthesis, 37% of normal after exposure to ultraviolet C or gamma irradiation.

## **Genes involved and proteins**

### **Note**

Mutations of RECQL4 gene, a member of the RECQ helicase gene family, have been identified in a limited number of RTS patients. Molecular testing is available on a research basis and useful in confirming the diagnosis in suspected RTS patients carrying RECQL4 mutations.

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