# **Brief Report**

## Syndrome-causing mutations in Werner syndrome

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Summary Complete loss of function in the WRN: RecQ3 DNA/RNA helicase gene causes Werner Syndrome (WS). WS patients with genetic instability manifest an early onset of age-related diseases including diabetes mellitus (DM), osteoporosis, atherosclerosis, and malignancy as well as early death. In 1,420 patients, WS was reported to be associated with chromosomal abnormality syndrome and other genetic diseases including Klinefelter syndrome in 2 patients, retinitis pigmentosa in 3, Wilson's disease in 1, xeroderma pigmentosum in 3, and porokeratosis Mibelli in 1. These clinical findings may support the concept of genetic instability in WS.

Keywords: Aging, Genetic instability, Mutation, Werner syndrome, Xeroderma pigmentosum

#### 1. Introduction

Half a century ago, Dr. Alex Comfort encouraged medical researchers to look for evidence of chromosomal abnormalities in many forms of constitutional disorders, the most important of which was Werner syndrome (WS: MIM#27770) (1). However, his theory faltered since normal chromosomes were found in WS (2-5). WS, the gene located at chromosome 8p11-12 (6) and caused by a recessively mutated WRN, is characterized by a variety of clinical manifestations mimicking features of advanced aging and thus may represent a typical progeroid syndrome (7). Wrn is a member of the RecQ helicase gene family (RecQ3) and may interact with a variety of DNA/RNA metabolism enzymes during repair, transcription, translation, recombination, replication, and chromosome segregation in the nucleus (8). Thus, actively proliferating cells may be affected by WRN dysfunction. Theories on the function of RecQ helicases and in vitro studies using WS fibroblasts and peripheral blood cells have suggested

genomic instability in WS cells (8-12). Patients with WS do not usually have apparent abnormalities before their teenage growth spurt, but they typically display hierarchical deterioration of a variety of connective tissue systems resulting in physical symptoms such as gray hair, alopecia, skin atrophy, skin sclerosis, skin hyper/hypo-pigmentation, vocal cord atrophy, osteoporosis, sarcopenia, bilateral cataracts, metastatic subcutaneous calcification, and atherosclerosis. Connective or supportive tissue may be a source of malignancies (and particularly sarcomas), and adversely affected systems include the endocrine system, resulting in type II diabetes mellitus (DM), hypogonadism, and thyroid disorders, and the metabolic system, resulting in hyperlipidaemia, hyperuricemia and hyaluronuria. Systems affected to a lesser degree include the immune system, resulting in excessive auto-antibody production, impaired cytokine response, and natural killer cell activity, and the nervous system, resulting in cognitive disorders and brain atrophy (5,7,13). Death due to malignancy or atherosclerosis-related conditions such as myocardial infarction typically occurs in the late 40s (13-15). In addition, in vivo mutation as may be associated with genetic instability of WS cells may induce other genetic diseases.

Since the first description of WS by the German family physician Otto Werner in 1904 (16), 1,420 cases of WS have been reported in total worldwide (17). Interestingly, 75% of WS patients are of Japanese

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descent (14), which is probably due to the relatively high frequency of consanguineous marriage in rural areas and an extremely high prevalence (1:100) of heterozygosity in the general Japanese population (18,19). Approximately 10 WS cases per year are regularly documented in Japan, while only 3.3 patients per year are reported outside the country. Adjusting roughly for population size, the frequency of WS in Japan is some 150-fold greater than in the rest of the world (Japan's population between 1966 and 2004 was about 113,000,000 with respect to a world population of about 5,000,000,000; that said, case reports are highly encouraged in Japan in comparison to some parts of the world).

Since the cloning of the WRN: RecQ3 helicase gene, the search for mutation-causing mutations in WS has been a matter of scientific/clinical interest (20). The current study looked for chromosomal abnormalities and genetic diseases associated with WS in WS patients.

#### 2. Materials and Methods

The first case of WS in Japan was reported in 1917 (21). Here, the clinical manifestations of WS as described in all papers published between 1917 and 2004 were analyzed. WS publications were selected through a citation index (Igaku-Chuo-Zasshi) and bibliographies of each report were extensively examined for additional references. For comparison of Japanese and foreign patients with WS, searches were performed using PubMed. Care was taken to thoroughly identify patient family details, personal histories, authors, institutions, and demographic characteristics to avoid the inclusion of duplicate patient data.

As most patients were diagnosed clinically with WS, diagnoses given by the original authors were carefully re-evaluated based on the presence of the following phenotypes: unusual body habitus, bilateral cataracts, skin sclerosis, painful corns, sarcopenia, metastatic subcutaneous calcification, skin ulcers, DM, and hyperlipidaemia (7,13,14).

#### 3. Results and Discussion

A total of 1,070 cases documented in 500 Japanese articles published between 1917 and 2004 were included in the analysis. Outside Japan, 350 cases of WS have been reported. As shown in Table 1, 1 case of WS associated with Klinefelter syndrome was diagnosed by chromosomal testing in Japan (22) and one was similarly diagnosed outside Japan (23). Several WS cases have been diagnosed as pseudo-Klinefelter syndrome because of several clinical similarities such as slender extremities with stocky trunk, gynecomastia, and atrophic testis (24, 25). However, chromosomal analysis of those cases indicated that they were normal males (26). The frequency of Klinefelter syndrome was

Table 1. Mutation causes mutations?

Klinefelter syndrome	27M; 46/XY, 47/XXY: Funayama 1976 24M; 48/XXYY: Ferramosca 1972
Extrachromosome	43 F; 46/XX, 49/XX; Tanihara 1986
Retinitis pigmentosa	58M; Tajima 1993 39M; Valero 1960 38M; Kleeberg 1949
Wilson disease	55M; Sakai 1990
Xeroderma pigmentosum	23F; Takahashi 1955 29F; Takahashi 1955 29F; Takahashi 1955
Porokeratosis Mibelli	49M; Machino 1984

about 1 in 700 live-born males in the general population, and the frequency among WS patients was about the same (27). In addition, one WS case of a patient with an extra chromosome was reported by Tanihara *et al.* in Japan (28).

In 1 Japanese case and 2 European cases of WS, patients had retinitis pigmentosa (29-31). Retinitis pigmentosa has several types of transmission and an overall frequency of about 1 in 3,700 (32). Thus, an incidence of 2:350-1:1070 among WS patients was probably higher than in the general population.

In 1 Japanese case of WS, the patient had Wilson disease (33). The frequency of Wilson disease was about 1 in 30,000-100,000 livebirths worldwide (32). Of particular interest, 3 female siblings with WS may have the variant form of xeroderma pigmentosum (X-P) (34,35). The frequency of all types of X-P in Japan was about 1 in 40,000, which was higher than in the rest of the world (27). However, both the cases of WS and of X-P were clinically diagnosed 50 years ago.

WS has been classified as a genetic instability syndrome, which includes Bloom syndrome (36,37), Rothmund-Thomson syndrome (38), Cockayne syndrome (37), ataxia telangiectasia (40,41), X-P (39), Fanconi anemia (Alter BP, NCI personal communication), and progeria (20). However, no association of additional chromosomal abnormalities or genetic diseases with genetic instability syndromes was noted except for WS and Bloom syndrome. Machino reported a case of WS with porokeratosis Mibelli, and Takemiya described a case of Bloom syndrome with porokeratosis Mibelli (42,43). The frequency of porokeratosis Mibelli in the general population is not known.

In all of the cases analyzed, WS and additional genetic diseases such as retinitis pigmentosa, Wilson disease, and X-P were clinically determined, which merely suggests that WS has genetic instability when encountered clinically.

Finally, some of the aging-associated phenotypes seen may relate directly to WRN dysfunction. Aging is believed to induce genetic instability leading to cancer (44,45), and thus the complete loss of WRN function may epigenetically and genetically impact other genes, promoters, or proteins related to agingassociated pathophysiology. It may also impact several disease-causing genes *via* acquired *in vivo* mosaicism or acquired *in vivo* mutation, as is reported in Rothmund-Thomson syndrome (*38*).

Medical researchers are encouraged to report cases of other genetic diseases or chromosomal abnormalities accompanying WS, as doing so may help to identify which diseases are associated with WS.

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