

# Cardiovascular Phenotype in Turner Syndrome—Integrating Cardiology, Genetics, and Endocrinology

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Cardiovascular disease is emerging as a cardinal trait of Turner syndrome, being responsible for half of the 3-fold excess mortality. Turner syndrome has been proposed as an independent risk marker for cardiovascular disease that manifests as congenital heart disease, aortic dilation and dissection, valvular heart disease, hypertension, thromboembolism, myocardial infarction, and stroke. Risk stratification is unfortunately not straightforward because risk markers derived from the general population inadequately identify the subset of females with Turner syndrome who will suffer events. A high prevalence of endocrine disorders adds to the complexity, exacerbating cardiovascular prognosis. Mounting knowledge about the prevalence and interplay of cardiovascular and endocrine disease in Turner syndrome is paralleled by improved understanding of the genetics of the X-chromosome in both normal health and disease. At present in Turner syndrome, this is most advanced for the *SHOX* gene, which partly explains the growth deficit.

This review provides an up-to-date condensation of current state-of-the-art knowledge in Turner syndrome, the main focus being cardiovascular morbidity and mortality. The aim is to provide insight into pathogenesis of Turner syndrome with perspectives to advances in the understanding of genetics of the X-chromosome. The review also incorporates important endocrine features, in order to comprehensively explain the cardiovascular phenotype and to highlight how raised attention to endocrinology and genetics is important in the identification and modification of cardiovascular risk. (*Endocrine Reviews* 33: 677–714, 2012)

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## I. Introduction

Turner syndrome (TS) occurs in 50 per 100,000 live-born females (1). The principally clinical diagnosis rests on the coexistence of partial or complete absence of an X-chromosome with classical traits that include reduced final height, estrogen deficiency, and infertility (2). The highly varied phenotype causes diagnostic delay as

well as nondiagnosis, with 30% of females with TS never obtaining correct diagnosis (1). Failure to diagnose TS is highly unfortunate because morbidity and mortality are a great deal higher than in the general population (1, 3, 4). Additionally, clinical assessment is poor even in diagnosed individuals (5), although multidisciplinary practice may improve this (6). Therefore, both appropriate identification of girls and women with TS and subsequent implementation of prophylactic measures to reduce morbidity and mortality leave much to be desired.

Risk assessment in TS is compromised by insufficient insight into the prevalence and causes of different syndrome-associated traits that may impact adversely on prognosis. This is especially the case for cardiovascular contributions to the excess all-cause mortality, where congenital and acquired heart diseases are felt to contribute to 8 and 41% of all-cause mortality, respectively (4). Cardiovascular risk in TS is predominantly stratified from evidence gathered on risk markers in the background population because insight into the cardiovascular phenotype in TS is limited, and previous guidelines have relied on expert consensus only (2, 7). This is unfortunate because complex patterns of not only cardiovascular but also endocrine diseases in TS render direct translation of evidence from other cohorts hazardous (8). The risk burden is so severe that TS is proposed as an independent risk factor for cardiovascular disease (9). More educated risk stratification and more appropriate clinical care can only be facilitated through a thorough delineation of the endocrine and cardiovascular phenotype in TS.

This review provides insight into the cardiovascular and endocrine phenotype in females with TS and presents an up-to-date condensation of current state-of-the-art knowledge of the cardiovascular phenotype, emphasizing the significance of not just congenital but also acquired pathologies. The aim is to provide an update on current insight into the pathogenesis of TS in relation to recent advances in the understanding of X-inactivation and its impact on female health. Moreover, the review provides an updated hypothesis on the genetic etiology of TS, highlighting our knowledge of the significance of X-chromosomal haploinsufficiency to congenital and acquired cardiovascular and endocrine traits. Finally, the review incorporates important endocrine features of TS, accounting for how genetics may explain the prevailing phenotype and that attention to endocrine factors is important in our efforts to identify and modify cardiovascular risk markers.

The full PubMed database was searched (without time restrictions) in December 2011 using the keyword “Turner syndrome” as MeSH term, as well as “Turner syndrome,” “Turner’s syndrome,” “Turner,” and “Turner’s” in titles and abstracts. Articles relevant to the topic were obtained

and reviewed, as well as older articles selected by the authors. Publications cited in this review were selected from those identified by the searches at the authors’ discretion.

## II. Morbidity and Mortality

Increased morbidity and mortality are cardinal traits in TS (1, 3, 4, 10). The risk of premature death is increased 3-fold [standardized mortality ratio (SMR), 3.0] (1, 4), and life expectancy is reduced by at least a decade (1, 4, 11). The early-life disease burden is high, with an inverse relation between the age achieved by a girl or woman with TS and her reduction in life expectancy (11). The inferior outcomes are attributed to endocrine, nervous, cardiovascular, respiratory, digestive, and genitourinary organ disease (Fig. 1) (1, 3, 4). Of these, cardiovascular disease is increasingly recognized as the principal component in the causation of decreased life expectancy (1, 3, 4) with half of the excess morbidity attributed to cardiovascular pathology (4) (Figs. 1 and 2). In TS, cardiovascular morbidity and mortality are attributed to:

- Congenital heart disease (SMR, 20.7)
- Aortic dilation and dissection (SMR, 23.6)
- Ischemic heart disease (SMR, 2.8)
- Cerebrovascular disease (SMR, 3.9) (1, 3, 4, 11)

### A. Karyotype-phenotype correlations

All-cause mortality is raised in 45,X when compared with mosaic karyotypes (1, 4). An exacerbated prognosis is also to some extent seen in the presence of an isochromosome, and most likely also in karyotypes with Y-chromosome material. Yet, all females with TS face increased morbidity compared with the general population (3), although patterns of specific pathologies may differ between karyotypes. An increased risk of cardiac congenital anomalies has been demonstrated in 45,X (3, 12, 13), and karyotypes with an isochromosome suffer from an increased risk of diabetes and hypothyroidism (4, 14). Karyotypes other than 45,X associate with an increased risk of noncongenital cardiovascular disease (3).

The patterns of morbidity and mortality appear to be changing over time, with a trend toward a reduction in mortality (1, 4). Changes in the composition of diagnosed karyotypes are likely to contribute to this because a larger proportion of less severe phenotypes with mosaic karyotypes is now more often being correctly diagnosed (15). Especially mosaic karyotypes encompass a larger fraction (16), where cell lines of normal karyotype may associate with more favorable profiles (17). Improvement in management in the diagnosed females may also contribute to a seemingly improved prognosis. Irrespective, the prognostic improvement is small, and it represents only a trend

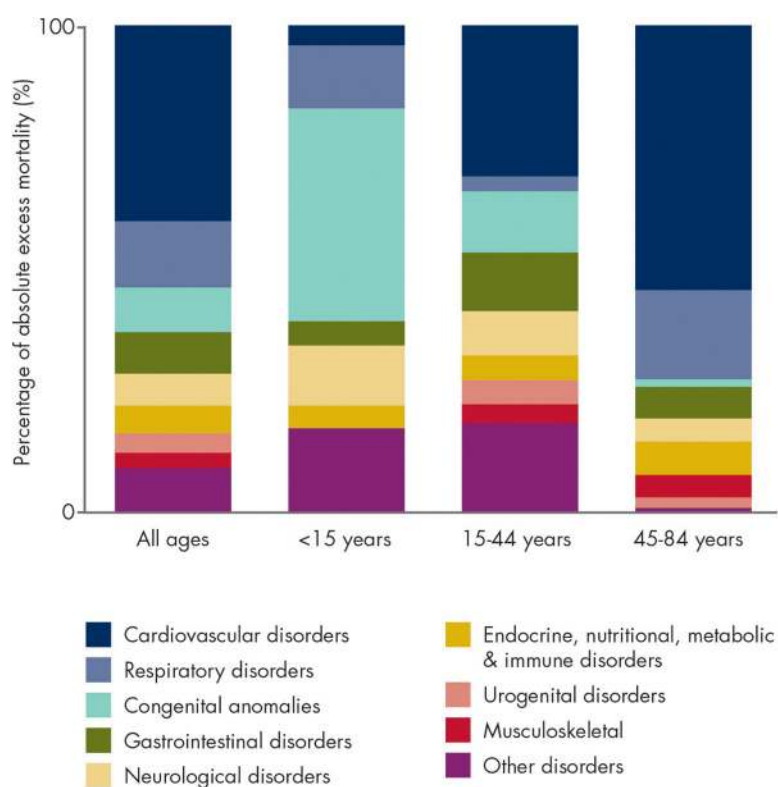
**Figure 1.**

Figure 1. Differentiated mortality in TS for all ages and according to age groups (4). Categories were defined according to International Classification of Diseases 9. Numbers are adapted to express the percentage of total absolute excess risk caused by the group of disorders in question.

in a cohort with grossly increased risk of premature death (1). Important to risk prophylaxis, many females with TS are diagnosed after a delay of many years (1).

## B. Genetic background

The advent of TS is caused by *de novo* meiotic or mitotic nondisjunction that results in a spectrum of karyotypes (Table 1) (3). There can be complete absence of an X-chromosome, a structurally abnormal X-chromosome (isochromosomes or ring chromosomes), different levels of abnormal karyotypes within the body tissues (mosaicism), and karyotypes with Y-chromosome material (Table 1). Maternal risk factors for offspring with TS have been proposed to include advanced maternal age and low maternal height, although predictive values of these factors are weak for a fetus with TS (18). Also, no correlation with maternal age or even higher risk in younger mothers has also been reported (16, 19). The risk of repeated pregnancy with a fetal karyotype compatible with TS is generally not considered raised, although rare familial occurrences have been described, and a recent case series did suggest an increased risk (20, 21).

The correct number of sex chromosomes (X- and Y-chromosomes) and their correct gene expression are essential for normal health (22). In keeping with this, increased morbidity and mortality are key features of structural and numerical disorders of the sex chromosomes (1, 23, 24). Structural or numerical abnormalities of the Y-chromosome are compatible with life, whereas complete lack of X-chromosome material is incompatible with fetal survival. In fact, even the complete lack of one X-chromosome in a female may not be feasible either, and it has been hypothesized that 45,X females are indeed low-level mosaics (25, 26). Concordantly, *in utero* lethality has been proposed to be especially increased for karyotypes with 45,X monosomy, with as many as one in 100 of all recognized early pregnancies being 45,X and more than 99% of these being spontaneously aborted (27). This process of natural selection occurs predominantly in early pregnancy and leads to a prevalence of karyotypes compatible with TS of 50 per 100,000 live births (1).

## C. X-Inactivation in Turner syndrome

The 1098 known genes on the X-chromosome by far exceed the 78 genes on the male Y-chromosome (28). This large difference in gene content exists despite the sex chromosomes stemming from a common progenitor autosome (29). Full transcription of all genes on both X-chromosomes in female somatic cells, as opposed to transcription from the single X- and Y-chromosomes in somatic cells of males, is detrimental. This delivers excess transcriptional products from the two X-chromosomes in females, when compared with the somatic male counterpart (30). Therefore, somatic cells in females are subject to gene dosage compensation in early embryonic development to equalize effective dosage of X-linked genes (31). This process is termed X-inactivation, and it involves packaging of transcriptionally active chromatin into “silenced” heterochromatin (31). The X-inactivation center is essential to gene dosage compensation and is comprised of a cluster of genes situated near the centromere of the long arm of the X-chromosome (Xq). This center encodes genes that are prerequisite to X-inactivation, including more types of noncoding RNA and regulatory protein binding sites (32). Among these, the X-inactive specific transcript (*XIST*) gene is critical to X-inactivation (33) because it transcribes noncoding *XIST* RNA that coats the X-chromosome and through methylation and histone modifications represses euchromatin to

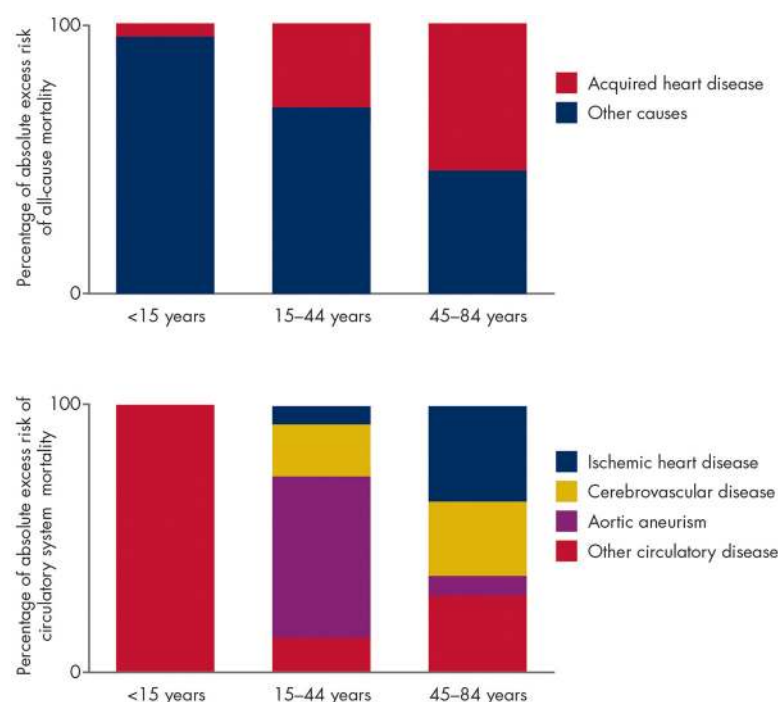
**Figure 2.**

Figure 2. Acquired heart disease in TS. Acquired heart disease accounts for a large proportion of the 3-fold excess mortality, increasing in relative contribution with age in TS (*top panel*). The contributions from ischemic heart disease, cerebrovascular disease, aortic aneurysm, and other heart diseases are shown *below* (4).

form the inactive X-chromosome (34). Initially, both X-chromosomes transcribe *XIST* RNA and an antagonistic noncoding *TSIX* RNA from the X-inactivation center (35). By general random selection of the maternal or paternal X-chromosome, the balance of noncoding RNA transcribed from the X-inactivation center shifts toward *XIST* in the X-chromosome that is subsequently inactivated. The other X-chromosome maintains *TSIX* transcription during the X-inactivation maintenance phase and avoids silencing (36). The random X-inactivation process introduces in the normal female an obligatory mosa-

icism for the paternal and maternal X-chromosome across somatic cell lineages (37).

The inactivated X-chromosome stays inactive in the specific cell clone (37). This equalizes the effective gene dosage to that of males, with only female germ cells having the potential to reactivate an inactivated X-chromosome (37). However, approximately 25% of genes on the inactivated X-chromosome escape transcriptional silencing to varying extents, depending on the tissue, individual characteristics, and age (38–40). Escapees include two pseudoautosomal regions (PAR1 and PAR2) located at the termini of Xq and the short arm of the Xp, which are common with the Y-chromosome (Fig. 3). Several genes outside these two regions also escape silencing, whereof 25 genes are X/Y homologous and others again have lost their Y-chromosomal counterpart (paralog genes) (28). The relative expression of nonsilenced genes on the inactive X-chromosome relative to the genes of the active X-chromosome has not been clarified, but expression of escape genes from the inactivated X-chromosome may be reduced compared with the active (38, 41).

In TS, complete or partial loss of activity of the entire or even parts of the X-chromosome detrimentally impacts both prenatal and postnatal survival (1, 27). The mechanisms of gene dosage compensation are complex even in normal health, and much still needs to be resolved. In TS, the prevailing traits may result from decreased expression of paralog genes or altered expression levels of homolog genes from within or outside PAR (or a combination of these). Additional variability between females with TS (and even between cell clones in the same females due to mosaicism) may be introduced through different patterns of X-inactivation. In 45,X the one normal X will inadvertently

**TABLE 1.** Postnatal and prenatal prevalence of TS

		Prenatal, % (n)	Postnatal, % (n)
n		383	1063
Monosomy	45,X	66% (254)	40% (426)
Mosaic	45,X/46,XX	23% (90)	24% (257)
Isochromosomes	45,X/46,X,i(Xq); 46,X,i(Xq); 45,X/46,X,i(Xq)/47,X,i(Xq),i(Xq), etc.	3% (11)	10% (107)
Deletions	45,X/46,X,del(X); 46,X,del(X)	3% (13)	6% (69)
Polyploidy	45,X/46,XX/47,XXX; 45,X/47,XXX; 45,X/46,XX/47,XXX/48,XXXX	3% (13)	11% (117)
Ring chromosomes	45,X/46,X,r(X)	<1% (2)	3% (34)
Y material	45,X/46,XY, other with Y material		5% (51)

National prevalence of TS updated in 2011 from the Danish Cytogenetic Central Register (previously unpublished data), where 72–82% chose termination of the pregnancy with a trend to fewer terminations in the last half of the study period. Note is made of a decrease in the proportion of 45,X postnatal karyotypes from the previously published data in Denmark: 47% (1996) (16) to 40% now. Prenatal prevalence was gathered from a screening program of recognized pregnancies using amniocentesis and chorionic villus sampling.

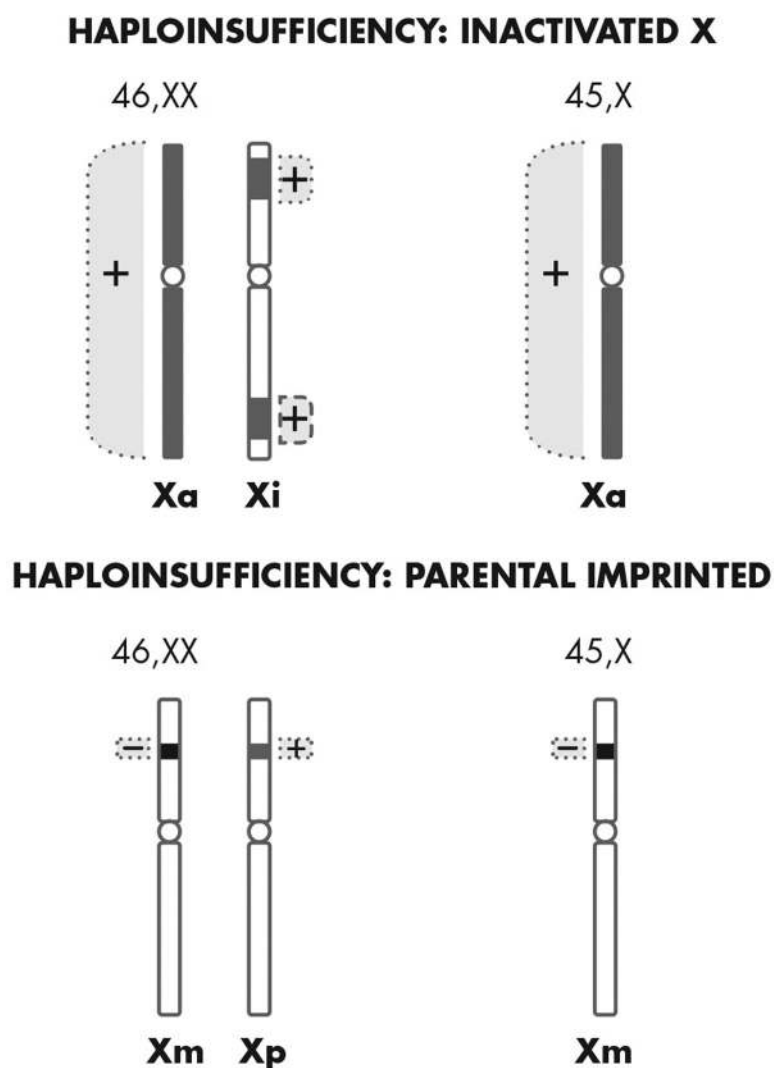
**Figure 3.**

Figure 3. Models of genetic etiology in TS. Haploinsufficiency for the X-chromosome in TS may result in an abnormal phenotype through different mechanisms. In the X-inactivation hypothesis (31), haploinsufficiency for genes that normally escape X-inactivation on the otherwise inactivated X-chromosome (Xi) result in abnormal gene dosage. These escapees are mainly located in PAR1 and -2 on the long and short arms of the X-chromosomes. Thus, only genes from the activated X-chromosome (Xa) are expressed in the haploinsufficient female with TS. In the imprinting hypothesis (59), genes on the X-chromosome are expressed in a monoallelic fashion depending on paternal or maternal origin. In TS, lack of activated allele will lead to loss of expression for that gene. This is shown here for the loss of the normally expressed paternal allele (Xp) with the nonexpressed maternal allele (Xm) left in TS.

be active, whereas in structural abnormalities such as isochromosomes or ring chromosomes, inactivation of the normal X-chromosome may further aggravate the phenotype (37). The evolutionary younger Xp is subject to more escapees from inactivation than Xq (29), and structural abnormalities involving the Xp may thus carry a more adverse impact, as has been proposed for aortic dilation in TS (42).

It has become evident that a sexual dimorphism in autosomal gene regulation is present, and that more than half of all genes in a tissue are differentially expressed, although the difference in expression in the majority of these genes is only in the order of a 1.2-fold difference (30). Historically, differences between the sexes were mainly attributed to differences in the level of sex hormones, but genetic factors seem also to be involved (43). New research shows that sexual dimorphism is determined directly by the sex chromosome complement, rather than the specific sex, in addition to a modulatory role of the male-determining *Sry* gene at least in mice (44). In this study, the authors also examined differences between male  $XX^{Y-}$  mice (male mice where the *Sry* gene is translocated to an autosome, thus with the same sex complement as a female) and XO mice (mice with “TS”), and found distinct differences in gene expression (44). It is not clear how X-chromosome complement causes this sexual dimorphism in gene expression, but candidate genes are genes that escape silencing (40), such as *UTX* (ubiquitously transcribed tetratricopeptide repeat gene on X-chromosome) and *JARID1C* (a histone methylase), which have previously been anticipated to play a role in the pathogenesis of TS (45, 46). Thus, if confirmed in humans, such a general mechanism of influence of sex chromosome complement on autosomal gene expression could explain features present in TS.

#### D. Candidate genes

Attempts to map candidate genes that escape gene dosage compensation in TS and other disorders of the sex chromosomes are ongoing. The short stature homeobox-containing gene on chromosome X (*SHOX*) situated in PAR1 on Xp has been implicated in growth retardation and bone changes (47) and is a homeodomain transcription factor. Brain natriuretic peptide and fibroblast growth factor receptor 3 are transcriptional targets of *SHOX* (48), and thus pose a link that may be of relevance to the etiology of cardiovascular disease in TS (49). Importantly, penetrance of gene defects in TS differs from X-linked recessive disorders. This is evident in a comparison with *SHOX* haploinsufficiency in Leri-Weill dyschondrosteosis, where a *SHOX* gene mutation (a range of single gene deletions) causes only 50–75% of the height retardation encountered in TS, and congenital heart disease is not a feature of Leri-Weill dyschon-

drosteosis (50). Putatively, the missing or structurally abnormal X-chromosome delivers a more profound impact on gene expression in TS than in monogenetic disorders through mechanisms that are unaccounted for (51). Hence, a range of features is known to associate with *SHOX* haploinsufficiency in TS:

- Reduced final height
- Sensorineural hearing impairment
- High-arched palate
- Short fourth metacarpal
- Madelung deformity of the radius
- Skeletal disproportionality (51)

Altered transcriptional expression of pseudoautosomal genes in PAR1 and PAR2 other than the *SHOX* gene is also likely to carry an impact in TS, with causative loci for various cardiovascular, endocrine, skeletal, or psychological traits awaiting discovery. Most PAR genes that escape X-inactivation are to be found on Xp, where 24 genes have been identified compared with only five on Xq (28). Thus, structural abnormalities involving unbalanced translocations or deletions of Xp are expected to more severely affect the phenotype than those involving Xq (37, 52). The consequences of haploinsufficiency for X/Y gene pairs beyond PAR1/PAR2 or the absence of X heterologous genes in TS remains undefined (30). Recently, work in pluripotent embryonic stem cells from amniocytes of a TS fetus showed that the pseudoautosomal genes *ASMTL* and *PPP2R3B* were expressed in lower levels than normal cells (53). *ASMTL* may be related to methyltransferase activity (and could thus be involved in proper methylation of the genome), whereas *PPP2R3B* is involved in cell cycle control (53). Insufficient up-regulation of another pseudoautosomal gene, *CSF2RA*, during embryoid body formation was also demonstrated, consistent with data showing lower expression of this gene in embryonic stem cells with TS (54). Decreased expression of *CSF2RA* points to insufficient placentation as an explanation for the increased early fetal lethality in TS. Interestingly, the TS-induced pluripotent embryonic stem cells differentiated normally into neural-like, cardiomyocyte-like, and hepatocyte-like cells, although the cardiomyocyte-like cells displayed a spontaneous action potential that was slightly longer compared with controls, but this was not statistically significant (53). These findings may help to explain the prolonged QT interval seen in TS (55).

The potential genetic mechanisms are complex, and disease may very well be the result of deregulation within different pathways that work in concert to cause abnormality. The phenotype of TS may result from loss of proteins involved in ordinary cellular function, which are normally expressed from the inactive X-chromosome. Al-

ternatively, loss of coding genes on an absent, inactive X-chromosome may alter gene expression from the active X-chromosomes. Loss of biallelic gene expression by either of the aforementioned mechanisms may directly impact the TS phenotype through altered expression of genes that regulate tissue genesis and maintenance, or more indirectly affect tissue phenotypes through loss of transcriptional up- or down-regulation of autosomal or even other X-linked genes. Several genes and critical regions have been proposed in TS, but the exact delineation of their roles in TS remains to be determined (54, 56–58). The same applies to the discovery of other tentative loci involved in the causation of the many cardiovascular, endocrine, skeletal, or psychological traits.

#### E. Parental imprinting of genes on the X-chromosome

Although haploinsufficiency for X-chromosomal genes that escape silencing may cause the abnormal phenotype of TS, an alternative hypothesis suggests that parental imprinting of gene expression profiles from the X-chromosome may be involved in the causation of TS. This hypothesis is based on the fact that whereas the majority of genes are expressed from either the paternal or maternal allele, some autosomal genes are expressed in a parent of origin-dependent fashion (59). Loss of the active (imprinted) allele has been implicated in diseases such as Prader-Willi syndrome (60). In TS, this process has been associated with the X-chromosome, and some patterns of morbidity may associate with loss of either the maternal or paternal X-chromosome (Fig. 3). In the imprinting hypothesis, loss of a homolog gene on the X-chromosome will lead to null expression of this gene when it is the active (imprinted) allele. The paternal X-chromosome is absent in approximately 60–80% of 45,X monosomy, with a more equal paternal *vs.* maternal X-chromosomal deficiency in mosaic and isochromosomal karyotypes (22, 61–63). The impact of parental X-chromosome imprinting in TS is not resolved. A determinant impact has been reported on aortic valve morphology and coarctation of the aorta (63), lipid levels, visceral and abdominal fat (62), white and gray matter cerebral volume (64), and neuropsychological profile (65). Conversely, other studies have not been able to reproduce these findings (22, 66). Thus, the genetics of TS is probably more complex than previously thought and could involve processes like:

- The X-inactivation process and X-chromosome dosage compensation
- Candidate genes other than *SHOX* are probably relevant in TS
- The role of haploinsufficiency of escape genes is not yet clear

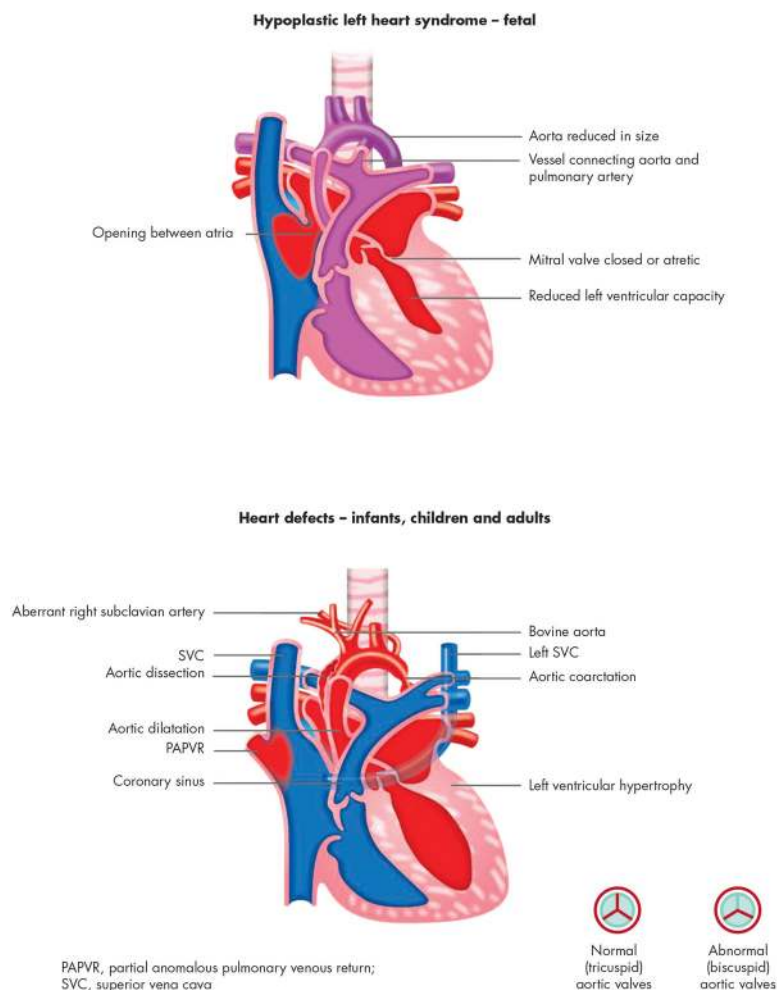
**Figure 4.**

Figure 4. The *upper figure* illustrates a fetal heart with hypoplastic left heart syndrome, a condition often seen during fetal life in TS and probably contributing to the high spontaneous abortion rate. The *lower figure* illustrates the most common congenital cardiovascular malformation seen in TS. The many different malformations and acquired conditions are not necessarily present at the same time in any given individual (see text for further information). PAPVR, Partial anomalous pulmonary venous return; SVC, superior vena cava.

- Genotype-phenotype relations are not yet fully uncovered
- Parental imprinting
- Differential methylation patterns

### III. Congenital Heart Disease

The risk of congenital heart disease is considerable [relative risk (RR), 8–50] (67–69). The disease spectrum spans from trivial defects to severe and highly complex disorders, and accordingly some lesions may be asymptomatic (67, 70), whereas others are incompatible with normal life expectancy or even fetal survival (4, 71). An estimated 22–70% of all females with TS have a form of congenital

heart disease (12, 67–69, 72) with congenital anomalies coexisting in other organs in 22% (12). Reports on the prevalence of congenital heart disease are highly dependent on the type of investigation used, spectrum of assessed anomalies, and karyotype distribution within the studied cohort. Excess early life mortality due to severe congenital heart disease may also cause selection bias into studies performed in adolescence and adulthood through survival with advancing age of less severe cardiac phenotypes (11).

#### A. Bicuspid aortic valve

A bicuspid aortic valve is found in 15–30% of TS patients (52, 67–69, 72), compared with 1–2% of the general population (Fig. 4) (73). The prevalence would appear to be increasing due to higher diagnostic sensitivity achieved with cardiac magnetic resonance when it comes to distinguishing the tricuspid from bicuspid aortic valves (52). Congenital aortic valve anomalies of an immediately severe nature may rarely present in the infant with TS, such as aortic valve dysplasia and congenital aortic stenosis (69, 74).

Outcomes with a bicuspid aortic valve are generally favorable in childhood, with the likelihood of associated morbidity increasing with age (73). In general, for the bicuspid valve, accelerated valve calcification often promotes premature valvular stenosis. Additionally, poor leaflet coaptation can lead to aortic regurgitation, which may become aggravated in a vicious circle of increased stroke volume and aortic root dilation (73). In view of this general population evidence, it is not surprising that aortic valve dysfunction is frequent in adults with TS; aortic valve stenosis is seen in 4–16% and regurgitation in 6–45% (52, 67, 72). For comparison, trivial and clinically irrelevant aortic regurgitation and stenosis may be seen in 11 and 6% of healthy females with tricuspid aortic valves, respectively (67). As expected, the risk of valvular dysfunction in TS links closely with the presence of bicuspid aortic valves (RR, 7.9 for stenosis; and RR, 4.2 for regurgitation), and the bicuspid valve morphology is the principal cause of all moderate and severe dysfunctions (52, 67). Pediatric studies of aortic valve function in TS are limited, but stenosis is observed in 28–46% and regurgitation in 42–50% of adult females with TS and a bicuspid aortic valve (67, 75). Unfortunately, functional grading has only been the focus of a few

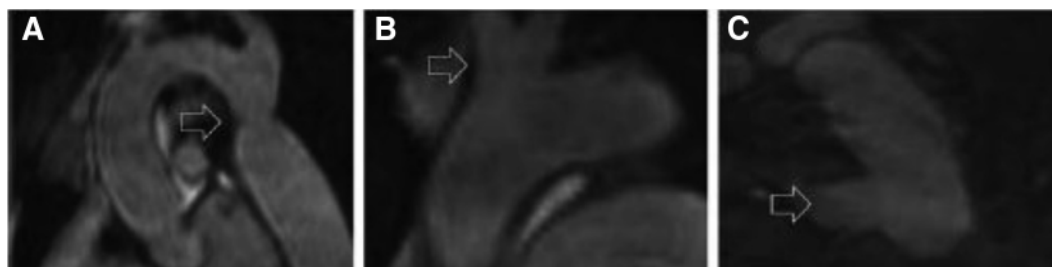
**Figure 5.**

Figure 5. Congenital abnormalities of the thoracic aorta in TS. Aortic coarctation (A), common origin of the innominate and left common carotid arteries (B), and an aberrant right subclavian artery (C) as seen by cardiac magnetic resonance in TS (67). [Reproduced from K. H. Mortensen et al.: Abnormalities of the major intrathoracic arteries in Turner syndrome as revealed by magnetic resonance imaging. *Cardiol Young* 20:191–200, 2010 (67) with permission. © Cambridge University Press.]

studies (52, 67). Hence, insight into the general state of the aortic valve in TS is scarce, which also includes our understanding of the onset of clinically significant disease, its optimal treatment, and to a large extent also the relationship between aortic valve function and aortic dilation.

Raphe configuration of the bicuspid aortic valve may be of prognostic importance for the development of aortic valve dysfunction and ascending aortic dilation (76). The right and left aortic leaflets are mainly fused in TS (82–95%) (52, 67), which, compared with leaflet fusions that involve the noncoronary cusp, has been indicated as a positive prognostic omen in some non-TS cohorts (73). Hitherto in TS, raphe configuration has nevertheless not been demonstrated to determine aortic valve function or aortic morphology (52, 67). Important to the clinician, the annual incidence of endocarditis in the general population is increased to 0.3–2% with bicuspid aortic valves (73). Therefore, due to the frequent occurrence of not just bicuspid aortic valves but also congenital heart disease in TS, the risk of endocarditis must also be assumed to be increased in these females, although the evidence hereof is limited (77).

### B. Coarctation of the aorta

Coarctation of the aorta is found in up to 17% of females with TS (13, 52, 67, 78), compared with 0.04% in the general population (Fig. 4) (79). The aortic obstruction often coexists with a bicuspid aortic valve (RR, 4.6) (52, 67), and coarctation of the aorta in TS is increasingly regarded as a part of an abnormal aortic phenotype (Figs. 5 and 6). This phenotype includes features such as an elongated and kinked transverse aortic arch (~50%), an aberrant right subclavian artery (~8%), a bovine aorta (~8%) (Fig. 4), and isolated cases of cervical aortic arch (13, 67, 80).

The severity of coarctation of the aorta in TS has received little attention. The prevalence of repaired coar-

tation of the aorta amounts to 5–12% in childhood and adulthood (13, 78, 81). In addition to this, undiagnosed coarctation of the aorta has been demonstrated by cardiac magnetic resonance in 5–8% (13, 67, 75). In reports of undiagnosed cases, the diagnosis is most often made by morphological analysis as a concentric narrowing with a posterior shelf-like morphology located typically in the descending aorta. These occult lesions have in all but one severe case been mild when assessed by cardiac magnetic resonance (neither arterial collaterals nor flow acceleration to warrant surgical intervention has been present) with very limited data from echocardiography or cardiac catheterization studies (13, 75, 80). Coarctation of the aorta may be seen in conjunction with an elongated aortic arch (13) or more adverse aortic arch phenotypes that include aortic arch hypoplasia and interrupted aortic arch (67, 80, 82).

Little is known about the optimal corrective technique of coarctation of the aorta in TS. Some surgical centers have hypothesized that aortic wall frailty increases the risk of perioperative hemorrhage and rupture of suture lines after end-to-end resections (observed in 25–38%) (83, 84). Favorable outcomes have been reported with patch angioplasty repair (83, 85). Percutaneous correction was encouraging in a very small series (86), although aortic dissection and pseudoaneurysm formation have been reported after balloon or stent repair (87–89). Most likely, the optimal surgical technique should be decided on a patient-to-patient basis and follow general experiences. Importantly, the risk of aortic dissection associated with the coarctation is not completely relieved with repair because dissections may occur even years after repair (90), which is in keeping with experiences in the correction coarctation of the aorta outside TS (91).

### C. Associated lesions

Several mild congenital heart defects can be found in TS (Fig. 3). These include:



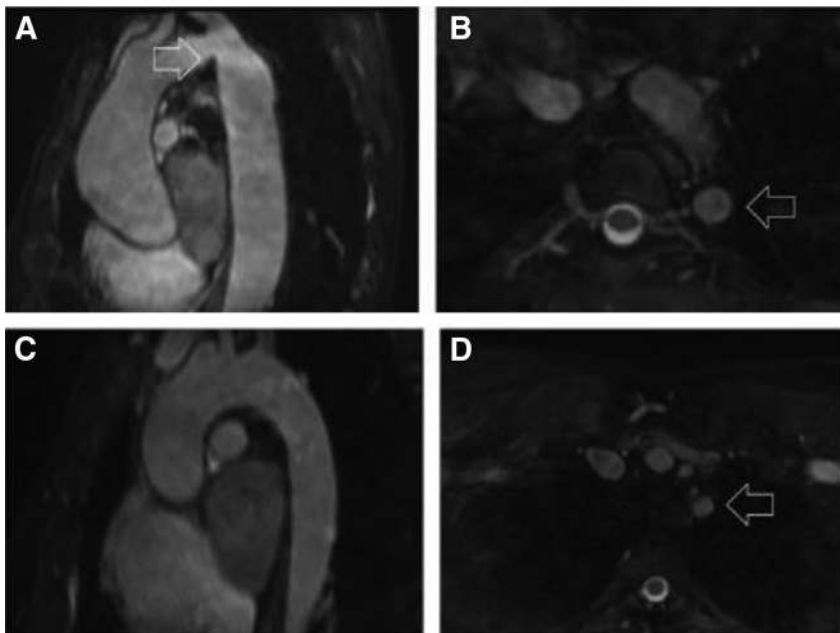
**Figure 6.**

Figure 6. The transverse aortic arch phenotype in TS. The normal and elongated transverse aortic arch as seen by cardiac magnetic resonance imaging in TS (67). The abnormality is defined by the coexistence of kinking of the inferior aortic contour at the aortic isthmus (A) and a left subclavian artery origin from a position posterior to the trachea in the horizontal plane (B). The normal transverse aortic arch (C) and normal origin of the left subclavian artery (D) are illustrated for comparison; kinking is absent, and the left subclavian artery departs anterior to the posterior aspect of the trachea. [Reproduced from K. H. Mortensen *et al.*: Abnormalities of the major intrathoracic arteries in Turner syndrome as revealed by magnetic resonance imaging. *Cardiol Young* 20:191–200, 2010 (67) with permission. © Cambridge University Press.]

- Atrial septal defects (in 1–2%; RR, 19) (12, 69, 75)
- Ventricular septal defects (in 1–4%; RR, 49) (12, 69, 75)
- Partial anomalous pulmonary venous drainage (in 13–15%; RR, 20) (13, 92) (Fig. 4)
- Persistent left superior vena cava (in 8–13%) (13, 92) (Fig. 4)
- Persistent arterial duct (72)
- Interrupted inferior vena cava with azygous continuation (80)
- Pulmonary valve stenosis (72)

Prenatally, severe left-sided cardiac malformations pave the way for 45,X conceptions comprising up to 10% of all recognized early miscarriages (27, 71, 93–95). This inferior outcome adds to a reported general inability to form an embryo proper with TS karyotypes (27, 53, 54). Spontaneous abortion occurs in 99% of all fetuses with 45,X TS, with the prevalence of TS dropping drastically from 1000 per 100,000 of all conceptions to 50 per 100,000 live births (1, 27). Fetal echocardiography performed in TS in gestational wk 11–15 has shown coarctation of the aorta (45%), hypoplastic left heart syndrome

(Fig. 4) (13%), and atrioventricular septal defect (one fetus) (71). These adverse findings were confirmed by an autopsy study, which additionally revealed the presence of unicuspid aortic valves (15%) or bicuspid aortic valves (85%), whereof all but one were 45,X karyotypes (94). Moreover, midgestation fetuses have myocardial hypoplasia (96), but this must be read in a context of the general state of small for gestational age in TS (18).

At birth, a range of complex congenital heart diseases can be found. Apparently, these severe lesions associate with an even more inferior prognosis than when occurring in the general population (97, 98). They include:

- Hypoplastic left heart syndrome (97) (Fig. 4)
- Congenital aortic valve stenosis (99)
- Neonatal coarctation of the aorta (12)
- Atrioventricular septal defects

Emphasizing the complexity of congenital heart disease in TS, lesions can be seen in both monosomy 45,X (12, 52, 67) and mosaic karyotypes (100). Furthermore, characteristic exterior morphological traits may or may not be

present to direct the attention toward congenital heart defects, and there is not necessarily complete concordance between external phenotype and congenital heart disease (12, 101).

#### D. Etiology of congenital heart disease

Early prenatal screening for TS relies on fetal sonography performed at gestational wk 11–14 (102). Nuchal translucency may often present in karyotypes compatible with TS, and this fetal appearance generally predicts chromosomal abnormalities and poor fetal outcome (including congenital malformations of the heart and skeleton) (103, 104). The remnant of this lymph accumulation in the lower neck region is appreciated postnatally as neck webbing in TS. Because this external feature cosegregates with congenital heart disease in TS (12, 52, 101), a causal relation between lymph accumulation and congenital heart disease has been proposed (105). Perturbations of flow and pressure are hypothesized to lead to left-sided defects by a mechanistic chain of events (106) that starts with abnormal fetal lymphangiogenesis (107, 108). The pres-

ence of severe edema in aborted TS fetuses with anomalies of the left ventricle and the aorta lends promise to this hypothesis (109). Yet, separate mechanisms have not been ruled out because postnatal neck webbing and congenital heart disease do not always coexist (12, 52). Furthermore, changes in the lymphatic vessels around the major vessels and the heart have not been demonstrated to be uniformly present in aborted fetuses with heart disease in TS (although most were examined after primary cardiac organogenesis) (109).

Insight into the genetic regulatory mechanisms behind formation and maturation of the thoracic aorta, the brachiocephalic vessels, the arterial duct, and the proximal pulmonary arteries from the five pharyngeal arch arteries may promote our understanding of congenital heart disease in TS. TGF- $\beta$ 2 could be of key importance in TS because knockout mice deficient of TGF- $\beta$ 2 suffer from obstructive lesions in the fourth pharyngeal arch artery (110) in a similar pattern to TS (94, 111). Also potentially perturbed in the fetus with TS, reduced TGF- $\beta$ 1 release from the endothelium in the context of low blood flow and pressure may adversely influence normal angiogenesis, which normally involves hormonal stimulation of migration and differentiation of cardiac neural crest cells into blood vessels to form vascular smooth muscle cells (112). Recent data indicate that TGF- $\beta$ 1 and its receptors (types I and II) activate several pathways, including SMAD-dependent and SMAD-independent pathways (113). SMADs are a family of transcriptional factors that transmit signals from receptors for TGF- $\beta$ 1 to the nucleus. These transmitters regulate multiple targets such as matrix proteins, collagen, fibronectin and matrix metalloproteinases, and members of the fibrinolytic system that includes plasminogen activator inhibitor-1 (110). In both syndromic (Marfan syndrome and other) and nonsyndromic (bicuspid aortic valves, degenerative aortic disease, and other) aneurysms of the aorta, SMAD2 activation and increased TGF- $\beta$ 1 activity have been documented in the arterial media layer, which indicates a final common pathway for thoracic aorta aneurysm syndromes (114). Molecules with potential roles in aortic wall weakening are many. A proteoglycan like biglycan was decreased in aneurysmatic aortic tissue (114), and the same biglycan was perturbed in fetuses with TS (115). Vascular endothelial growth factor has been implicated as important to early angiogenesis as well as in postnatal maintenance of aortic wall integrity in TS (57). In contrast to these potential contributors, the angiotensin type 2 receptor appears to have been ruled out in the etiology of congenital heart disease in TS (116).

A close link has been hypothesized between arteriogenesis involving neural crest cell invasion and differentiation

to vascular smooth muscle cells and peripheral neurons (117). This association infers that the lifelong perturbed autonomic nervous system in TS (49, 118) could share a common origin with congenital heart disease. Moreover, normal lymph vessel development is guided by vascular endothelial growth factor C produced by blood vessels (119), and with blood vessels and peripheral nerves evolving alongside (117), a common origin of the abnormalities of these three components could explain the phenotype seen in TS with disturbances of the neural, the arterial, and the lymphatic systems. Collectively, further studies of not only angiogenesis but also genetic models and signaling molecules in TS are essential in order to identify the causative mechanisms. Unfortunately, relevant animal models of TS are not available due to different patterns of X-inactivation as well as difficulties in creating the relevant karyotypes for TS in animals (120).

#### IV. Acquired Heart Disease

Acquired heart disease is a major cause of morbidity and mortality in TS (Fig. 2). An increased risk of acute aortic dissection is present from as early as the second decade of life, stroke incidence is increased from the third decade, and myocardial infarction usually appears from the fifth decade (1, 3, 4). Although acquired heart disease with age contributes increasingly to all-cause excess mortality (Fig. 2), it may often be impossible to distinguish congenital from acquired lesions. Many adverse features such as aortic dilation (67, 78), proarrhythmic traits (55, 121), and hypertension (122) appear to be interrelated, and to some extent present throughout the entire lifetime in TS.

##### A. Aortic dissection

Aortic dissection occurs in 1 or 2 of 100 females with TS (123). Incidence rates are increased up to 100-fold when compared with the female background population (Fig. 4) (42, 123). This often devastating event occurs 40 yr before expected and with peak incidences in the third to fifth decades of life (Fig. 7) (123). Aortic dissection is not restricted to adulthood in TS. Incidence rates are increased even in the below 19 yr age group with 14 per 100,000 yr (123), and aortic dissection may occur as early as the first decade of life (90).

The outcome of aortic dissection is unknown in TS. The prognosis is likely to be at least equally as grave as in the general population, where 22 in 100 cases die before reaching a hospital (124). In general, prognosis depends on the location and extent of the dissection and is improved when limited to the descending aorta (Stanford B/DeBakey III) compared with both isolated ascending

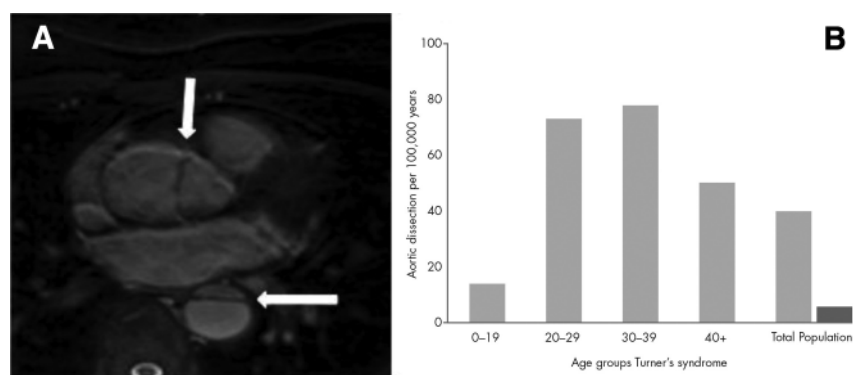
**Figure 7.**

Figure 7. Aortic dissection in TS. A, Chronic Stanford type A dissection in a 50-yr-old woman with TS with 45,X/46,Xr(X) mosaic karyotype and no known risk factors for aortic dissections but moderate aortic regurgitation (81, 123). B, Aortic dissections occur at increased incidence throughout the entire life span of individuals with TS (123). [Panel B was reproduced from C. H. Gravholt *et al.*: Clinical and epidemiological description of aortic dissection in Turner's syndrome. *Cardiol Young* 16:430–436, 2006 (123) with permission. © Cambridge University Press.]

aortic involvement (Stanford A/DeBakey II) and those propagating from the descending aorta into the ascending aorta (Stanford A/DeBakey I) (125). In TS, aortic dissections predominantly include the ascending aorta (63%), with isolated descending aortic involvement occurring less frequently (37%) (123). The causative lesions are not well characterized in TS, be it intimal tear, intramural hematoma, or penetrating ulcer (126). Preponderance to Stanford type A dissection over type B is in accordance with a high prevalence of both ascending aortic dilation and

bicuspid aortic valves (42, 81). Descending aortic dissection is most likely to occur in the presence of coarctation of the aorta (also after repair) (88, 90, 123).

Risk factors for aortic dissection in TS are not easily defined. Although the risk of aortic dissection and rupture is 100-fold increased (42), the event occurs on the backdrop of a syndrome prevalence of 1 in 2000 live-born girls (1), considerable diagnostic delay, a high extent of nondiagnosis (1), and aortic dissection affecting 1 or 2 of 100 females with TS over the course of an entire lifetime (123). Hence, the risk of aortic dissection may very well be severely increased, yet aortic dissections are infrequent encounters in clinical practice. Therefore, risk stratification in TS is mainly based on numerous case

reports and one register-based survey (123, 127). One prospective study has assessed aortic dissection in TS (42). However, the total number of years at risk was relatively low, and it has remained difficult to conclude anything beyond a severely increased risk of dissection (42). In TS, the currently acknowledged risk markers for aortic dissection are hypertension, karyotype 45,X, aortic dilation, and left-sided obstructive lesions including bicuspid aortic valves, coarctation of the aorta, and other obstructive arch

**TABLE 2.** Risk markers for aortic dissection in TS

General population risk factors	Evidence level	Special concerns in TS	Unresolved	Current risk factors in TS
Age (124)	Case reports, case series (127), one register study (123), one cohort study (42), and more cross-sectional studies of associations between aortic diameter and phenotype traits (78, 81)	Karyotype (127)	ERT?	Hypertension
Hypertension (125)	Expert consensus (Level C) (130)	Pregnancy (148)	Diabetes? GH?	Aortic caliber Aortic growth Bicuspid aortic valve Aortic coarctation Aortic valve dysfunction Iatrogenic trauma Age Pregnancy Karyotype External trauma Drugs
Aortic diameter (135)				
Aortic growth (130)				
Bicuspid aortic valve (73)				
Aortic coarctation (347)				
Aortic valve dysfunction (140)				
Iatrogenic trauma (348)				
Aortitis (126)				Aortitis
External trauma (349)				
Drugs (126)				
Pregnancy (126)				
Atherosclerosis (126)				

Risk markers for future aortic dissection in TS are currently based on evidence gathered in the general population and extrapolated onto TS as well as on limited evidence specific to TS. Hence, any conclusions on risk markers in TS are at evidence level C. ERT, Estrogen replacement therapy and estrogen deficiency; GH, GH therapy and GH deficiency.

**TABLE 3.** Maximum aortic diameter in TS using cardiac magnetic resonance

First author (Ref.)	Age (yr)	n	Assessed	Aortic diameter in comparison with controls	Aortic dilation: TS	Association <sup>a</sup>
Cleemann (78)	<24	41	9 positions	Absolute: smaller in TS except for similar in proximal descending compared to controls (n = 50); BSA-indexed: similar except for arch, isthmus, and descending where smaller in TS	15%, total <sup>b</sup>	BSA, +; CoA, –
Hjerrild (81)	>18	97	8 positions	Absolute: comparable in TS to controls (n = 24) for all, except smaller at distal arch and isthmus	23%, total <sup>c</sup>	BSA, +; BAV, +; blood pressure, +; age, +
Ostberg (75)	>18	128	2 positions	Absolute: mid-ascending and mid-descending in TS comparable to controls (n = 36); height-indexed: both aforementioned positions enlarged after adjustment for height and BSA	16%, mid-ascending <sup>b</sup>	BAV, +; age, +
Matura (289)	>18	166	2 positions	Absolute: mid-ascending similar and mid-descending smaller compared to controls (n = 26); BSA-indexed, larger mid-ascending aorta in TS and similar descending aorta	24%, mid-ascending <sup>c</sup>	BSA, +; BAV, +; ETA, +; karyotype, +
Kim (92)	<26	50	9 positions		2–30%, all positions, highest for aortic sinus and sinotubular junction <sup>d</sup>	Age, +; PAPVR, –; ETA, +
Dawson-Falk (80)	<35	40	1 position		13%: ascending aorta	

BSA, Body surface area; ETA, elongated transverse aortic arch; BAV, bicuspid aortic valve; PAPVR, partial anomalous pulmonary venous return; CoA, coenzyme A.

<sup>a</sup> Associations indicate positive (+) or inverse (–) correlation between aortic diameter and the variable in question.

<sup>b</sup> Aortic dilation was defined as 2 SD above the mean of the control group.

<sup>c</sup> Aortic dilation was defined as a maximum aortic diameter exceeding 3.4 cm after height adjustment.

<sup>d</sup> Aortic dilation was defined as two standardized z-scores above the normal values of children and adolescents.

<sup>e</sup> Aortic dilation was defined as >95% CI of nomograms for BSA and aortic diameter derived from x-ray computed tomography.

lesions (Table 2). Furthermore, a risk of aortic dissection is introduced in association with diagnostic cardiac and aortic catheterizations (87, 88, 123). Aortitis as a cause of dissection has been reported in TS, but only in one case (128). With insight into the causative mechanism of aortic dissection being scarce, much is left to the attending clinician on a case-to-case basis. This is especially true because the presently acknowledged risk factors are likely to fail to identify at least 10 of 100 acute aortic syndromes (127, 129).

### 1. Aortic diameter

Aortic diameter is the principal risk stratification tool for aortic dissection (130). Assessed by cardiac magnetic resonance, absolute aortic diameter is smaller in young girls with TS compared with healthy peers (Fig. 4) (78). Conversely, absolute aortic diameter is comparable to normal healthy peers in adulthood (allowing for smaller diameter in the arch, isthmus, and proximal descending aorta) (42, 81). Hence, aortic diameter seemingly catches up from childhood to adulthood with regard to aortic size, and this occurs despite a persistently smaller height and weight in TS. Aortic diameter in adults with TS is noted to have a larger range than healthy female peers, emphasizing

the essential clinical issue with separating a normal aortic diameter from an abnormal one in TS (Table 3) (75, 78). Echocardiographic studies generally support these findings (52, 81), with one study inferring a very early enlargement of aortic diameter in children with TS (9).

Physical stature is a determinant of aortic diameter in TS (42, 81). Even when normalizing aortic diameter for body surface area, aortic dilation is evident in adults with TS in the form of larger indexed diameters at all levels (except the distal aortic arch and adjacent to the classical site of coarctation of the aorta) (42, 81). In younger females the same applies around the coarctation site, whereas other diameters are comparable to controls (78). Nevertheless, the prevalence of dilation approaches 30% when using z-scores to normalize aortic diameter in young cohorts (92). Aortic dilation is also common in adulthood, and certain factors have been found to contribute to the variation in aortic diameter in TS. Much still needs to be clarified in relation to factors determining aortic diameter, and aortic dilation may occur in the absence of such factors (67, 131). The identified determinants of aortic size are:

- Body surface area (42, 78)
- Increasing age (9, 81)

- Systemic blood pressure (81, 132)
- Karyotype (67)
- Coarctation of the aorta (67, 78)
- Elongated transverse aorta (67)
- Bicuspid aortic valve (42, 81)

Insight into aortic dilation in TS is improving with better imaging, allowing pervasive characterization of the entire thoracic aorta and expanding our knowledge of an aortic disease that extends beyond the ascending aorta. At echocardiography, acoustic windows are often restricted due to a “barrel-shaped” chest (81, 133). Magnetic resonance and x-ray computed tomography circumvent this issue and may with orthogonal postprocessing of three-dimensional imaging provide an opportunity to minimize measurement error due to angulations in two-dimensional imaging (134). In keeping with this, echocardiography in TS underestimates magnetic resonance diameters in the ascending and descending aorta (75, 80, 133). As in the general population (130), cardiac magnetic resonance is emerging as a “gold standard” for delineation of anatomical aortic variants and aortic diameter when these occur downstream from the aortic sinus. Importantly, cardiac magnetic resonance has limitations beyond cost and availability, being contraindicated in the presence of ferromagnetic objects that even with magnetic resonance-compatible devices may create significant artifact. Yet, approximately 90% of girls and women with TS will successfully complete magnetic resonance studies (75, 133).

## 2. Risk stratification for aortic events

An enlarged aortic diameter increases the risk of aortic dissection in TS (42). However, aortic dissection may occur with aortic diameters that are considered “normal” even after normalizing to the smaller physical stature in TS (127). Aortic dilation appears to offer good positive predictive value for future aortic dissection (42), whereas the negative predictive value is less encouraging (127). This is in accordance with evidence from the general population, where on one hand absolute aortic diameters associate with a nonlinear increment in risk (135), whereas on the other hand 20% of all aortic dissections strike at diameters below 45 mm (136). The proposal to normalize aortic diameter using body surface area arose from this shortcoming (137), but the implications to risk need further assessment; outcome studies in the general population using indexed diameters are limited, and guidelines base surgical cutoff values on nonindexed numbers (130). In keeping with this, statements on critical aortic diameter have remained conservative in TS (2) because there is no firm evidence for a specific cutoff value, where the risk asso-

ciated with prophylactic aortic root replacement is superseded by the risk of acute aortic events (130).

Aortic dilation over time provides important prognostic information in populations that face a high risk of aortic dissection and rupture (130). Hence, aortic growth is a potential clinical marker that may assist in the complex risk assessment in TS. Changes in aortic diameter in TS were first prospectively studied using echocardiography. The ascending aorta grew 0.24–1.22 mm per year during 37 months in a mixed pediatric and adult cohort (age range, 3–39 yr) (133). A subsequent magnetic resonance study of a purely adult cohort showed that ascending aortic diameters increased over 29 months by 0.1–0.4 mm per year, and bicuspid aortic valves (but no other congenital anomaly or blood pressure) predicted aortic growth rates (134). For comparison, estimated aortic growth (using different imaging modalities) is 0.07 mm/yr in normal females (138), 0.1 mm/yr in thoracic aortic aneurysm populations (139), and 0.2 mm/yr in the setting of a bicuspid aortic valve (140, 141). The clinical significance of the reported aortic growth rates in TS is not known because the growth rates have not been linked with outcomes. Nevertheless, growth rates appear to be at least equivalent to other states of thoracic aortic disease, and growth rates may when used cautiously add weight to risk stratification also in TS.

Other measures than indexing to body surface area have been suggested in order to avoid over- or underestimation of thoracic aortic disease in TS (75). This includes a patient-specific ratio of ascending to descending aortic diameter (142), which is unfortunately inherently flawed in TS, where ascending aortic dilation will be underestimated in the context of frequent descending aortic dilation (75, 81). Attempts to quantify thoracic aortic disease using indices other than size have been made in TS. The aortic wall appears to be stiffened compared with healthy peers, although this is not a consistent finding, and the implications in terms of stratification for the risk of aortic dissection remain to be defined (132, 143–145).

## 3. Pregnancy and aortic dissection

The risk of aortic dissection and rupture during pregnancy in TS may be as high as two in 100 pregnancies (146–148), with maternal mortality approaching 86% (127). An increased occurrence of systemic arterial hypertension, if not overt eclampsia, may contribute to this risk (149, 150). However, some report gestational hypertensive disorders in TS, equally common in TS and in assisted pregnancies in general (146, 151). There are no outcome-driven criteria for absolute and relative contraindications to pregnancy (or the risk to mother and fetus, and mode of birth), although cardiovascular investigation before and

during pregnancy clearly is crucial (147). Aortic diameter exceeding 2.0 cm/m<sup>2</sup> has been proposed as an absolute contraindication for pregnancy in TS (152). Then again, aortic diameter may not accurately define and monitor the risk of dissection associated with pregnancy because aortic size was comparable between nulliparous and parous females with TS (147). While waiting for further evidence, we may have to refrain from categorical statements and limit advice to “severe” aortic dilation and complex congenital heart disease weighing against pregnancy. Therefore, counseling regarding pregnancy in TS needs to be given on a case-by-case basis, and preferably with cardiologists experienced in congenital heart disease and thoracic aortic disease taking center stage. On the note of counseling, unassisted pregnancy occurs in 2–5 per 100 females with TS (147, 153), and even females with 45,X monosomy or karyotypes with Y-material can become pregnant (146). Hence, careful information about birth control and hormonal contraceptives as well as the cardiovascular risk associated with assisted and unassisted pregnancy should be intrinsic to clinical care in potentially fertile age groups in TS (154).

#### 4. Etiology of aortic dilation

Insight into the causative mechanism of aortic dilation, dissection, and rupture is scarce. Intima-media changes with cystic medial degeneration have been reported in 42–72% of aortic dissections in TS (123, 155). Collagen fiber composition may also be altered (123). These microscopic changes in the aortic wall composition compare well with other states of thoracic aortic aneurysm formation, where changes in vascular smooth muscle cells, elastin, collagen, and other extracellular matrix components are encountered (156). Normal tissue homeostasis within the intima-media layer results mainly from a complex interplay of contractile, secretory, and proliferative activities of vascular smooth muscle cells (156). These cells maintain the extracellular matrix in delicate interaction with extracellular matrix components, which in turn are regulated by biochemical and mechanical stimuli of the aortic wall microenvironment (156). Interactions with vascular smooth muscle cells include factors such as wall stress from the propagating aortic pulse wave (157) and blood flow perturbations in the setting of bicuspid aortic valves (158, 159). Vascular smooth muscle cells may also be regulated by cytokines secreted by immune and endothelial cells (156). In thoracic aortic aneurysms, this equilibrium of the media layer is shifted toward structural degeneration, and this process is felt secondary to a primary intrinsic wall defect leading to mechanical and elastic aortic failure (156). The culprit lesion for these adverse changes can be increased synthesis of degenerative matrix metalloprotei-

nases (or reduced inhibitory molecules) from vascular smooth muscle cells (160). This is the case for Marfan and Loays-Dietz syndromes (161, 162), where TGF- $\beta$ 1-mediated stimulation of vascular smooth muscle cell synthesis of matrix-degrading metalloproteinases is increased. These proteins then cause lyses of extracellular matrix proteins with secondary loss of structural integrity (160, 163). Other mechanisms may also be involved in increasing local protease activity, such as stimulation by immune cells within the adventitia layer (164). In theory, aortic disease in TS could result from disruption of the same or parts of these disrupted pathways of synthesis and degradation. There is a call for further studies in TS, where a myriad of factors could be causative as primary lesions or exacerbating components. Hypertension, tachycardia, or flow perturbation in relation to bicuspid aortic valves may, for instance, provide the substrate for an adverse mechanical stimulus within the aortic medial layer (49). Alternatively, immunological hyperreactivity could evoke an adverse shift in the extracellular matrix components within the aortic wall (165, 166).

Aortic wall disease may also result from a primary defect in the structural components of the extracellular matrix in TS, such as in vascular Ehlers-Danlos syndrome (167), or a primary conformational change of vascular smooth muscle cells (168). Furthermore, an ontological component has been inferred in the pathogenesis of thoracic aortic aneurysm in general (126). Aneurysm in the thoracic aorta is now considered a nonatherosclerotic process in contrast to the mainly atherosclerotic abdominal aortic aneurysms (although atherosclerosis may be superimposed in thoracic aortic disease) (130, 169). A hitherto undetermined fetal process may also be the source of cystic medial degeneration in TS, as seen in aortic dissection in this cohort (123, 127). Intriguingly, such aortic wall pathology was also observed in a 20-wk-old 45,X fetus (Fig. 8) (94). Although seen in only one case, this finding certainly infers a primary defect that could then potentially be exacerbated by the presence of other factors both ante- and postnatally. Conclusively, delineation of the precise chain of events leading to thoracic aortic aneurysm formation leading to a more complete understanding, awaits discovery both for TS and other states of syndromic and nonsyndromic aortic dilation. Certainly, relevant points of focus might include the TGF- $\beta$  system or activation of SMAD pathways in TS (112), as well as a more in-depth delineation of aortic wall changes to ascertain other potentially involved pathways.

There are concerns that recombinant human GH treatment and estrogen replacement therapy could induce unfavorable aortic wall changes. Treatment with GH may directly influence the homeostasis within the media layer

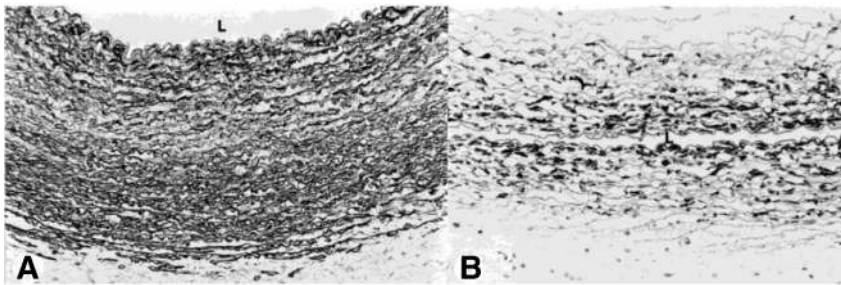
**Figure 8.**

Figure 8. Changes within the tunica media are present *in utero* in TS. Aortic wall pathology may be present from an early stage, as seen in an autopsy of a 20-wk-old 45,X fetus that presented with markedly reduced numbers of smooth muscle cells and elastic fibers in a thin-walled and hypoplastic aorta (A). The appearance of the aortic wall was consistent with degeneration of the aortic media layer (A) and differed markedly from the normally developed aortic wall of a 21-wk-old normal fetus (B) (94). Illustrations are from a photomicrograph where smooth muscle cells are demonstrated with anti- $\alpha$ -smooth muscle actin antibody, and elastic fibers are stained with Victoria blue. [Reproduced from S. Miyabara *et al.*: Developmental analysis of cardiovascular system of 45,X fetuses with cystic hygroma. *Am J Med Genet* 68:135–141, 1997 (94) with permission. © Wiley-Liss, Inc.]

of the aortic wall, as seen in animal and human cell models (170, 171). Alternatively, such treatment may impose mechanical stress on the aortic wall as a part of a hyperkinetic syndrome comparable to that observed in acromegalic individuals (172). Hitherto, there has been no overtly adverse impact of GH treatment on aortic diameter in TS, although conclusions are limited by primarily retrospective designs without adequate controls or pretreatment comparison (144, 173). Estrogen replacement therapy appears to reduce central arterial stiffness (174), improve endothelial function (175), and decrease carotid intima-media thickness (176) in short-term studies in TS. Hence, an improved risk profile for aortic dissection has been inferred. The precise relevance of these findings to the risk of aortic events in TS is undefined. On the note of treatments that aim to decrease morbidity and mortality in TS, the extension of the experiences with prevention of dilation in Marfan syndrome (177, 178) to TS would be highly interesting. However, specific evidence is undoubtedly needed for this cohort in light of the comorbidities that renders direct comparisons with non-TS studies hazardous.

## B. Stroke

Stroke incidence in TS is increased from the second decade of life onward, overall causing increased morbidity (RR, 2.7) and mortality (SMR, 3.9) (3, 4). Only one cohort study has attempted to ascertain the etiology of stroke in TS (4). With the limitations of death certificates in identifying causes of death, an estimated 90% of events were hemorrhagic (4). Case reports have described numerous causes, which are more or less common in the general population:

- Atherosclerosis (179)
- Hemorrhage (180)
- Fibromuscular dysplasia (181)
- Moyamoya disease (182)
- Hemophilia (183)
- Congenital carotid hypoplasia (184)
- Hemangioma (185)
- Dissection of head and neck vessels (186)

Naturally, deriving causative mechanisms from solitary case sources is a dubious act due to likely publication bias. The verdict is pending, and the etiology will only be clarified with a systematic assessment of both morbidity and mortality related especially to the contribution of ischemia, intracerebral hematoma and subarachnoid hemorrhage to stroke in TS (common causes in the background population) (187).

### 1. Cerebrovascular arteriopathy

The burden of general population risk markers for stroke of any cause (188) is increased from a young age in TS, and among other markers includes hypertension, insulin resistance, and visceral adiposity (122, 189). These markers may influence the cardiovascular system toward early aging (190), which in turn may predispose to ischemic or hemorrhagic stroke (187). Early cardiovascular aging (191) may potentially superimpose onto any adverse *in utero* programming of the cardiovascular system (192) and be aggravated by metabolic complications (193) related to the low birth weight in TS (18). Estrogen and GH deficiencies (or equally incorrectly administered replacement therapies) could further deliver a negative impact on the vasculature (194, 195).

Carotid intima-media thickness is increased as early as the second decade of life in TS (143, 196). This surrogate risk marker quantifies the combined thickness of media and intima layer, which for the elastic carotid arteries predominantly reflects intimal pathology (principally atherosclerosis) (197). Intima-media thickness predicts the risk of stroke and coronary atherosclerosis in the general population, adding value beyond traditional risk markers (198, 199). Consequently, increased intima-media thickness in young females with TS may indicate a premature atherosclerosis. In a pediatric cohort with TS, intima-media thickness was not abnormal, but there were correlations with atherogenic factors (200), which does align with a hypothesis of early atherosclerosis. In adulthood, intima-media thickness continually associates with ath-

erosclerotic risk markers, including blood pressure, C-reactive protein, clotting factors, lipids, and age (143, 196, 201). Outcome studies are completely lacking for TS, but note can be made that increased intima-media thickness is generally considered an even more adverse prognostic omen when encountered in the young (202). Perturbations in other indices that may also reflect premature atherosclerosis have been identified in TS, such as aortic stiffness, augmentation index, and ambulatory arterial stiffness index (143, 196, 203).

Arterial dilation extends beyond the aorta and into the large arteries in TS (67, 196). Therefore, a primary and intrinsic, generalized arteriopathy may include the cerebral vasculature and potentially cause small-vessel disease and stroke (187) or promoting dissections within regional arteries (186, 204). Case reports of stroke have accounted for vascular smooth muscle cell deregulation in fibromuscular dysplasia and moyamoya disease in TS (205, 206). These arterial disorders are characterized by luminal narrowing rather than dilation (207, 208). However, their presence could be seen as indicative of disequilibrium of vascular smooth muscle cell activity and extracellular matrix components. Vascular smooth muscle cells of the head and neck arteries notably share embryonic origin with aortic vascular smooth muscle cells, stemming from pharyngeal arch arteries (I–III). The cerebral arteries are subject to the same regulatory processes during angiogenesis (112), which could support a common primary defect for arterial disease in TS. Intriguingly, other states of media layer deregulation can have a systemic component, as seen in skin biopsies of cervical artery dissections (209). Conversely, the observed association of thoracic aortic disease and conduit artery dilation in TS (67) could be a hemodynamic effect rather than marking a universal arterial disorder.

## 2. Thromboembolism

Atrial fibrillation could increase the risk of thromboembolic stroke in TS, although reports of atrial tachycar-

dia are limited to date (210–212). However, P-wave dispersion is increased in TS (213) as a potential substrate for atrial fibrillation (214). Risk factors for atrial fibrillation (hypertension, type 2 diabetes, *etc.*) are overrepresented in TS, and arguably atrial arrhythmia should occur at least with the general population prevalence of one in every 200 (215). Systematic assessment is warranted, especially with atrial fibrillation often remaining subclinical until associated events unmask the arrhythmia. Other possible sources of thromboembolic stroke in TS include congenital cardiac lesions such as intracardiac shunts and thrombus (216), or calcifications and vegetations on abnormal mitral and aortic valves (217). Thromboembolism may also develop in the setting of heart failure (218), which is an anticipated feature in TS in association with the increased incidence of ischemic heart disease and type 2 diabetes, or even in normotensive and euglycemic young females with cardiomyopathy (70).

## 3. The coagulation system

Procoagulant disorders of the clotting system can lead to thromboembolic stroke, which is especially relevant for stroke in the young (217), and this may be a contributing factor for often premature cerebral events in TS (4). No outcome studies for thrombotic disease are available in TS. Nevertheless, the risk of thrombus formation seems increased even in the absence of functional or morphological cardiovascular substrates (Table 4). Clotting factors and times may be normal for cohorts with TS when assessed as a whole (212, 219). On the individual level, many will, however, have procoagulant levels of clotting and fibrinolytic factors (201, 219). This has been shown for C-reactive protein (40%), fibrinogen (15%), fibrin D-dimer (15%), factor VIII (25%), and von Willebrand factor (15%) (201). Proteins C and S may also be reduced (220). Inflammatory markers and clotting factors correlate with carotid intima thickness, heart rate, and blood pressure (201), which support a prognostic significance of clotting abnormalities. Of procoagulant mutations, only

**TABLE 4.** Reported events related to thrombosis in TS

First author (Ref.)	Event	Age (yr)	Karyotype	Clotting system abnormality	Other
Jobe (350)	Deep venous thrombosis	17	45,X/46,XY (q11.2)	Prothrombin G20210 mutation	Ongoing ERT
Donal (351)	Aortic thrombus	56			
Kopacek (352)	Portal vein thrombosis	3	45,X	↑ Factor VIII, ↑ von Willebrand factor	
	Portal vein thrombosis	1	45,X	↑ Factor VIII, ↑ von Willebrand factor	
	Portal vein thrombosis	12	45,X	↑ Factor VIII, ↑ von Willebrand factor	
Ureten (353)	Atraumatic osteonecrosis	38	45,X	↓ Protein S	Ongoing ERT
Pinto (354)	Portal vein thrombosis	1		↓ Protein S, ↓ Protein C, heterozygote for MTHFR mutation	
	Portal vein thrombosis	2		↓ Protein C; heterozygote for MTHFR mutation	

Case reports of thrombotic and related disease in TS, occurring in the absence of reported morphological and functional disturbance to explain the lesions beyond perturbations in the clotting system. ERT, estrogen replacement therapy.



factor V Leiden G1691A gene polymorphism heterozygosity has been found more commonly in TS (13%) compared with the background population (2%) (201). An in-depth characterization of disturbances of the clotting system, their etiology, and any implications for morbidity is warranted in TS. This also applies to tendencies toward increased bleeding as a cause of hemorrhagic stroke in TS patients (221, 222), where X-linked recessive hemophilia could hypothetically manifest as frequently as in their male counterparts.

### C. Ischemic heart disease

Ischemic heart disease is more frequent in TS than in general (SMR, 3.5) (1). These events mainly raise mortality rates from the fifth decade of life in TS, occurring with incidences skewed toward older age groups when compared with other acquired heart diseases (4).

The origin of coronary artery disease in TS has not been formally assessed. A plethora of causative mechanisms conceivably contribute to causing these events as early as in the fourth decade of life. Reported causes of myocardial infarction in TS include atherosclerosis, embolism from intracardiac thrombus or coronary artery involvement in aortic dissection (223–225). However, the etiology has in many cases not been identified (226), and causative mechanisms are not easily characterized by case reports. However, looking toward atherosclerosis, in the capacity of being most common in the background population, there is ample possibility for premature atherosclerosis. Markers for atherosclerotic heart disease are frequent in TS:

- Hypertension (81)
- Insulin resistance (227) and overt diabetes (1)
- Dyslipidemia (228)
- Obesity (229)
- Estrogen deficiency (229)
- GH deficiency (229)
- Hypercoagulable clotting system (201)

Coronary artery anomalies have also been reported in TS. These include coronary arterial dilation (230), single coronary ostium (231), coronary fistulas (232), and aberrant origin from the descending thoracic aorta (233). The prevalence of these anomalies is completely unexplored. From an embryonic perspective, the proximal coronary arteries share origin with the ascending aorta and aortic valve, and adequate migration of neural crest cells and appropriate differentiation into vascular smooth muscle cells are essential to all of these structures (112). Hence, and in line with the aortic and cerebral vascular territories in TS, the delineation of connections between coronary heart disease with ontological programming of an intrinsic

arterial wall defect and the interplay with comorbidities awaits further scientific progress.

### D. Hypertension

Arterial blood pressure is raised in all ages of TS patients (9, 132, 212). Hypertension affects 21–40% of children and adolescents (9, 122) and 50–58% of adults (81, 132). Characteristically for TS, hypertension affects systolic, diastolic, and pulse pressures, and there is a blunted nocturnal dipping pattern (122, 234). Morbidity (RR, 2.9) and mortality (SMR, 6.0) directly attributed to hypertension are substantial (3, 4), and there is an unfortunate degree of nondiagnosis with only 4–22% receiving treatment (9, 212).

#### 1. Essential hypertension

Hypertension is principally described as being essential by nature in TS (212, 235), and a multifactorial etiology thus prevails. Relative sympathetic hyperactivity in TS provides a possible foundation for high output hypertension (49, 118). Increased activation of the renin-angiotensin-aldosterone system has also been indicated (235, 236) with potential secondary salt and water retention, arteriolar constriction, and potentiated sympathetic drive. Conversely, lower peripheral vascular resistance has been demonstrated in young girls with TS, but this finding could relate to ongoing recombinant GH treatment (237). Abnormal levels of renin associate with abnormal captopril challenges in TS (236), but outright renovascular hypertension is rare (238). Conversely, normal renin and aldosterone have been reported for TS (49).

Oral contraceptives and postmenopausal hormone therapy have been claimed to induce hypertension (239). For TS, the verdict on this matter is unsettled (Table 5). Estrogen replacement therapy induced abnormal renin levels and captopril responses (236), and a cross-sectional study found elevated blood pressure in females with TS on oral estrogen replacement therapy (240). As opposed to these studies, renin levels were not affected by estrogen in another cohort with TS (49). Furthermore, it has been implied that estrogen counter-regulates the renin-angiotensin-aldosterone and sympathetic nervous systems in animal and human models (241). In a mixed population of patients with premature ovarian failure that included TS, transdermal 17- $\beta$ -estradiol (and vaginal progesterone) reduced 24-h blood pressure and serum angiotensin II when compared with baseline and treatment with synthetic ethinyl estradiol (and norethisterone), with unchanged serum aldosterone (242). So the general appearance is that estrogen deficiency leads to a slight elevation of blood pressure, and that appropriate estrogen treatment could lower

**TABLE 5.** Cardiovascular, metabolic, and clotting factors during estrogen replacement therapy compared with no treatment in TS

	Gravholt (234, 276)	Papagianni (309)	Ostberg (176)	Elsheik (174)	Chan. (175)	Host (312)	Lanes (219)
Participants							
n	26	12	14	21	7	8	5
Mean age (yr)	33	15	32	32	29	29	17
Intervention							
Duration	4-month pause	12-month treatment		3-month pause	6-wk pause	2-month pause	6-month treatment
Route	Oral/transdermal	Oral	Oral	Oral	Oral	Oral	Oral
Status when on ERT							
Hemodynamic	↓ 24-h diastolic AMBP, →AASI		↓ IMT, →FMD, →PWV, →AIX	↓ AIX, →clinic BP	↑ FMD, →clinic BP		
Metabolic	↓ Glucose tolerance, ↓ insulin, →glucose, ↑ FFM, →BMI, →WHR, ↑ HDL	↑ HDL, →other lipids	↑ HDL, ↓ glucose	↓ Insulin, ↓ glucose, ↓ WHR, →BMI, →lipids	→Lipids		→Lipids
Clotting system		→APTT, →fibrinogen, →TRC	→Fibrinogen				
Other	↑ VO <sub>2</sub> max		→IL-6, →CRP			↓ Adiponectin	→PAI-I

AASI, Ambulatory arterial stiffness index; AIX, Aortic augmentation index; AMBP, ambulatory BP; APTT, activated partial thromboplastin time; BMI, body mass index; BP, blood pressure; CRP, C-reactive protein; ERT, estrogen replacement therapy; FFM, fat free mass, FMD, forearm flow-mediated dilation; HDL, high-density lipoprotein; IMT, carotid intima-media thickness; VO<sub>2</sub> max, maximum oxygen absorption capacity; WHR, waist-to-hip ratio; TRC, platelet count; PAI-I, plasminogen activator inhibitor-I; PWV, pulse wave velocity.

blood pressure to a similar degree as antihypertensive monotherapy.

Other causes of hypertension are common in TS, such as insulin resistance [seen in some (227, 243) but not all studies (189, 244)], visceral adiposity (245, 246), and GH deficiency or resistance (8). Altered cardiovascular auto-regulation due to adverse imprinting in fetal life or poor thriving early in antenatal life (247) may very well raise the risk of hypertension in TS also. Intriguingly, metabolic syndrome appears to have a component of prenatal programming in TS (248), and hypertension is a well-established component of this syndrome (249). Further study of a putative prenatal programming component is required because findings are unequivocal with birth weight and blood pressure unassociated in another study of TS (122).

## 2. Secondary hypertension

Classical secondary hypertension is uncommon in TS. Nonetheless, hypertension does associate with vascular pathology in some females, such as coarctation of the aorta (repaired and unrepaired) (250) or, less commonly, hypoplastic transverse aortic arch (13). Still, congenital heart disease and hypertension do not always coexist (235). Renovascular (238) and endocrine hypertension (181, 251) are rare in TS, although various renal malformations occur in a third of all TS (252). Despite this, hypertension may develop secondary to impaired renal function in TS (132).

## 3. Sympathovagal dysfunction

Sympathetic adrenergic drive is relatively increased in TS (49), which is present in utero and evident as sinus

tachycardia (253). A primary developmental component is therefore likely, as is also indicated in a recent stem cell model of TS (53). However, this does not preclude a contribution for acquired factors. Certainly, changes in glucose metabolism are common in TS (189), and complications analogous to those seen in diabetic autonomous neuropathy could be of importance (254). Estrogen also contributes to normal autonomous regulation. This is appreciated by QT interval fluctuations and arrhythmic potential fluctuating over the menstrual cycle as well as by the relative tachycardia and heart rate-corrected QT-interval elongation induced by female puberty (194). Hence, estrogen deficiency in TS may play a role, although the sympathetic drive relative to parasympathetic activity was not affected by short-term estrogen replacement therapy (49). Another relevant factor in TS is the potential disturbance to the renin-angiotensin-aldosterone system (236) and its link to the overly active sympathetic nervous system.

## 4. Prognostic impact of hypertension

Hypertension contributes to aortic dilation and dissection in TS (81, 127). In addition, elevated blood pressure marks an increased risk of ischemic heart disease and stroke (188, 255). Outcome studies in TS are lacking, but several studies have demonstrated an association between blood pressure elevation and intermediate cardiovascular markers such as carotid intima-media thickness, aortic pulse wave velocity, and aortic diameter (81, 132, 143, 196). Hypertension also causes left ventricular failure in the general population (256). Presumably, a causal rela-

**TABLE 6.** The left ventricle of the heart in TS

First author (Ref.)	No. of participants (age)	Selection criteria	Comparative abnormalities	Putative involved mechanisms
Andersen (70)	33 (>18 yr)	+, ERT; ÷, HTN, diabetes, IGTT, LVEF <50%, CHD (excl. bicuspid aortic valve)	Diastolic dysfunction (27%), systolic dysfunction (normal LVEF), <sup>a</sup> left atrial dilation	Hypertension, sympathovagal dysfunction, aortic arch abnormalities, aortic valve dysfunction, intrinsic cardiac defect, estrogen deficiency, GH deficiency, insulin resistance, thyroid disease
Sozen (355)	31 (>18 yr)	÷, HTN, <sup>b</sup> CHD, ERT, all medicines	Diastolic dysfunction, left ventricular hypertrophy, left ventricular dilation, left atrial dilation	
Tancredi (261)	50 (<38 yr)	÷, Aortic dilation <sup>c</sup>	Diastolic dysfunction, left ventricular hypertrophy, systolic dysfunction (normal LVEF), left ventricular dilation	

Abnormal left ventricular function and geometry is a common finding on echocardiography in TS, although many abnormalities may only be appreciated in their true extension when taking the smaller physical stature of women with TS into account. Overall, the pathologies balance toward hypertrophy and enlarged chamber dimensions, contrasting indications of cardiac hypoplasia in fetal life. CHD, Congenital heart disease including aortic valve disease; ERT, estrogen replacement therapy; HTN, hypertension; IGTT, impaired glucose tolerance; LVEF, left ventricular ejection fraction; +, all received ERT; ÷, none had HTN, diabetes, IGTT, LVEF < 50%, and CHD.

<sup>a</sup> Systolic dysfunction was seen as decreased myocardial strain rate with normal ejection fraction when compared to controls (n = 33).

<sup>b</sup> Diagnosed hypertension was an exclusion criterion, but systolic blood pressure was higher in TS than in controls (n = 30).

<sup>c</sup> Hypertension was not an exclusion criterion, and diastolic blood pressure was higher in TS than in controls (n = 56).

tion may also be present in TS between the common hypertensive disorder and widespread dysfunction and abnormal geometry of the left ventricle in TS, although such association remains to be assessed in unselected cohorts in TS (Table 6).

Appropriate thresholds for antihypertensive treatment are not available in TS. However, the burden of cardiovascular morbidity and mortality certainly justifies tight blood pressure control. When a considerable burden of cardiovascular disease, including aortic dilation and risk of dissection, is present, lowering blood pressure to the lowest tolerated levels is probably paramount (130). There is no evidence to support specific treatments in TS, and the choice of therapeutic agent should follow general guidelines where reduction in blood pressure is the principal goal (257). However, some antihypertensive medicines may offer secondary benefits beyond lowering of blood pressure such as reducing progression of disease within the aortic wall in TS. Again, evidence in TS is non-existent, and clinicians have to look to other cohorts with increased risk of aortic dilation and dissection, bearing in mind that the aortic pathology has not been disclosed in TS. The general recommendation for Marfan syndrome has been to use  $\beta$ -adrenergic receptor blockade as an aortic wall-stabilizing agent. Angiotensin receptor antagonists now challenge this practice by potentially superior prophylaxis for aortic dissection through TGF- $\beta$ 1 antagonism (177, 178, 258). Then again, combined  $\alpha$ - and  $\beta$ -adrenergic receptor blockade reduced the incidence of aortic dissection in type IV vascular Ehlers-Danlos syndrome (259). Although some lessons may be learned from Mar-

fan syndrome, clinicians must bear in mind that causative mechanisms behind aortic dilation are unidentified in TS, and unequivocal evidence of superiority of one agent over the other is lacking, with prospective trials highly awaited. To date, only one study has showed an actual reduction in blood pressure during a generally increased attention to cardiovascular risk prophylaxis in TS (134). Unfortunately, this study allows no conclusions other than that blood pressure can be controlled through the use of antihypertensive medicines, as also indicated with short-term estrogen replacement (234).

### E. Arrhythmia

Sinus tachycardia is a lifelong phenomenon in TS (55, 260, 261). Conduction of the electrical impulse through the atria and atrioventricular node is accelerated (121), and the risk of atrial tachycardia may be increased (213), whereas bradycardia is rare (262). Furthermore, the myocardial action potential is prolonged with delayed repolarization, with pathological prolongation present in 33–36% of children and 21% of adults with TS (*i.e.*, heart rate-corrected QT interval >440 msec) (55, 121). These abnormalities of excitability and conduction are negative prognostic omens in the general population (263, 264), whereas their predictive capacity in TS remains to be determined. So far in TS, only one case of sudden, unexplained death has been related to known prolongation of the QT interval (265).

The cause of altered cardiac electrophysiology is unknown, but several aspects suggest an inherited defect:

- The 45,X karyotype has an adverse impact on the heart rate-corrected QT interval (although not consistently in all studies) (55, 121), and
- The heart rate-corrected QT interval prolongation is not associated with traditional indices of heart rate-corrected QT interval elongation (*i.e.*, age, electrolytes, left ventricular hypertrophy, medicines, coronary heart disease, and thyroid disorders) (55, 121).

It remains to be resolved whether the X-chromosomal haploinsufficiency in TS results in altered transcription of proteins involved directly in the activity of ion channels as structural or regulatory components, such as is seen in congenital long QT syndrome (266). Alternatively, altered expression of X-linked genes may function as transcriptional regulators of autosomal genes involved in ion channel activity. Interestingly, the action potential of the cardiomyocyte-like differentiated pluripotent stem cells seems prolonged in TS (53), which points toward a developmental abnormality. Further insight is necessary into the etiology, also including the role played by the relatively increased sympathetic drive seen in TS (49, 118), and likely increases automaticity, excitability, and conductivity of cardiomyocytes and cardiac conduction tissues.

## V. Endocrine and Metabolic Risk Factors

### A. Growth hormone deficiency or resistance

*SHOX* haploinsufficiency causes reduced final height in 95–99% of females with TS (267, 268). In addition to this genetic perturbation of growth, there is an unresolved imbalance in the “GH–IGF-I–IGF binding protein” axis. This disturbance may involve relative resistance to GH actions (269, 270), pituitary production failure of GH, or increased serum binding (8). Spontaneous and/or stimulated secretion of GH has been found to be diminished by some (271–273), whereas others have demonstrated normal GH secretion (274, 275). GH secretion is reduced by 50% in adulthood, with plausible influences from body composition (276). The bioactivity of circulating GH may also be reduced (277), and there are low levels of IGF-I as well as free and bioactive IGF-I (245). Moreover, proteolysis of IGF binding proteins is increased, which possibly impedes normal tissue delivery of IGF-I (278, 279). In view of this available evidence on GH signaling in TS and with normal final height being achievable with supraphysiological doses of recombinant human GH (280, 281), treatment should commence in infancy. Starting treatment early facilitates appropriate catch-up of skeletal growth, and it should be terminated upon reaching final height or when growth potential is no longer available (2).

Disorderly signaling within the GH–IGF-I axis may very well extend beyond the realm of physical stature in TS, affecting other organ systems such as the cardiovascular system. Comparatively, cardiovascular prognosis is impaired in overt GH deficiency (282, 283), and low levels of IGF-I increase the risk of stroke and myocardial infarction, even in the absence of a clearly pathological deviation of the GH–IGF-I axis (284, 285). The precise cause for this adverse association remains to be defined (195). However, GH stimulates hepatic IGF-I production through auto- and paracrine actions, which in turn regulates a myriad of cardiovascular cells, including cardiac myocytes, fibroblasts, and vascular smooth muscle cells (195). It is well established that many cardiovascular and metabolic features of TS are present in other states of GH deficiency (195). Additional study is needed to assess the role of a deviation in the GH–IGF-I axis to these common features that include insulin resistance (227) and visceral adiposity (245) as well as unfavorable changes in intima-media thickness (143), left ventricular function (70), blood pressure (212), sympathetic drive (49), exercise performance (261), and the clotting system (201). Recombinant GH treatment in adulthood is not advocated at present, and the importance of relative GH deficiency or resistance for morbidity awaits elucidation, with perturbed GH signaling continuing into adulthood in TS (286).

Increasing the activity of the GH–IGF-I axis from low normal levels benefits cardiovascular risk markers such as blood pressure, vascular resistance, and insulin sensitivity, as well as levels of the immune system (195). Conversely, hypersecretion of GH increases cardiovascular mortality as seen in acromegaly (287). In uncorrected acromegaly, an initial hyperkinetic syndrome with increased cardiac output is followed by diastolic left ventricular failure due to biventricular myocardial hypertrophy (fibrosis and cardiomyocyte hypertrophy), and ultimately dilated cardiomyopathy and heart failure ensue (288). Excess GH may also cause mitral and aortic valve calcifications, arrhythmia, and hypertension, just as glucose resistance is an issue (195). Therefore, unfavorable cardiovascular effects are a concern when supraphysiological doses of recombinant human GH are prescribed to girls and adolescents with TS over several years.

Studies of the cardiovascular impact of recombinant human GH in TS are limited by a lack of phenotype characterization before treatment (Table 7) as well as a scarcity of insight into normal age-associated, cardiovascular development in children with TS. As is the case for estrogen replacement therapy, placebo-controlled studies are not likely to take place because these treatments are now recommended universally in TS (2). Somewhat reassuringly, available evidence suggests that recombinant human GH

**TABLE 7.** The impact of GH treatment on cardiovascular phenotype in TS

	Sas (356)	van den Berg (357)	van den Berg (144)	Matura (289)	Bondy (173)	Radetti (237)
Participants (n)	68	31	38	67	53	26
GH duration	≈7 yr	≈9 yr	≈9 yr	≈4 yr	≈5 yr	≈5 yr
Design	Prospective	Retrospective (5 yr)	Retrospective (5 yr)	Retrospective <sup>c</sup>	Retrospective	Cross-sectional (ongoing)
Comparison group	Healthy peers	Healthy peers	Healthy peers	TS <sup>d</sup>	TS <sup>d</sup>	Healthy peers
Cardiac outcomes <sup>a</sup>	→LVM	→LVM, →RVM ↓ Biventricular volumes →RV EF, →LV EF, →CI		→LVM →Biventricular volumes →LV FS, →E/A ratio		→LVM →LV EF, ↑ LV FS, →CI ↓ Diastolic filling
Blood pressure	↓ Diastolic BP →Hypertension	→BP, ↑ HR	↑ Diastolic BP, ↑ mean BP			↓ SVR ↑ Diastolic BP, ↑ systolic BP, ↑ HR
Aortic outcomes <sup>a</sup>			↑ Aortic diameter, ↓ aortic distensibility <sup>b</sup>		→Aortic diameter	

CI, Cardiac index; BP, blood pressure; EF, ejection fraction; FS, fractional shortening; HR, heart rate; LV, left ventricle; LVM, left ventricular mass; RVM, right ventricular mass; RV, right ventricle; SVR, systemic vascular resistance; E/A ratio, early/late (atrial) ventricular filling velocity.

<sup>a</sup> Cardiac and aortic measurements indexed for body surface area.

<sup>b</sup> Aortic distensibility was decreased in the mid-ascending aorta and at the level of the diaphragm.

<sup>c</sup> Mainly retrospective study with 65% having terminated their GH treatment.

<sup>d</sup> GH-treated individuals were compared to age-comparable individuals with TS who had never received GH treatment.

delivered over several years does not induce an exaggerated trophic stimulus on cardiomyocytes in TS (Table 7). Likewise, recombinant human GH does not seem to increase aortic diameter compared with nontreated females with TS (173). The impact of dose and duration of recombinant human GH is unresolved, although retrospective reports imply no adverse impact of the current doses on the aorta (144, 173, 289).

Mapping of any adverse or advantageous influences of GH treatment is essential because it is delivered to an age group where the incidence of aortic dissections is raised to 14–19 per 100,000 population years (123, 290). The cardiovascular system undergoes rapid changes with altered signaling in the GH–IGF-I axis in other states of abnormal production (288). Although evidence is limited, there are no clear indications of such influences in TS (237). Furthermore, some effects of recombinant human GH are likely to revert toward pretreatment levels even after long-term delivery. This has been noted for diastolic blood pressure in TS (291), although more convincingly for glucose metabolism (292, 293). Metabolic change may also impact the circulatory system during recombinant human GH treatment, where insulin sensitivity has mainly been reported to decrease (292–294). By contrast, lipoproteins either are unaffected (295) or undergo favorable changes (296), and visceral and total adipose tissues decrease (294, 297). The verdict is pending for metabolic influences during long-term delivery of GH treatment in childhood and

adolescence. Prospective controlled studies focusing on metabolism as well as the cardiovascular system during recombinant human GH would be of great value.

## B. Estrogen deficiency

Estrogen deficiency is common in TS, where approximately 70% fail to undergo natural pubarche or menarche (298). Premature ovarian failure often ensues in adolescence, or early adulthood and is caused by apoptosis of ovarian follicles that commences in fetal life and progresses at varying rates (299, 300). The etiology of this gonadal demise is undefined. Umbilical cord blood samples have shown increased spontaneous apoptosis as well as higher levels of apoptosis mediated by TNF and CD95 pathways (301), and overexpression of Müllerian-inhibiting substance in ovarian granulosa cells of TS may also be involved (302). Due to this loss of ovarian function, lifelong deficiency of estrogen and progestin often prevails in TS, with a mere 2–5% being able to achieve unassisted pregnancy (147, 153). Hence, estrogen replacement therapy is necessary for feminization, achievement and maintenance of normal bone mineral density, and stimulation of neurocognitive development (2). Initial concerns about early introduction of estrogen as a cause of growth plate closure and reduced final height are being rejected in TS. For final height, impressive effects are shown with recombinant human GH despite early commencement of estrogen treatment (280, 303). After introduction, estrogen

replacement therapy should continue throughout adulthood, aiming for a total exposure time comparable to that in normal sex hormone synthesis (2).

Endogenous estrogen synthesis by the ovaries is crucial to cardiovascular health, as appreciated by the adverse prognostic impact of declining ovarian function in the normal menopause and premature ovarian failure (304, 305). Estrogen exerts direct cardiovascular effects on nuclear receptors of vascular smooth muscle cells, endothelial cells, and cardiomyocytes (194). Additional indirect effects are also important. Endogenous estrogen regulates hepatic synthesis of blood clotting factors and lipoproteins and promotes loss of visceral fat and increases insulin sensitivity (194). At the same time, estrogen down-regulates the immune system (194) and antagonizes the renin-angiotensin-aldosterone system (241). The precise mechanisms of action are in many instances unknown, but overall the effects of endogenous estrogen appear to be advantageous.

There may be complex and individualized influences of estrogen replacement therapy in different states of estrogen deficiency (306). The cardiovascular benefits are thought to inversely relate to the duration of estrogen deficiency before estrogen replacement therapy introduction (194). Exogenous estrogen is conceptualized as a pleiotropic hormone that may both improve and aggravate cardiovascular prognosis, depending on the atherosclerotic substrate of the individual and with timing of treatment being crucial (306). Therefore, lessens of estrogen delivery in TS should only be learned from the study of states with similar states of pronounced and early-life estrogen deficiency, rather than from cohorts with late-life deficiency after decades of normal exposure (postmenopausal women) or even normal levels (the contraceptive setting).

The effect of estrogen replacement therapy on hard cardiovascular endpoints has not been elucidated in TS. Short-term studies of intermediate risk markers suggest a beneficial impact of estrogen replacement therapy during the second to fourth decades of life in TS (Table 5). Overall, findings are congruent with known effects of estrogen on cardiovascular cells, causing acute nitrogen monoxide release and up-regulation of nitrogen monoxide synthase expression in the endothelium, which in turn relaxes vascular smooth muscle cells and has antiinflammatory effects (194). Estrogen also appears to beneficially influence the extracellular matrix synthesis and the vascular smooth muscle cell proliferation (194). For these reasons, estrogen replacement therapy in appropriate dosages could protect against atherosclerosis in TS, and may even antagonize a tentative up-regulation of the TGF- $\beta$ 1–vascular smooth muscle cell–matrix metalloproteinase axis in the aortic wall. However, estrogen replacement therapy may also

promote a procoagulant state and contribute to an apparently increased risk of venous thrombus formation in TS. This adverse effect is seen in the contraceptive and postmenopausal setting (307, 308), and oral estrogen has been proposed to cause a procoagulant state secondary to hepatocytes that respond to high estrogen levels through increased protein synthesis (194). This may theoretically be avoided using transdermal or sc administration (194). In TS, findings of exogenous estrogen as an up-regulator of the clotting system are equivocal. Some studies report stationary levels of clotting factors (176, 219, 309), whereas others find increased fibrinogen levels during estrogen treatment (220). Although the transdermal route may offer a more physiological profile (310), route of administration did not have any influence on other cardiovascular risk markers in TS (234). Beyond administration route, the progestin component of sex hormone replacement therapy may also be of relevance to the cardiovascular effects; some progestins activate androgen and mineralocorticoid receptors (311) and potentially oppose advantageous effects of estrogens.

Metabolic changes during estrogen replacement therapy may impact cardiovascular disease. Estrogen delivery in TS has been inferred to improve metabolism with decreased visceral adipose tissue (174), increased high-density lipoprotein cholesterol (176, 234, 309), and improved levels of circulating adipokines (312). However, findings during correction of the premature ovarian failure in TS are ambiguous. Glucose tolerance may be reduced by estrogen replacement therapy during the short term, with no change in insulin sensitivity (234). Speculatively, delivery of estrogen during the longer term will improve glucose homeostasis and metabolism in TS.

### C. Androgens

Sex steroid derangement in TS includes androgen deficiency (313–315). Androgen replacement therapy in TS is controversial and far from routine, and any association between correction of androgen deficiency and cardiovascular disease is unresolved (316). Delivery of androgens in other states of female androgen deficiency benefits intermediate cardiovascular markers (317). Oxandrolone, a nonaromatizable (to estradiol) derivative of testosterone, can be used as an accelerating adjuvant to recombinant human GH in pronounced growth failure or late introduction of GH treatment in TS (2). Cardiovascular effects of oxandrolone also remain controversial, and there are concerns that delivery may worsen any adverse impact of recombinant human GH on glucose metabolism in addition to deferring feminization (318–321).

#### D. Diabetes and metabolism

Both type 1 (RR, 11.6) and type 2 (RR, 4.4) diabetes occur with increased frequency in TS and directly aggravate prognosis (1, 3, 4, 10). Glucose tolerance is impaired in 25–78% of adult TS when assessed by an oral glucose tolerance test (234, 322), with higher glucose levels, whereas insulin levels are comparable (323), higher (322), or even lower (324) in comparison with controls. The insulin response has been described as delayed compared with healthy controls (227, 322). Fasting levels of glucose are generally comparable to controls (234, 325), whereas insulin levels have been reported as increased in some cohorts (323, 325) but not in other cohorts (234, 322). Using the hyperinsulinemic euglycemic clamp, some early studies indicated increased insulin resistance in adolescents with TS (243, 325). However, these early reports are challenged by more recent studies that show no such difference when compared with controls that are meticulously matched for age and body composition (189, 227).

The cause for diabetes in TS is unexplained. Insulin release after an iv glucose tolerance test has been shown to be reduced, especially at the early time-points (first-phase insulin response) (189). This infers that impaired  $\beta$ -cell function, rather than decreased insulin sensitivity, could be the foundation of impaired glucose tolerance in TS (244). Reduced gastric inhibitory polypeptide in young girls with TS who had impaired glucose tolerance, compared with age-matched girls with TS with normal glucose tolerance, suggests that an imbalance in incretin- $\beta$ -cell cross talk might contribute to disturbances in insulin secretion (326). Moreover, insulin sensitivity was normal in young females with TS compared with controls closely matched for age and body mass index, whereas discrete perturbations in glucose handling were present (189). In the same study, blood glucose levels were increased during oral and iv glucose tolerance test, with a poor compensatory increase in insulin and a reduced insulin-to-glucose ratio (189).

Haploinsufficiency of genes on Xp increases the risk of type 2 diabetes to 18–23%, and haploinsufficiency of Xp combined with trisomy for Xq genes (karyotypes with isochromosome Xq) further increases this risk (327). This was shown in a large cohort of TS females with a mean age of 35 yr and 25% of these TS females had type 2 diabetes (327), where the distribution was:

- del Xq: type 2 diabetes rate of 9% (compares with the general population)
- 45,X: type 2 diabetes rate of 18%
- del Xp: type 2 diabetes rate of 23%
- Isochromosome Xq: type 2 diabetes rate of 43%

The discovery of specific causative genes on the X-chromosome is likely to follow. Gene array data suggest that overexpression of Xq transcription factors is involved in altered pancreatic islet and  $\beta$ -cell function, as well as in proinflammatory action in TS, which was mirrored by increased levels of circulating C-reactive protein, IGF-II, and anti-glutamic acid decarboxylase 65 antibodies (327). Clinical studies have documented raised C-reactive protein (326, 328) as well as glutamic acid decarboxylase 65 antibodies (166), especially linked to isochromosome Xq (166). Such data infer a link between autoimmunity and diabetes in TS. Collectively, oral and iv glucose tolerance tests and clamp studies suggest that a number of factors might interact and in combination explain the perturbed  $\beta$ -cell function.

With premature ovarian failure affecting most adult females with TS, it is essential to evaluate the impact of estrogen replacement on glucose metabolism. Glucose tolerance assessed by oral glucose tolerance test was impaired in 78% after 6 months of estrogen replacement therapy compared with 50% before such treatment (234). However, free fat mass and physical fitness increased during exogenous estrogen treatment, lending promise of further improvements in glucose metabolism with extended treatment duration. A meta-analysis evaluated the effect of estrogen replacement therapy in the postmenopausal setting and found that muscle strength was increased during active treatment (329), as has also been indicated in TS (234, 330). Route of administration of estrogen replacement therapy (oral or transdermal) does not seem to influence glucose metabolism in TS (234, 240, 310). Incorporating data from conditions other than TS, which also include young women with surgically induced menopause, it seems that 17- $\beta$ -estradiol in the normal range is associated with beneficial effects on insulin sensitivity, whereas both a hypo- and hyperestrogenic milieu is associated with insulin resistance (331, 332). Indeed, 17- $\beta$ -estradiol and its  $\alpha$  and  $\beta$  receptors have profound beneficial effects on energy homeostasis, skeletal muscle, adipose tissue, liver, pancreas ( $\beta$ -cells), and the cardiovascular system (333).

Increased liver enzymes are common in TS (334), and cirrhosis is 5-fold more common (3). Deficiency of sex hormones, especially 17- $\beta$ -estradiol, seems to play an essential role in the hepatic abnormalities because treatment with estrogen replacement therapy reduces or even normalizes the liver enzymes (276, 335). The liver primarily expresses estrogen receptor  $\alpha$ , and to some extent estrogen receptor  $\beta$  also, and recent evidence shows that stimulation of estrogen receptor  $\alpha$  protects against inflammation and hypercholesterolemia, and that it is essential for normal glucose homeostasis (333). The histopathology of liver biopsies in females with TS who had persistently el-

evated liver enzymes showed a range of diseases, which included cirrhosis associated with obliterative portal venopathy, multiple focal nodular hyperplasia, and nodular regenerative hyperplasia (336). Less severe changes were also present, such as portal fibrosis, inflammatory infiltrates, and nonalcoholic fatty liver disease. The overall conclusion was that the principal causative mechanisms were congenital vascular disorders and nonalcoholic fatty liver disease (336). 17- $\beta$ -Estradiol directly mediates transcriptional regulation of vascular endothelial growth factor through estrogen receptor  $\alpha$  and  $\beta$ , activating receptors 1 and 2 for this growth factor on hepatic sinusoidal endothelial cells (337). This leads to paracrine release of growth and survival factors protecting hepatocytes from toxins and promoting hepatocyte proliferation. In this way, a lack of estrogen could lead to accelerated hepatocyte apoptosis and leakage of liver enzymes and vice versa (337). In a recent study, an abnormality of hepatic lipid storage was suggested because females with TS with high cholesterol and high body mass index had higher  $\gamma$ -glutamyl transferase, and magnetic resonance imaging confirmed intra-hepatocellular lipid deposition in TS (338). With the available evidence in mind, it can be concluded that estrogen replacement has a prominent role in maintaining normal liver metabolism in TS.

### E. Autoimmunity

Autoimmunity is increased in TS (165, 166), resulting not only in type 1 diabetes but also thyroiditis (RR, 16.6) (3). Thyroid autoimmune disease principally manifests as hypothyroidism, which increases in prevalence with age to affect 23–34% of adolescents and adults with TS (166, 339). No links have been made between hypothyroidism and cardiovascular morbidity in TS. However, inadequate diagnosis and suboptimal substitution may aggravate the prevailing metabolic and cardiovascular phenotype (340).

Haploinsufficiency of X-chromosome material has been proposed as a possible primary cause for autoimmunity (335), although lack of estrogen and up-regulation of proinflammatory cytokines (245) as well as accelerated apoptosis of T-cell subsets may also be involved (301). Estrogen has numerous effects on the immune system encompassing modulation of T cells (341) and inhibition of inflammation (342), although studies in humans and animal models are inconclusive. From periovulatory to pregnancy levels, 17- $\beta$ -estradiol has been shown to have a stimulatory effect on IL-4, IL-10, and interferon  $\alpha$ , but it inhibits TNF from CD4+ T cells, indicating a shift toward down-regulation of T-cell autoimmunity (342). At the same levels, 17- $\beta$ -estradiol has been shown to stimulate antibody secretion by CD5+ B cells while suppressing bone marrow B-cell lineage precursors

(342). The exact role of estrogen in autoimmunity in TS is unknown. Estrogen could induce an increased B-cell autoimmunity, but at the same time decrease T-cell autoimmunity (342). Other causes for the breakdown of self-tolerance such as fetal microchimerism, skewing of X-chromosome inactivation, gene duplication, or up-regulation of proinflammatory cytokines are being investigated. Androgen predominantly exerts an inhibitory effect on the immune system (342) and could therefore also be implicated through the relative androgen insufficient state in TS (314).

### VI. Future Perspectives

*SHOX* haploinsufficiency is currently the only genetic mechanism that explains part of the genotype and phenotype of TS (343). Here, we have presented a host of other putative mechanisms, which may or may not play a role in cardiovascular disease among TS patients. For many years, the pathogenesis of TS has eluded researchers, and the discovery of *SHOX* gene and its role in TS has naturally led researchers to believe that another gene or several genes play similar roles to partially or completely explain the spectrum of phenotypes. Such genes could be situated in one of the two PAR of the X-chromosome, although genes outside of this area may also play a role in the pathogenesis of TS. Interestingly, pseudoautosomal genes such as *ASMTL* and *PPP2R3B* in experimental models were expressed at lower levels than in normal cells (53). Because *ASMTL* is a methyltransferase, it can be speculated that improper methylation of the genome plays a role in TS. At present, the involvement of *CSF2RA* points toward insufficient placentation as the most plausible explanation for the high fetal lethality in TS (53, 344). However, these findings need to be confirmed in additional studies before any firm conclusions can be drawn. Likewise, the process and consequences of X-chromosome inactivation (or the lack hereof) also need to be studied in further detail. Another perspective comes from recent scientific advances indicating that sexual dimorphism between males and females plays a role in transcriptional regulation of autosomal genes and that this leads to differential expression between males and females of more than half of the coding genome (44). This must be studied in further detail, and the exact role of sexual dimorphism and the possible changes inflicted by the lack of one X-chromosome in TS need to be explored. Suffice it to say, the genetic mechanisms behind TS are far from understood, and the interaction between different genetic mechanisms may even complicate matters further. TS may actually be the result of deregulation within many different pathways that work



in concert to cause deviation from normality. Thus, linking genetics with phenotypic traits should prove extremely interesting in future studies, and we may very well have to revise our understanding of the genetics behind TS in the coming years. At present, we can probably divide the phenotypic traits of TS into (Fig. 9): 1) congenital cardiovascular malformations of unresolved pathogenesis; 2) decreased intrauterine viability, where haploinsufficiency for X-linked pseudoautosomal genes operating in the placenta has been suggested to be involved (*CSF2RA* and possibly other genes); 3) ovarian dysgenesis, where many genes have been implicated, but none proven; 4) predilection to autoimmune disease of unknown genes; and 5) altered brain development, especially social-cognitive development, which is altered in many cases, often in a more “male-like” direction.

Today, we can conclude that females with TS suffer from a condition with a high risk of congenital and acquired heart diseases, diabetes, metabolic disorders, and autoimmune disease (Fig. 9), which necessitates multidisciplinary care with involvement of many specialties (2). In many centers (6, 155), guidelines for multidisciplinary practice have been implemented, but clinical care has remained suboptimal elsewhere (5). The prevalence of associated diseases has been well detailed, but we lack a clear understanding of the pathophysiological mechanisms, including details on the genetic and molecular events that take place and lead to the diverse phenotypes. Life expectancy is decreased (11), and conditions related to the heart and great vessels are involved in about half of all excess deaths (4). Mounting evidence supports the notion that the specific X-chromosomal deficit and the tissue affected

**Figure 9.**

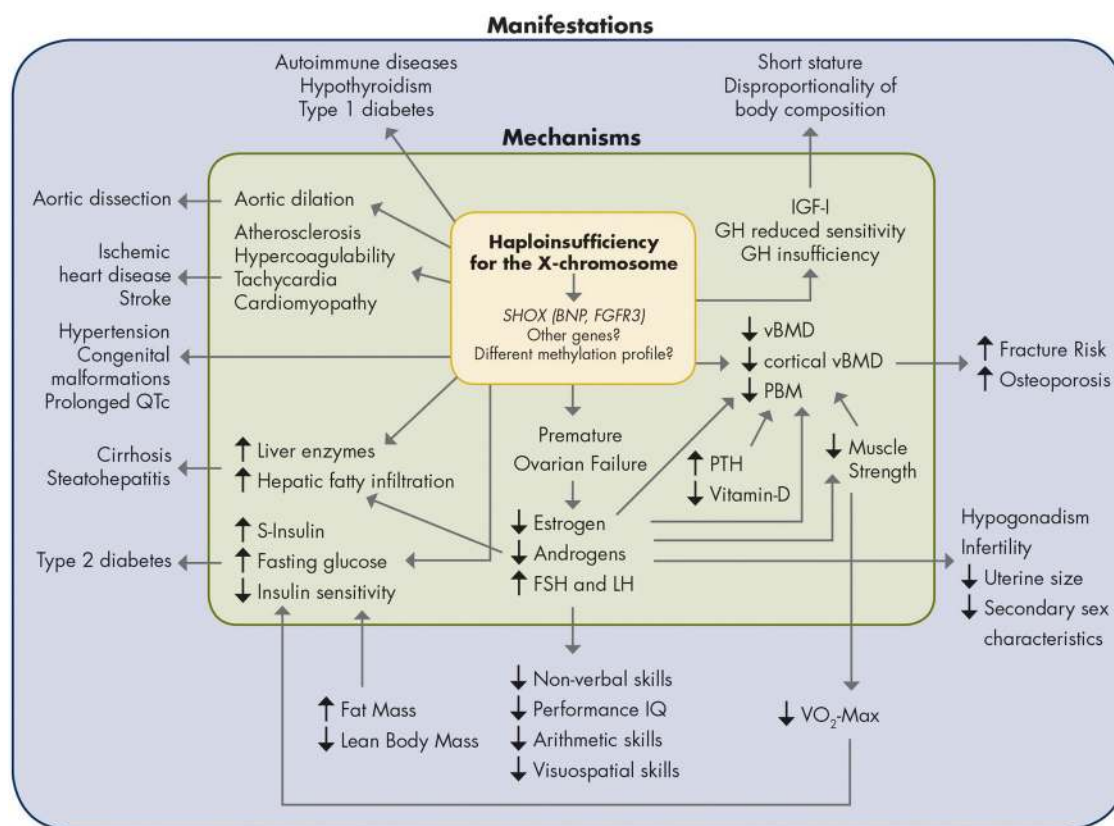


Figure 9. The serious effect of haploinsufficiency of genes on the X-chromosome (and possibly other genetic mechanisms working in concert) is at the center of the current understanding of pathogenesis in TS. This leads to a range of effects for the cardiovascular system (congenital and acquired diseases), the endocrine system (especially with premature ovarian failure and thus female hypogonadism), body composition, as well as the brain and other organs. Hypogonadism has pervasive effects, affecting: 1) different hormone levels; 2) cardiovascular features; 3) metabolic features; and 4) features related to sex hormones, such as infertility. In addition, mounting evidence suggests that hypogonadism in TS leads either directly or indirectly to a reduced quality of life. Haploinsufficiency of genes on the X-chromosome has been implicated in the presence of an increased risk of congenital malformations, although no specific genes have been identified so far. Arrows indicate possible consequences—not all interactions have been shown in scientific studies. VO<sub>2</sub>max, Maximum capacity to transport and utilize oxygen during incremental exercise; BNP, brain natriuretic peptide; FGFR3, fibroblast growth factor receptor 3; BMD, bone mineral density; PBM, peak bone mineral mass; IQ, intelligence quotient.

by this karyotype will determine the pathology encountered by the affected girl or woman. Although TS is not primarily considered a syndrome of premature aging (345), a number of characteristics point to some form of accelerated process—for instance, premature occurrence of aortic dissection, stroke, diabetes, autoimmune disease, and sensorineural deafness similar to presbycusis (346). So with these aspects in mind, a more thorough understanding of the total task that females with TS pose to the clinician is expected to evolve, facilitating better clinical care and more targeted development of new medications and translation from basic research to future treatments.

## VII. Conclusion

Females with TS suffer a multifaceted syndrome that manifests within the realm of several organ systems, whereof many especially cardiovascular components are only just being unraveled (Figs. 4 and 9). There is ample opportunity for progress because our understanding of the causation behind the plethora of associated cardiovascular traits remains patchy. An in-depth description of the genetic mechanisms leading to the advent of this X-chromosomal disorder should facilitate fundamental changes in diagnosis, treatment, and monitoring. Treatment of TS comes at a high cost for society, with an estimated 78,000 females affected in the United States and another 125,500 females affected in the European Union. If we can pinpoint new genetic mechanisms leading to the abundance of clinical manifestations and traits related to TS, we will have provided the important framework for development of new and targeted therapy.

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