

## RESEARCH

# Disorders of sex development: timing of diagnosis and management in a single large tertiary center

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## Abstract

**Background:** We describe the phenotypic spectrum and timing of diagnosis and management in a large series of patients with disorders of sexual development (DSD) treated in a single pediatric tertiary center.

**Methods:** DSD patients who had visited our tertiary center during the survey period (between 2004 and 2014) were identified based on an ICD-10 inquiry, and their phenotypic and molecular genetic findings were recorded from patient charts.

**Results:** Among the 550 DSD patients, 53.3% had 46,XY DSD; 37.1% had sex chromosome DSD and 9.6% had 46,XX DSD. The most common diagnoses were Turner syndrome (19.8%, diagnosed at the mean age of  $4.7 \pm 5.5$  years), Klinefelter syndrome (14.5%,  $6.8 \pm 6.2$  years) and bilateral cryptorchidism (23.1%). Very few patients with 46,XY DSD (7%) or 46,XX DSD (21%) had molecular genetic diagnosis. The yearly rate of DSD diagnoses remained stable over the survey period. After the release of the Nordic consensus on the management of undescended testes, the age at surgery for bilateral cryptorchidism declined significantly ( $P < 0.001$ ).

**Conclusions:** Our results show that (i) Turner syndrome and Klinefelter syndrome, the most frequent single DSD diagnoses, are still diagnosed relatively late; (ii) a temporal shift was observed in the management of bilateral cryptorchidism, which may favorably influence patients' adulthood semen quality and (iii) next-generation sequencing methods are not fully employed in the diagnostics of DSD patients.

## Key Words

- ▶ disorders of sex development
- ▶ Turner syndrome
- ▶ Klinefelter syndrome
- ▶ cryptorchidism
- ▶ children

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## Background

Disorders of sexual development (DSD) are congenital conditions in which the chromosomal, anatomic or gonadal sex development is atypical (1). Today, DSD is classified into three major categories by the patient's karyotype: sex chromosome DSD, 46,XY DSD and 46,XX DSD (1). The phenotypic spectrum of DSD is wide, and it can manifest as a complete sex reversal, a solitary genital abnormality or it can be a part of a syndrome affecting other organ systems (2). The etiology of DSD is multifaceted and can be caused by genetic and environmental factors;

especially, hypospadias and undescended testes have been linked to the actions of environmental agents (3). A family history with genital abnormalities, delayed puberty, infertility, stillbirths or multiple miscarriages should lower the threshold for molecular genetic investigations (4). A newborn with ambiguous genitalia requires evaluation by a multidisciplinary team that aims to determine the sex of rearing, and eventually designs a long-term management plan in close co-operation with the patient and the family (1). It is generally agreed that

timely diagnosis and management are important as they potentially prevent adverse health outcomes and improve long-term, health-related quality of life (5, 6, 7, 8, 9, 10).

There are currently only a handful of systematic and comprehensive reports on the etiology and management of DSD patients (11, 12, 13, 14, 15, 16, 17). At the same time, there is a vivid ongoing global debate on the treatment policies of these patients (4, 14, 18, 19, 20). The first step toward evidence-based, optimal care of this patient group would be to describe the current status of diagnostics and management in pediatric tertiary centers. In this study, we describe the phenotypic spectrum and molecular genetic findings of DSD patients treated at the Helsinki University Hospital between 2004 and 2014, with special attention on the timing of diagnosis and management of the most prevalent DSDs.

## Patients and methods

We performed an ICD-10 code-based inquiry covering the years from 2004 to 2014 to the electronic patient information system of the Helsinki University Hospital, the tertiary center of the Helsinki metropolitan area (catchment area 1.2 million people in 2014, 0.3 million of which were children and adolescents 0–20 years old (21)) also nationally responsible for treating severe and rare illnesses. We included diagnoses describing phenotypes with deviations in chromosomal, gonadal or phenotypic sexual development. To examine the relation of minor anatomic deviations to actual DSDs, we accepted a wide range of diagnoses (i.e. distal hypospadias and/or unilateral cryptorchidism) in the search (Table 1). We identified altogether 3206 patients who had visited the Pediatric, Pediatric Endocrine and/or Pediatric Surgical Clinics for the evaluation of treatment of DSD or a minor anatomic deviation. The selection of these outpatient clinics was based on their clinical practice: every DSD patient visits either pediatrician, pediatric endocrinologist or pediatric surgeon. Data from other departments (i.e. genetics) were utilized, if available. In the more thorough examination, we included patients with ICD-10 diagnoses describing DSD phenotypes as defined by the LWPES/ESPE consensus group in 2006 (1), and hence, also accepted bilateral cryptorchidism and penoscrotal or perineal hypospadias. We reviewed the electronic patient records of these patients ( $n=885$ ) for clinical features, karyotype and molecular genetics to verify their diagnoses and record their family anamnesis (Fig. 1). The date of diagnosis was defined as the first remark by a pediatric endocrinologist

or a pediatric surgeon in the patient data system on either clinical or molecular genetic DSD diagnosis. Patients over 20 years old and patients with unilateral cryptorchidism, distal hypospadias, epispadias, chordee, cloaca, congenital rectovaginal fistula, imperforate hymen, bifid clitoris or fusion of labia were excluded from the analyses. Boys with congenital adrenal hyperplasia (CAH) were not considered DSD patients since no DSD phenotype was present.

We paid special attention to patients who had either Turner syndrome (TS) or Klinefelter syndrome (KS). The ICD-10 inquiry returned originally 134 patients with TS diagnosis. Manual verification of the patient records revealed, however, that 25 of them did not meet the definition for TS (9) and therefore 109 TS patients were included in the analyses. Similarly, we excluded 5 of the 85 KS patients identified by the original ICD-10 inquiry, as they were found not to meet the criteria for KS (22). The age at diagnosis was available for 90% ( $n=98$ ) of TS patients and 85% ( $n=68$ ) KS patients. Approximately half of TS (51%) and KS (55%) patients were diagnosed during the study period.

To study the management of bilateral cryptorchidism, we evaluated all 46,XY DSD patients retrieved by the ICD-10 search (Table 1). The patient records of boys referred for isolated bilateral cryptorchidism or a more complex phenotype accompanied by bilaterally undescended testes were reviewed. The age at bilateral orchiopexy or the first of the two unilateral orchiopexies was recorded for each boy and used in subsequent analyses.

## Statistical analyses

Correlations between variables were assessed with Spearman rank correlation analysis. Mann–Whitney  $U$  test and chi-square test were used to compare the ages of boys with bilateral cryptorchidism operated between 2004 and 2007 or 2008 and 2014. The data in the text are presented with a mean  $\pm$  s.d. and the data in the tables as median (range) unless otherwise stated.  $P<0.05$  was accepted to indicate statistical significance.

## Results

### Phenotypic spectrum of DSD patients and timing of diagnosis

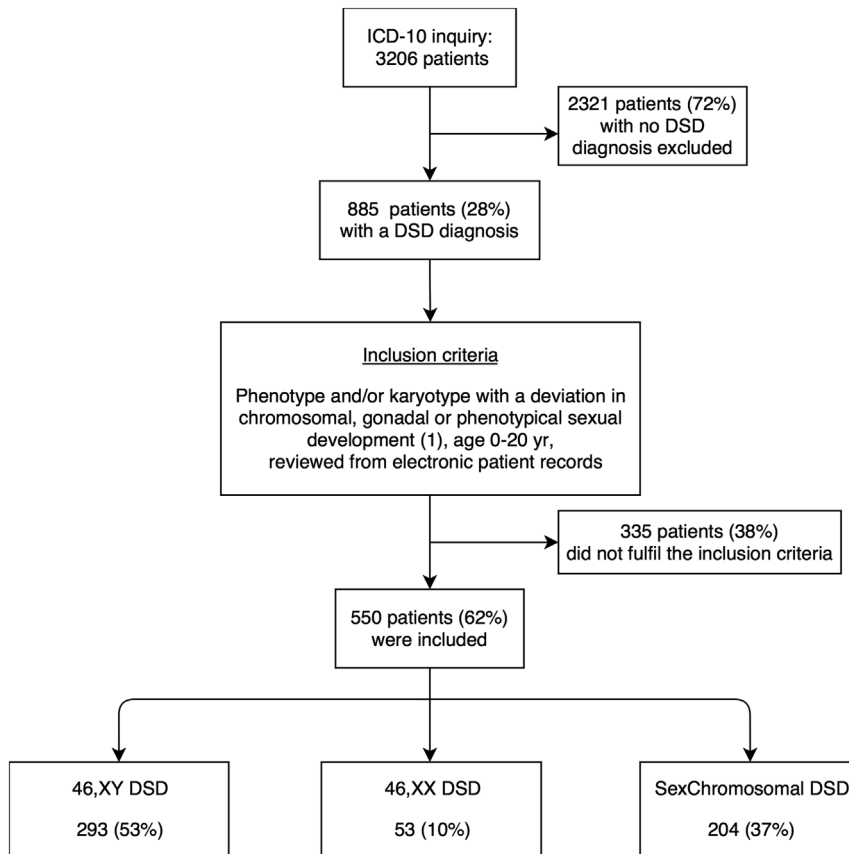
We identified 3206 patients who were evaluated for one or multiple ICD-10 diagnosis of interest in either Pediatric, Pediatric Endocrine and/or Pediatric Surgery Outpatient

**Table 1** ICD-10 diagnoses included in the inquiry.

<b>46,XY DSD</b>	
E34.5	Androgen insensitivity syndrome
Q53.2	Undescended testicle, bilateral
Q54.2	Hypospadias, penoscrotal
Q54.3	Hypospadias, perineal
Q55.00	Absence and aplasia of testis
Q55.01	Anorchia
Q56.1	Male pseudohermaphroditism, not elsewhere classified
Q97.3	Female with 46,XY karyotype
<b>46,XX DSD</b>	
Q52.0	Congenital absence of vagina
Q52.1	Doubling of vagina
Q56.2	Female pseudohermaphroditism, not elsewhere classified
Q98.3	Other male with 46,XX karyotype
E25.00	Salt-losing congenital adrenal hyperplasia
E25.01	Congenital adrenal hyperplasia
<b>Sex chromosome DSD</b>	
Q96.0-Q96.9	Turner syndrome and variants
Q97.0-Q97.9	Other sex chromosome abnormalities, female phenotype, not elsewhere classified
Q98.0,Q98.1	Klinefelter syndrome and variants
Q98.2,Q98.4	
Q98.5-Q98.9	
<b>Unspecified DSD</b>	
E25.9	Adrenogenital disorder, unspecified
Q56.0	Hermaphroditism, not elsewhere classified
Q56.3	Pseudohermaphroditism, unspecified
Q56.4	Indeterminate sex, unspecified
<b>Other diagnoses</b>	
Q53.0	Ectopic testis
Q53.1	Undescended testicle, unilateral
Q53.9	Undescended testicle, unspecified
Q54.0	Hypospadias, balanic
Q54.1	Hypospadias, penile
Q54.4	Congenital chordee
Q54.8	Other hypospadias
Q54.9	Hypospadias, unspecified
Q55.1	Hypoplasia of testis and scrotum
Q55.20	Retractile testis
Q55.28	Unspecified congenital malformations of testis and scrotum
Q55.6	Other congenital malformations of penis
Q55.8	Other specified congenital malformations of male genital organs
Q55.9	Congenital malformation of male genital organ, unspecified
Q52.2	Congenital rectovaginal fistula
Q52.3	Imperforate hymen
Q52.4	Other congenital malformations of vagina
Q52.5	Fusion of labia
Q52.8	Other specified congenital malformations of female genitalia
Q52.9	Congenital malformation of female genitalia, unspecified
Q64.1	Exstrophy of urinary bladder
Q43.7	Persistent cloaca

Clinics at the Helsinki University Hospital between 2004 and 2014. Over two-thirds of these patients (72%) had minor deviations of the genitourinary tract that are not classified as DSDs (Fig. 1 and Table 1). Patients with a DSD diagnosis ( $n=550$ ) were divided into three karyotype-based subgroups as suggested in the Chicago consensus statement (1); 53.3% had 46,XY DSD; 37.1% had sex chromosome DSD and 9.6% had 46,XX DSD (Fig. 1).

Only 21% of 46,XX DSD patients and 7% of 46,XY DSD patients had a molecular genetic diagnosis, and none of those were made with next-generation sequencing methods. Even in patients with a distinct phenotype, such as persistent Müllerian duct syndrome, the diagnosis was seldom confirmed genetically. Age at diagnosis was available for 75% of 46,XY DSD patients, 86% of 46,XX DSD patients and 87% of sex chromosome DSD



**Figure 1** Inclusion criteria and outcomes. Patients retrieved from the ICD-10 inquiry ( $n=3206$ ) and DSD patients ( $n=550$ ) included in the analyses, divided into three subgroups based on 'Consensus statement on management of intersex disorders' (1).

patients. Table 2 shows detailed distribution and timing of diagnoses among these DSD patients; most common diagnoses were bilateral cryptorchidism (23.1%), TS (19.8%) and KS (14.5%). The mean age at diagnosis was  $2.2 \pm 3.8$  years for 46,XY DSD,  $2.4 \pm 4.6$  years for 46,XX DSD and  $5.3 \pm 5.8$  years for sex chromosome DSD. In the majority of patients with 46,XY DSD or 46,XX DSD the diagnosis, based on phenotype or molecular genetic finding, was made immediately after birth, apart from 46,XX DSD patients expressing SRY and those presenting DSD as part of a syndrome.

Consanguinity was seldom reported in our data: only in two families, the parents were either first or second cousins. Two families with multiple DSD cases were identified. One family presented with several cases of persistent Müllerian duct syndrome. In the other family, there were two brothers with a DSD phenotype of unknown etiology (one brother had a micropenis with proximal hypospadias and a vaginal remnant and the other proximal hypospadias). In addition, three families where the DSD patient's sibling had a minor anatomic deviation were identified.

We next investigated the annual number of new DSD patients between 2004 and 2014 ( $n=290$ ), excluding

patients who had been diagnosed before 2004 and had follow-up visits during the survey period. The annual number of new DSD patients remained stable ( $26 \pm 7$  patients per year; mean  $\pm$  s.d., range 10–40) and did not correlate with time ( $r=0.18$ ,  $P=0.6$ ,  $n=11$ ). No significant trends in the annual number of patients were observed either when estimated separately in subgroups of 46,XY DSD (range 5–24,  $r=0.31$ ,  $P=0.4$ ,  $n=11$ ), sex chromosome DSD (range 5–15,  $r=-0.43$ ,  $P=0.18$ ,  $n=11$ ) and 46,XX DSD (range 0–6,  $r=-0.50$ ,  $P=0.12$ ,  $n=11$ ).

### Timing of diagnosis in sex chromosome DSDs: Turner syndrome and Klinefelter syndrome

Our data allowed us to estimate the age at diagnosis of (TS) and (KS) (Fig. 2). For the 80 boys with KS, the diagnosis was set at the mean age of  $6.8 \pm 6.2$  years mostly due to specific language impairment and/or learning difficulties (40%), delayed motor development (13%) or small testicles (11%). Twelve boys were diagnosed already before birth (Fig. 2). Boys referred for evaluation due to developmental issues were aged  $8.0 \pm 3.5$  years, and boys detected due to small testicular size  $15.8 \pm 1.3$  years. Klinefelter boys diagnosed after birth were most frequently referred from

**Table 2** Classification of DSD patients ( $n=550$ ) by molecular diagnosis or phenotype, age at diagnosis and rate of molecular genetic diagnoses.

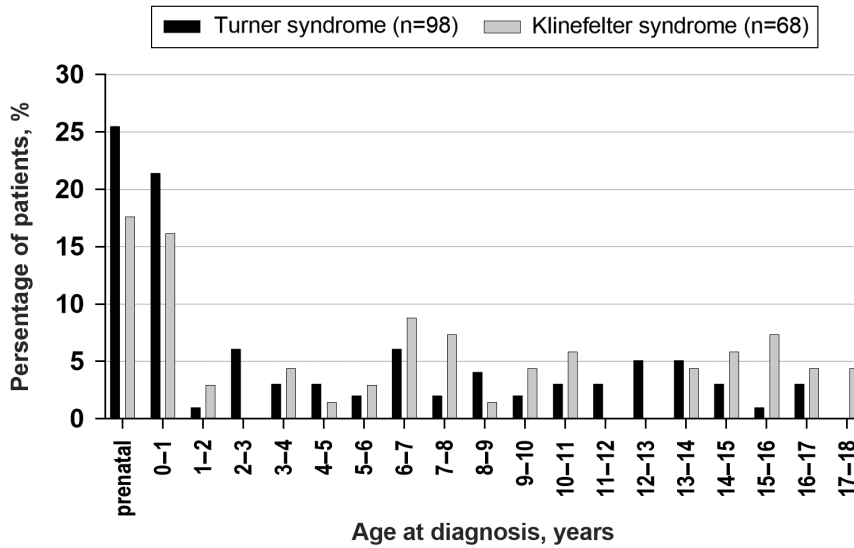
	<i>n</i> (%)	Median (range) age at diagnosis, years	Molecular genetic diagnosis (%)
<b>46,XY DSD</b>			
(A) Disorders of gonadal development			
1. Gonadal regression	27 (4.9)	0.84 (0.01–14.9)	0
2. Ovotesticular DSD	2 (0.4)	7.9 (0.04–15.7)	0
(B) Disorders in androgen synthesis or action			
1. Androgen biosynthesis defect (3BHSD)	1 (0.2)	NA	0
2. Defect in androgen action (AIS)	5 (0.9)	7.7 (0–15.3)	20
3. Disorders of AMH of AMH-receptor (persistent Müllerian duct syndrome)	7 (1.3)	0.7 (0–2.4)	29
(C) Other			
1. Unknown male undermasculinization			
Bilateral cryptorchidism	125 (22.7)	0.9 (0–16.1)	9
Severe hypospadias	55 (10.0)	0.01 (0–1.7)	0
Severe hypospadias and bilateral cryptorchidism	16 (2.9)	0.02 (0–0.04)	0
Micropenis	5 (0.9)	0.01 (0–10.6)	0
Micropenis and bilateral cryptorchidism	6 (1.1)	0.1 (0–10.5)	20
Micropenis and severe hypospadias	8 (1.5)	0.01 (0–0.4)	0
Micropenis, bilateral cryptorchidism and severe hypospadias	4 (0.7)	0 (0–0.1)	0
2. Part of a syndrome (e.g. Prune Belly, Prader–Willi, Noonan)	19 (3.5)	0.02 (0–11.5)	42
3. Cloacal extrophy	13 (2.4)	0.0	0
<b>46,XX DSD</b>			
(A) Disorders of gonadal development			
1. Ovotesticular DSD	1 (0.2)	0.0	0
2. SRY +	4 (0.7)	6.3 (0.3–18.1)	100
(B) Androgen excess			
1. CAH	31 (5.6)	0.0 (0–8.8)	23
(C) Other			
1. Congenital structural abnormalities (e.g. vagina duplex, vaginal atresia, clitoris hyperplasia)	5 (0.9)	0.05 (0–10.6)	0
2. Part of a syndrome (e.g. MRKH, HFGS)	6 (1.1)	6.4 (0–15.4)	0
3. Cloacal extrophy	6 (1.1)	0.0	0
<b>Sex chromosome DSD</b>			
(A) Turner syndrome and variants	109 (19.8)	2.1 (0–17.0)	
(B) Klinefelter syndrome and variants	80 (14.5)	6.7 (0–17.4)	
(C) 45,X/ 46,XY mixed gonadal dysgenesis, ovotesticular DSD	14 (2.5)	0.0 (0–10.1)	
(D) Other			
Structural anomaly of Y chromosome and testis retention	1 (0.2)	0.07	

the local healthcare center (24%), maternity hospital (9%) or department of pediatric neurology (9%).

The girls with TS ( $n=109$ ) were diagnosed at the mean age of  $4.7 \pm 5.5$  years. The age distribution was skewed as twenty-five girls were diagnosed prenatally, and, in 47% of the cases, diagnosis was reached by the age of one year (Fig. 2). Turner girls diagnosed after birth were most frequently referred by child health care clinic (19%), maternity hospital (16%) or another hospital district (6%). The main complaint (41% of all cases) was delayed growth. There were no significant trends in the age at diagnosis in either KS or TS patient groups between 2004 and 2014 (KS range 0–17 year,  $r=0.05$ ,  $P=0.75$ ,  $n=44$ ; TS range 0–17 year,  $r=0.003$ ,  $P=0.98$ ,  $n=56$ ).

### Timing of management in bilateral cryptorchidism

Finally, we reviewed the changes in the management of cryptorchidism by estimating the age at operation for bilateral cryptorchidism before (years 2004–2007) and following the release of the Nordic consensus (years 2008–2014) on treatment of undescended testes (8). Altogether, during the whole study period (2004–2014), 142 patients were operated and the average age of operation for bilateral cryptorchidism was  $45.1 \pm 44.8$  months. The annual number of operated patients (range 8–17) did not increase significantly during the survey period ( $R=0.41$ ,  $P=0.21$ ,  $n=11$ ). However, the age at operation declined after the release of the consensus from  $54.5 \pm 43.0$  months to  $41.1 \pm 45.3$  months ( $P=0.0001$ ).



**Figure 2** Percentage of TS (n=98) and KS (n=68) patients diagnosed per age group. Data obtained from patients visiting the outpatient clinics of a single tertiary center between 2004 and 2014 whose age at diagnosis was available.

Indeed, before the Nordic consensus statement, all bilateral cases (n=42) were operated after the age of 1 year but after 2007, almost one-third (28 of 100 boys) was operated before the age of 1 year (P=0.0001, Fig. 3). To exclude boys with acquired cryptorchidism and cases of congenital cryptorchidism missed in early childhood, we limited the analysis to those boys operated before the age of three (23). In this subanalysis, the age at operation had declined from 24.5±5.4 months in boys operated between 2004 and 2007 to 13.7±5.3 months in those operated after 2007 (P<0.0001).

### Discussion

Attempts to estimate the etiological distribution of DSDs according to the 2006-launched DSD classification system

are few. In our series, approximately half of the patients had 46,XY DSD, whereas only one-tenth (9.6%) had 46,XX DSD. The proportion of 46,XY DSD patients is similar to the results reported from Turkey (47%) (12), South Africa (57%) (13) and North India (52.5%) (17), and somewhat lower than has been reported from Indonesia (68%) (16). On the other hand, our series contained more cases with sex chromosome DSDs than the previously mentioned reports, which may reflect differences in prenatal screening policies. Of note, in a recent Danish nationwide study, the estimated prevalence of androgen insensitivity syndrome was 2.3 per 100,000 live born females (24), which appears close to our frequency estimate based on the current catchment area of our tertiary center. In our study, congenital adrenal hyperplasia (CAH) was expectedly the most common cause of 46,XX DSD. After the implementation of the newborn screening in Finland



**Figure 3** Age at orchiopexy. Age at bilateral orchiopexy or first of the two unilateral orchiopexies in a single tertiary center in 142 boys operated between 2004 and 2014. 42 boys were operated between 2004 and 2007 before the release of the Nordic consensus on treatment of undescended testes (8) and 100 boys thereafter.

in 2015, six new CAH patients have been diagnosed by it. We anticipate that screening leads to better diagnostic yield and conceivably better care for this patient group in the future.

Since early diagnosis of DSD is considered important (7, 25, 26), the efficacy of the diagnostic procedures of most frequent patient groups merits evaluation. The mean age at diagnosis of TS patients in our series was 4.7 years, a result similar to the findings reported from the United States and Spain (26, 27). A quarter of our TS patients was diagnosed prenatally, based on the voluntary, combined first trimester screening (a nuchal translucency scan and maternal serum screening followed by a prenatal chromosome test, if necessary). After birth, the most frequent symptom guiding to diagnosis was short stature (41%), the leading clinical cue also in other studies (26, 27). The diagnosis of KS patients in our series was even further delayed, and the diagnosis was typically suspected during childhood due to developmental problems, and during adolescence due to small testicular size. If the frequency of KS was 1/667 male births in our area (28), we could roughly estimate that, between 2004 and 2006, ~20–30% of the boys with KS were diagnosed before the age of 8 years. This low diagnostic yield is in agreement with the Danish data suggesting that only 10% of Danish KS patients are diagnosed before the age of 14 years (29). One possibility to expedite diagnostics would be to include sex chromosome abnormalities in the national newborn screening programs, an idea that has been well-taken by parents of KS patients and pediatricians (10, 30). Also peer groups of TS (e.g. Turner Syndrome Society of the United States) are in favor of diagnosing TS as early as possible in order to anticipate the possible need of services required for variable defects in childhood (31). On the other hand, many pregnancies, in which TS is prenatally diagnosed, are currently terminated, although the severity of the TS phenotype is hard to predict and a postnatal re-evaluation of chromosomes is needed for an accurate diagnosis (9, 31).

The timing of orchiopexy is highly variable, and for example, 95% of surgeries were performed after the first year of life in German hospitals (32), and in Singapore, the corresponding number was 70% (33). Nordic guidelines for the treatment of cryptorchidism were renewed at 2007 and accordingly the age for orchiopexy was advanced to 6–12 months (8). We tested whether implementation of the guideline in our tertiary center functions as an indicator for standard of care. Between 2004 and 2007, before the release of the Nordic consensus statement (8), not a single

boy with bilateral cryptorchidism was operated before the age of one year in our hospital, whereas thereafter, the age at orchiopexy has decreased at least by 10 months. Our finding is in line with the results published from Sweden and Norway (34, 35).

In this cohort, 21% of patients with 46,XX DSD and only 7% of 46,XY DSD were diagnosed genetically. In the largest patient group of 46,XX DSD, CAH, the current standard of diagnostics is still based on steroid hormone and other biochemical analyses instead of molecular genetics (36). For 46,XY DSD boys, it seems that the etiology remains largely unknown in the clinical setting. In the research setting, however, the overall genetic diagnostic rate achieved with a targeted DSD gene panel, can increase up to 43% (37). An accurate molecular genetic diagnosis is probably valuable in planning the long-term treatment of the patient, although more research is needed to show if the ever more precise diagnostics is associated with beneficial changes in clinical practice from the patient's perspective (38). While awaiting such evidence, it is still reasonable to conclude that the molecular genetic diagnostic rate should be improved by applying NGS methodology.

Our retrospective study has certain limitations. The ICD-10 classification has multiple different diagnosis codes describing the same phenomenon. To overcome this, we evaluated the patient records and chose the diagnostic code, which best described the patient's phenotype. Moreover, often there are no specific ICD-10 codes reserved for rare genetic diseases.

## Conclusions

We present data from one of the largest series of DSD patients to date. Analysis of patients with chromosomal DSD shows that patients with TS or KS are diagnosed relatively late and often the initiative for the diagnostic workup is related to special needs of the patients (KS) or deceleration of growth (TS). It is feasible that age at diagnosis in patients with TS or KS and the age at operation of bilateral cryptorchidism could serve as indices for monitoring the diagnostic efficiency and management of DSD patients within and between different tertiary centers providing care for this vulnerable patient group.

### Declaration of interest

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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### Author contribution statement

All authors have provided substantial input to the concept of the study and drafting the article. E K, P M and T R designed the study. E K and A T collected the data. P M, S T, M H and T R advised in interpreting the patient data. E K analyzed the data and wrote the first draft of the manuscript. All authors contributed in revising and editing the manuscript.

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