

RESEARCH

Open Access

Holt Oram syndrome: a registry-based study in Europe

Ingeborg Barisic^{1*}, Ljubica Boban¹, Ruth Greenlees², Ester Garne³, Diana Wellesley⁴, Elisa Calzolari⁵, Marie-Claude Addor⁶, Larraitz Arriola⁷, Jorieke EH Bergman⁸, Paula Braz⁹, Judith LS Budd¹⁰, Miriam Gatt¹¹, Martin Haeusler¹², Babak Khoshnood¹³, Kari Klungsoyr¹⁴, Bob McDonnell¹⁵, Vera Nelen¹⁶, Anna Pierini¹⁷, Annette Queisser-Wahrendorf¹⁸, Judith Rankin¹⁹, Anke Rissmann²⁰, Catherine Rounding²¹, David Tucker²², Christine Verellen-Dumoulin²³ and Helen Dolk²

Abstract

Background: Holt-Oram syndrome (HOS) is an autosomal dominant disorder characterised by upper limb anomalies and congenital heart defects. We present epidemiological and clinical aspects of HOS patients using data from EUROCAT (European Surveillance of Congenital Anomalies) registries.

Methods: The study was based on data collected during 1990–2011 by 34 registries. The registries are population-based and use multiple sources of information to collect data on all types of birth using standardized definitions, methodology and coding. Diagnostic criteria for inclusion in the study were the presence of radial ray abnormalities and congenital heart disease (CHD), or the presence of either radial ray anomaly or CHD, with family history of HOS.

Results: A total of 73 cases of HOS were identified, including 11 (15.1%) TOPFA and 62 (84.9%) LB. Out of 73 HOS cases, 30.8% (20/65) were suspected prenatally, 55.4% (36/65) at birth, 10.7% (7/65) in the first week of life, and 3.1% (2/65) in the first year of life. The prenatal detection rate was 39.2% (20/51), with no significant change over the study period. In 55% (11/20) of prenatally detected cases, parents decided to terminate pregnancy. Thumb anomalies were reported in all cases. Agenesis/hypoplasia of radius was present in 49.2% (30/61), ulnar aplasia/hypoplasia in 24.6% (15/61) and humerus hypoplasia/phocomelia in 42.6% (26/61) of patients. Congenital heart defects (CHD) were recorded in 78.7% (48/61) of patients. Isolated septal defects were present in 54.2 (26/48), while 25% (12/48) of patients had complex/severe CHD. The mean prevalence of HOS diagnosed prenatally or in the early years of life in European registries was 0.7 *per* 100,000 births or 1:135,615 births.

Conclusions: HOS is a rare genetic condition showing regional variation in its prevalence. It is often missed prenatally, in spite of the existence of major structural anomalies. When discovered, parents in 45% (9/20) of cases opt for the continuation of pregnancy. Although a quarter of patients have severe CHD, the overall first week survival is very good, which is important information for counselling purposes.

Keywords: Holt Oram syndrome, Congenital anomalies, Prenatal diagnosis, Epidemiology, Europe

* Correspondence: ingeborg.barisic@kdb.hr

¹Zagreb Children's Hospital, Medical School, University of Zagreb, Klaićeva 16, Zagreb 10 000, Croatia

Full list of author information is available at the end of the article

Background

Holt-Oram syndrome (HOS, OMIM 142900) is a rare autosomal dominant multiple malformation syndrome characterised by high penetrance and variable expression of upper limb abnormalities, congenital heart defects (CHD) and/or conduction abnormalities [1,2].

Sequence variants of *TBX5* gene, a member of the T-box family of transcription factors, have been identified to affect function in 75% of HOS cases [3-5]. Most are truncating alterations that result in haploinsufficiency, but occasionally sequence variations can lead to extension of the *TBX5* protein [6,7]. Different types of sequence variants can cause complete loss or reduction of *TBX5* protein function by affecting nuclear localisation of the protein or by disrupting its interaction with other transcriptional cofactors and downstream target genes. Some sequence variants may result in the gain-of-function causing a similar phenotype [8-13]. Since under- and overexpression cause the same phenotype, *TBX5* function is considered to be gene dosage sensitive [13].

The carriers of the *TBX5* allelic variants affecting function show high intra- and interfamilial variability of clinical presentation. These variations can be due to the type and location of the sequence alteration, but also to other modifier factors, e.g., sequence variations in enhancers regulating *TBX5* expression during heart development [14].

Clinical diagnostic criteria of HOS include pre-axial radial ray malformations in at least one upper limb and CHD and/or conduction defects [15,16]. When a heart anomaly is not present, there should be a family history of HOS consistent with an autosomal dominant type of inheritance [2,5,15]. The spectrum of skeletal upper limb defects ranges from an abnormal carpal bone or triphalangeal/fingerlike thumb to bilateral phocomelia. CHD is present in 75% of individuals with HOS. Ostium secundum atrial septal defect (ASD) and ventricular septal defect (VSD) are the most common, but other more severe heart anomalies have been reported as well [2,17]. Conduction heart disease may occur in the absence of structural anomaly. No correlation between the severity of heart and limb defects has been established.

Several large clinical series of HOS patients have been published examining the clinical presentation, differential diagnosis and diagnostic criteria [16,18,19], but population-based epidemiological studies are rare because of the need for large populations and standardised data collection. The prevalence of HOS is estimated to be 0.95/100,000 births, based on a single epidemiological study from Hungary [20]. Most clinical reports refer to live born (LB), children or adults, while there is little information on fetal deaths (FD), terminations of pregnancy after prenatal ultrasound detection of severe anomaly/

anomalies (TOPFA) and on patients diagnosed in the neonatal period.

The aim of this study was to investigate the epidemiological and clinical aspects of HOS patients diagnosed prenatally or in the early years of life, using data notified to the European Surveillance of Congenital Anomalies (EUROCAT) database.

Methods

The study was based on data routinely collected in the period from 1990 to 2011 by 34 EUROCAT registries in 16 countries covering approximately 30% of Europe's annual birth population and extracted from the central database in June 2013. The EUROCAT registries are population-based and use multiple sources of information to collect data on all major structural congenital anomalies, chromosomal abnormalities, and other genetic and environmental conditions presenting with structural defects among LB, FD with gestational age (GA) ≥ 20 weeks, and TOPFA using standardized definitions and coding. Six registries include cases diagnosed up to one week after birth, three up to one month, and 25 registries include cases diagnosed up to at least one year of age. A detailed description of registries, methods of case ascertainment, data collection and processing is available elsewhere [21,22].

The central EUROCAT database was searched for the International Classification of Diseases (ICD)/British Paediatric Association (BPA) version 9 (759.842, 759.84), ICD10/BPA (Q87.20) and OMIM (142900) codes assigned to cases of HOS. Diagnostic criteria for inclusion in the study were the presence of radial ray abnormalities and CHD, or the presence of either radial ray anomaly or CHD, with family history of HOS [2,4]. Minor anomalies, including those detectable exclusively by x-ray (e.g., carpal bone defects, hypoplasia of hand/arm/shoulder muscles, sloping shoulders), functional limb defects (e.g., minor limitation of movements of thumbs, elbows, or shoulders) or heart defects (e.g., conduction abnormalities) were not systematically recorded, as the EUROCAT focus is on collecting data on major structural anomalies [21].

In two-thirds of EUROCAT registries, clinical geneticists take part in the examination and diagnosing of all patients with dysmorphic features and congenital anomalies, and in the remaining registries clinical geneticists participate in the examination of selected cases [23]. This genetic expertise allows EUROCAT to collect and analyse data on rare genetic syndromes that manifest a characteristic pattern of anomalies prenatally, at birth or in early infancy [23-25]. A medical geneticist reviewed all records including textual description in order to include all relevant clinical information. Local registries were contacted to provide additional/missing information.

The variables included in the analysis were the time of diagnosis (pre- or postnatal), birth outcome (proportion of LB, FD at GA ≥ 20 weeks, TOPFA), type of congenital anomalies (ICD9/ ICD10 code and written text), infant sex, survival up to 1 week of age, maternal and paternal age at delivery, birth weight (BW), GA at birth or termination of pregnancy, use of assisted reproductive techniques (ART), multiple pregnancy (twin or triplet), and family history. Until very recently, data on genetic testing were not systematically collected and therefore were not included in this study.

Statistics

Descriptive data are presented as numbers and percentages for categorical data. Means and 95% confidence intervals, based on Poisson distribution, were used to calculate prevalence and a Generalized Linear/Nonlinear model based on Poisson distribution was used for statistical testing of time trends. A difference between two proportions test was used to test prevalence differences between registries and the total prevalence rate. A chi-square test was performed to determine differences in maternal age distribution between HOS and other EUROCAT cases. Maternal age comparisons were restricted to the registries/time period with available maternal age denominator data (63% of EUROCAT total birth population). Statistical analysis was performed using STATISTICA 6.1 StatSoft Inc. 1983–2003 (serial number AGA304B211928E61).

Results

Between January 1990 and December 2011, a total of 73 cases of HOS were identified in the European population covered by 34 European registries included in the EUROCAT network. There were 11 (15.1%) TOPFA, no FD, and 62 (84.9%) LB. Two of the 62 LB children died in the neonatal period (3.2%). Among the patients for which the time of diagnosis was known ($n = 65$), 30.8% were diagnosed prenatally, 55.4% at birth, 10.7% in the first week of life, and 3.1% in the first year of life.

The prenatal detection rate was 39.2% (20/51) with no significant change between the 1990–2000 and 2001–2011 periods ($P > 0.05$) (Table 1). The mean GA at prenatal diagnosis by obstetric ultrasound was 18.6 ± 4.6 (range 14–26) gestational weeks. In 55% (11/20) of cases

with prenatally diagnosed severe anomaly/anomalies, parents decided to terminate the pregnancy. The mean GA at termination was 21.1 ± 1.8 (range 14–35) gestational weeks. The diagnosis was verified by *post mortem* examination in 10 of 13 cases (eight TOPFA and two neonatal deaths). In 31 out of 51 cases with available data (60.8%), prenatal ultrasound was performed but did not detect any anomaly, although major anomalies (e.g., absence of forearm, bilateral radial aplasia, tetralogy of Fallot, pentalogy of Fallot, etc.) were present in 10/31 patients detected postnatally. Three undetected cases were familial.

Characteristics of the HOS patients are shown in Table 2. The male-to-female ratio was 1.1:1. The mean GA in LB was 39.3 ± 3.1 (range 33–42) in males and 38.4 ± 2.9 (range 27–42) in females. The mean BW was 2998 ± 370 g (range 2240–3800 g) for males and 3241 ± 458 g (range 2400–4200 g) for females. Of 62 live born cases, 96.8% survived the first week of life.

Eleven families had more than one affected member. Maternal age distribution did not differ significantly from that of the total EUROCAT population ($P = 0.06$). Multiple pregnancies were not noted. Three of 34 patients with known information on ART were conceived by induced ovulation. There were no cases of *in vitro* fertilization (IVF) or intracytoplasmic sperm injection (ICSI). Data on karyotyping were available for 46.1% of cases and all results were normal.

Description of the type and frequency of major congenital anomalies was available for 61 HOS patients. The results are presented in Table 3, together with those on the three large series of HOS patients published so far. Thumb anomalies were present in all patients. Agenesis/hypoplasia of radius was present in 49.2% (30/61), ulnar aplasia/hypoplasia in 24.6% (15/61), and humerus hypoplasia/phocomelia in 26/61 (42.6%) patients.

CHD was recorded in 78.7% (48/61) of patients. Isolated septal defects were present in 54.2%, ASD + VSD in 8.3% and various other CHD in 12.5% of cases (Table 4). There were 25% of patients with complex/severe heart defects: tetralogy of Fallot (1), pentalogy of Fallot (1), pulmonary valve atresia (1), atrial septal defect and tricuspid atresia (1), VSD and tricuspid atresia (1), AVSD with multiple ventricular septal defects (2), AVSD with multiple ASD (1), common arterial truncus (1),

Table 1 Outcome of pregnancies and prenatal detection of Holt Oram syndrome in the EUROCAT registries, 1990-2011

Monitored period	Total no. of patients	LB	FD	TOPFA	Prenatally detected/number of patients with available data	Prenatal detection rate% (95% CI)
1990-2000	42	38	0	4	11/28	39.3 (95% CI 21.2-57.4)
2001-2011	31	24	0	7	9/23	39.1 (95% CI 19.2-59.1)
1990-2011	73	62	0	11	20/51	39.2 (95% CI 25.8-52.6)

Abbreviations: LB live born, FD fetal deaths, TOPFA termination of pregnancy for fetal anomaly, 95% CI 95% Confidence Interval.

Table 2 Descriptive epidemiological data on patients with Holt Oram syndrome, EUROCAT registries, 1990-2011

Characteristics	Number of HOS patients N = 73 (%)	
Sex		
Male	38 (52.8)	
Female	34 (47.2)	
Unknown	1	
Gestational age, wk	N = 59	
	(live births with known gestational age and sex)	
	M	F
	n = 27	n = 32
<37	5 (18.5)	4 (12.5)
37-41	20 (74.1)	25 (78.1)
≥42	2 (7.4)	3 (9.4)
Birth weight (g)	N = 47	
	(live births with known gestational weight and sex)	
	M	F
	N = 20	N = 27
<1500	0	0
1500-1999	3 (15.0)	1 (3.7)
2000-2499	6 (30.0)	8 (29.6)
2500-2999	10 (50.0)	10 (37.1)
3000-3499	1 (5.0)	7 (25.9)
>3500	0	1 (3.7)
Positive family history	15.1% (11/73)	
Mean maternal age (range)	28.3 ± 4.6 (17-41)	
Mean paternal age (range)	32.7 ± 1.3 (19-62)	

wk: weeks.
g: grams.

pulmonary valve atresia (1), aortal valve insufficiency ASD and VSD (1), and double outlet right ventricle (DORV) (1).

CHD was present in all but one familial case (90%). In this maternal case, there was aplasia of the thumb and radial hypoplasia without CHD. Distribution of CHD types among familial and sporadic cases is shown in Table 4.

The prevalence of HOS was restricted to data from 16 registries with an above average prevalence of genetic syndromes and microdeletions (EUROCAT average is 4.8 *per* 10,000 births) according to the Data Quality Indicators developed by EUROCAT (<http://www.eurocat-network.eu/content/DQI-2013.pdf>). This was done in order to ensure a more homogeneous ascertainment of cases. During the 1990-2011 period, a total population of 5,017,781 births were monitored in these registries and 37 cases of HOS identified. Therefore, the mean

prevalence of HOS diagnosed in the early years of life was 0.7 *per* 100,000 births or 1: 135,615 births. The prevalence rates for 1990-2000 and 2001-2011 were 1.1 and 0.4 *per* 100,000 births, respectively ($P = 0.03$) (Table 5). The number of HOS patients and prevalence rates *per* registry are shown in Table 6.

Discussion

The Holt-Oram syndrome is an autosomal dominant condition associated with defective development of the radial ray and cardiac structures, resulting in a wide spectrum of phenotypes. It needs to be emphasized that there is no intellectual impairment which is important information for genetic counselling of affected families. The diagnosis is based on established clinical criteria and can be confirmed by molecular genetic testing in a proportion of cases. The diagnosis is not difficult in the presence of typical clinical features and positive family history. In isolated cases, the differential diagnosis will include hand-heart syndromes type II (Tabatznik) and III (Spanish type), other genetic syndromes with upper limb anomalies (e.g., thrombocytopenia-absent radius syndrome, Fanconi anaemia, *SALL4*-related disorders, ulnar mammary, Kaufmann McKusick, Roberts or Nager syndromes), chromosomal anomalies, associations such as VACTERL and rare teratogenic embryopathies (thalidomide, valproate) [2,26]. In most cases, these conditions can be excluded without major difficulties by careful clinical examination and appropriate diagnostic evaluation including cytogenetic and molecular testing. These tests, including genetic testing for *TBX5*, 22q11.2 microdeletion and Fanconi anaemia, were also performed in the studied patients as a part of the routine clinical evaluation for establishing the final diagnosis. We are not able to present these data, as we have only recently started to collect data on genetic testing in a systematic way.

Prenatal diagnosis

As HOS is rare and most cases are sporadic, it is mostly reported in familial cases when a more detailed ultrasound examination is performed rather than by prenatal screening [27,28]. The radius and ulna are easy to see at 13-16 weeks, and most cardiac anomalies, with exception of ASD and small VSD, are clearly visible on ultrasound screening for anomalies at 18-20 weeks. Our study showed that over 60% of cases were not suspected prenatally, although many presented with major anomalies that could have been easily detected by prenatal ultrasound and although some cases were familial. In addition, the prenatal detection rate did not improve over time.

Parents of the 9 of 20 patients diagnosed prenatally decided to continue the pregnancy. Two cases were

Table 3 Major congenital anomalies in Holt Oram syndrome: EUROCAT data and in previously published studies

TYPE OF ANOMALY	EUROCAT STUDY	Newbury-Ecob et al., 1996 [16]	Hurst et al., 1991 [19]	Smith et al., 1979 [18]	
	N = 61 (%)	N = 55 (%)	N = 43 (%)	N = 39 (%)	
Congenital heart anomalies*					
Atrial septal defect	28 (45.9)	20 (36.4)	20 (46.5)	14 (35.9)	
Ventricular septal defect	23 (37.7)	15 (27.2)	3 (7.0)	9 (23.1)	
Atrioventricular septal defect	3 (4.9)			1 (2.6)	
Pulmonary atresia/stenosis	2 (3.3)	1 (1.8)		1 (2.6)	
Double outlet right ventricle	1 (1.6)			1 (2.6)	
Aortic valve insufficiency	1 (1.6)			1 (2.6)	
Aortic valve stenosis		2 (3.6)			
Tricuspid atresia	2 (3.3)				
Mitral valve abnormality		3 (5.5)	1 (2.3)	2 (5.1)	
Patent ductus arteriosus	3 (4.9)	1 (1.8)	1 (2.3)	1 (2.6)	
Pentalogy of Fallot	1 (1.6)				
Tetralogy of Fallot	1 (1.6)	2 (3.6)			
Common arterial truncus	1 (1.6)				
Dextrocardia	1 (1.6)	1 (1.8)	1 (2.3)	1 (2.6)	
Right aortic arch				1 (2.6)	
Congenital heart disease, NOS	1 (1.6)			4 (10.3)	
LIMBS					
Thumbs					
	Triphalangeal/digital thumb	24 (39.3)	10 (18.2)	10 (23.3)	24 (61.5)
	Hypoplasia of thumb	16 (26.2)	21 (38.2)	9 (20.9)	13 (33.3)
	Absence of thumb	30 (49.1)	27 (49.1)	10 (23.3)	16 (41.0)
	Accessory/bifid thumb	4 (6.6)			1 (2.6)
	Syndactyly of thumb	3 (4.9)	7 (12.7)		6 (15.4)
Fingers					
	Aplasia of hand and/or fingers	8 (13.1)	6 (10.9)		8 (20.5)
	Hypoplasia of hand and/or fingers	8 (13.1)	7 (12.7)		8 (20.5)
	Syndactyly	20 (32.8)	6 (10.9)		16 (41.0)
Lower arm					
	Agensis radius	14 (23.0)	13 (23.6)		8 (20.5)
	Hypoplastic radii, short forearms	16 (26.2)	24 (43.6)	9 (20.9)	12 (30.8)
	Ulnar hypoplasia	11 (18.0)	20 (36.4)		7 (17.9)
	Ulnar aplasia	4 (6.6)	4 (7.3)		1 (2.6)
	Synostosis of radius and ulna	12 (19.7)	32 (58.1)		5 (12.8)
Upper arm					
	Humerus hypoplasia	24 (39.3)	28 (50.9)		10 (25.6)
	Humerus aplasia/Phocomelia	2 (3.3)	1 (1.8)	3 (7.0)	1 (2.6)
	Clavicles, abnormalities	8 (13.1)	40 (72.2)		11 (28.2)

Table 3 Major congenital anomalies in Holt Oram syndrome: EUROCAT data and in previously published studies (Continued)

THORAX			
	Rib anomalies	2 (3.3)	5 (12.8)
SPINE			
	Hemivertebra	1 (1.6)	1 (2.6)
	Fusion of vertebrae		1 (2.6)

*Congenital heart anomalies are presented *per* type of anomaly and not *per* patient.
 NOS = non specified.

diagnosed late (at 32 and 36 gestational weeks). Among the remaining seven, there were three familial cases suggesting that families accept and tolerate well the clinical consequences of the genetic disorder in their family, as there is no intellectual impairment.

Clinical manifestations

The distribution of gender was equal. The intrauterine growth and development was not affected. Over two-thirds of HOS patients were born at term and of 50 patients born after 37 weeks of gestation, only four (8%) weighed less than 2,500 g. Likewise, the first week survival was excellent, taking into account a high rate of severe congenital heart anomalies.

By definition, all HOS patients have upper extremity anomalies. Bone abnormalities are usually bilateral and asymmetric, with left side often more affected than the right side. Most skeletal manifestations are visible on inspection but some patients have only functional abnormalities or carpal/metacarpal bone anomalies visible on x-ray. These functional and subclinical skeletal manifestations of HOS were not systematically reported in our dataset, as EUROCAT records major anomalies, while recording of minor anomalies and functional symptoms is optional and would be included in the database only if associated with a major defect. In addition, some of these manifestations are difficult to assess prenatally, at *post mortem*, or in the neonatal period/infancy, when most of our patients were diagnosed.

Table 4 Type of congenital heart defect in sporadic and familial patients with Holt Oram syndrome (N = 48)

Type of CHD	Sporadic	Familial	TOTAL
ASD	13 (34.2%)	5 (50.0%)	18 (37.5%)
VSD	7 (18.4%)	1 (10.0%)	8 (16.7%)
ASD + VSD	2 (5.3%)	2 (20.0%)	4 (8.3%)
Complex	12 (31.6%)	-	12 (25%)
Other	4 (10.5%)	2 (20.0%)	6 (12.5%)
TOTAL	38	10	48/61* (78.7%)

*Number of HOS patients with available data on type and frequency of major congenital anomalies.

CHD - congenital heart defect, ASD - atrial septal defect, VSD - ventricular septal defect.

The most common thumb anomalies were absence of thumbs, followed by triphalangeal/digital thumbs and thumb hypoplasia. Radial hypoplasia was only slightly more common than radial agenesis. The involvement of the ulna was less frequent than of the radius. Upper arm involvement was present in over 40% of patients. These findings are largely in agreement with previous clinical reports [16,18,19]. Due to the previously mentioned methodology issues, synostosis of radius and ulna and subtle shoulder anomalies (narrow shoulders, short clavicles/hypoplasia of head of humerus, and hypoplastic musculature of the shoulder girdle) are probably under-reported. Contrary to the observation of Newbury Ecob et al. [16], no difference between isolated and familial cases in the severity of thumb aplasia, radial and ulnar involvement was observed. The predominant ulnar involvement and atypical findings in isolated cases were not noted. Only 1 in 30 patients had phocomelia, which is somewhat less than previously observed [16,29]. Smith et al. [18] found phocomelia to be present more often in familial cases and in females. Both cases of phocomelia found in our series were recorded in males and were sporadic. Some studies noted that skeletal defects were more severe in females than in males [18,29-31], but that males had a greater number of bones involved than females [18]. In this study, we did not observe any sex difference in the type and number of skeletal defects.

CHD occur in approximately 75% of HOS patients. A variety of structural heart anomalies are seen, with ASD and VSD being the most common. The proportion of CHD found in this study was 78.7%. In particular, we observed a higher rate of VSD and a higher rate of

Table 5 Prevalence of Holt Oram syndrome patients in 16 selected EUROCAT registries, 1990-2011

Monitored period	Total births	Total no. of patients	Birth prevalence per100 000 (95% CI)
1990-2000	2 387 019	26	1.1 (0.69-1.51)
2001-2011	2 630 762	11	0.4 (0.2-0.6)
1990-2011	5 017 781	37	0.7 (0.32-1.08)

Abbreviations: 95% CI, 95% Confidence Interval.

Note: the registries were selected according to the <http://www.eurocat-network.eu/content/DQI-2013.pdf>.

Table 6 Prevalence of Holt Oram syndrome patients per 16 selected EUROCAT registries, 1990–2011

REGISTRY	Population	No. of HOS patients (confirmed familial cases)	Prevalence per 100000
Styria (Austria)	229506	2 (2)	0.9
Antwerp (Belgium)	341573	2	0.6
Hainaut (Belgium)	277204	4 (2)	1.4
Dublin (Ireland)	489614	1	0.2
Odense (Denmark)	121532	1	0.8
Paris (France)	703650	3	0.4
Mainz (Germany)	75496	1	1.3
Cork And Kerry (Ireland)	141421	1	0.7
N Netherlands (NI)	421026	7 (2)	1.7*
Vaud (Switzerland)	166950	4 (1)	2.4*
Glasgow (UK)	122803	1	0.8
N W Thames (UK)	661527	2	0.3
Wessex (UK)	492559	5	1.0
Thames Valley (UK)	291759	1 (1)	0.3
Northern England (UK)	382900	1	0.3
Malta	98261	1 (1)	1.0
TOTAL	5017781	37	0.7

*Registries with statistically significantly higher prevalence rates compared to the total average prevalence rate ($P < 0.05$).

severe CHD than in previous reports [16–19]. Our series of patients included a more severely affected subset of patients compared with the clinical series that describe mostly adult patients coming from affected families. HOS patients in our study were diagnosed mostly at birth or prenatally, and pregnancies with severe CHD that resulted in TOPFA were also included. Additionally, some of the VSDs diagnosed at birth would eventually resolve and will not be recorded later in childhood or in adulthood. Although the number of familial cases in our series is small, it is of note that none of them had complex CHD.

Cardiac conduction abnormalities are commonly found in HOS. They are more likely to occur in those with a structural heart defect, but it is reported that about 40% of cases have conduction abnormalities alone [16]. We were not able to study this manifestation of HOS, as recording of functional abnormalities is optional and reported data are not complete. It is therefore possible that some patients with upper limb anomalies and only conduction heart defects will be diagnosed later in adolescence or adulthood when the symptoms of heart disease develop.

Other associated anomalies have also been described in HOS patients and may include craniofacial, tracheal, pulmonary, vertebral, renal and lower limb anomalies

[5,32–36]. These can be incidental findings or may be atypical cases of HOS due to specific sequence variants of the *TBX5* gene [11,36,37]. We observed single cases with cleft uvula, brain cyst, spleen anomaly, pylon duplex, ectopic kidney and hemivertebra. Lower limb anomalies were found in 3 patients. The types of anomalies (one bilateral hip dislocation, one lower limb shortening, one club foot) would suggest random association rather than the clinical spectrum of HOS.

Two cases with renal anomalies and a case with hemivertebra were evaluated for the possible diagnosis of VATER, but this was not conclusive. The patient with hemivertebra had ASD type secundum, radial and thumb aplasia on the right hand and phocomelia on the left side, which is more indicative of HOS. Data on molecular tests for cases with associated anomalies were not available.

The proportion of familial cases was 15.1%, which is consistent with the results of the only population study on HOS [20] and contrasts with clinical reports citing that up to 85% of cases are familial due to the ascertainment bias [16,20,23]. The mean parental age did not differ from the general EUROCAT population, although a wide range (19–62 years) of paternal ages was observed.

ART methods are known to be associated with a higher risk of congenital anomalies, especially congenital heart defects [38,39] and are a risk factor for some genetic syndromes [25,40]. In addition, ART could facilitate propagation of pre-existing mutations that are associated with impaired fertility, e.g., in the sperm of older men [41]. Our sample, although small, did not show correlation between the ART techniques and HOS.

Prevalence

The results of this European study show that HOS is a very rare condition with an average prevalence of 0.7 per 100,000 births and a high regional variation [range 0.3 (N W Thames)–2.4 (Vaud) or 1: 330,763 to 1: 41,737]. The prevalence is higher in registries where familial cases are recorded. The mean prevalence of 1: 135,615 births represents a minimum figure and refers to a group of patients with obvious clinical presentation.

The only population-based study on HOS was conducted in Hungary covering the 1975–1988 period [20]. The established prevalence of 0.95 per 100,000 births is in agreement with the prevalence of 1.0 per 100,000 births recorded in our dataset for 1990–2000. Interestingly, there is a significant decrease in the prevalence rate for 2001–2011 period, for which there is no obvious explanation and requires further monitoring.

Strengths and limitations of the study

This is the largest population-based study on HOS to date. The results are based on the same methodology

and case description and there is sufficient genetic expertise to have confidence in the clinical diagnosis. All cases met strict diagnostic criteria as proposed by McDermott [2005, 2013], but a proportion of mild cases without overt anomalies within the radial ray or those with only conduction heart defects are missed. Another study limitation is that we have data on genetic testing only for a minority of patients, which does not allow genotype phenotype analysis or confirmation of cases with associated anomalies.

Conclusions

HOS is a rare genetic condition with regional variability. It is often missed prenatally in spite of the presence of major anomalies. When discovered, half of the parents will opt to continue the pregnancy even in the presence of severe limb defects and CHD. Our data indicate that the number of severe CHD in HOS is underestimated since they are present in one quarter of HOS patients. In spite of this, the overall first week survival is good, which is important information for genetic counselling of affected families.

Abbreviations

HOS: Holt Oram syndrome; EUROCAT: European surveillance of congenital anomalies; LB: Live born; FD: Fetal deaths; SB: Stillborn; BW: Birth weight; GA: Gestational age; TOPFA: Termination of pregnancy for fetal anomaly; CHD: Congenital heart defects; ASD: Atrial septal defect; VSD: Ventricular septal defect; AVSD: Atrioventricular septal defect; ICD: International classification of diseases; BPA: British paediatric association; OMIM: Online mendelian inheritance in man; ART: Assisted reproductive techniques; IVF: *in vitro* fertilization; ICSI: Intracytoplasmic sperm injection.

Competing interests

The authors declare that they have no competing interest.

Authors' contributions

IB conceived, and wrote the manuscript. LB performed the statistical analysis. RG extracted the dataset from the database and checked for accuracy. EG, DW, EC, MCA, LA, JEHB, PB, JLSB, MG, MH, BK, KK, BMCD, VN, AP, AQL, JR, AR, CR, DT, CVD participated in the study design, contributed to the interpretation of the results and revised the manuscript. HD participated in the design and coordination of the study and critically revised the manuscript. All authors read and approved the final manuscript.

Acknowledgements

This study is part of the EUROCAT Joint Action funded by the EC, under the framework of EU Health Programme 2008–2013, Grant Agreement 20102204 (Executive Agency for Health & Consumers). All registers in England are funded by Public Health England. We thank doctors Sebastiano Bianca, Jean Chapple, Berenice Doray, Christine Francannet, Anna Latos-Bielenska, Carmen Mosquera-Tenreiro, Mary O'Mahony, Joaquin Salvador, Gioacchino Scarano, David Stone, Romano Tenconi for providing data for the present study. We also thank the many people throughout Europe involved in providing and processing information, including affected families, clinicians, health professionals, medical record clerks and registry staff.

Author details

¹Zagreb Children's Hospital, Medical School, University of Zagreb, Klaićeva 16, Zagreb 10 000, Croatia. ²EUROCAT Central Registry, University of Ulster, Jordanstown Campus, Room 12 L09 Shore Road, Newtownabbey, Co. Antrim, Northern Ireland BT37 0QB, UK. ³Pediatric Department, Hospital Lillebaelt, Skovvangen 2-6, Kolding DK 6000, Denmark. ⁴Wessex Clinical Genetics Service, Princess Anne Hospital, Coxford Road, Southampton SO16

5YA, UK. ⁵Registro IMER, Azienda Ospedaliero-Universitaria di Ferrara, Corso Giovecca, Ferrara 202 44121, Italy. ⁶Division of Medical Genetics, CHUV, Lausanne, Switzerland. ⁷Registro Anomalías Congénitas CAPV. Dirección de Salud Pública. Departamento de Sanidad. Instituto BIO-Donostia, Basque Government CIBER Epidemiología y Salud Pública - CIBERESP, Donostia-San Sebastian 1, Vitoria Gasteiz 01010, Spain. ⁸Department of Genetics, University of Groningen, University Medical Center Groningen, Hanzeplein 1, Groningen 9700 RB, The Netherlands. ⁹Departamento de Epidemiologia, Instituto Nacional de Saude Doutor Ricardo Jorge, Av. Padre Cruz, Lisbon 1649-016, Portugal. ¹⁰Department of Health Sciences, University of Leicester, 22 28 Princess Road West, Leicester LE1 6TP, UK. ¹¹Malta Congenital Anomalies Registry, Directorate of Health Information and Research, G'mangia Hill, G'mangia PTA 1313, Malta. ¹²Styrian Malformation Registry, Medical University of Graz, Auenbruggerplatz 14, Graz AT 8036, Austria. ¹³Paris Registry of Congenital Malformations, INSERM U953, Maternité de Port-Royal, 53 av de l'Observatoire, Paris 75014, France. ¹⁴Medical Birth Registry of Norway, University of Bergen, Kalfareien 31, Bergen N-5018, Norway. ¹⁵Health Information Unit, Health Service Executive, Dr Stevens Hospital, Dublin 8, Dublin, Ireland. ¹⁶Provinciaal Instituut voor Hygiene, Kronenbrgstraat 45, Antwerp B-2000, Belgium. ¹⁷CNR Institute of Clinical Physiology, Via Moruzzi 1, Pisa I 56124, Italy. ¹⁸Universitätskinderklinik Mainz, Langenbeckstrasse 1, Postfach 3960, Mainz D 55101, Germany. ¹⁹Institute of Health & Society, Newcastle University, Newcastle upon Tyne NE2 4AA, UK. ²⁰Malformation Monitoring Centre Saxony-Anhalt, Medical Faculty Otto-von-Guericke University, Leipzigerstrasse, Haus 39, Magdeburg D 39120, Germany. ²¹National Perinatal Epidemiology Unit, University of Oxford, Old Road, Headington, Oxford OX3 7LF, UK. ²²Congenital Anomaly Register & Info Service Public Health Level 3 West Wing, Singleton Hospital, Sketty Lane, Swansea, Wales SA2 8QA, UK. ²³Institut de Pathologie et de Génétique, Avenue Georges Lemaître 25, Charleroi (Gosselies) 6041, Belgium.

Received: 11 June 2014 Accepted: 30 September 2014

Published online: 25 October 2014

References

- Holt M, Oram S: **Familial heart disease with skeletal malformations.** *Br Heart J* 1960, **22**:236–242.
- McDermott DA, Fong JC, Basson CT: **Holt-Oram syndrome.** In *Gene Reviews at Gene Tests: Medical Genetics Information Resource (database online)*. Copyright, Seattle: University of Washington; 1997–2008. (Updated 4/4/13). <http://www.genetests.org>.
- Basson CT, Bachinsky DR, Lin RC, Levi T, Elkins JA, Soultis J, Grayzel D, Kroumpouzou E, Traill TA, Leblanc-Straceski J, Renault B, Kucherlapati R, Seidman JG, Seidman CE: **Mutations in human TBX5 [corrected] cause limb and cardiac malformation in Holt-Oram syndrome.** *Nat Genet* 1997, **15**:30–35.
- Basson CT, Huang T, Lin RC, Bachinsky DR, Weremowicz S, Vaglio A, Bruzzone R, Quadrelli R, Lerone M, Romeo G, Silengo M, Pereira A, Krieger J, Mesquita SF, Kamisago M, Morton CC, Pierpont ME, Müller CW, Seidman JG, Seidman CE: **Different TBX5 interactions in heart and limb defined by Holt-Oram syndrome mutations.** *Proc Natl Acad Sci U S A* 1999, **96**:2919–2924.
- McDermott DA, Bressan MC, He J, Lee JS, Aftimos S, Brueckner M, Gilbert F, Graham GE, Hannibal MC, Innis JW, Pierpont ME, Raas-Rothschild A, Shanske AL, Smith WE, Spencer RH, St John-Sutton MG, van Maldergem L, Waggoner DJ, Weber M, Basson CT: **TBX5 genetic testing validates strict clinical criteria for Holt-Oram syndrome.** *Pediatr Res* 2005, **58**:981–986.
- Böhm J, Heinritz W, Craig A, Vujic M, Ekman-Joelsson BM, Kohlhasse J, Froster U: **Functional analysis of the novel TBX5 c.1333delC mutation resulting in an extended TBX5 protein.** *BMC Med Genet* 2008, **9**:88.
- Muru K, Kalev I, Teek R, Sönajalg M, Kuuse K, Reimand T, Ounap K: **A Boy with Holt-Oram syndrome caused by novel mutation c.1304delT in the TBX5 gene.** *Mol Syndromol* 2011, **1**:307–310.
- Fan C, Duhagon MA, Oberti C, Chen S, Hiroi Y, Komuro I, Duhagon PI, Canessa R, Wang Q: **Novel TBX5 mutations and molecular mechanism for Holt-Oram syndrome.** *J Med Genet* 2003, **40**:e29.
- Fan C, Liu M, Wang Q: **Functional analysis of TBX5 missense mutations associated with Holt-Oram syndrome.** *J Biol Chem* 2003, **278**:8780–8785.
- Postma AV, van de Meerakker JB, Mathijssen IB, Barnett P, Christoffels VM, Ilgun A, Lam J, Wilde AA, Lekanne Deprez RH, Moorman AF: **A gain-of-function TBX5 mutation is associated with**

- atypical Holt-Oram syndrome and paroxysmal atrial fibrillation. *Circ Res* 2008, **102**:1433–1442.
11. Porto MP, Vergani N, Carvalho AC, Cernach MC, Brunoni D, Perez AB: **Novel mutations in the TBX5 gene in patients with Holt-Oram syndrome.** *Genet Mol Biol* 2010, **33**:232–236.
 12. Boogerd CJ, Dooijes D, Ilgun A, Mathijssen IB, Hordijk R, van de Laar IM, Rump P, Veenstra-Knol HE, Moorman AF, Barnett P, Postma AV: **Functional analysis of novel TBX5 T-box mutations associated with Holt-Oram syndrome.** *Cardiovasc Res* 2010, **88**:130–139.
 13. Patel C, Silcock L, McMullan D, Brueton L, Cox H: **TBX5 intragenic duplication: a family with an atypical Holt-Oram syndrome.** *Eur J Hum Genet* 2012, **20**:863–869.
 14. Smemo S, Campos LC, Moskowitz IP, Krieger JE, Pereira AC, Norbrega MA: **Regulatory variation in a TBX5 enhancer leads to isolated congenital heart disease.** *Hum Mol Genet* 2012, **21**:3255–3263.
 15. Basson CT, Cowley GS, Solomon SD, Weissman B, Poznanski AK, Traill TA, Seidman JG, Seidman CE: **The clinical and genetic spectrum of the Holt-Oram syndrome (heart-hand syndrome).** *N Engl J Med* 1994, **330**:885–891.
 16. Newbury-Ecob R, Leanage R, Raeburn JA, Young ID: **The Holt-Oram syndrome: a clinical genetic study.** *J Med Genet* 1996, **33**:300–307.
 17. Sletten LJ, Pierpont MEM: **Variation in severity of cardiac disease in Holt-Oram syndrome.** *Am J Med Genet* 1996, **65**:128–132.
 18. Smith AT, Sack GH, Taylor GJ: **Holt-Oram syndrome.** *J Pediatr* 1979, **95**:538–543.
 19. Hurst JA, Hall CM, Baraitser M: **The Holt-Oram syndrome.** *J Med Genet* 1991, **28**:406–410.
 20. Elek C, Vitéz M, Czeizel E: **Holt-Oram syndrome.** *Orv Hetil* 1991, **132**:73–74, 77–78.
 21. **EUROCAT 'Guide 1.4'** [http://www.eurocat-network.eu/aboutus/datacollection/guidelinesforregistration/guide1_4]
 22. Greenlees R, Neville A, Addor MC, Amar E, Arriola L, Bakker M, Barisic I, Boyd P, Calzolari E, Doray B, Draper E, Vollset SE, Garne E, Gatt M, Haeusler M, Kallen K, Khoshnood B, Latos-Bielenska A, Martinez-Frias ML, Materna-Kiryluk A, Dias CM, McDonnell B, Mullaney C, Nelen V, O'Mahony M, Pierini A, Queisser-Luft A, Randrianaivo-Ranjatoelina H, Rankin J, Rissman A, et al: **Paper 6: EUROCAT Member Registries: Organization and Activities.** *Birth Defects Res A Clin Mol Teratol* 2011, **91**:S51–S100.
 23. Barisic I, Tokic V, Loane M, Bianchi F, Calzolari E, Garne E, Wellesley D, Dolk H, and EUROCAT Working Group: **Descriptive epidemiology of Cornelia de Lange syndrome in Europe.** *Am J Med Genet* 2008, **146A**:51–59.
 24. Barisic I, Odak L, Loane M, Garne E, Wellesley D, Calzolari E, Dolk H, Addor MC, Arriola L, Bergman J, Bianca S, Boyd PA, Draper ES, Gatt M, Haeusler M, Khoshnood B, Latos-Bielenska A, McDonnell B, Pierini A, Rankin J, Rissmann A, Queisser-Luft A, Verellen-Dumoulin C, Stone D, Tenconi R: **Fraser syndrome: epidemiological study in a European population.** *Am J Med Genet* 2013, **161A**:1012–1018.
 25. Barisic I, Odak L, Loane M, Garne E, Wellesley D, Calzolari E, Dolk H, Addor MC, Arriola L, Bergman J, Bianca S, Doray B, Khoshnood B, Klungsoyr K, McDonnell B, Pierini A, Rankin J, Rissmann A, Rounding C, Queisser-Luft A, Scarano G, Tucker D: **Prevalence, prenatal diagnosis and clinical features of oculo-auriculo-vertebral spectrum: a registry-based study in Europe.** *Eur J Hum Genet* 2014, doi:10.1038/ejhg.2013.287.
 26. Stoll C, Dott B, Alembik Y, Roth MP: **Associated malformations among infants with radial ray deficiency.** *Genet Couns* 2013, **24**:223–234.
 27. Brons JT, van Geijn HP, Wladimiroff JW, van der Harten JJ, Kwee ML, Sobotka-Plojhar M, Arts NF: **Prenatal ultrasound diagnosis of the Holt-Oram syndrome.** *Prenat Diagn* 1988, **8**:175–181.
 28. Law KM, Tse KT: **Prenatal sonographic diagnosis of familial Holt-Oram syndrome associated with type B interrupted aortic arch.** *Hong Kong Med J* 2008, **14**:317–320.
 29. Holmes LB: **Congenital heart disease and upper-extremity deformities. A report of two families.** *N Engl J Med* 1965, **272**:437–44.
 30. Gall JC, Stem AM, Cohen MM, Adams MS, Davidson RT: **Holt-Oram syndrome: clinical and genetic study of a large family.** *Am Hum Genet* 1966, **18**:187–200.
 31. Poznanski AK, Gall JC Jr, Stern AM: **Skeletal manifestations of the Holt-Oram syndrome.** *Radiology* 1970, **94**:45–54.
 32. Li QY, Newbury-Ecob RA, Terrett JA, Wilson DI, Curtis AR, Yi CH, Gebuhr T, Bullen PJ, Robson SC, Strachan T, Bonnet D, Lyonnet S, Young ID, Raeburn JA, Buckler AJ, Law DJ, Brook JD: **Holt-Oram syndrome is caused by mutations in TBX5, a member of the Brachyury (T) gene family.** *Nat Genet* 1997, **15**:21–29.
 33. Lehner R, Goharkhay N, Tringler B, Fasching C, Hengstschläger M: **Pedigree analysis and descriptive investigation of three classic phenotypes associated with Holt-Oram syndrome.** *J Reprod Med* 2003, **48**:153–159.
 34. Brassington AM, Sung SS, Toydemir RM, Le T, Roeder AD, Rutherford AE, Whitby FG, Jorde LB, Bamshad MJ: **Expressivity of Holt-Oram syndrome is not predicted by TBX5 genotype.** *Am J Hum Genet* 2003, **73**:74–85.
 35. Tseng YR, Su YN, Lu FL, Jeng SF, Hsieh WS, Chen CY, Chou HC, Peng SS: **Holt-Oram syndrome with right lung agenesis caused by a de novo mutation in the TBX5 gene.** *Am J Med Genet A* 2007, **143A**:1012–1014.
 36. Garavelli L, De Brasi D, Verri R, Guareschi E, Cariola F, Melis D, Calcagno G, Salvatore F, Unger S, Sebastio G, Albertini G, Rivieri F, Soli F, Superti-Furga A, Gentile M: **Holt-Oram syndrome associated with anomalies of the feet.** *Am J Med Genet A* 2008, **146A**:1185–1189.
 37. Faria MH, Rabenhorst SH, Pereira AC, Krieger JE: **A novel TBX5 mutation (V263M) in a family with atrial septal defects and postaxial hexadactyly.** *Int J Cardiol* 2008, **130**:30–35.
 38. Davies MJ, Moore VM, Willson KJ, Van Essen P, Priest K, Scott H, Haan EA, Chan A: **Reproductive technologies and the risk of birth defects.** *N Engl J Med* 2012, **366**:1803–1813.
 39. Tararbit K, Lelong N, Thieulin AC, Houyel L, Bonnet D, Goffinet F, Khoshnood B, EPICARD Study Group: **The risk for four specific congenital heart defects associated with assisted reproductive techniques: a population-based evaluation.** *Hum Reprod* 2013, **28**:367–374.
 40. Vermeiden JP, Bernardus RE: **Are imprinting disorders more prevalent after human in vitro fertilization or intracytoplasmic sperm injection?** *Fertil Steril* 2013, **99**:642–651.
 41. Kochanski A, Merritt TA, Gadzinowski J, Jopek A: **The impact of assisted reproductive technologies on the genome and epigenome of the newborn.** *J Neonat Perinat Med* 2013, **6**:101–108.

doi:10.1186/s13023-014-0156-y

Cite this article as: Barisic et al.: Holt Oram syndrome: a registry-based study in Europe. *Orphanet Journal of Rare Diseases* 2014 **9**:156.

Submit your next manuscript to BioMed Central and take full advantage of:

- Convenient online submission
- Thorough peer review
- No space constraints or color figure charges
- Immediate publication on acceptance
- Inclusion in PubMed, CAS, Scopus and Google Scholar
- Research which is freely available for redistribution

Submit your manuscript at
www.biomedcentral.com/submit

