# Ten-year survival of children with trisomy 13 or trisomy 18: a multi-registry European cohort study

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# ABSTRACT

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**Objective** To investigate the survival to 10 years of age of children with trisomy 13 (T13) and children with trisomy 18 (T18), born 1995-2014.

Design Population-based cohort study that linked mortality data to data on children born with T13 or T18, including translocations and mosaicisms, from 13 member registries of EUROCAT, a European network for the surveillance of congenital anomalies.

Setting 13 regions in nine Western European countries. Patients 252 live births with T13 and 602 with T18. Main outcome measures Survival at 1 week, 4 weeks and 1, 5 and 10 years of age estimated by randomeffects meta-analyses of registry-specific Kaplan-Meier survival estimates.

**Results** Survival estimates of children with T13 were 34% (95% CI 26% to 46%), 17% (95% CI 11% to 29%) and 11% (95% CI 6% to 18%) at 4 weeks, 1 and 10 years, respectively. The corresponding survival estimates were 38% (95% CI 31% to 45%), 13% (95% CI 10% to 17%) and 8% (95% CI 5% to 13%) for children with T18. The 10-year survival conditional on surviving to 4 weeks was 32% (95% CI 23% to 41%) and 21% (95% CI 15% to 28%) for children with T13 and T18, respectively.

**Conclusions** This multi-registry European study found that despite extremely high neonatal mortality in children with T13 and T18, 32% and 21%, respectively, of those who survived to 4 weeks were likely to survive to age 10 years. These reliable survival estimates are useful to inform counselling of parents after prenatal diagnosis.

defects, chromosomal and genetic syndromes,

affect about 2%–3% of births in Europe<sup>1</sup> and in the

USA,<sup>2</sup> and are a leading cause of infant mortality.<sup>3 4</sup> They are also a growing contributor to mortality

of children under 5 years of age<sup>5</sup> and of older

children.<sup>6</sup> Survival of children with major CAs

beyond 1 year has substantially improved during

the last few decades due to advances in neonatal

care and surgical interventions.7 8 As shown in

our recent multi-centre European study, 10-year

survival exceeded 90% for most major structural

anomalies and the most common chromosomal

anomaly, Down syndrome (trisomy 21).<sup>9</sup> Trisomy

13 (T13) (Patau syndrome) and trisomy 18 (T18)

(Edwards syndrome) are the most common

#### **INTRODUCTION** Congenital anomalies (CAs), including structural

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# WHAT IS ALREADY KNOWN ON THIS TOPIC

- $\Rightarrow$  Children with trisomv 13 or trisomv 18 have extremely high neonatal and infant mortality.
- $\Rightarrow$  A recent Canadian population-based study reported that about 13% of children with trisomy 13 and 10% with trisomy 18 may survive to age 10 years.
- $\Rightarrow$  Long-term follow-up population-based studies of survival in children with trisomy 13 or trisomy 18 are lacking.

# WHAT THIS STUDY ADDS

- $\Rightarrow$  The majority of children born alive with trisomy 13 or trisomy 18 between 1995 and 2014 in 13 Western European regions died during the first 28 days of life: 66% of children with trisomy 13 and 62% with trisomv 18.
- $\Rightarrow$  Survival at age 5 and 10 years was 16% (95%) CI 10% to 26%) and 11% (95% CI 6% to 18%), respectively, for children with trisomy 13, and 10% (95% CI 7% to 14%) and 8% (95% CI 5% to 13%), respectively, for children with trisomy 18.
- $\Rightarrow$  Ten-year survival conditional on surviving the first 28 days of life was 32% (95% CI 23% to 41%) and 21% (95% CI 15% to 28%) for trisomy 13 and trisomy 18, respectively.

# HOW THIS STUDY MIGHT AFFECT RESEARCH, **PRACTICE OR POLICY**

 $\Rightarrow$  This study demonstrates that reliable survival estimates can be obtained for children with rare anomalies by linking administrative mortality data to data on live births from European population-based congenital anomaly registries and combining results across registries. The results are important for counselling parents after prenatal diagnosis of these conditions.

autosomal trisomies after Down syndrome and are characterised by multiple structural anomalies and intellectual disability in survivors. The combined total prevalence including pregnancies resulting in a termination of pregnancy for fetal anomaly (TOPFA), stillbirths and live births varies from 5 to 10 per 10 000 births.<sup>10-12</sup> Children with T13 or T18 have a high mortality risk during the first weeks of life and the majority die during the first year.<sup>11 13–17</sup> Recent population-based US and Canadian studies



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reported median survival time of  $5^{14}$ –12. $5^{15}$  days for T13 and  $8^{14}$ –9<sup>15</sup> days for T18, while 5-year survival was 9.7% (95% CI 7.2% to 12.5%) for T13 and 12.3% (95% CI 10.1% to 14.8%) for T18 in the USA<sup>14</sup> and 15% (95% CI 10% to 21%) for T13 and 11% (95% CI 8% to 16%) for T18 in Canada.<sup>15</sup> In Canada, conditional 10 year survival for children who survived to 1 year, was 65% (95% CI 46% to 79%) for T13 and 77% (95% CI 56% to 89%) for T18.<sup>15</sup> Recent population-based information on longer-term survival of European children with T13 and T18 is lacking.<sup>7</sup>

The aim of this multi-registry European study was to investigate the survival up to 10 years of age of children born alive with T13 or T18 by linking data from 13 EUROCAT (European network for the surveillance of CAs) population-based registries in nine Western European countries to their local mortality data sources. This study was part of the wider EUROlinkCAT data linkage project that investigated the survival, health and educational outcomes to 10 years of age of European children born with a major CA.<sup>18</sup>

#### **METHODS**

## Design, population and data linkage

We conducted a European, population-based linked cohort study. The full cohort included all live births with a major CA collected and validated by population-based CA registries that are members of EUROCAT (https://eu-rd-platform.jrc.ec. europa.eu/eurocat\_en).

Each registry has ethics permissions and procedures for routine surveillance, data collection and transmission of anonymised individual-level data to a central database according to national guidelines. For the EUROlinkCAT study, local ethics approvals or other permissions to link registry data with local mortality data sources were obtained by 12 registries; one registry (Norway) obtained permission to use data that were already linked.

Data on all children with a major CA born alive between 1 January 1995 and 31 December 2014 recorded in the 13 registries in nine Western European countries were linked to administrative mortality data sources up to the child's 10th birthday or to 31st December 2015, whichever was earlier, so that all children have at least 1 year of follow-up information. Registries linked their CA data to either national/vital statistics (11 registries) or to mortality records only (two registries) (table 1). Linkage to national/vital statistics that included both birth and death registration data provided information on the vital status for all linked children (dead or alive) including those who moved to other country areas; in contrast, linkage to mortality records can identify deaths only, and hence, children with no death record were assumed to be alive. A detailed description of the linkage process and accuracy of the linked data for each registry together with an analysis of the survival data validity is provided elsewhere.<sup>19</sup> The included birth year periods differed between registries due to different years of EUROCAT membership or due to inclusion of the years with high quality linked data only (table 1 and online supplemental table 1). There was no standard approach to neonatal treatment of children with T13/T18 across participating regions.

The inclusion criteria were all liveborn children with a diagnostic code (*International Statistical Classification of Diseases* and Related Health Problems, Ninth Revision or Tenth Revision (ICD-9 or ICD-10)) 758.1 (ICD-9) or Q914-Q917 (ICD-10) (karyotype 47,XX+13 or 47,XY+13 and translocations/

 Table 1
 Participating EUROCAT‡ registries, birth years, population covered, total and live birth (LB) prevalence of cases with trisomy 13 (T13) and trisomy 18 (T18) (per 10 000 births) by registry

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			Trisomy 13		Trisomy 18	
Participating registries	Included birth years	Birth population covered*	Total prevalence per 10 000 (95% CI)*	LB prevalence per 10 000 (95% CI)*	Total prevalence per 10 000 (95% CI)*	LB prevalence per 10 000 (95% CI)*
Registries that linked to nat	tional/vital statistics†					
Denmark: Funen	1995–2014	105 570	1.9 (1.2 to 2.9)	0.1 (0.0 to 0.5)	5.2 (3.9 to 6.8)	1.0 (0.5 to 1.9)
Finland	1995–2014	1 174 727	2.4 (2.1 to 2.7)	0.7 (0.5 to 0.8)	6.8 (6.4 to 7.3)	1.4 (1.2 to 1.6)
France: Paris	1995–2014	597 822	3.9 (3.4 to 4.4)	0.3 (0.2 to 0.4)	11.4 (10.6 to 12.3)	0.7 (0.5 to 0.9)
Italy: Emilia Romagna	2008–2014	282 094	1.0 (0.7 to 1.4)	0	3.9 (3.2 to 4.7)	0.4 (0.2 to 0.7)
Italy: Tuscany	2005–2014	299 869	1.7 (1.3 to 2.2)	0.2 (0.1 to 0.4)	5.1 (4.4 to 6.0)	0.4 (0.2 to 0.7)
Netherlands: Northern	1995–2014	372 192	1.5 (1.1 to 2.0)	0.5 (0.3 to 0.8)	5.5 (4.8 to 6.3)	1.2 (0.9 to 1.6)
Norway	1999–2014	956 939	1.9 (1.6 to 2.2)	0.5 (0.4 to 0.7)	4.4 (4.0 to 4.9)	1.2 (1.0 to 1.4)
UK: East Midlands and South Yorkshire	2003–2012	717 264	2.3 (2.0 to 2.7)	0.4 (0.3 to 0.6)	5.4 (4.9 to 6.0)	0.8 (0.6 to 1.0)
UK: Thames Valley	2005–2013	270 327	3.4 (2.8 to 4.2)	0.5 (0.3 to 0.8)	8.0 (6.9 to 9.1)	0.9 (0.6 to 1.3)
UK: Wales	1998–2014	569 341	2.1 (1.8 to 2.6)	0.4 (0.2 to 0.6)	5.4 (4.9 to 6.1)	1.0 (0.8 to 1.3)
UK: Wessex	2004–2014	325 339	2.9 (2.4 to 3.6)	0.3 (0.2 to 0.6)	7.6 (6.7 to 8.6)	0.9 (0.6 to 1.3)
Registries that linked to mo	ortality records†					
Malta	1995–2014	84 737	0.7 (0.3 to 1.5)	0.7 (0.3 to 1.5)	3.7 (2.5 to 5.2)	3.1 (2.0 to 4.5)
Spain: Valencian Region	2007–2014	403 099	1.5 (1.2 to 2.0)	0.2 (0.1 to 0.4)	4.1 (3.5 to 4.8)	0.5 (0.3 to 0.7)
Total		6 159 520				

The registers in Finland, Norway, Wales and Malta are national, while other registries are regional.

In Malta, termination of pregnancy is illegal that explains similar total and live birth prevalence of T13 and T18 in Malta.

\*Extracted from the EUROCAT website: https://eu-rd-platform.jrc.ec.europa.eu/eurocat/eurocat-data/prevalence\_en (accessed on 6 Jan 2022). Total prevalence includes

terminations of pregnancy for fetal anomaly, fetal deaths/stillbirths from 20 week' gestation and live births per 10 000 registered live and stillbirths.

†National/vital statistics include birth and death registration data and all live births will have a record; mortality records only include death registration and live births who remain alive will not have a record.

‡EUROCAT (European network of population-based registries for the epidemiological surveillance of congenital anomalies).

mosaicism) for T13 and 758.2 (ICD-9) or Q910-Q913 (ICD-10) (karyotype 47,XX+18 or 47,XY+18 and translocations/mosaicism) for T18, meaning that children with less severe forms of T13 and T18 were also included. At a later stage, the registries reported the karyotype for infant deaths and for children who survived beyond 1 year where possible to confirm long-term survival results.

## **Statistical analysis**

The study included the development of a common data model to standardise the local variables available in the national/vital statistics or mortality databases (https://www.eurolinkcat.eu/wp2-buildingresultsrepository/eurolinkcatpubliccommondatam odels).<sup>18</sup> This formed the basis for the development of centrally written syntax scripts used for checking the data linkage quality and for the local analyses to be run by the participating registries.<sup>18</sup> <sup>19</sup>

Each registry calculated the survival probability of children with T13 and T18 at prespecified ages by running Kaplan-Meier survival analysis on the individual case data to account for censoring, as not all children reached their 10th birthday during the study period. The registry-specific Kaplan-Meier survival estimates with 95% CIs (all 13 registries), the number at risk (alive at the beginning of each age point) and the number of deaths at each age (all registries, except Netherlands: Northern) were then uploaded to the Central Results Repository at Ulster University (UK) using a secure web platform. The Netherlands: Northern registry rounded the number of deaths to the nearest 0 or 5 after age 4 weeks due to the national small number restrictions; therefore, their data could not be included in the meta-analysis.

No individual case data were shared.

The registry-based Kaplan-Meier survival estimates were combined centrally in random-effects meta-analyses of the survival at five ages (1 week, 4 weeks and 1, 5 and 10 years) to estimate the overall survival of children with T13 and T18. The meta-analytic approach applied to these data involved modifying a method proposed by Combescure *et al*<sup>20</sup> and is described in detail elsewhere<sup>9</sup> and in online supplemental box 1.

Kaplan-Meier survival analyses were performed using Stata V.16 (StataCorp LLC, 2019). Meta-analyses were performed using R software.

#### RESULTS

Table 1 shows the data from 13 EUROCAT population-based contributing registries covering a population of 6 159 520 births in 1995–2014. The live birth prevalence of T13 and T18 was much lower than the total prevalence, as total prevalence also includes TOPFAs and stillbirths. Overall, the live birth/total prevalence ratio decreased by about 40% between 1995–2004 and 2005–2014 (from 0.26 to 0.15 for T13 and from 0.22 to 0.13 for T18), which may have resulted from improvement in prenatal diagnosis and higher TOPFA rates.

Figure 1 shows the Kaplan-Meier survival estimates with 95% CIs at age 1 week, 4 weeks and 1 year for infants with T13 and T18 by contributing registries and the pooled survival provided by the meta-analysis. The heterogeneity between registries was high at 1 week (T13:  $I^2=54\%$ ; T18:  $I^2=63\%$ ) and lowest at 1 year (T13:  $I^2=24\%$ ; T18:  $I^2=23\%$ ). The variation of the survival estimates and the width of the 95% CIs were relatively high as a result of the different sizes of the population covered by each registry and the rarity of T13.

Table 2 reports pooled survival estimates with 95% CI at age 1 week, 4 weeks, 1, 5 and 10 years for the 252 children born with T13 (total deaths=226) and the 602 with T18 (total deaths=535). Forty-five per cent of children with T13 and 41% with T18 died within the first week of life, 66% of children with T13 and 62% with T18 died within the first 4 weeks. Although the majority of these children died in infancy, 10.8% (95% CI 5.7% to 17.8%) of children with T13 and 8.0% (95% CI 5.0% to 12.8%) of children with T18 survived to age 10 years.

Pooled survival estimates produced by the sensitivity analysis that included 11 registries with more reliable linkage results (linked to vital/national statistics) were very similar to the survival estimates based on 13 registries (less than one percentage point difference).

Nine of the 11 registries with survivors beyond 1 year of age provided additional karyotype information for some children, which suggests that the percentage of children with less severe trisomy forms was relatively higher among survivors than among infant deaths. We do not report the exact figures as karyotype information was missing in up to 22% of survivors, with substantial variation across registries.

The overall survival at 10 years conditional on surviving to 4 weeks (a third of children with either trisomy survived 28 days) was 32% (95% CI 23% to 41%) for children with T13 and 21% (95% CI 15% to 28%) for children with T18 (table 2).

## DISCUSSION

This multi-registry population-based European linked cohort study of liveborn infants delivered in 1995–2014 with T13 and T18 reported that over 60% of these infants died during the first 28 days of life and over 80% did not survive to their first birthday. Despite such high infant mortality, 16% and 10% of children with T13 and T18, respectively, survived to 5 years and 11% (T13) and 8% (T18) survived to 10 years. The 10-year survival conditional on surviving to 28 days was 32% for children with T13 and 21% for children with T18. The survival estimates were relatively consistent between the contributing registries at 1 year, but there was a substantially higher heterogeneity at 1 week.

Due to very high infant mortality of live births with T13 and T18, earlier studies reported survival during infancy only. However, more recent population-based studies have demonstrated that approximately 6% to 20% of these children survived the first year<sup>11</sup> <sup>13–17</sup> <sup>21</sup> <sup>22</sup> and around 10% survived up to 10 years<sup>15</sup> <sup>22</sup> (table 3). Our study's survival estimates at 1 month, 1 year and 5 years for European children with trisomy 18 are mostly in agreement with large recent international studies that included any trisomy variants<sup>11 14 15</sup> (table 3). Ten-year survival is also comparable with that in a Canadian study covering a similar birth year period.<sup>15</sup> For children with T13, there is slightly more inconsistency in survival estimates between the published studies, in particular for longer term survival. For example, 5-year survival of children with T13 is similar in our European study and the mentioned Canadian study,<sup>15</sup> while it is higher than in other large recent studies<sup>11 14</sup> (table 3). As expected, the 1-year, 5-year and 10-year survival in our study that included children with any cytogenetic variants was higher than in studies reported for children with full trisomies<sup>13 17 21</sup> (table 3), as partial and mosaic variants are associated with a higher survival. In addition, improved survival in more recent years may be associated with a wider use of neonatal intensive care in infants with T13 and T18 than previously and surgical interventions<sup>15 23-25</sup> in some infants who survived the first week/month. For example,



**Figure 1** Registry-specific Kaplan-Meier survival estimates with 95% CIs and the combined survival: (A) at 1 week, (B) 4 weeks and (C) 1 year for children with trisomy 13 and trisomy 18. The numbers given in square brackets for each registry at 1 week (A) are the numbers alive at birth (number at risk), and the numbers at risk for age 4 weeks and 1 year are suspended due to small number of cases. The registries are ordered in descending order of survival estimates. The number of presented registries differs depending on the data available at certain age; for example, there were no live births with trisomy 13 in the Italy: Emilia Romagna registry; the 1 week and 4 week survival for children with trisomy 13 is not presented for the Denmark: Funen registry as there were  $\leq$ 5 live births; the survival estimates after 4 weeks of age are not presented for the Netherlands: Northern registry due to the national small number restrictions. DK, Funen, Denmark: Funen; Fr, Paris, France: Paris; IT, E Romagna, Italy: Emilia Romagna; IT, Tuscany, Italy: Tuscany; North Neth, Netherlands: Northern; SP, Valencian R, Spain: Valencian region; UK, CAROBB, UK: Thames Valley; UK, EMSYCAR, UK: East Midlands and South Yorkshire; UK, WANDA, UK: Wessex; Wales, UK: Wales.

a recent single-centre Japanese study reported improvement in 3-year survival of children with T18 from 13.8% in 2008–2012 to 44.4% in 2013–2017, likely resulting from increased surgical interventions in the later period in infants with T18 admitted to a paediatric tertiary centre within the first 7 days of life.<sup>25</sup> Our study confirmed that children who survived the first 28 days of life had a higher likelihood of survival to age 10 years: 32% for children with T13 and 21% for children with T18 compared with 30% (95% CI 20% to 41%) for T13 and 28% (95% CI 19% to 38%) for T18 in a Canadian study conditional on surviving to 30 days.<sup>15</sup>

Despite accumulating evidence of improvement in survival as a result of neonatal intensive treatment and surgical interventions

in children with T13 and T18,<sup>15 24-28</sup> there is still some controversy regarding treatment strategies including cardiac surgery for patients with T13 and T18 due to poor prognosis for more vulnerable patients, significant neurodevelopmental disability in survivors, sparse information on quality of life of the children and families, high individual and societal costs and a number of ethical issues involved.<sup>29–35</sup> Although the approaches to care of live births with T13 and T18 may differ between countries, with reports on neonatal intensive care and surgical interventions mostly from North America and Japan,<sup>15 23 25 26 28 30 33</sup> current medical expert's view is developing towards evidence-based individualised medical care of these children<sup>24 28 30</sup> with careful consideration of condition severity and comorbidities

Table 2Pooled survival estimates (with 95% CIs) for five age points up to 10 years of age and 10-year survival conditional on surviving to 4weeks for children born with trisomy 13 or trisomy 18 in 13 EUROCAT registries in nine Western European countries, 1995–2014

			Survival estimates %	95% CI)				
Trisomy type	No. of live births	No. of deaths up to 10 vears	1 week	4 weeks	1 vear	5 years	10 vears	10 years conditional on surviving to 4 weeks
-71		<b>, .</b>			· <b>,</b>	- ,		
Trisomy 13	252	226	55.1 (43.2 to 70.1)	34.3 (25.7 to 45.7)	17.4 (10.6 to 28.6)	16.1 (10.0 to 25.8)	10.8 (5.7 to 17.8)	32 (23 to 41)
l <sup>2</sup>			54%	38%	24%	45%	37%	
Trisomy 18	602	535	59.1 (51.4 to 67.9)	37.6 (31.4 to 45.1)	12.8 (9.5 to 17.3)	10.0 (6.9 to 14.4)	8.0 (5.0 to 12.8)	21 (15 to 28)
l <sup>2</sup>			63%	41%	23%	46%	0%	

There was no complete follow-up for all registries and all birth years to age 10 years; hence, 10-year survival cannot be calculated as deaths/births. Therefore, Kaplan-Meier survival analysis that accounts for censoring was used to estimate registry-specific survival.

The number of deaths from the Netherlands: Northern registry was rounded to the nearest 0 or 5 after age 4 weeks to follow the national restrictions in relation to small numbers and therefore could not be included in the meta-analysis.

 $l^2$  statistic was used as a measure of the observed between-registry heterogeneity (with  $l^2 > 50\%$  indicating significant heterogeneity<sup>38</sup>) calculated by a random effect metaanalysis.

and discussions with parents taking into account their wishes and values and respecting their informed decisions.  $^{29\,31\,32\,36\,37}$ 

The main study strength was the follow-up of children with T13 and T18 to 10 years of age to determine the pooled survival estimates of these children using linked data between high-quality population-based specialised CA registries from 13 regions across nine Western European countries and their mortality data sources, including high quality linked data from national/vital statistics for 11 of 13 registries. This resulted in the creation of a large European cohort of children with T13 and T18 with 10-year survival data, which increased the study's statistical power and the reliability of its survival estimates. A

further strength was a combination of standardised approaches to data collection, coding and classification in EUROCAT registries and standardising the linked mortality data to a EUROlinkCAT common data model, development of standardised syntax scripts and production of standardised analytic results.

This study was limited to survival data only for children with T13 or T18, and therefore, no information on morbidity, hospitalisation or surgical interventions was available to explore their association with survival. Although the EUROCAT registries collect information on cytogenetic variants of these chromosomal syndromes and associated structural anomalies in live births, for this study, we did not request that level of detail for

Rasmussen et al (2003) <sup>13</sup> Niedrist et al (2006) <sup>21</sup> Wang et al (2013) <sup>17</sup> Weyer et al (2016) <sup>14</sup> Nelson et al (2016) <sup>15</sup> Schneuer et al (2019) <sup>16</sup> Goel et al (2019) <sup>16</sup> Current study           Trisomy 13           Study period         1968–1999         —         1983–2006         2004–2011         1999–2007         1991–2012         2004–2009         1974–2014         1995–2014           Geographical region         Georgia, USA         —         New York State, USA         England and Wales         USA, multi-state         Ontario, Canada         NSW, Australia         Multi-registry         Western Europe           Sample size         70         —         525         120         693         174         25         2537         252           Trisomy variant included         Mosaicism excluded         —         Any         Full trisomy         Any         Any         Any         Any         Any         Any           1 month         30.0         38.1         29         25.5         42         40.0         NR         34.3           1 year         8.6         —         21.3         8.0         11.5         19.8         LN         13         17.4           5 years         1
Trisomy 13         Study period       1968–1999       –       1983–2006       2004–2011       1999–2007       1991–2012       2004–2009       1974–2014       1995–2014         Geographical region       Georgia, USA       –       New York State USA       England and Wales       USA, multi-state       Ontario, Canada       NSW, Australia       Multi-registry       Western Europe         Sample size       70       –       525       120       693       174       25       2537       252         Trisomy variant included       Mosaicism excluded       –       Any       Hul trisomy       Any
Study period         1968–1999          1983–2006         2004–2011         1999–2007         1991–2012         2004–2009         1974–2014         1995–2014           Geographical region         Georgia, USA          New York State, USA         England and Vales         USA, multi-state         Ontario, Canada         NSW, Australia         Multi-registry         Western Europe           Sample size         70          525         120         693         174         25         2537         252           Trisomy variant included         Mosaicism excluded          Sa1         29         693         174         25         2537         252           Age         Proportion survive (%)         Full trisomy         Any
Geographical regionGeographical regionGeographical sourceGeographical USAGeographical USAGeographical USAMew York State, USAEngland and WalesUSA, multi-state officialOntario, Canada NSW, AustraliaMulti-registryWestern EuropeSample size70525120693174252537252Trisomy variant includedMosaicism excludedAnyFull trisomy Full trisomyAnyAnyAnyAnyAnyAgeProportion surviving (%)38.12925.54240.0NR34.31 year8.621.38.011.519.8LN1317.45 years118.439.715LN716.110 yearsNRNR*NRNR12.9NRNR10.8Trisomy 18
Sample size         70          525         120         693         174         25         2537         252           Trisomy variant included         Mosaicism excluded          Any         Full trisomy         Any         A
$ \begin{array}{ c c c c c c c c c c c c c c c c c c c$
Age         Proportion surviving (%)           1 month         30.0         38.1         29         25.5         42         40.0         NR         34.3           1 year         8.6          21.3         8.0         11.5         19.8         LN         13         17.4           5 years         1          18.4         3         9.7         15         LN         7         16.1           10 years         NR         -         NR*         NR         NR         12.9         NR         NR         10.8
1 month         30.0         38.1         29         25.5         42         40.0         NR         34.3           1 year         8.6          21.3         8.0         11.5         19.8         LN         13         17.4           5 years         1          18.4         3         9.7         15         LN         7         16.1           10 years         NR          NR*         NR         NR         12.9         NR         NR         10.8
1 year         8.6          21.3         8.0         11.5         19.8         LN         13         17.4           5 years         1          18.4         3         9.7         15         LN         7         16.1           10 years         NR          NR*         NR         NR         12.9         NR         NR         10.8           Trisomy 18
5 years         1          18.4         3         9.7         15         LN         7         16.1           10 years         NR          NR*         NR         NR         12.9         NR         NR         10.8           Trisomy 18 <td< td=""></td<>
10 years         NR         NR         NR         12.9         NR         NR         10.8           Trisomy 18                    10.8                10.8 </td
Trisomy 18
Study period 1968–1999 1964–2003 1983–2006 2004–2011 1999–2007 1991–2012 2004–2009 1974–2014 1995–2014
Geographical         Georgia, USA         Switzerland         New York State, USA         England and Wales         USA, multi-state         Ontario, Canada         NSW, Australia         Multi-registry         Western Europe
Sample size         114         161         773         309         1113         254         34         6122         602
Trisomy variant         Mosaicism         Mosaicism         Any         Full trisomy         Any
Age Proportion surviving (%)
1 month 38.6 22.4 46.8 39 37.2 35 35.3 NR 37.6
1 year 8.4 6.2 18.8 8.0 13.4 12.6 20.6 12 12.8
5 years NR 2 15.2 NR 12.3 11 17.6 7.7 10.0
10 years NR 1.2 NR* NR NR 9.8 NR NR 8.0

The study by Niedrist et al, 2006 was restricted to trisomy 18 only, therefore, no data for trisomy 13 could be provided; 1 month can differ between 28 and 30 days in different studies, for example, 28 days in Meyer *et al.*, Wang *et al.* and in our study.

\*Ten-year survival not reported; 15-year survival was 16.2% and 13.2% for children with trisomy 13 and 18, respectively.

LN, low number (less than 5 cases at risk at that time interval); NR, not reported; NSW, New South Wales.;

## **Original research**

practical reasons (expecting very small numbers by cytogenetic variant per registry), which prevented reporting pooled survival by trisomy variant and comorbidities. However, an examination of trisomy variants among long-term survivors suggested a relatively higher percentage of children with mosaicism/translocation among survivors compared with infant deaths, as expected. The survival results for the Netherlands: Northern registry were included for the first 4 weeks of life only as after this age the number of survivors was too small and could not be included in the meta-analysis due to the national small number restrictions. Although the survival data were combined from 13 registries, the relatively low number of survivors beyond 1 year did not allow analysing the association with demographic/infant risk factors.

In conclusion, we confirmed that 1-year survival of children born with T13 or T18 remains low. However, we found that 16% and 10% of children born in 1995–2014 with T13 and T18, respectively, survived to 5 years and 11% and 8%, respectively, survived to 10 years. Reliable information on longer-term survival of live births with T13 and T18 in Western Europe is important for health professionals when counselling parents following prenatal diagnosis of these conditions and would help parents to make informed decisions in relation to termination of pregnancy. It is also valuable for parents of liveborn children with T13 and T18 to choose the treatment approach optimal for their child in consultation with health professionals.

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Data availability statement Data are available on reasonable request. The data that support the findings of this study are available from the contributing registries of congenital anomalies but restrictions apply to the availability of these data, which were used under license for the current study, and so are not publicly available. Limited data are however available from the authors for scientifically valid requests and with permission of the contributing registries. To apply for the data please complete the data request form available on https://www.eurolinkcat.eu/contacti nformationanddatarequests. The survival results for various congenital anomalies can be explored using EUROlinkCAT Results Explorer on the website: http://www.EUROlinkCAT.eu/eurolinkcatresultsexplorer.

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