

### 2.3. Template for Associate Member Registry Data Transmission

The forms below describes the data that should be sent to the Central Registry by associate registries (aggregate data) for a given birth year. The aggregate data should ideally be sent in a .csv file.

The following columns should be used in the .csv file for transmitting aggregate data. The file will be used by the Central Registry for importing the data in DMS.

- Centre
- Year
- anomalyId
- live
- still
- topfa
- live\_ec
- still\_ec
- topfa\_ec
- completed\_data

Centre Number:	
<b>Denominator Data Year XXXX:</b>	
Live births:	
Stillbirths:	
Total Births:	
<b>Births (live and still) by Maternal Age:</b>	
<20	
20-24	
25-29	
30-34	
35-39	
40-44	
45+	
Not known	









	Includes Chromosomal Cases				Excludes Chromosomal Cases			
	No of LB	No of FD	No of TOPFA	Total cases	No of LB	No of FD	No of TOPFA	Total cases
Pierre-Robin sequence								
Caudal regression sequence								
Sirenomelia								
Septo-optic dysplasia								
Vascular disruption anomalies								
Laterality anomalies								
Teratogenic syndromes resulting in major malformations								
Valproate syndrome								
Maternal infections resulting in major malformations								
<b>Genetic disorders</b> <i>(genetic syndromes, hereditary skin disorders, skeletal dysplasias and chromosomal anomalies)</i>								
Skeletal dysplasias								
Down syndrome /trisomy 21								
Patau syndrome/trisomy 13								
Edwards syndrome/trisomy 18								
Turner syndrome								
Triploidy and polyploidy								

LB = Live births

FD = Fetal deaths / Still births from 20 weeks gestation

TOPFA = Termination of pregnancy for fetal anomaly following prenatal diagnosis

\* All Anomalies = all cases with a major structural congenital anomaly, excluding cases with only minor anomalies as defined in Section 3.2 in Guide 1.5 for cases born post-2005. Cases with more than one anomaly are only counted once in the "All Anomalies" subgroup.

Last update 31.05.2022