EUROCAT Data Management Software Analyse data Statistical surveillance - Trends



Introduction

- DMS performs the trends analysis according to the methodology described in the <u>Statistical monitoring protocol</u>.
- The analysis can run on one or several centres. If two or more centres are selected, the program also outputs pan-centre/pan-European results, based on the aggregation of the data from the individual centres selected.

- <u>Section 1</u> (respectively <u>section 2</u>) shows how the trends analysis run for 1 registry selected (resp. 2 or more registries selected)
- The trends analysis requires to install the « *R distribution for Eurocat DMS.msi* » and to save the appropriate collection of R libraries. Please, follow instructions in <u>section 3</u>.

Table of contents

1. Run trends analysis for 1 registry

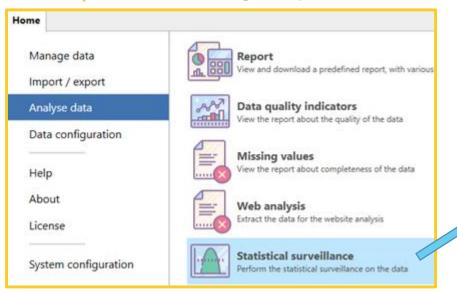
2. Run trends analysis for several registries

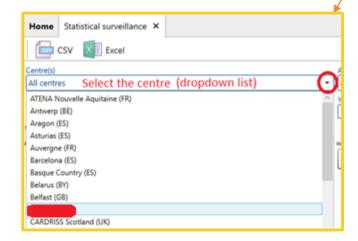
- 3. R distribution for DMS
 - a. Install the R distribution for Eurocat DMS.msi
 - b. New R libraries for DMS

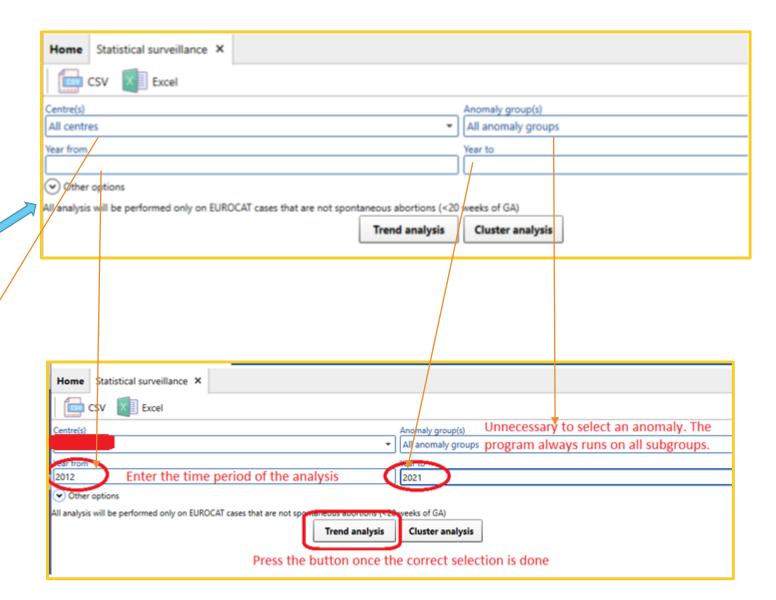
1. Run trends analysis for 1 registry



1 local registry selected

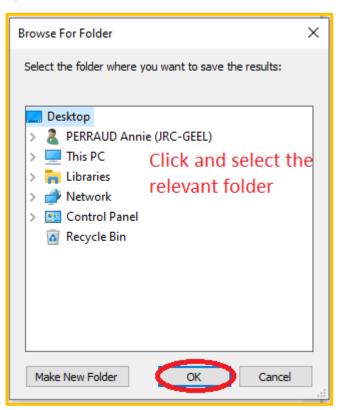




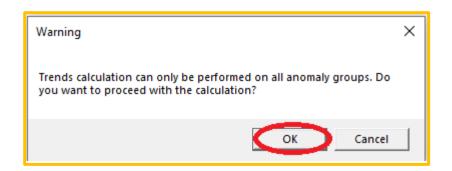


1 local registry selected

1. A pop-up window asks to locate the folder where you want to save the outputs of the R script (graphs and csv).



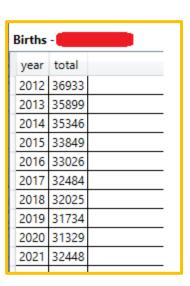
2. A pop-up message warns that the analysis will run on all the anomaly subgroups.



1 local registry selected

Once the analysis has run, DMS displays:

1. a table with the total number of births in the selected registry



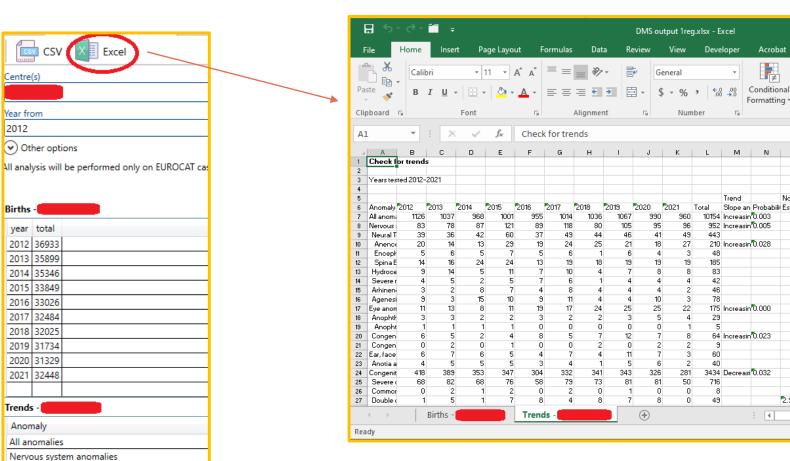
2. a summary of the detected trends in the selected registry

Trends -												
Anomaly	2012	2013	2014	2015	2016	2017	2018	2019	2020	2021	Total	SlopeAndPercentageChange
All anomalies	1126	1037	968	1001	955	1014	1036	1067	990	960	10154	Increasing: 1.0% (95% CI 0.4% to 1.7%
Nervous system anomalies	83	78	87	121	89	118	80	105	95	96	952	Increasing: 3.2% (95% CI 1.0% to 5.5%
Neural Tube Defects	39	36	42	60	37	49	44	46	41	49	443	
Anencephaly and similar	20	14	13	29	19	24	25	21	18	27	210	Increasing: 5.4% (95% CI 0.6% to 10.4
Encephalocele and meningocele	5	6	5	7	5	6	1	6	4	3	48	
Spina Bifida	14	16	24	24	13	19	18	19	19	19	185	
Hydrocephaly	9	14	5	11	7	10	4	7	8	8	83	
Severe microcephaly	4	5	2	5	7	6	1	4	4	4	42	
Arhinencephaly / holoprosencephaly	3	2	8	7	4	8	4	4	4	2	46	
Agenesis of corpus callosum	9	3	15	10	9	11	4	4	10	3	78	
Eye anomalies	11	13	8	11	19	17	24	25	25	22	175	Increasing: 13.1% (95% CI 7.3% to 19.
Anophthalmos / microphthalmos	3	3	2	2	3	2	2	3	5	4	29	
Anophthalmos	1	1	1	1	0	0	0	0	0	1	5	
Congenital cataract	6	5	2	4	8	5	7	12	7	8	64	Increasing: 10.5% (95% CI 1.4% to 20.
Congenital glaucoma	0	2	0	1	0	0	2	0	2	2	9	
Ear, face and neck anomalies	6	7	6	5	4	7	4	11	7	3	60	
Anotia and atresia / stenosis / stricture of external auditory canal	4	5	5	5	3	4	1	5	6	2	40	
Congenital Heart Defects	418	389	353	347	304	332	341	343	326	281	3434	Decreasing: -1.3% (95% CI -2.4% to -
Severe congenital heart defects	68	82	68	76	58	79	73	81	81	50	716	
Common arterial truncus	0	2	1	2	0	2	0	1	0	0	8	
Double outlet right ventricle	1	5	1	7	8	4	8	7	8	0	49	
N 11 11 11 11 11 11 11 11 11 11 11 11 11	1.	_	I		_	_	_	^	^	_	^	

1 local registry selected

Neural Tube Defec

The tables can be exported to Excel → 1 spreadsheet per table

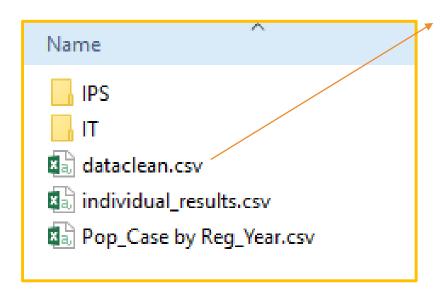


If you choose to export to csv, DMS will create one csv file per table (i.e. one for the total number of births and one for the summary of the trends).

Excel may give an error when opening the .xlsx generated. Please ignore it.

1 local registry selected

In the folder you selected in step 2 (see <u>page 6</u>), the program has created 2 folders and 3 csv files.

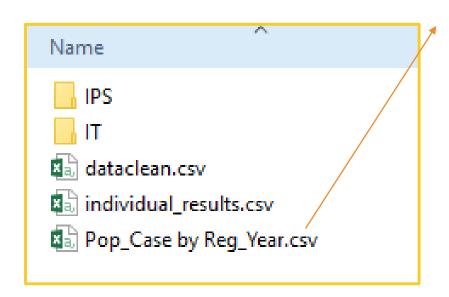


dataclean.csv: input data prepared for analysis

\square	Α		В	С	D	Е	F	G	Н	I	J
1	reg	_[anom	year	case	рор	ganom	nyears	minyr	descanom reg	gna
2		٦	1	2012	1126	36933	0	10	2012	All anoma	
3			1	2013	1037	35899	0	10	2012	All anoma	
4			1	2014	968	35346	0	10	2012	All anoma	
5			1	2015	1001	33849	0	10	2012	All anoma	
6			1	2016	955	33026	0	10	2012	All anoma	
7			1	2017	1014	32484	0	10	2012	All anoma	
8			1	2018	1036	32025	0	10	2012	All anoma	
9			1	2019	1067	31734	0	10	2012	All anoma	
10			1	2020	990	31329	0	10	2012	All anoma	
11			1	2021	960	32448	0	10	2012	All anoma	
12			2	2012	83	36933	1	10	2012	Nervous s	
13		J	2	2013	78	35899	1	10	2012	Nervous s	

1 local registry selected

In the folder you selected in step 2 (see <u>page 6</u>), the program has created 2 folders and 3 csv files.

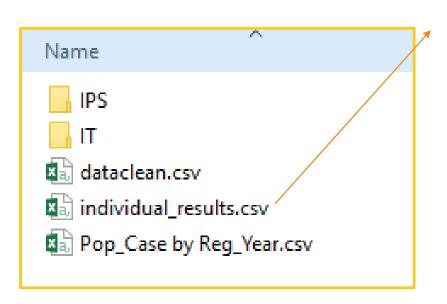


Pop_Case by Reg_year.csv: total number of cases by year and by registry

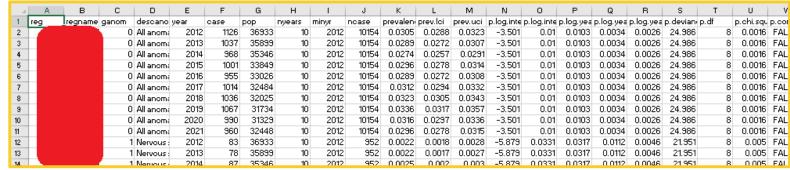
4	А	R		C	U	
1	reg	year		pop	case	
2		2	012	3877965	4012	
3		2	013	3769395	3781	
4		2	014	3711330	3599	
5		2	015	3554145	3675	
6		2	016	3467730	3562	
7		2	017	3410820	3684	
8		2	018	3362625	3725	
9		2	019	3332070	3847	
10		2	020	3289545	3554	
11		2	021	3407040	3498	
12						

1 local registry selected

In the folder you selected in step 2 (see <u>page 6</u>), the program has created 2 folders and 3 csv files.



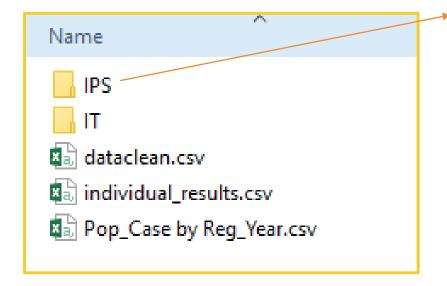
individual_results.csv: detailed results of the linear and spline models, for the local registry selected in each anomaly group

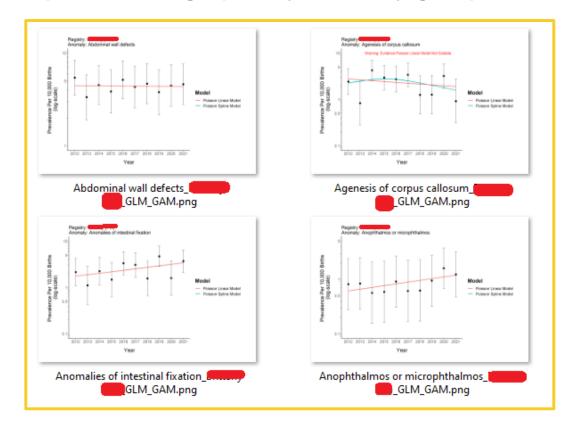


1 local registry selected

In the folder you selected in step 2 (see <u>page 6</u>), the program has created 2 folders and 3 csv files.

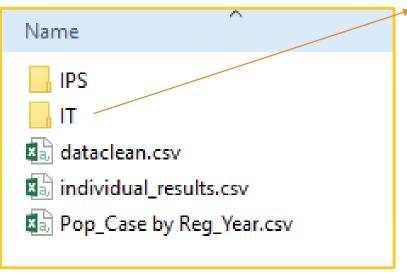
IPS: Fitted prevalence graphs by anomaly group





1 local registry selected

In the folder you selected in step 2 (see <u>page 6</u>), the program has created 2 folders and 3 csv files.



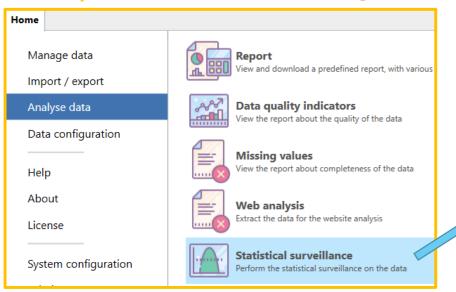
IT: Forest plot of decreasing/increasing trends + list of anomaly groups excluded (e.g. too few cases)

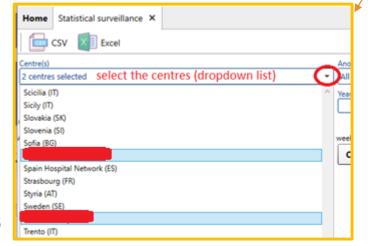


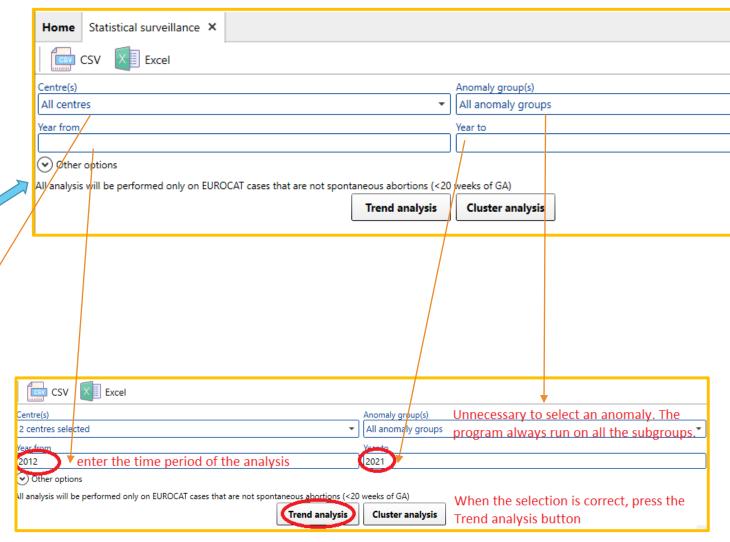
2. Run trends analysis for 2 or more registries



2 or more local registries selected

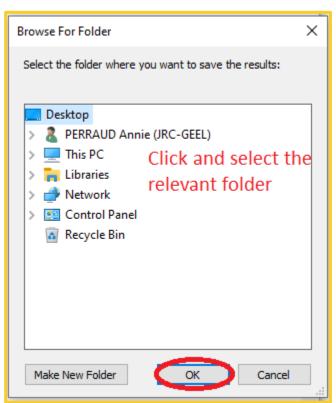




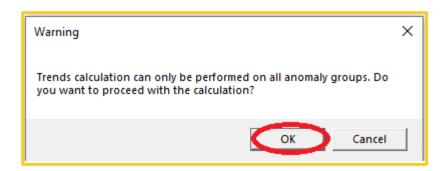


2 or more local registries selected

1. A pop-up window asks to locate the folder where you want to save the outputs of the R script (graphs and csv).



2. A pop-up message warns that the analysis will run on all the anomaly subgroups.



2 or more local registries selected

Once the analysis has run, DMS displays:

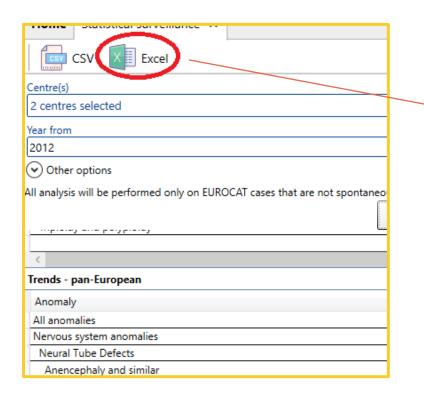
1 – a table with the total number of births in each of the selected registry

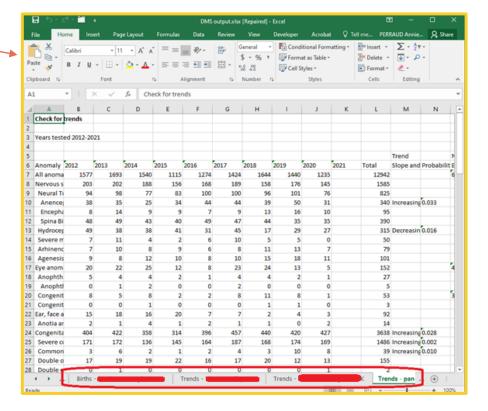
2 – a summary of the detected trends in each of the selected registry and at « pan-European » level

Anomaly	2012	2013	2014	2015	2016	2017	2018	2019	2020	2021	Total	SlopeA	ndPercentageCha
All anomalies	516	572	508	357	492	507	607	508	462		4529	Increasi	ng: 1.8% (95% CI
Nervous system anomalies	69	55	59	58	66	72	57	60	46		542		
Neural Tube Defects	30	24	18	27	36	42	36	32	25		270	Increas	ng: 5.2% (95% CI
Anencephaly and similar	12	11	4	14	15	20	14	15	11		116		
Encephalocele and meningocele	2	6	4	2	5	4	6	7	6		42		
Spina Bifida	16	7	10	11	16	18	16	10	8		112		
Hydrocephaly	20	14	17	17	11	19	8	10	8		124	Decreas	ing: -7.1% (95% (
rends -								_ _					
Anomaly		2012	2013	2014	201	201	6 201	7 20	18 20	19 20	020 2	021 To	tal SlopeAndPe
All anomalies		1061	1121	1032	758	782	917	103	7 93	2 77	3	84	13
Nervous system anomalies	/stem anomalies				98	102	117	101	11	6 99		104	13
Neural Tube Defects		64	74	59	56	64	58	60	69	51		55	5
Anencephaly and similar		26	24	21	20	29	24	25	35	20)	224	1
Encephalocele and meningocele		6	8	5	7	2	5	7	9	4		53	
Spina Bifida		32	42	33	29	33	29	28	25	27		278	3
Hydrocephaly		29	24	21	24	20	26	9	19	19		19	1
Frends - pan-European					2045	2015		2042		2020	2024		
Anomaly	_										2021	Total	SlopeAndPerce
All anomalies		_	_	1540	\rightarrow	-	_	_		1235	_	12942	
Nervous system anomalies	_	-			-	_			176	145		1585	
Neural Tube Defects	9.	-	_						101	76		825	
Anencephaly and similar	3	_	_		_	\rightarrow	_	_	50	31		340	Increasing: 4.6%
Encephalocele and meningocele	8	_	_		_	_	_	-	16	10	_	95	
Spina Bifida	4		\rightarrow	-	_				35	35		390	
Hydrocephaly	4	9 [3	8 3	88	41	31	45	17 I	29	27	I	315	Decreasing: -5.2

2 or more local registries selected

The tables can be exported to Excel → 1 spreadsheet per table

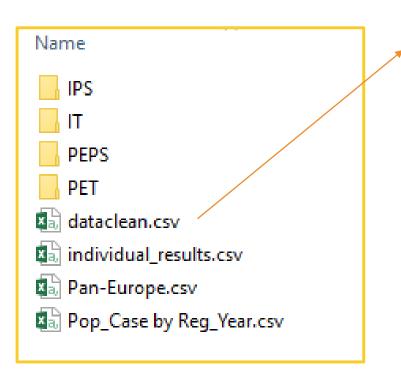




If you choose to export to csv, DMS will create one csv file per table (i.e. one for the total number of births and one for the summary of the trends).

2 or more local registries selected

In the folder you selected in step 2 (see <u>page 16</u>), the program has created 4 folders and 4 csv files.



dataclean.csv: input data prepared for analysis

4	А	D	C	U	С	г	U	п	1
1	reg	anom	year	case	pop	ganom	nyears	minyr	descanom re
2		1	2012	516	31179	0	9	2012	All anoma
3		1	2013	572	29640	0	9	2012	All anoma
4		1	2014	508	29573	0	9	2012	All anoma
5		1	2015	357	29777	0	9	2012	All anoma
6		1	2016	492	29480	0	9	2012	All anoma
7		1	2017	507	28398	0	9	2012	All anoma
8		1	2018	607	27690	0	9	2012	All anoma
9		1	2019	508	26767	0	9	2012	All anoma
10		1	2020	462	26135	0	9	2012	All anoma
11		2	2012	69	31179	1	9	2012	Nervous s
12		2	2013	55	29640	1	9	2012	Nervous s
13		2	2014	59	29573	1	9	2012	Nervous s

2 or more local registries selected

In the folder you selected in step 2 (see <u>page 16</u>), the program has created 4 folders and 4 csv files.

Pop_Case by Reg_Year.csv: total number of cases by year

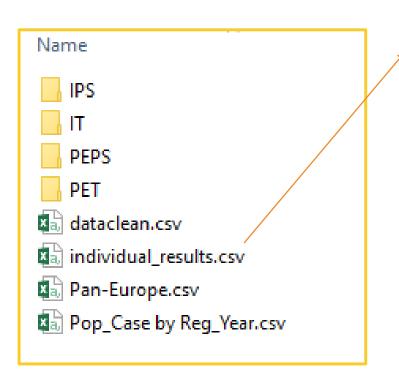
and by registry

Name
<mark>∏</mark> IPS
☐ IT
PEPS /
PET /
dataclean.csv
individual_results.csv
Pan-Europe.csv
Pop_Case by Reg_Year.csv

	A	В	C	D	E
1	reg	year	рор	case	
2		2012	3273795	2113	
3		2012	5485830	4074	
4		2013	3112200	2258	
5		2013	5247480	4194	
6		2014	3105165	1917	
7		2014	5236875	3838	
8		2015	3126585	1570	
9		2015	5204745	3194	
10	1	2016	3095400	2061	
11		2016	5145420	3462	
12		2017	2981790	2052	
13		2017	5004825	3737	
14		2018	2907450	2353	
15		2018	4881555	4092	
16		2019	2810535	2027	
17	•	2019	4747995	3682	
18		2020	2744175	1788	
19		2020	4526130	3116	
20					

2 or more local registries selected

In the folder you selected in step 2 (see <u>page 16</u>), the program has created 4 folders and 4 csv files.

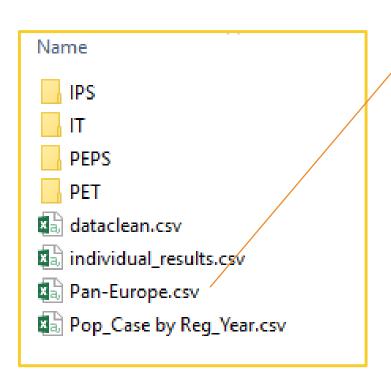


individual_results.csv: detailed results of the linear and spline models, by individual registry for each anomaly group

91				-	-		9	- 0						191	1.5
ı	reg	regname	ganom	descanom	year	case	рор	nyears	n	ninyr	ncase	prevalenc	prev.lci	prev.uci	p.log.inte
8				0 All anoma	2012	516	31179		9	2012	4529	0.01655	0.015192	0.018026	-4.04364
		- 10		0 All anoma	2013	572	29640		9	2012	4529	0.019298	0.017793	0.020928	-4.04364
				0 All anoma	2014	508	29573		9	2012	4529	0.017178	0.015758	0.018723	-4.04364
			70 Ye	0 All anoma	2015	357	29777		9	2012	4529	0.011989	0.010814	0.01329	-4.04364
			83	0 All anoma	2016	492	29480		9	2012	4529	0.016689	0.015289	0.018216	-4.04364
			V Y	0 All anoma	2017	507	28398		9	2012	4529	0.017853	0.016377	0.01946	-4.04364
8				0 All anoma	2018	607	27690		9	2012	4529	0.021921	0.020262	0.023713	-4.04364
			10	0 All anoma	2019	508	26767		9	2012	4529	0.018979	0.017412	0.020684	-4.04364
			(i 19	0 All anoma	2020	462	26135		9	2012	4529	0.017677	0.016149	0.019347	-4.04364
				1 Nervous s	2012	69	31179		9	2012	542	0.002213	0.001749	0.0028	-6.16778
				1 Nervous s	2013	55	29640		9	2012	542	0.001856	0.001426	0.002414	-6.16778
				1 Nervous s	2014	59	29573		9	2012	542	0.001995	0.001547	0.002572	-6.16778
Ø.		. 3		1 Nervous s	2015	58	29777		9	2012	542	0.001948	0.001507	0.002517	-6.16778

2 or more local registries selected

In the folder you selected in step 2 (see <u>page 16</u>), the program has created 4 folders and 4 csv files.

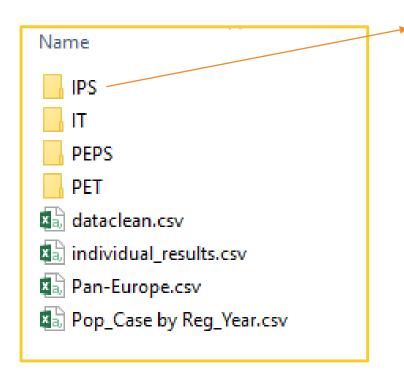


Pan-Europe.csv: detailed results of the linear and spline models, at pan-European level for each anomaly group

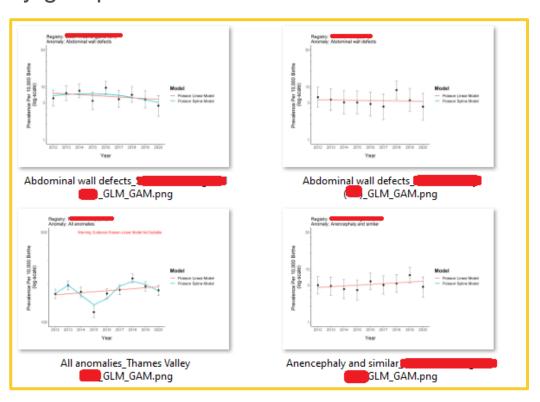
- 4	A	В	L	U	E	-	G	Н	l l	J	K	L	IAI	IV	U	۲	Ų	н	5	1	U	
1	year	ganom	descano	case	рор	minyr	nyears	ncase	prevalen	prev.lci	prev.uci	prev.con	p.log.inte	p.log.inte	p.log.yea	p.log.yea	p.log.yea	p.re.inter	p.re.slop	p.AIC	p.link.fit	p.fit
2	2012	0	All anoma	1577	83425	2012	9	12942	184.09	159.72	212.17	NA	-3.992	0.0362	0.0052	0.0093	0.5775	0.0024	0.0001	389.81	-4.013	0
3	2017	0	All anoma	1424	76063	2012	9	12942	187.21	177.74	197.2	NA	-3.992	0.0362	0.0052	0.0093	0.5775	0.0024	0.0001	389.81	-3.987	0.
4	2014	0	All anoma	1540	79448	2012	9	12942	189.41	166.46	215.51	NA	-3.992	0.0362	0.0052	0.0093	0.5775	0.0024	0.0001	389.81	-4.003	0.
5	2015	0	All anoma	1115	79346	2012	9	12942	136.09	114.94	161.13	NA	-3.992	0.0362	0.0052	0.0093	0.5775	0.0024	0.0001	389.81	-3.998	0.
6	2016	0	All anoma	1274	78484	2012	9	12942	162.33	153.65	171.49	NA	-3.992	0.0362	0.0052	0.0093	0.5775	0.0024	0.0001	389.81	-3.992	0.
7	2019	0	All anoma	1440	71986	2012	9	12942	199.92	188.75	211.76	NA	-3.992	0.0362	0.0052	0.0093	0.5775	0.0024	0.0001	389.81	-3.977	0.
8	2020	0	All anoma	1235	69241	2012	9	12942	178.36	168.69	188.59	NA	-3.992	0.0362	0.0052	0.0093	0.5775	0.0024	0.0001	389.81	-3.972	0.
9	2013	0	All anoma	1693	79616	2012	9	12942	209.11	188.39	232.1	NA	-3.992	0.0362	0.0052	0.0093	0.5775	0.0024	0.0001	389.81	-4.008	0.
0	2018	0	All anoma	1644	74181	2012	9	12942	221.62	211.21	232.53	Model fai	-3.992	0.0362	0.0052	0.0093	0.5775	0.0024	0.0001	389.81	-3.982	0.
1	2017	1	Nervous:	189	76063	2012	9	1585	24.847	21.545	28.654	NA	-6.095	0.049	-0.009	0.0118	0.4232	0.0034	7E-05	145.98	-6.105	0.0
2	2014	1	Nervous:	188	79448	2012	9	1585	23,431	19.579	28.041	NA	-6.095	0.049	-0.009	0.0118	0.4232	0.0034	7E-05	145.98	-6.076	0.0
3	2013	1	Nervous:	202	79616	2012	9	1585	23,803	17.281	32.787	NA	-6.095	0.049	-0.009	0.0118	0.4232	0.0034	7E-05	145.98	-6.067	0.0

2 or more local registries selected

In the folder you selected in step 2 (see <u>page 16</u>), the program has created 4 folders and 4 csv files.

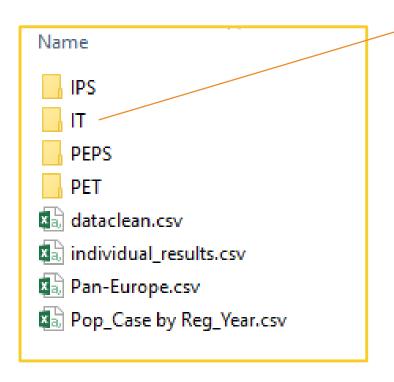


IPS: Fitted prevalence graphs for each individual registry by anomaly group



2 or more local registries selected

In the folder you selected in step 2 (see <u>page 16</u>), the program has created 4 folders and 4 csv files.

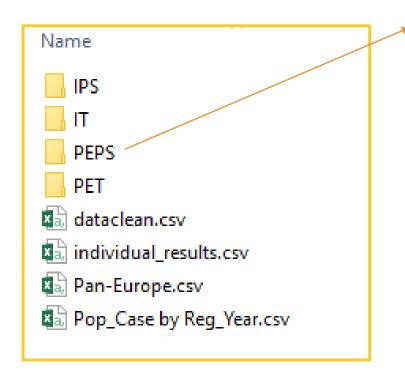


IT: Forest plot of decreasing/increasing trends for each individual registry + list of anomaly group excluded (e.g. too few cases)

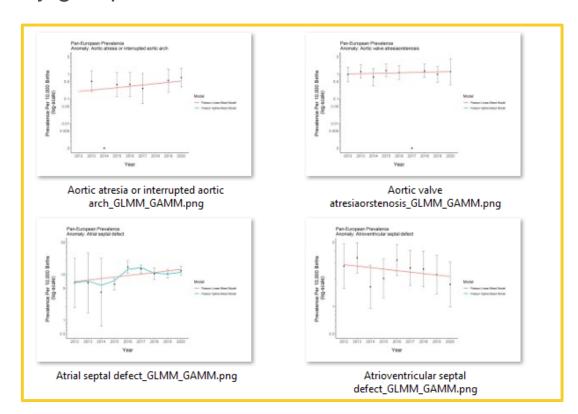


2 or more local registries selected

In the folder you selected in step 2 (see <u>page 16</u>), the program has created 4 folders and 4 csv files.

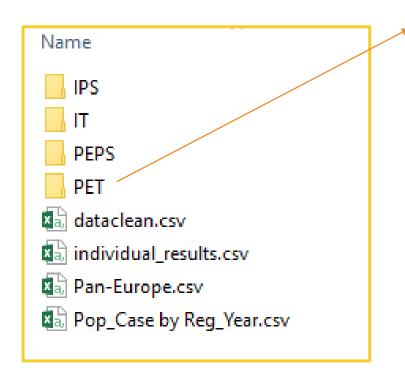


PEPS: Fitted prevalence graphs, at Pan-European level, by anomaly group

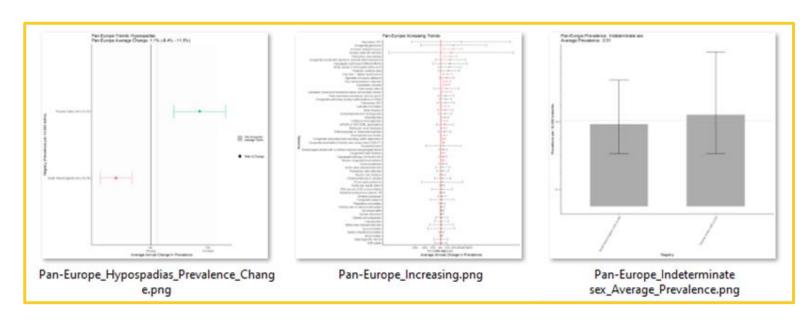


2 or more local registries selected

In the folder you selected in step 2 (see <u>page 16</u>), the program has created 4 folders and 4 csv files.



PET: Forest plot of decreasing/increasing trends at Pan-European level + average prevalence by anomaly group + % change by registry



3. R distribution for DMS



Install the R distribution for Eurocat DMS.msi

Accessing the statistical moniroting functionnalities in the DMS implies to install a distribution of the free R software that is recognised by the DMS.

To request the installation files, please contact the <u>JRC-EUROCAT central registry</u>.



The *.msi* must be run on the user Windows profile, after the main structure of the software (*.exe*) has been installed. An IT administrator of your organisation may need to enter his credentials to allow the installation.

New R libraries for DMS - download

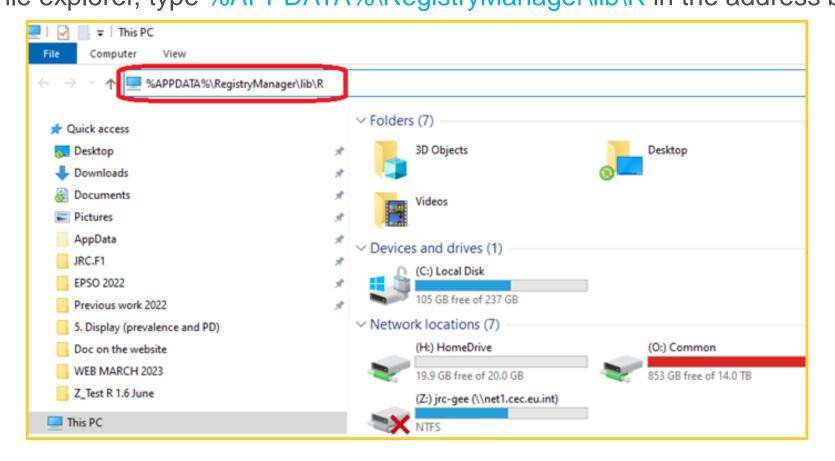
Once the R distribution has been installed (see <u>page 28</u>), the R libraries need to be updated. Please, proceed as indicated in the following slides.

1. Please contact the <u>JRC-EUROCAT central registry</u> to get the complete list of libraries.

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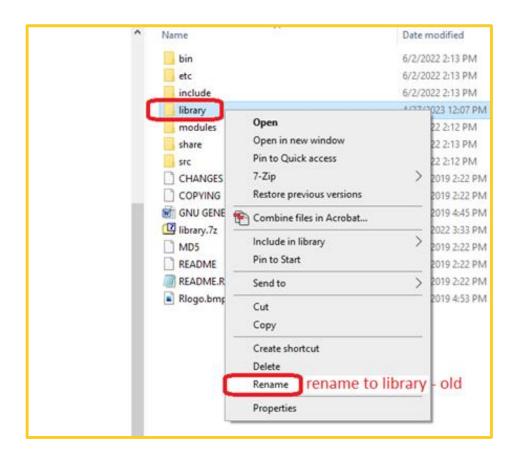
New R libraries for DMS – locate the folder where the libraries will be saved

2. In your File explorer, type %APPDATA%\RegistryManager\lib\R in the address bar and press ENTER:



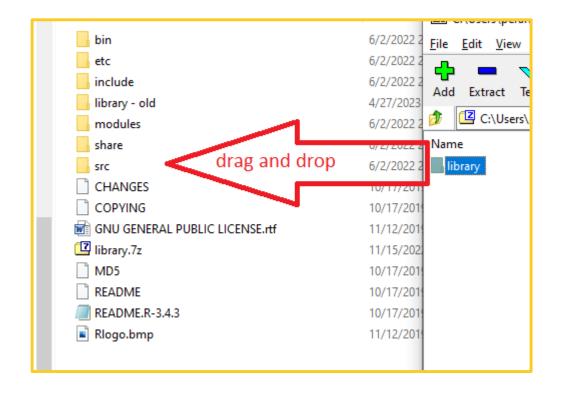
New R libraries for DMS – rename existing folder

3. Rename the folder library to library - old



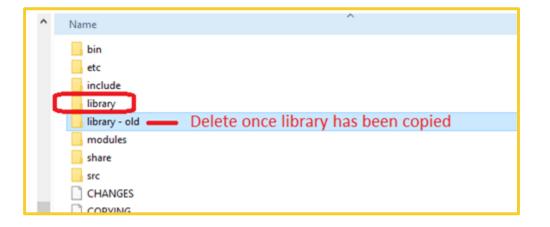
New R libraries for DMS – copy new R libraries

- 4. Open the .zip file you downloaded in step 1 (see page 29)
- 5. Drag and drop the folder named library in this file to the RegistryManager\lib\R



New R libraries for DMS – delete old R libraries

6. Wait until the folder is copied, then you can delete the folder library - old





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