Leishmaniasis Country Profile Generator

User Manual

The Leishmaniasis Country Profile Generator, from now LCPG, retrieves country data from several sources for a specific year and puts it in an HTML country profile format ready to print in PDF.

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1. Changelog

Version	Date	Changes
0.23	2019.11.19	Charts aligned. Charts size and font size reduced to adjust to two pages.
0.22	2019.11.18	Title on yearly incidence and new cases charts
0.21	2019.11.08	Introduction about N/A and No data added to 3.2 "Country profile" chapter.
		Total population and Population at risk description improved. (B1, C10, D5).
		Bugfix: new cases and incidence chart. Gap between years and data fixed.
0.20	2019.11.07	Gender and Population age group data is now retrieved from UN_WPP_POP indicators.
		Life expectancy at birth is now retrieved from GHO, not WB.
0.19	2019.11.05	All texts reviewed and adapted to 2015 CP texts.
		Country General Information and Control and Surveillance sections show now in one column.
		Printing the CP chapter of the manual updated
		Long numbers are now converted to locale format (e.g. 1,203,103 instead 1203103)
		Life expectancy at birth and GDP values are now rounded to the nearest integer. (comments
		updated in this manual)
0.18	2019.11.04	Country names are now shortNames
		Incidence rates indicators updated
		Bugfix: Control & Surveillance and Treatment and medicines sections were being only filled
		with VL data.
		antiCache added for PDF manual
0.17	2019.11.04	New chart configuration section
		Version and date now available on loading page
		Bugfix: User was not able to select first map or legend in the map configuration section.
0.16	2019.11.01	Bugfix: App crashed when trying to retrieve footnotes from countries not having children.

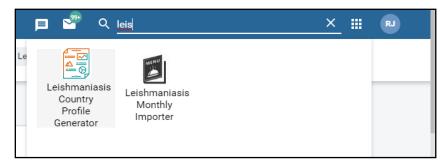
2. Metadata and permissions requirements

LCPG and the user using it need metadata (and eventually) data read access to the following metadata:

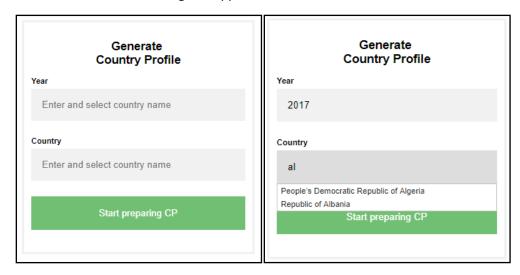
Туре	ID	Name	Comments
program	w9hSFsNr3Vh	CL_cases_by provenance	
program NVUlJzIakuO Fo		Footnotes for Report Generator RG_	Needs to be assigned to the country
program	Jd8gnElt8uT	Leishmaniasis endemicity	Needs to be assigned to the country
program	i5JSf4ffFl2	VL_cases_by provenance	,
dataSet	Uc3j0vpsfSB	Cutaneous Leishmaniasis - ACL/ZCL - Detailed aggregated - Annual	
dataSet	Sn0dExPzQqW	Cutaneous Leishmaniasis - ACL/ZCL - Simple aggregated - Annual	
dataSet	tnek2LjfuIm	Cutaneous Leishmaniasis - Detailed aggregated - Annual	
dataSet	zna8KfLMXn4	Cutaneous Leishmaniasis - Simple aggregated - Annual	
dataSet	NKWbkXyfO5F	General information	Needs to be assigned to the country
dataSet	p0NhuIUoeST	GHO indicators for NTDs	Needs to be assigned to the country
dataSet	fdBM4sWSuPR	Visceral Leishmaniasis - Detailed aggregated - Annual	
dataSet	SHw2zOysJ1R	Visceral Leishmaniasis - Simple aggregated - Annual	
sqlViews	mejiVo59hWs	categoryOptionCombos in DS	
sqlViews	oQdIVqkVlxC	data elements in dataSet	
sqlViews	IrawAndH02Y	data elements used in program	
legendSet	clwSlrqvmMx	ACL Incidence	
legendSet	TnU2O8YxH51	CL Incidence	
legendSet	gUOjExXros1	VLIncidence	
legendSet	TbrqpLWzLS8	ZCL Incidence	
indicatorGroup	nozEoB0uRq9	NTD_Leish_CP_INC_charts_IG	It doesn't need the IG itself but its indicators.
indicatorGroup	VvTNYst2QCW	NTD_Leish_CP_maps_IG	It doesn't need the IG itself but its indicators.
indicatorGroup	KUdeVRtIK45	NTD_Leish_CP_popAtRisk_IG	It doesn't need the IG itself but its indicators.
indicatorGroup	Wp7ZgcxoAwM	NTD_LSH_EPI_NEW_UNS_IG	It doesn't need the IG itself but its indicators.
indicatorGroup	OxgkCeNyVVm	NTD_LSH_TREAT_completed_IG	It doesn't need the IG itself but its indicators.
indicatorGroup	jCYF44Wq3r7	NTD_LSH_SCREEN_passive_IG	It doesn't need the IG itself but its indicators.
indicatorGroup	VbB8TCGqmH5	UN_WPP	It doesn't need the IG itself but its indicators.

3. Generating a country profile

Look for the app called "Leishmaniasis Country Profile Generator" in the apps bar of the WIDP instance.

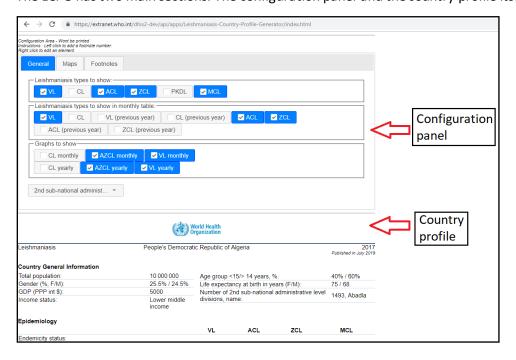


A form box like the following will appear



Start typing a year and a country. Once it starts appearing in the list, select it. This selection is needed, otherwise, the system won't take in account the text you typed.

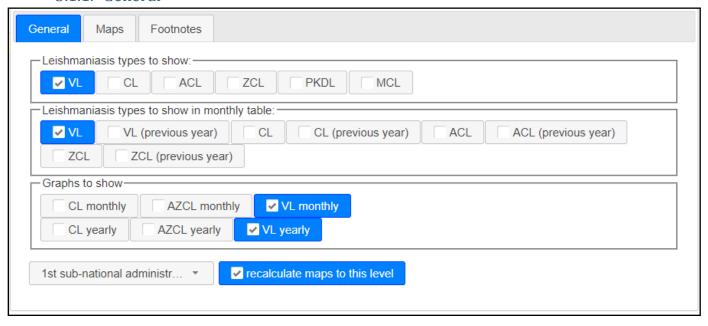
The LCPG has two main sections: The configuration panel and the country profile itself.



3.1. Configuration panel

In this panel you can configure which elements are showing in the CP and how. It has three tabs: General, to manage which types of the disease will appear in the tables, the graphs and the subnational level taken in account; maps to configure maps, legends and "notas bene"; and Footnotes, to manage which footnotes will appear in the CP.

3.1.1. General



Leishmaniasis types to show are checked if, at least, one leishmaniasis dataset of this type (VL, CL or ACL/ZCL) is assigned to the country. PKDL and MCL are unchecked by default.

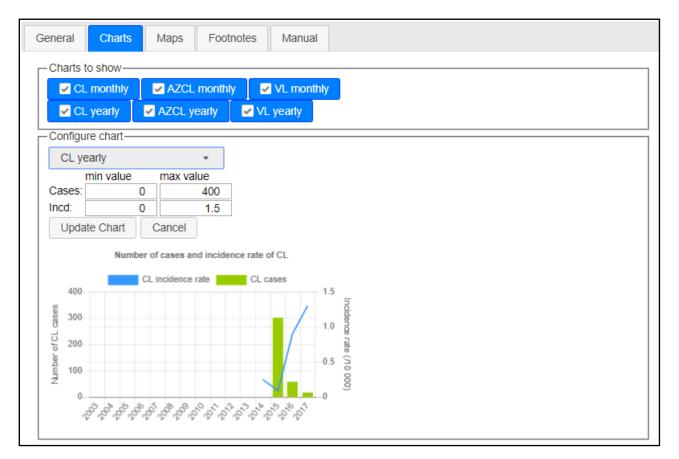
The behavior is the same in the monthly table checkboxes except for previous years, whose checkboxes are by default unchecked.

The graphs checkboxes follow also the same logic.

The subnational level dropdown menu is set, by default, to the first subnational level. Changing it to 2nd or 3rd subnational level will update:

- The "Number of endemic X sub-national..." row text-and-value in the Country General Information section.
- The "Number of endemic X sub-national..." row text-and-value in the Epidemiology section.
- All the maps if the "recalculate maps to this level" checkbox is checked.

3.1.2. Charts



The active chart checkboxes are checked by default. Check or uncheck a checkbox to, respectively, make a chart appear or disappear.

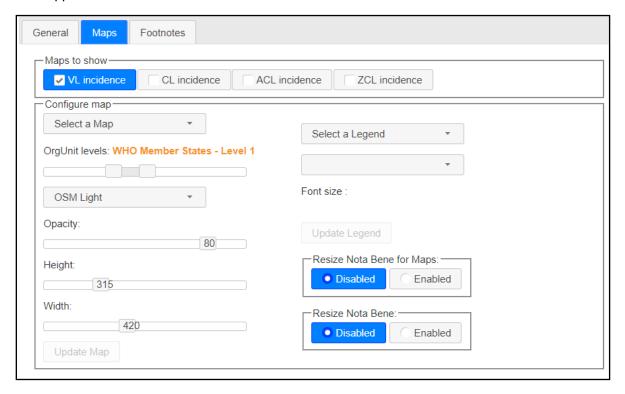
You can select a chart to configure it (if its checkbox is unchecked it will appear as greyed out).

You can adapt the lower and upper bounds of the "number of cases" axis and (if available) the incidence axis. Just type or use the arrows in the correspondent input field to modify the values. You will be able to see changes in real time.

Click on "Update chart" to update it with the new bounds or Cancel to leave it as it was before.

3.1.3. Maps

The active map checkboxes are checked by default. Check or uncheck a checkbox to, respectively, make a chart appear or disappear.

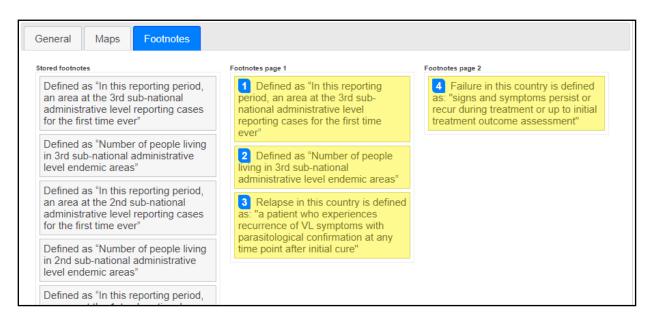


To update a map, select a map name on the dropdown. Select the deepest subnational level you want to see drawn in the map. You can select also the background layout, the opacity, the height and the width. Click "Update Map" once you finished the configuration.

You can relocate and resize the legend within the map. To do that, select a legend on the right dropdown menu. You will see appear the legend at right. You can resize it, edit the text or select the corner where it will be shown. Click "Update Legend" to apply changes.

Size of the two "Nota Bene" in the CP are locked by default. You may enable this feature to adapt the size of the box to the content or the location. Remember putting them back to "disabled" once you have finished.

3.1.4. Footnotes



In this section you organize the footnotes stored in and got from the footnote program.

To move one footnote from one page to another or to remove from the CP, just drag and drop it in the correct box. You can reorder them within the box. The footnote index will be accordingly updated. However, the index you put in the CP text are not "linked" to these footnotes: It's up to you to make sure the reference and the index match.

3.2. Country profile

LPCG usually shows N/A when Not Applicable (for example, data was not requested to the country¹) and No data when blank data (no 0's) is found in the system. Other complex cases are explained in their own section.

3.2.1. Texts to update before printing

Two texts in CP are generated but must be verified before printing. Those text are highlighted in red and, as described in 2.2.2 section, can be modified and its color changed to black. The texts are:

Name of the division levels: The system, takes, as example, the first subdivision level found, but this text should be updated to its specific category name (region, district, department, upazilla... etc).

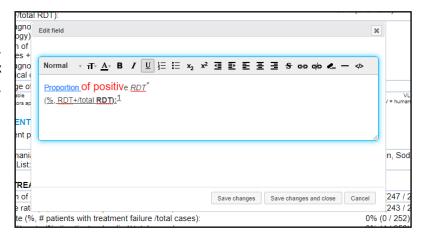


Title on maps: The text in red is automatically changed when you check or uncheck the maps, checkboxes. Just, verify the title is correct and change the color to blue.

Distribution of VL and CL cases per 10 000 population

3.2.2. Edit elements in the CP

You can edit almost any text in the CP. To do that, just right click on the element and an edit field box will pop up. You can change the text, color, size, text style, add hyperlinks, etc.



3.2.3. Footnote elements in the CP

You can footnote almost any element of the CP by left clicking on the element. A footnote index (1 for first clicked element) will be added to the element. The next element will be footnoted with the number "2" and so on. If you want to

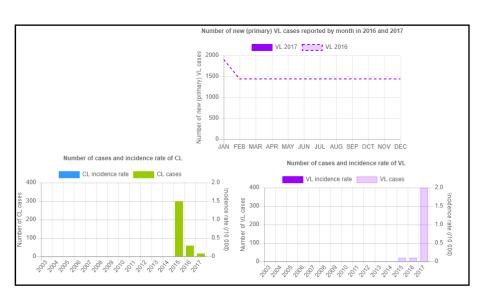
Was there any outbreak?¹
Number of new foci:²
N/A not applicable

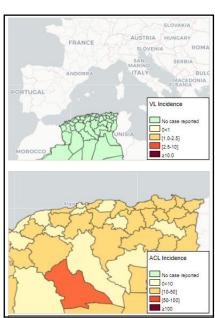
¹ Currently, that's only valid when data comes directly from dataElements and not from DHIS2 indicators.

remove a footnote index, just left click again on it. It's better to remove all higher footnote indexes first, to keep a logic sequence of indexes.

3.2.4. Arrange maps and charts

Charts and maps can be moved by clicking on them and moving the mouse to the desired location. In addition, maps can be zoomed in and its content moved up, down, left or right. However, they cannot be zoomed out for the instance. If you need to reset a map, just go to the maps section in the configuration zone, select the map and click on "Update map".





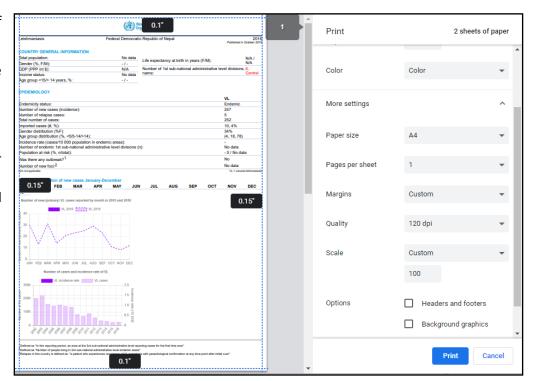
3.2.5. Printing the CP or saving it as PDF

To generate the PDF version of the CP, click [Ctrl]+[P].

You can adjust the final with the parameters:

- paper size (A4)
- scale (normally 100%)
- margins (minimum or custom).
- Uncheck "Headers and footers"

Select your printer or the option "Save as PDF".



4. Source of information in country profile sections

4.1. Country general information section

	COUNTRY GENERAL INFORMATION				
B1	Total population:	41,320,000	В5	Age group <15/> 14 years, %:	0% / 0%
B2	Gender (%, F/M):	0% / 0%	В6	Life expectancy at birth in years (F/M):	75 / 68
В3	GDP (PPP int \$):	5000	D 7	Number of 2nd sub-national administrative level	1493. Abadla
В4	Income status:	Lower middle	D/	divisions, name:	1495, Abadia
		income			

CODE	DataSet / Program	DataElement / Indicator	CatCombos / comments						
B1	DS_GeneralInformation	GEN_UN_WPP_Pop_Tot_1	It shows "No data" if no data value found.						
		000 * 1000							
			Total population (GEN_UN_WPP_Pop_Tot_1000) is						
			used in B1, C10 and D5.						
D2	DC Conoral Information	UN_WPP_POP_GENDER_FEMALE_%							
B2	DS_GeneralInformation	UN_WPP_POP_GENDER_MALE	Z_%						
В3	DS_GeneralInformation	NY.GDP.PCAP.PP.CD	Value is rounded to the nearest integer.						
B4	DS_GeneralInformation	GEN_WB_IncomeGroup							
B5	DS GeneralInformation	UN_WPP_POP_AGE_U15_%							
	DS_GeneralInformacion	UN_WPP_POP_AGE_OVER15_							
В6		WHOSIS 000001 FMLE							
	DS_GeneralInformation	WHOSIS_000001_MLE	Value is rounded to the nearest integer.						
В7	Number of subdivisions in the orgUnitTree for the current country at the selected level. The name is the first occurrence found in the								
	orgUnitTree.								

4.2. Epidemiology section

	EPIDEMIOLOGY						
		VL	CL	ACL	ZCL	PKDL	MCL
C1	Endemicity status:	Non endemic	Previously endemic	Endemic	Endemic	Error!	Error!
C2	Number of new cases (incidence):	25	15	100	50	No data	105
C3	Number of relapse cases:	No data	1	No data	No data	N/A	N/A
C4	Total number of cases:	25	18	100	50	No data	105
C5	Imported cases (#, %):	No data, No data	21, 33%	No data, No data	No data, No data	N/A	N/A
C6	Gender distribution (%F):	No data	33%	No data	No data	No data	99%
C7	Age group distribution (%, <5/5-14/>14):	No data	(43, 47, 10)	No data	No data	No data	(100, No data, No data).
C8	Incidence rate (cases/10 000 population in endemic areas):		0	-	-	N/A	N/A
C9	Number of endemic 1st sub-national administrative level divisions (n):	9	47	No data	No data	N/A	N/A
C10	Population at risk (%, n/total):	0% 1800 / 41320000	0% 9200 / 41320000	0% 0 / 41320000	0% 0 / 41320000	N/A	N/A
C11	Was there any outbreak?	No data	Yes	No data	No data	N/A	N/A
C12	Number of new foci:	No data	987	No data	No data	N/A	N/A
	N/A not VL = visceral CL = cutaneous ACL = anthropo applicable leishmaniasis leishmaniasis leishmaniasis		ZCL = zoonotic cutaneou eishmaniasis	s PKDL = po leishmania	st-kala-azar dermal sis		CL = mucocutaneous shmaniasis

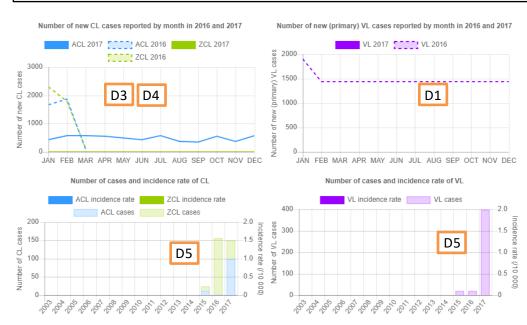
See detailed descriptions on table on next page. Codes are used when possible. Names or names and UIDs otherwise.

CODE	DataSet	DataElement / Indicator	CatCor	mbos / Comments			
C1	GHO_NTDs	NTD_LEISHVEND NTD_LEISHCEND NTD_LEISHACEND NTD_LEISHZCEND NTD_LEISHMCEND NTD_LEISHPKDLEND	-	It replaces the numeric code (1,3 or 5) by "Endemic", "Previously endemic" or "Non endemic". It shows "Error!" if other code is found.			
C2	DS_VL_Detailed_Annual DS_VL_Simple_Annual GHO_NTDs	VL_EPI_Type	New (default for	It shows "No data" if no data found in the system.			
	DS_CL_Detailed_Annual DS_CL_Detailed_Monthly DS_CL_Simple_Annual GHO_NTDs	CL_EPI_Type MCL_GEN_EPID_cases	PKDL and MCL)				
	DS_ACL/ZCL_Detailed_Annual	ACL_EPI_Type ZCL_EPI_Type					
	DS_VL_Detailed_Annual DS_VL_Simple_Annual	PKDL_GEN_EPID_cases					
C3	*As C2 for each DE	VL_EPI_Type CL_EPI_Type ACL_EPI_Type ZCL_EPI_Type	Relapse (N/A for PKDL and MCL)	It shows "No data" if no data found in the system.			
C4	*As C2 for each DE	VL_EPI_Type CL_EPI_Type ACL_EPI_Type ZCL_EPI_Type PKDL_GEN_EPID_cases MCL_GEN_EPID_cases	New Relapse Type unspecified (default for PKDL and MCL)	It shows "No data" if no data found in the system.			
C5	DS_VL_Simple_Annual DS_VL_Detailed_Annual	VL_EPI_Type_Origin	New, Autochthone Relapse, Autoch	thonous			
	DS_CL_Detailed_Annual DS_CL_Detailed_Monthly DS CL Simple Annual	CL_EPI_Type_Origin	New, Imported Relapse, Importe Type unspecified				
	DS_ACL/ZCL_Detailed_Annual	ACL_EPI_Type_Origin ZCL_EPI_Type_Origin		unknown d, Origin unknown			
			not able to ca	instead XY% if it was alculate percentage.			
C6	DS_VL_Detailed_Annual	VL_EPI_Type_Gender	name="New, Female"				
	DS_CL_Detailed_Monthly DS_CL_Detailed_Annual	CL_EPI_Type_Gender	name="New, Gender Uname="New, Male" id=	Jnknown" id="FaYhAlKLX16" -"GnO7H8hC7iV"			
	DS ACL/ZCL Detailed Annual	ACL_EPI_Type_Gender		ed, Female" id="wGED4K5Bs37"			
	_ ,	ZCL_EPI_Type_Gender		ecified, Gender Unknown"			
			id="zkKbllarKWM" name="Type unspecifi	ed, Male" id="aWWYWv6buzp"			
				instead XY% if it was cloulate percentage.			
	DS_VL_Detailed_Annual	PKDL_EPID_sex		id="V2LdgcGgFQt"			
	DS_CL_Detailed_Monthly DS_CL_Simple_Annual DS_CL_Detailed_Annual	MCL_EPID_sex	name="Gender Unknown" id="j name="Male" id	NbFhhnUsQv" l="Z2hvpF7mhh7"			
			LCPG shows - instead XY% if it was not able to calculate percentage.				
C7	DS_VL_Detailed_Annual	VL_EPI_Type_Age		over" id="DDIiBAHqwGV"			
	DS_CL_Detailed_Monthly DS_CL_Detailed_Annual	CL_EPI_Type_Age		nown" id="dVuOzmU4xbI"			
	DS_ACL/ZCL_Detailed_Annual	ACL_EPI_Type_Age	name="New, Under 5y" id="hKq5WASZw8q" name="Type unspecified, 15 y a				
		ZCL_EPI_Type_Age	id="UQMTeRPY2U0" name="Type unspecifi	ed, 5 to 14 y" id="P6R9XEaqQbz"			
			name="Type uns id="nlbrdHlIMKh"	pecified, Age Unknown"			

DS_VL_Detailed_Annual PKDL_PID_age name="15 y and over"id=*TMSELIVE(pr) name="15 y and o				name="Type unspecifie	ed, Under 5y" id="rZwYGlqR8GG"		
DS_CL_Detailed_Monthly DS_CL_Simple_Annual DS_CL_Detailed_Montal DS_CL_Detailed_Montal DS_CL_Detailed_Montal DS_CL_Detailed_Annual DS_CL_Detailed_Annual DS_CL_Detailed_Annual NTD_ISH_AC_EPI_NEW_UNS_I NTD_ISH_AC_EPI_NE		DS VL Detailed Annual	PKDL EPID age				
DS_CL_Detailed_Annual NND_ISH_VL_FPL_NEW_UNS_IND_ST_CL_EPL_NEW_IND_ST_CL_EPL				name="5 to 14 y" id="moktBQGym51"			
NPT LSH VL EPI NEW UNS.] NPT LSH VL EPI NEW		DS_CL_Simple_Annual					
MND LSH ALE PLANE WIND NOT LSH ALE PLANE W		DS_CL_Detailed_Annual		name="Under 5y" id="F	IDXcEOGT2s1"		
NPD_SH_ACL_FN_NW_UNS_ *10000 / population at risk (numerator at C10) *10000 / population (in part at the selected subnational level in CODEHERE having "1" as value for the dataElement and year. *10000 / population (in part at the selected subnational level in CODEHERE having "1" as value for the dataElement and year. *10000 / population (in part at the selected subnational level in CODEHERE having "1" as value for the dataElement and year. *10000 / population (in part at the selected subnational level in CODEHERE having "1" as value for the dataElement and year. *10000 / population (in part at the selected subnational level in CODEHERE having "1" as value for the dataElement and year. *10000 / population (in part at the selected subnational level in CODEHERE having "1" as value for the dataElement and year. *10000 / population (in part at the selected subnational level in CODEHERE having "1" as value for the dataElement and year. *10000 / population (in part at the selected subnational level in CODEHERE having "1" as value for the dataElement and year. *10000 / population (in part at the selected subnational level in CODEHERE having "1" as value for the dataElement and year. *10000 / population (in part at the selected subnational level in code subnational level in code subnational level in code subnational feet of the dataElement and year. *10000 / population (in part at the selected subnational level in code subnational level in code subnational level in code subnational feet of the dataElement and year. *10000 / population (in part at the selected subnational level in code subnational level in code subnat	C8	-	NTD_LSH_VL_EPI_NEW_UNS_I	If population at risk is (), the incidence text shows N/A.		
Total population (GEN_UN_RPP_Pop_Tot_1000) is used in 81, Cl0 and DS. Leishmaniasis endemicity DET_VL_endemicity WHD DET_CL_endemicity WHO DET_CL_endemicity WHO DET_CL_endemicity WHO DET_CL_endemicity WHO DET_CCL_endemicity WHO DET_CCL_endemicity WHO DET_CCL_endemicity WHO DET_CCL_endemicity WHO DET_CCL_endemicity WHO DET_CCL_endemicity WHO Total population (GEN_UN_RPP_Pop_Tot_1000) is used in 81, Cl0 and MCL. VL_EOR_AT_RISK_I CL_POR_AT_RISK_I CEN_POR_AT_RISK_I CEN_P				N/A for PKDL and MCL.			
* 10000 / population at risk (numerator at C10) DET VL endemicity WHO DET CL endemicity V WHO DET ACL ENDEMISS I ACL POP AT RISK I ACL POP AT RISK I CEN POP Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the corresponding program indicator value: GEN pop Leish if the value: GEN pop Leish if the value: GEN pop Leish if the value: GEN pop Leish in the selected substance v							
C10 DS VL_Simple_Annual DS CL_Simple Annual DS ACL/ZCL_Detailed_Annual DS CL_Simple_Annual DS CL_S			NTD_LSH_ZCL_EPI_NEW_UNS_I	Total population (GEN	UN WPP Pop Tot 1000) is		
C10 DS VL_Simple_Annual DS CL_Simple Annual DS ACL/ZCL_Detailed_Annual DS CL_Simple_Annual DS CL_S				used in B1. C10 and D5			
C9				,			
Subnational level in CODERER DET CL endemicity WHO DET ACL endemicity WHO DET ACL endemicity WHO DET ZCL endemicity WHO Factorial ZCL POP AT RISK I ACL POP AT RISK I ACL POP AT RISK I ZCL POP AT RISK I DET ZCL		- 1 1 1 1 1 1 1 1	,				
DET_CL_endemicity WHO DET_ACL_endemicity WHO DET_ACL_endemicity WHO DET_ACL_endemicity WHO DET_ACL_endemicity WHO DET_ACL_endemicity WHO DET_ACL_endemicity WHO DET_ACL_ENDEMINER DET_ACL_EN	C9	Leishmaniasis endemicity		Gets the count of			
C10 DS VL Simple Annual DS CL Semple Annual DS CL Simple Annual DS CL Simple Annual DS ACL/ZCL_Detailed Annual DS CL Simple Annual Scus DS CL Simple Annual S			_				
C10 C10 C10 C10 C10 C10 C10 C10				_			
C10			_		_		
C10 C10 C10 C10 C11 C11 C12 C12				N/A for PKDL and MCL.			
C10							
VL_POP_AT_RISK_I CL_POP_AT_RISK_I CL_POP_AT_RISK_I ACL_POP_AT_RISK_I CL_POP_AT_RISK_I CL_POP_AT_RISK_I CL_POP_AT_RISK_I CCL_POP_AT_RISK_I CCL_POP_AT			DET_ZCL_endemicit				
CI_POP_AT_RISK_I ACL_POP_AT_RISK_I ACL_POP_AT_RISK_I ACL_POP_AT_RISK_I ACL_POP_AT_RISK_I CI_POP_AT_RISK_I CEN_POP_AT_RISK_I CEN_POP_A			y WHO				
CI_POP_AT_RISK_I ACL_POP_AT_RISK_I ACL_POP_AT_RISK_I ACL_POP_AT_RISK_I ACL_POP_AT_RISK_I CI_POP_AT_RISK_I CEN_POP_AT_RISK_I CEN_POP_A							
ACL_POP_AT_RISK_I ACL_POP_AT_RISK_I ACL_POP_AT_RISK_I CEN_pop_Lelsh if the corresponding program indicator	C10	-	VL_POP_AT_RISK_I				
ZCL_POP_AT_RISK_I ZCL_POP_AT_RISK_I Program indicator XXX_endemicity WHO factorl_PI equals 1. 0 otherwise. Denominator: GEN_UN_WPP_Pop_Tot_1000 * 1000 LCGEN_UN_WPP_Pop_Tot_1000 * 1000 LCGEN_UN_WPP_Pop_Tot_1000) is used in B1, C10 and D5. N/A for PKDL and MCL. C11 DS_VL_Simple_Annual DS_CL_Detailed_Annual DS_CL_Simple_Annual DS_ACL/ZCL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_CL_GEN_EPID_new focus CCL_GEN_EPID_new			CL POP AT RISK I	The POP AT RIS	K I indicator value:		
Denominator: GEN_UN_WPP_Pop_Tot_1000 * 1000			ACL POP AT RISK I				
equals 1. 0 otherwise. Denominator: GEN_UN_WPP_Pop_Tot_1000 * 1000 LCPG shows '-' instead XY% if it was not able to calculate percentage. Total population (GEN_UN_WPP_Pop_Tot_1000) is used in B1, C10 and D5. N/A for PKDL and MCL. C11 DS_VL_Simple_Annual			ZCL POP AT RISK I				
Denominator: GEN_UN_WPP_Pop_Tot_1000 * 1000 LCPG shows '-' instead XY% if it was not able to calculate percentage. Total population (GEN_UN_WPP_Pop_Tot_1000) is used in B1, C10 and D5. N/A for PKDL and MCL. DS_VL_Simple_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_ACL/ZCL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Sim				XXX endemicity	WHO factor1 PI		
Denominator: GEN_UN_WPP_Pop_Tot_1000 * 1000 LCPG shows '-' instead XY% if it was not able to calculate percentage. Total population (GEN_UN_WPP_Pop_Tot_1000) is used in B1, C10 and D5. N/A for PKDL and MCL. DS_VL_Simple_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_ACL/ZCL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Sim				equals 1. 0 ot	herwise.		
C11 DS_VL_Simple_Annual DS_CL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual Sccus DS_CL_GEN_EPID_new Sccus DS_CL_GEN_EPID_DS_CL_GEN_EPID_DS_CL_GEN_EPID_DS_CL_GEN_EPID_DS_CL_GEN_EPID_D							
LCPG shows '-' instead XY% if it was not able to calculate percentage. Total population (GEN_UN_WPP_Pop_Tot_1000) is used in B1, C10 and D5. N/A for PKDL and MCL. DS_VL_Simple_Annual DS_VL_Detailed_Annual DS_CL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_VL_Simple_Annual DS_ACL_GEN_EPID_outbreak ZCL_GEN_EPID_outbreak ZCL_GEN_EPID_outbreak ZCL_GEN_EPID_outbreak ZCL_GEN_EPID_new Focus DS_CL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_GEN_EPID_new Focus ZCL_GEN_EPID_new Focus ZCL_GEN_EPID_new Focus ZCL_GEN_EPID_new Focus ZCL_GEN_EPID_new				Denominator:			
LCPG shows '-' instead XY% if it was not able to calculate percentage. Total population (GEN_UN_WPP_Pop_Tot_1000) is used in B1, C10 and D5. N/A for PKDL and MCL. DS_VL_Simple_Annual DS_VL_Detailed_Annual DS_CL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_VL_Simple_Annual DS_ACL_GEN_EPID_outbreak ZCL_GEN_EPID_outbreak ZCL_GEN_EPID_outbreak ZCL_GEN_EPID_outbreak ZCL_GEN_EPID_new Focus DS_CL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_CL_GEN_EPID_new Focus ZCL_GEN_EPID_new Focus ZCL_GEN_EPID_new Focus ZCL_GEN_EPID_new Focus ZCL_GEN_EPID_new					Tot 1000 * 1000		
was not able to calculate percentage. Total population (GEN_UN_WPP_Pop_Tot_1000) is used in B1, C10 and D5. N/A for PKDL and MCL. DS_VL_Simple Annual DS_CL_Detailed Annual DS_CL_Simple Annual DS_CL_Simple Annual DS_CL_Simple Annual DS_ACL/ZCL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_CL_Simple Annual DS_CL_S				0210110p			
was not able to calculate percentage. Total population (GEN_UN_WPP_Pop_Tot_1000) is used in B1, C10 and D5. N/A for PKDL and MCL. DS_VL_Simple Annual DS_CL_Detailed Annual DS_CL_Simple Annual DS_CL_Simple Annual DS_CL_Simple Annual DS_ACL/ZCL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_CL_Simple Annual DS_CL_S				LCPG shows '-'	instead XY% if it		
DS_VL_Simple_Annual							
Total population (GEN_UN_WPP_Pop_Tot_1000) is used in B1, C10 and D5. N/A for PKDL and MCL. C11 DS_VL_Simple Annual							
used in B1, C10 and D5. N/A for PKDL and MCL. C11 DS_VL_Simple_Annual DS_VL_Detailed_Annual DS_CL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_ACL/ZCL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_VL_Simple_Annual DS_VL_Simple_Annual DS_VL_Simple_Annual DS_VL_Simple_Annual DS_VL_Simple_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Simple							
used in B1, C10 and D5. N/A for PKDL and MCL. C11 DS_VL_Simple_Annual DS_VL_Detailed_Annual DS_CL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Simple_Annual DS_ACL/ZCL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual DS_VL_Simple_Annual DS_VL_Simple_Annual DS_VL_Simple_Annual DS_VL_Simple_Annual DS_VL_Simple_Annual DS_VL_Detailed_Annual DS_VL_Detailed_Annual DS_CL_Simple_Annual DS_CL_Simple				Total population (GEN	I UN WPP Pop Tot 1000) is		
C11 DS_VL_Simple_Annual VL_GEN_EPID_outbr eak DS_CL_Detailed_Annual CL_GEN_EPID_outbr eak DS_CL_Simple_Annual ACL_GEN_EPID_outbr reak C12 DS_VL_Simple_Annual PCL_GEN_EPID_new focus DS_CL_Simple_Annual Focus DS_CL_Simple_Ann				used in B1, C10 and D5.			
C11 DS_VL_Simple_Annual vL_GEN_EPID_outbr eak DS_CL_Detailed_Annual eak DS_CL_Simple_Annual eak DS_ACL/ZCL_Detailed_Annual ACL_GEN_EPID_outbreak C12 DS_VL_Simple_Annual vL_GEN_EPID_outbreak DS_VL_Simple_Annual eak C13 DS_VL_Simple_Annual vL_GEN_EPID_outbreak DS_VL_Detailed_Annual focus DS_CL_Detailed_Annual focus DS_CL_Detailed_Annual focus DS_CL_Simple_Annual focus DS_CL_Simple_Annual focus DS_ACL/ZCL_Detailed_Annual focus DS_ACL/ZCL_Detailed_Annual focus DS_ACL_ZCL_Detailed_Annual Focus DS_ACL_							
DS_VL_Detailed_Annual							
DS_VL_Detailed_Annual	C11	DS VL Simple Annual	VL GEN EPID outbr	default	Converts the boolean value to		
DS_CL_Detailed_Annual							
DS_CL_Simple_Annual		DS CL Detailed Annual	CL GEN EPID outbr		N/A for PKDL and MCL.		
DS_ACL/ZCL_Detailed_Annual							
reak ZCL_GEN_EPID_outb reak C12 DS_VL_Simple_Annual VL_GEN_EPID_new DS_VL_Detailed_Annual focus DS_CL_Detailed_Annual focus DS_CL_Simple_Annual focus DS_ACL/ZCL_Detailed_Annual ACL_GEN_EPID_new focus ZCL_GEN_EPID_new focus ZCL_GEN_EPID_new			ACL GEN EPID outb				
C12 DS_VL_Simple_Annual VL_GEN_EPID_new focus DS_CL_Detailed_Annual focus DS_CL_Simple_Annual focus DS_CL_Simple_Annual focus DS_ACL/ZCL_Detailed_Annual ACL_GEN_EPID_new focus ZCL_GEN_EPID_new focus ZCL_GEN_EPID_new							
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DS_VL_Detailed_Annual focus DS_CL_Detailed_Annual CL_GEN_EPID_new DS_CL_Simple_Annual focus DS_ACL/ZCL_Detailed_Annual ACL_GEN_EPID_new focus ZCL_GEN_EPID_new	C12	DS VI Simple Annual		default	N/A for PKDL and MCL		
DS_CL_Detailed_Annual	CIZ			uciauit	INA IOI FINDE ATTU IVICE.		
DS_CL_Simple_Annual focus DS_ACL/ZCL_Detailed_Annual ACL_GEN_EPID_new focus ZCL_GEN_EPID_new				1			
DS_ACL/ZCL_Detailed_Annual ACL_GEN_EPID_new focus ZCL_GEN_EPID_new							
focus ZCL_GEN_EPID_new							
ZCL_GEN_EPID_new		DS_ACL/ZCL_Detailed_Annual					
focus							
			focus				

4.3. Monthly distribution of new cases January-December section

ſ	Monthly distribution of new cases January-December												
		JAN	FEB	MAR	APR	MAY	JUN	JUL	AUG	SEP	OCT	NOV	DEC
D1	VL												
۲	VL (previous year)	1909	1441	1441	1441	1441	1441	1441	1441	1441	1441	1441	1441
D2	CL	323	473	439	445	552	593	412	417	451	445	443	421
02	CL (previous year)	1661	1597	276									
D3	ACL	427	581	569	554	498	438	569	371	350	549	380	570
	ACL (previous year)	1664	1865	101									
D4	ZCL	0	0	0	0	0	0	0	0	0	0	0	0
<i>D</i> 4	ZCL (previous year)	2300	1794	122									



CODE	Program	DataElement
D1	VL_cases_by provenance	VL_cases_byProvenance_T
D2		CL_cases_byProvenance_T
D3	CL_cases_by provenance	ACL_cases_byProvenance_T
D4		<pre>ZCL_cases_byProvenance_T</pre>

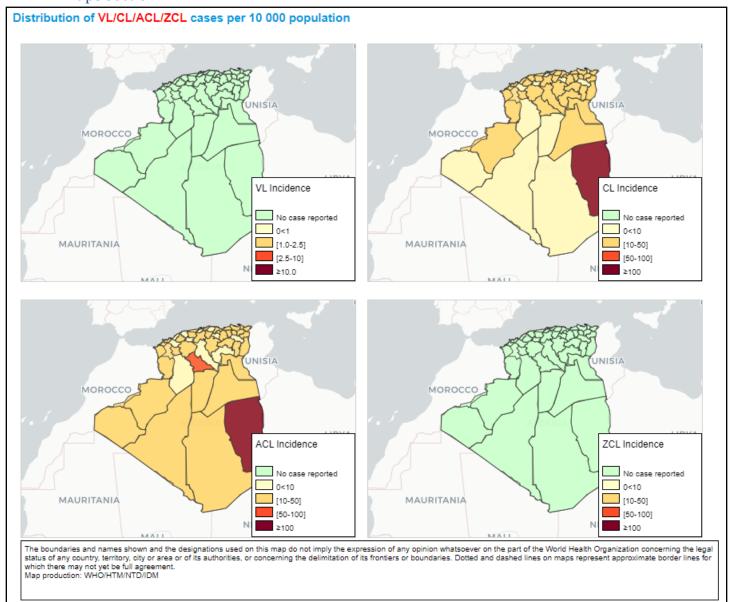
Number of cases (D5)

INDICATOR	Numerator	den	Comments
NTD_LSH_VL_EPI_NEW_UNS_I	VL_EPI_Type New + VL_EPI_Type Type unspecified	1	indicatorType:
NTD_LSH_CL_EPI_NEW_UNS_I	CL_EPI_Type New + CL_EPI_Type Type unspecified		number
NTD_LSH_ACL_EPI_NEW_UNS_I	ACL_EPI_Type New + ACL_EPI_Type Type unspecified		
NTD LSH ZCL EPI NEW UNS I	ZCL EPI Type New + ZCL EPI Type Type unspecified		

Incidence rates (D5)

INDICATOR	Numerator	denominator	Comments
IA_VL_EPI_INC_PopUN_10000	VL_EPI_Type New + VL_EPI_Type Type	GEN_UN_WPP_P	indicatorType:
	unspecified	op_Tot_1000	Per ten
IA_CL_EPI_INC_PopUN_10000	CL_EPI_Type New + CL_EPI_Type Type	* 1000	thousand
	unspecified		
IA_ACL_EPI_INC_PopUN_10000	ACL_EPI_Type New + ACL_EPI_Type Type		
	unspecified		
IA_ZCL_EPI_INC_PopUN_10000	ZCL_EPI_Type New + ZCL_EPI_Type Type		
	unspecified		

4.4. Maps section



INDICATOR	Numerator	denominator	Comments
VL_EPI_INC_PopData_LSH_10000	<pre>VL_cases_byProvenance_T</pre>	GEN_pop_Leish	indicatorType:
CL_EPI_INC_PopData_LSH_10000	CL_cases_byProvenance_T	(In Population data	Per ten
ACL_EPI_INC_PopData_LSH_10000	ACL_cases_byProvenance_T	dataset)	thousand
ZCL EPI INC PopData LSH 10000	ZCL cases byProvenance T		

LEGENDSET name	Legend Name	startValue	endValue	Color
	No case reported	0.0	0.001	#CCFFCC
	0<1	0.001	1.0	#FFFFCC
VL_INCIDENCE_LEGEND_0_10	[1-2.5]	1.0	2.5	#FED976
	[2.5-10]	2.5	10.0	#FC4E2A
	≥10	10.0	10000.0	#800026
	No case reported	0.0	0.001	#CCFFCC
CL INCIDENCE LEGEND 0 100	0<10	0.001	10.0	#FFFFCC
ACL_INCIDENCE_LEGEND_0_100	[10-50]	10.0	50.0	#FED976
ZCL_INCIDENCE_LEGEND_0_100	[50-100]	50.0	100.0	#FC4E2A
	≥100	100.0	10000.0	#800026

4.5. Control and surveillance section

CONTROL AND SURVEILLANCE				
G1 Year Leishmaniasis National Control Programme (LNCP) was established:	2001	G5	Year latest national guidelines (CL / VL):	No data / No data
G2 Type of surveillance (CL / VL):	No data / Integrated	G6	Is leishmaniasis notifiable (mandatory report)? (CL / VL):	No data / No data
G3 Is there a vector control programme?	Yes	G7	Is there a reservoir host control programme?	Yes
Type of insecticide used for Indoor residual Spraying (IRS):	101	G8	Number of leishmaniasis health facilities (CL / VL):	No data / No data

CODE	DataSet	DataElement / Indicator	Comments
G1	DS_VL_Detailed_Annual	Leish_GEN_LNCP_year	It shows "No data" when no entry found in the
	DS_CL_Detailed_Annual		system.
	DS_ACL/ZCL_Detailed_An		
	nual		
G2	DS_CL_Detailed_Annual	CL_GEN_Surv_Type	Converts codes into texts:
	DS_VL_Detailed_Annual	VL_GEN_Surv_Type	1: Vertical
			2: Integrated
			7: Other
			8: Non-applicable
			9: Unknown
G3	DS_CL_Detailed_Annual	Leish_GEN_VectorControl	Converts codes into texts:
	DS_ACL/ZCL_Detailed_An		1: Yes
	nual		2: No
	DS_VL_Detailed_Annual		9: Unknown
G4		Leish_GEN_VectorControl	It shows "No data" when no entry found in the
	,	_Insecticide	system.
G5	DS_ACL/ZCL_Detailed_An	CL_GEN_Guidelines_year	It shows "No data" when no entry found in the
	nual		system.
	DS_CL_Detailed_Annual		
	VL_GEN_Guidelines_year	VL_GEN_Guidelines_year	
G6	DS_CL_Detailed_Annual	CL_GEN_Surv_Notif	Converts codes into texts:
	DS_ACL/ZCL_Detailed_An		1: Yes
	nual		2: No
	DS_VL_Detailed_Annual	VL_GEN_Surv_Notif	9: Unknown
G7	DS_CL_Detailed_Annual	Leish_GEN_ReservoirCont	Converts codes into texts:
	DS_ACL/ZCL_Detailed_An	rol	1: Yes
	nual		2: No
	DS_VL_Detailed_Annual		9: Unknown
G8	DS_CL_Detailed_Annual	CL_GEN_Surv_HF	It shows "No data" when no entry found in the
		VL GEN Surv HF	system.

4.6. Diagnosis section

	DIAGNOSIS						
114		VL	CL	ACL	ZCL	PKDL	MCL
	Number of people screened actively for: Number of people screened passively for:	No data No data	No data N/A	N/A N/A	N/A N/A	N/A N/A	N/A N/A
Н3	VL cases diagnosed by RDT [*] (%, RDT+/total VL cases):	86% (216 / 252)	N/A	N/A	N/A	N/A	N/A
	Proportion of positive RDT* (%, RDT+/total RDT):	100% (216 / 216)	N/A	N/A	N/A	N/A	N/A
H5	Cases diagnosed by direct exam (parasitology) (%, # slides +/total cases):	15% (38 / 252)	No data	N/A	N/A	N/A	N/A
Н6	Proportion of positive slides (%, # slides +/total slides):	100% (38 / 38)	No data	No data	No data	N/A	N/A
H7	Cases diagnosed clinically (%, # clinical cases/total cases):	0% (0 / 252)	No data	N/A	N/A	N/A	N/A
Н8	Percentage of cases with HIV-VL coinfection:	0% (0 / 252)	N/A	N/A	N/A	N/A	N/A
	N/A not VL = visceral CL = cutaneous ACL = anthroponotic cutane applicable leishmaniasis leishmaniasis leishmaniasis *These indicators apply only for primary VL cases	ous ZCL = zoonotic cutaneous leishmaniasis RDT = rapid diagnostic rest	PKDL = leishman	post-kala-azar de niasis	leish	= mucocutan maniasis an immunode	

CODE	DataSet	DataElement / Indicator	CatCombos / Comments	
H1	DS_VL_Detailed_Annual	VL_SCREEN_active	-	N/A for PKDL and MCL
	DS_CL_Detailed_Monthly	CL_SCREEN_active		
	DS_CL_Detailed_Annual			
	DS_ACL/ZCL_Detailed_An	ACL_SCREEN_active		
	nual	ZCL_SCREEN_active		
H2	DS_VL_Detailed_Annual	NTD_LSH_VL_SCREEN_pass	=	The related DE is assigned to the
		ive_I		dataset but it's not in the form!
	DS_CL_Detailed_Monthly	NTD_LSH_CL_SCREEN_pass	-	The related DE is not assigned to
	DS_CL_Detailed_Annual	ive_I		the dataset!
	DS_ACL/ZCL_Detailed_An	NTD_LSH_ACL_SCREEN_pas	=	The related DEs are assigned to
	nual	sive_I		the dataset but they are not in
		NTD_LSH_ZCL_SCREEN_pas		the form!
		sive_I		N/A for PKDL and MCL
Н3	DS_VL_Detailed_Annual	VL_Lab_RDT_results_type	N/A for CL (All types) and PKDL.
		/		
		NTD_LSH_VL_EPI_NEW_UNS		
		I		
H4	DS_VL_Detailed_Annual	VL Lab RDT tested type	name="New"	VL Lab RDT results type
			id="psVSPLcly	(New + Unsp.)
			Fj"	/
			name="Type	VL_Lab_RDT_tested_type
			unspecified"	(New + Unsp.)
			id="IRW4YrOtk	(New - Shisp.)
			5q "	N/A for CL (All types) and
		VL_Lab_RDT_results_type	name="New,	PKDL.
			Positive"	PRDL.
			id="jRcT6HVKb	
			2t"	
			name="Type	
			unspecified,	
			Positive"	
			id="YXktM46Yi	
			Xo"	
H5	DS_VL_Detailed_Annual	VL_Lab_parasito_tested	New	Direct exam diagnosed
		_type	Relapse	/
				Total cases (C4)

	DS_CL_Detailed_Annual DS_ACL/ZCL_Detailed_Mo nthly DS_ACL/ZCL_Detailed_An nual	CL_LAB_parasito_Suspects ACL_Lab_Parasito_Results ZCL_Lab_Parasito_Results	Type unspecified	
H6	DS_VL_Detailed_Annual DS_CL_Detailed_Annual DS_ACL/ZCL_Detailed_Mo nthly	VL_LAB_parasito_result _type CL_LAB_Parasito_Result s	name="New, Positive" id="jRcT6HVKb 2t" name="Relapse , Positive"	Direct exam diagnosed / Direct exam diagnoses (numerator on H5)
	DS_ACL/ZCL_Detailed_An nual	ACL_Lab_Parasito_Results ZCL_Lab_Parasito_Results	<pre>id="QKqVJ13mG ZI" name="Type unspecified, Positive" id="YXktM46Yi Xo"</pre>	
H7	DS_VL_Detailed_Annual DS_CL_Detailed_Annual DS_ACL/ZCL_Detailed_Mo nthly DS_ACL/ZCL_Detailed_An	NO DATA ELEMENT NO DATA ELEMENT	New Relapse Type unspecified	Clinical cases / Total cases (C4)
	nual	NO DATA ELEMENT	1137	
Н8	DS_VL_Detailed_Annual	VL_LAB_HIVstatus_Type	name="New, Positive" id="jRcT6HVKb 2t" name="Relapse , Positive" id="QKqVJ13mG ZI" name="Type unspecified, Positive" id="YXktM46Yi Xo"	VL_LAB_HIVstatus_Type (New Positive + Relapse Positive + Unsp. Positive) / Total cases (C4)

4.7. Treatment and medicines and Treatment Outcome section

11	TREATMENT AND MEDICINES Is treatment provided for free in the public sector? (CL /	N/A / Yes				
	VL): Antileishmanial medicines included in the National	Amphotericin B deoxychola	te, Miltefosine, Paromomy	cin, Sod	ium	
12	Medicine List:	stibogluconate (SSG)				
	INITIAL TREATMENT OUTCOME FOR NEW CASES		VL	CL	ACL	ZCL
13	Proportion of cases treated (%, # treated cases/ total cases):		98% (247 / 252)	N/A	N/A	N/A
14	Initial cure rate (%, # cases initially cured /total cases):		96% (243 / 252)	N/A	N/A	N/A
15	Failure rate (%, # patients with treatment failure /total cases):		0% (0 / 252)	N/A	N/A	N/A
16	Case fatality rate (%, # patients who died/ total cases):		2% (4 / 252)	N/A	N/A	N/A

CO DE	DataSet	DE / Indicator	Comments
11	DS_VL_Detailed_Annual	VL_GEN_TxFree	Converts codes into texts:
	DS_CL_Detailed_Annual	CL_GEN_TxFree	1: Yes
	DS_ACL/ZCL_Detailed_Annu		2: No
	al		9: Unknown
12	DS_VL_Detailed_Annual	Leish_GEN_EML_AmphotericinB	LCPG retrieves ids and replaced by hardcodes names:
	DS_CL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual	Leish_GEN_EML_LiposomalAmp Leish_GEN_EML_Meglumine	Amphotericin B deoxycholate
	bs_, tely zee_betailed_, tillidal	Leish_GEN_EML_Miltefosine	Liposomal amphotericin B
		Leish_GEN_EML_Paromomycin	Meglumine antimoniate Miltefosine
		Leish_GEN_EML_Pentamidine	Paromomycin
		Leish_GEN_EML_SSG	Pentamidine
			Sodium stibogluconate (SSG)
13	DS_VL_Detailed_Annual	VL_TREAT_completed	
		/	
	DC CL Data lad Accord	NTD_LSH_VL_EPI_NEW_UNS_I	
	DS_CL_Detailed_Annual	CL_TREAT_completed	
		, NTD_LSH_CL_EPI_NEW_UNS_I	
	DS_ACL/ZCL_Detailed_Annu	NTD_LSH_ACL_TREAT_completed_I	
	al	NTD_LSH_ZCL_TREAT_completed_I	
		NTD ISH ACLEDI NEW LINE I	
		NTD_LSH_ACL_EPI_NEW_UNS_I NTD_LSH_ZCL_EPI_NEW_UNS_I	
14	DS_VL_Detailed_Annual	VL_INIT_ITxO_Drug_Type	Ambisome, New, Initial Cure
	DS_CL_Detailed_Monthly	CL_ITxO_Tx-route	Ambisome, Type unspecified, Initial Cure Antimonials, New, Initial Cure
	DS_CL_Detailed_Annual DS_ACL/ZCL_Detailed_Annual	ACL_ITxO_Tx-drug ZCL_ITxO_Tx-drug	Antimorials, Type unspecified, Initial Cure
	D3_ACL/2CL_Detailed_Affilial	ZCL_ITXO_TX-drug	Meglumine Antimoniate (glucantime), New, Initial Cure
		/	Meglumine Antimoniate (glucantime), Type unspecified, Initial Cure Other CL drug, New, Initial Cure
			Other CL drug, Type unspecified, Initial Cure
		NTD_LSH_VL_EPI_NEW_UNS_I	Other VL drug, New, Initial Cure Other VL drug, Type unspecified, Initial Cure
		NTD_LSH_CL_EPI_NEW_UNS_I NTD_LSH_ACL_EPI_NEW_UNS_I	SSG, New, Initial Cure
		NTD_LSH_ZCL_EPI_NEW_UNS_I	SSG + Paramomycin, New, Initial Cure
			SSG + Paramomycin, Type unspecified, Initial Cure SSG, Type unspecified, Initial Cure
			Treatment Drug Unknown, New, Initial Cure
			Treatment Drug Unknown, Type unspecified, Initial Cure VL drug unspecified, New, Initial Cure
			VL drug unspecified, Type unspecified, Initial Cure
15			Ambisome, New, Failure
			Ambisome, Type unspecified, Failure Antimonials, New, Failure
			Antimonials, Type unspecified, Failure
			Meglumine Antimoniate (glucantime), New, Failure Meglumine Antimoniate (glucantime), Type unspecified, Failure
			Other CL drug, New, Failure
			Other CL drug, Type unspecified, Failure Other VL drug, New, Failure
			Other VL drug, New, Failure Other VL drug, Type unspecified, Failure
			SSG, New, Failure
			SSG + Paramomycin, New, Failure SSG + Paramomycin, Type unspecified, Failure
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	SSG, Type unspecified, Failure
	Treatment Drug Unknown, New, Failure
	Treatment Drug Unknown, Type unspecified, Failure
	VL drug unspecified, New, FailureVL drug unspecified, Type unspecified, Failure
16	Ambisome, New, Death
	Ambisome, Type unspecified, Death
	Antimonials, New, Death
	Antimonials, Type unspecified, Death
	Meglumine Antimoniate (glucantime), New, Death
	Meglumine Antimoniate (glucantime), Type unspecified, Death
	Other CL drug, New, Death
	Other CL drug, Type unspecified, Death
	Other VL drug, New, Death
	Other VL drug, Type unspecified, Death
	SSG, New, Death
	SSG + Paramomycin, New, Death
	SSG + Paramomycin, Type unspecified, Death
	SSG, Type unspecified, Death
	Treatment Drug Unknown, New, Death
	Treatment Drug Unknown, Type unspecified, Death
	VL drug unspecified, New, DeathVL drug unspecified, Type unspecified, Death