BacLink – EARS-Vet Data Import



WHO Collaborating Centre for Surveillance of Antimicrobial Resistance

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Background

EARS-Vet stands for the "European Antimicrobial Resistance Surveillance Network in Veterinary Medicine". WHONET can assist users with exporting their AMR data to the EARS-Vet file format, which can be uploaded to their data collection platform. For more information on EARS-Vet, please see the following URL: https://anses.hal.science/anses-04685234v1/document

About this document

This document provides information on how to import data from an existing electronic data source into the WHONET data structure compatible with exportation to the EARS-Vet file format.

Obtain a sample data file from your existing data source

Please follow the guidance in the section "BacLink – Exporting data from desktop applications, laboratory instruments, and laboratory information systems" found near the bottom of the following support page. <u>https://whonet.org/training.html</u>

The steps will be different depending on the type of information system your facility uses, but the goal is to generate a file which can be subsequently processed with the BacLink software.

How to create a new EARS-Vet-specific BacLink configuration

To import existing data stored in an export from your LIS, laboratory instrument, or other electronic data source, you first need to create the corresponding BacLink configuration file tailored for EARS-Vet.

1. From the main screen of BacLink, press the "New format" button highlighted below.

BacLink 2024		-		\times
File Select language Help				
	original data file. v data file. Click on 'Begin conversion'. ot appear on the list, choose 'New format'.			
File format C:\WHONET\	Test configuration	New	format	
		Edit	format	
Test configuration-TEXT.cfg		Delet	e format	
File name	C:\WHONET\Data*.txt	Br	owse	
New data file				
File name	C:\WHONET\Data*.sqlite	Br	owse	
Table name	For Access files only			
File format	WHONET (SQLite) ~			
	Begin conversion		Exit	

- 2. Fill in the first three questions regarding your country, laboratory description, and laboratory code.
- 3. Press the "File structure" button and select from the list of options depending on the type of existing data files you have.
 - a. Please see the "BacLink Getting started" document at the following URL for more information. <u>https://whonet.org/WebDocs/BacLink.1_Getting_started_Introduction.pdf</u>
 - b. You may also wish to review the links in the section entitled "BacLink Exporting data from desktop applications, laboratory instruments, and laboratory information systems" found near the bottom of the following support page. <u>https://whonet.org/training.html</u>
 - c. You may also need to configure the antibiotics for your data file. Please follow the general guidance available on our support page at the links above for more information.
 - d. While configuring the antibiotics, you will answer questions about the specific way your antibiotic results appear in the file, such as whether the antibiotics appear horizontally in their own columns, or whether they appear vertically with one antibiotic per row. Because of the many possible responses to these questions, we cannot provide a comprehensive set of instructions here which will cover every scenario.
 - e. If you are unable to answer the questions on the "Antibiotics" screen, then you should consult the documentation mentioned above, or contact the WHONET group for additional support.
- 4. One of the questions on the "Antibiotics" form is related to the antibiotic guidelines which were used in performing the tests in your data file. Some laboratories use CLSI, EUCAST, or SFM exclusively, while others use a mixture of these. If your laboratory uses mixed guidelines, please choose "Multiple guidelines" from the list of shown below.

Configure antibiotics				>
File format T		TEXT (DELIM	ITED)	ок
Does your file include antibiotic	s results?	O Yes	⊖ No	Cancel
Guidelines	Multiple guidelines		~	Configure guidelines
The antibiotics of one isolate req	EUCAST CA-SFM Other Multiple guidelines		w an one row	
In what sequence do the antibiot	ics appear?	0	Fixed antibiotic sequences of the sequences of the second	
a. Currently, BacLink c	only supports multip	le guidelines fo	or data files wh	ich have antibiotics

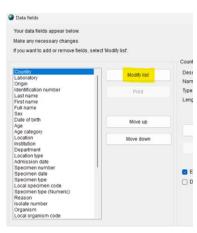
- on separate rows of the data file.b. If you have a mixture of guidelines, but your antibiotics are horizontally in separate
- b. If you have a mixture of guidelines, but your antibiotics are horizontally in separate columns, please contact the WHONET team for more support with your configuration.
- c. When "Multiple guidelines" is selected, there will be an additional data field added to your configuration called "Antibiotic test guideline" which should be matched to one of your data fields which serves this purpose.
- 5. After answering the questions on the "Antibiotics" form, you must now add the EARS-Vet data fields to your configuration. Press the "Data fields" button near the bottom of the "File structure" form shown below.

Country	France	~ F	RA
Laboratory name	EARS-Vet test laborator		
Laboratory code	TST		F
Maximum 10 letters			
_	File structure		
File structure		(
Codes and dates	File structure	Text (Delimited)	~
	Field delimiter	Tab 🗸	
New data file	File location	W:\WHONET.Source\BacLink\bin\x64\Debug\D:	ata\ Browse
	File name	*.bxt	Browse
Data filter			
Save as	File origin	Unicode ~	
Cave as	Character set	Unicode (UTF-8) V	
	Antibiotics	Enter information about the antibiotics in yo	our data file
	Guidelines	No answer	
	Number of rows of da	ta for each isolate No answer	
	Antibiotic sequence	No answer	
	Test methods	No answer	
	Number of test metho	ds in one row of data No answer	
	Does the first row of t	ne data file have the names of the data fields?	
		No	

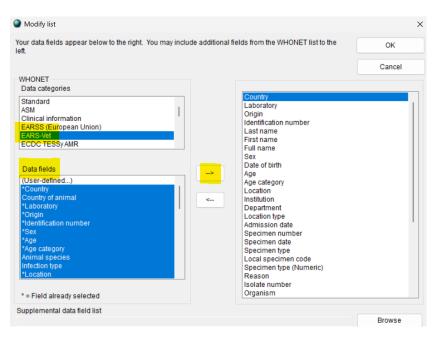
6. Press the "Modify the list of data fields" button.

Data fields			
Select a sample data file			
Data fields in the new file Identification number = «None» Last name = «None» Full name = «None» Sex = «None» Date of birth (DM/Y) = «None»			Data fiel
Age = «None» Location = «None» Department = «None» Specimen durb = «None» Specimen durb = «None» Isolate number = «None» Comment = «None» Comment = «None» Antibiotic result 1 (Undefined) = «Non	e>	=	
Modify the list of data fi	elds		
Antibiotics			
Add	Delete		
Define	Insert		

7. On the "Data fields" form which will appear, press the "Modify list" button.



- BacLink has a default set of data fields included with all new configurations, shown in the box on the right. For the purposes of EARS-Vet, you should remove all the optional default data fields by highlighting each data field and pressing the arrow which points to the left near the center of the form.
 - a. There is a small number of data fields which cannot be removed, for which you will receive a message. Please remove all other data fields.
- 9. Now that you have cleared the default data fields from your configuration, you can proceed to include the EARS-Vet set.
- 10. Choose "EARS-Vet" from the "Data categories" list and add all the EARS-Vet data fields to your configuration by selecting one or more data fields (using the shift key and your mouse) and pressing the arrow which points to the right.
 - a. If you attempt to add a data field which is already part of your configuration, you will receive a message indicating that this data field was skipped.



- 11. The list of data fields shown in the column on the right should now include the EARS-Vet data fields. Press "OK" on the "Modify list" form to return to the "Data fields" form.
- 12. It is now time to match the data fields in your sample input file with those you've just added to your configuration. Begin by pressing the "Select a sample data file" button and choosing your sample AMR data file you've extracted previously from your facility's information system.

Data fields		×
Select a sample data file		ОК
Data fields in the new file Identification number = <none></none>	Data fields in the original file Select a sam	nle data file
Last name = <none></none>		

- 13. After selecting your sample file, BacLink will read the first isolate and display it on the screen along with the data field header names found in your sample file as shown below.
- 14. You must now match each of your data fields in the left column with the corresponding column shown on the right by highlighting the fields in both columns and pressing the equals sign in the center of the screen.

Click on a WHONET data field and on the corresponding	g field from your data file.	OK
Click '=' to match the two fields.		
Data fields in the new file	Vitek2-demo-data.txt	
Specimen number = Lab ID	<none below="" of="" the=""></none>	
Specimen date (M/D/Y) = Collection Date	Lab ID = 371173143	
Specimen type = Specimen Type	Isolate Number = 1 Patient Name = John Smith	
Isolate number = Isolate Number	Patient ID = 2044259	
Organism = Organism Name	Patient Location = ICU	
Comment = <none></none>	Specimen Type = Blood	
Specimen Source = Specimen Source	Specimen Source = Collection Date = 2014-06-02	
Testing Date (M/D/Y) = Testing Date	Testing Date = 2014-06-01	
Bio Number = Bio Number	Organism Name = Staph.haemol	yticus
Country of animal = <none></none>	Organism Code = MHA Bio Number = 010002042660031	

15. There are several differences between the WHONET and EARS-Vet terminology used to describe the same concepts. Please refer to the following table.

EARS-Vet name	WHONET/BacLink name	
Isolate ID	Isolate number	
Bacterial species	Organism	
AMPc phenotype	AMPc production	
ESBL profile	ESBL	
AST technique*	Antibiotic test method	
AST standard*	Antibiotic test guideline	

* An important note: partners who have only one AST method or guideline used in their data set (specified in the "file structure" configuration) will not see the corresponding "AST technique" or "AST standard" respectively. In all cases, the columns will be present in the WHONET export to EARS-Vet.

- 16. Some of the EARS-Vet data fields may not have a corresponding field in your input file. For example, if you do not have a data field for "Year of sampling", then you should use a "Fixed value", for example "2024". To do this, type the fixed value into the corresponding text box and press the "equals" sign in the center of the screen.
 - a. You may be missing other data fields, such as the AST instrument, etc. You can use a fixed value to provide these as well.
 - b. If you needed to set a fixed value for the "Year of sampling", you will need to adjust this fixed value each year in your configuration.

Year of testing = '2024' Laboratory = 'JEJE' Animal identification number = <none> Sex = <none> Location = <none> Specimen number = <none> Specimen date (D/M/Y) = <none> Age category = Age category Infection type = Infection type AST instrument = AST instrument AST standard (other) = AST standard other</none></none></none></none></none>	AST standard - other = AST standard - other = Antimicrobial agent = amoxicillin Disk diameter or MIC value = 12 Virulence testing technique = pcr be Virulence testing technique_other = Virulent strain = - ESBL confirmation technique = EUVSEC2 panel be ESBL confirmation technique_other = ESBL profile = + AmpC confirmation technique = AmpC confirmation technique = AmpC phenotype = - PCR mecA = +
Modify the list of data fields	PCR mecC = - Fixed value 2024

- 17. Once you have completed your data field matches, you may press "OK" to return to the "File structure" screen.
- 18. Press "OK" on the "File structure" screen to return to the "BacLink Configuration" screen.
- 19. Press "Save" on this screen and return to the main menu, where your new configuration file should be listed.

Importing data files with BacLink

The procedure above for generating your BacLink configuration must only be performed once on your system. Once you have an EARS-Vet configuration, you can simply select it from the list, choose your corresponding data file, and convert the data into the WHONET data file structure using the procedure below in a few clicks. While the configuration process can be tedious due to the number of details that must be managed in describing your input files, the data export process itself is very simple. After you have successfully created a configuration for your data files, this is the only section of this document that you will need to repeat on an ongoing basis, unless your files or other details change which would necessitate a corresponding change to your configuration.

- 1. Choose your EARS-Vet configuration file from the list of options on the BacLink main screen.
- 2. Press the "Browse" button associated with the file name, highlighted in the image below.

•				_	
BacLink 2024			-		×
File Select language Help					
	original data file. v data file. Click on 'Begin conversion'. t appear on the list, choose 'New format'.				
File format	Boston General Hospital (SILAB)		New	format	
W:\WHONET.Source\BacLink\bin \x64\Debug\	Brigham and Women's Hospital EARS-Vet test laboratory		Edit	format	
EARS-Vet test laboratory- VITEK.cfg	CARG-VELTEST REUTING		Delet	e format	
File name	W:\WHONET.Source\BacLink\bin\x64\Debug\Da	ata*.bd	Br	owse	
New data file					
File name	W:\WHONET.Source\BacLink\bin\x64\Debug\Da	ata*.sqlite	Br	owse	
Table name	For Access files only				
File format	WHONET (SQLite)	~			
		Begin conversion		Exit	

3. Choose your input data file using the file browser and press OK to select it.

4. A default output file name will be provided in the "new data file" section once you select your data file above. You may choose another name or location for the output data or accept the default.

BacLink 2024			_		\times
File Select language Help					
	original data file. v data file. Click on 'Begin conversion'. vt appear on the list, choose 'New format'.				
File format	Boston General Hospital (SILAB)		Nev	v format	
W:\WHONET.Source\BacLink\bin \x64\Debug\	Brigham and Women's Hospital EARS-Vet test laboratory		Edi	t format	
EARS-Vet test laboratory- VITEK.cfg			Dele	te format	
File name	C:\WHONET\Data\test.txt		В	rowse	
New data file					
File name	C:\WHONETIData\FRA-TST-test.sqlite		В	rowse	
Table name	For Access files only				
File format	WHONET (SQLite)	~			
		Begin conversion		Exit	

- 5. Press "Begin conversion" near the bottom of the form.
- 6. The first three isolates will be shown sequentially so that you can make a brief visual inspection. Some fields will be translated into the WHONET code set, others will be copied as-is. The system should also recognize your antibiotics, which are shown in the lower window. If you do not see antibiotics on this screen, then the system may need to finish processing the file to request that you match these with the known WHONET antibiotics.

Field name	Local value	WHONET value
Identification number		
Location		
Department		
Specimen number		
Specimen date		
Specimen type	CECUM	cecum
Local specimen code	CECUM	CECUM
Isolate number		
Organism	ESCCOL	eco
Local organism code	ESCCOL	ESCCOL
Comment		
Data year		
Data representativeness		
Surveillance program		
Animal species	PIC	
Animal use	MEAT	mea
Markat oota aan	DOM	4
AMP_EM = 2	CHL_EM = 64	CIP_EM = 0.015
COL_EM = 1	CTX_EM = 0.25	CAZ_EM = 0.5
GEN_EM = 0.5	MEM_EM = 0.03	NAL_EM = 4
SMX_EM = 1024	TCY_EM = 64	TGC_EM = 0.25
TMP_EM = 32		

- 7. For each isolate, press "Next" once you have finished looking over the record.
- 8. After the third isolate, the remainder of the data file will be processed. You can monitor the conversion on the progress screen. Newly discovered codes will appear, as well as any conversion problems, such as dates with unrecognized formats, etc.
- 9. Once the conversion has completed, you will be presented with a dialog box allowing you to either continue back to the BacLink main screen, or you can view the entire database's contents if you choose "View database".
 - a. If there are undefined codes, please follow the standard BacLink documentation regarding code mapping found on the training page of our website.
 - b. If one of the date fields was incorrectly formatted, you can change this in the "Data fields" configuration area. This is also covered in the standard BacLink documentation.
- 10. If you have defined new codes or made any other modifications to the configuration you should rerun the BacLink conversion again using the updated configuration. Once the data appears correct, and no further changes are required, you may use the data files generated by BacLink with WHONET analyses and the EARS-Vet export.
- 11. For general BacLink questions not covered in this document, please refer to the documentation found on your computer at C:\WHONET\Documents\ or online at the following URL:
 - a. <u>https://whonet.org/training.html#bacLinkResources</u>