

WHONET

Resistance Profiles

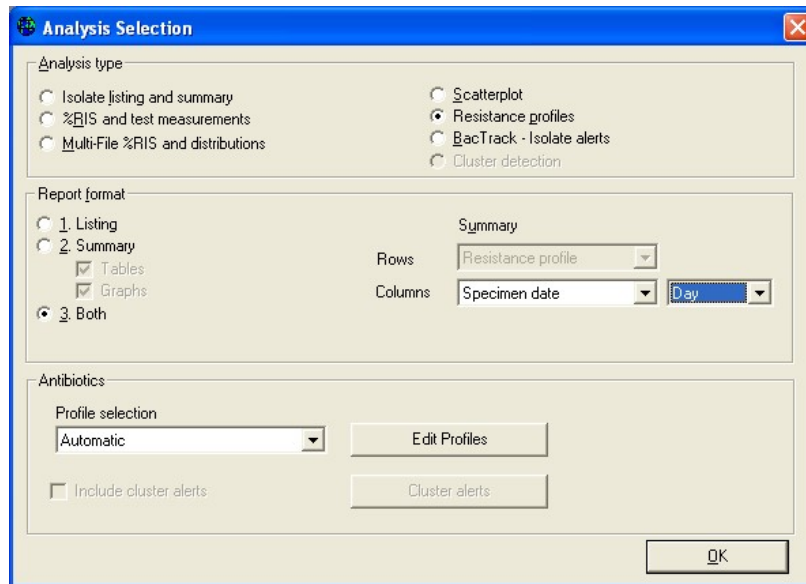


**WHO Collaborating Centre for
Surveillance of Antimicrobial Resistance**

Boston, July 2022

Part 1. Antimicrobial resistance profiles

The final analysis that covered in this tutorial is the study of multi-resistance patterns using the analysis “Resistance profiles”. Click on “Analysis type”, and select “Resistant profiles”. Because you are analyzing only one month of data, change the option “Month” in the middle of the screen to “Day”.

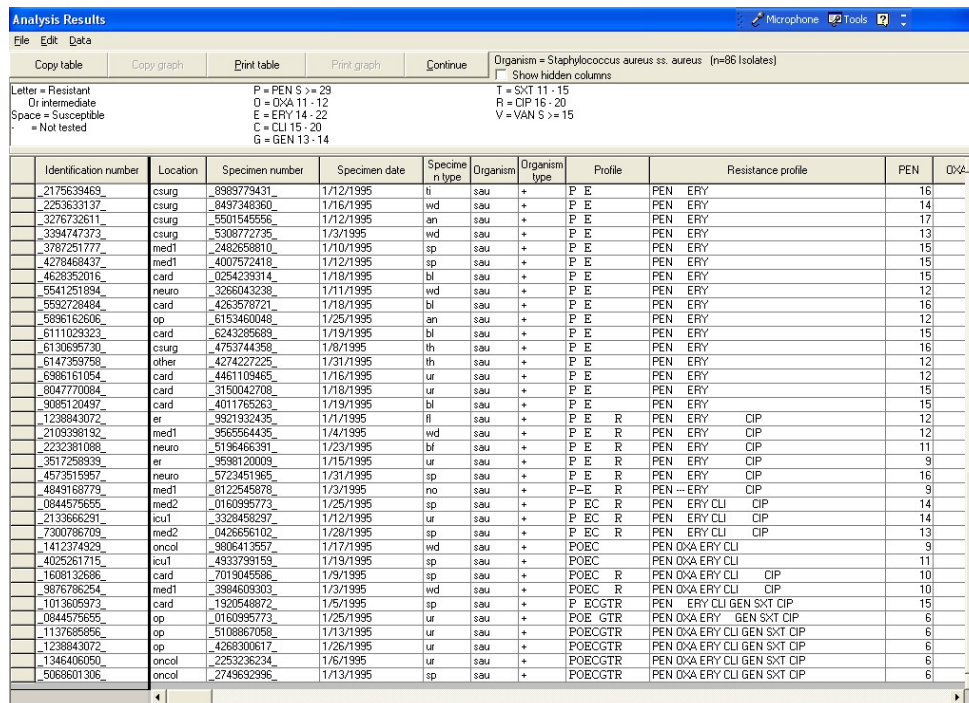


The "Analysis Selection" dialog box is shown. It has a blue title bar with the text "Analysis Selection" and a close button. The dialog is divided into several sections:

- Analysis type:** Contains radio buttons for "Isolate listing and summary", "%RIS and test measurements", "Multi-File %RIS and distributions", "Scatterplot", "Resistance profiles" (selected), "BacTrack - Isolate alerts", and "Cluster detection".
- Report format:** Contains radio buttons for "1. Listing", "2. Summary", and "3. Both" (selected). Below these are checkboxes for "Tables" and "Graphs". To the right, there are dropdown menus for "Rows" (set to "Resistance profile") and "Columns" (set to "Specimen date"), and a "Day" dropdown menu.
- Antibiotics:** Contains a "Profile selection" dropdown menu (set to "Automatic"), an "Edit Profiles" button, and a "Cluster alerts" button. There is also an unchecked checkbox for "Include cluster alerts".

An "OK" button is located at the bottom right of the dialog.

Then “OK” and “Begin analysis”. The following isolate listing shows the bottom portion of the output.



The "Analysis Results" window displays the output of the analysis. It has a menu bar with "File", "Edit", and "Data". Below the menu bar are buttons for "Copy table", "Copy graph", "Print table", "Print graph", and "Continue". The window title is "Organism = Staphylococcus aureus ss. aureus (n=86 Isolates)".

Legend:

- Letter = Resistant
- Or intermediate
- Space = Susceptible
- = Not tested

Summary statistics:

- P = PEN S >= 29
- D = OXA 11 - 12
- E = ERY 14 - 22
- C = CLI 15 - 20
- G = GEN 13 - 14
- T = SXT 11 - 15
- R = CIP 16 - 20
- V = VAN S >= 15

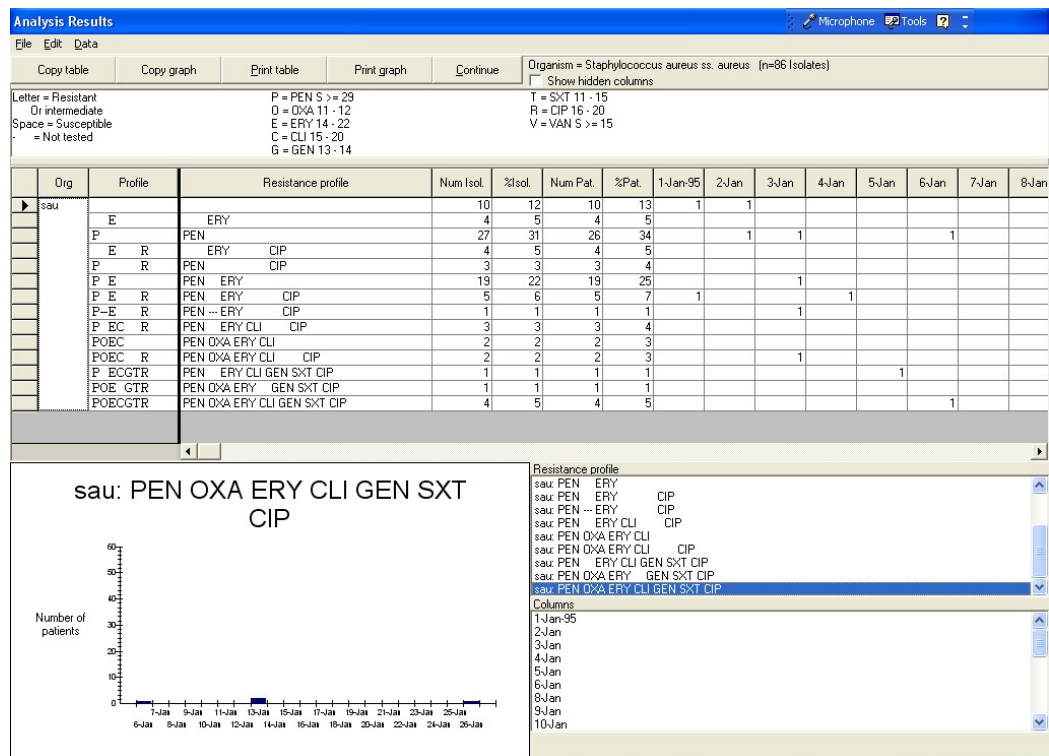
Table columns: Identification number, Location, Specimen number, Specimen date, Specimen type, Organism, Organism type, Profile, Resistance profile, PEN, OXA.

Identification number	Location	Specimen number	Specimen date	Specimen type	Organism	Organism type	Profile	Resistance profile	PEN	OXA
2175639469	csurg	9989779431	1/12/1995	ti	sau	+	P E	PEN ERY	16	
2253633137	csurg	9497348360	1/16/1995	wd	sau	+	P E	PEN ERY	14	
3276732611	csurg	5501545556	1/12/1995	an	sau	+	P E	PEN ERY	17	
3394747373	csurg	5308772735	1/3/1995	wd	sau	+	P E	PEN ERY	13	
3787251777	med1	2482658810	1/10/1995	sp	sau	+	P E	PEN ERY	15	
4278468437	med1	4007572418	1/12/1995	sp	sau	+	P E	PEN ERY	15	
4628352016	card	0254239314	1/18/1995	bl	sau	+	P E	PEN ERY	15	
5541251894	neuro	3266043238	1/11/1995	wd	sau	+	P E	PEN ERY	12	
5552729484	card	4263578721	1/18/1995	bl	sau	+	P E	PEN ERY	16	
588616206	op	6153460040	1/25/1995	an	sau	+	P E	PEN ERY	12	
6111029323	card	6243285689	1/19/1995	bl	sau	+	P E	PEN ERY	15	
6130695730	csurg	4753744358	1/8/1995	th	sau	+	P E	PEN ERY	16	
6147359758	other	4274227225	1/31/1995	th	sau	+	P E	PEN ERY	12	
6986161054	card	4461109465	1/16/1995	ur	sau	+	P E	PEN ERY	12	
8047770084	card	3150042708	1/18/1995	ur	sau	+	P E	PEN ERY	15	
9085120497	card	4011765263	1/19/1995	bl	sau	+	P E	PEN ERY	15	
1238843072	er	9921932435	1/1/1995	fl	sau	+	P E R	PEN ERY CIP	12	
2109381132	med1	966564435	1/4/1995	wd	sau	+	P E R	PEN ERY CIP	12	
2232381080	neuro	5196463391	1/23/1995	bl	sau	+	P E R	PEN ERY CIP	11	
3517258939	er	9598120009	1/15/1995	ur	sau	+	P E R	PEN ERY CIP	9	
4573515957	neuro	5723451965	1/31/1995	sp	sau	+	P E R	PEN ERY CIP	16	
4849168779	med1	8122545878	1/3/1995	no	sau	+	P E R	PEN ERY CIP	9	
0844575655	med2	0160995773	1/25/1995	sp	sau	+	P EC R	PEN ERY CLI CIP	14	
2133666291	icu1	3328458297	1/12/1995	ur	sau	+	P EC R	PEN ERY CLI CIP	14	
7300786709	med2	0426656102	1/28/1995	sp	sau	+	P EC R	PEN ERY CLI CIP	13	
1412374529	oncol	9806413557	1/17/1995	wd	sau	+	POEC	PEN OXA ERY CLI	9	
4025261715	icu1	4933799159	1/19/1995	sp	sau	+	POEC	PEN OXA ERY CLI	11	
1608132686	card	7018046586	1/3/1995	sp	sau	+	POEC R	PEN OXA ERY CLI CIP	10	
9876786254	med1	3984609303	1/3/1995	wd	sau	+	POEC R	PEN OXA ERY CLI CIP	10	
1013605973	card	1320548872	1/5/1995	sp	sau	+	P ECGR	PEN ERY CLI GEN SXT CIP	15	
0844575655	op	0160995773	1/25/1995	ur	sau	+	POE GTR	PEN OXA ERY GEN SXT CIP	6	
1137685856	op	5108867058	1/13/1995	ur	sau	+	POECGTR	PEN OXA ERY CLI GEN SXT CIP	6	
1238843072	op	4268300617	1/26/1995	ur	sau	+	POECGTR	PEN OXA ERY CLI GEN SXT CIP	6	
1346406050	oncol	2253236234	1/6/1995	ur	sau	+	POECGTR	PEN OXA ERY CLI GEN SXT CIP	6	
5068601306	oncol	2749632996	1/13/1995	sp	sau	+	POECGTR	PEN OXA ERY CLI GEN SXT CIP	6	

This listing is very similar to the one you did in Data analysis 1 with the analysis option “Isolate listing and summary”. But there are two additional columns: “Profile” (using one-letter antibiotic codes) and “Resistance profile” (using three-letter antibiotic codes). These columns indicate the multi-resistance pattern of the isolates. The profiles indicate the drugs to which the isolate is either resistant or intermediate (*i.e.* non-susceptible). For example “PE” = “PEN ERY” = isolates non-susceptible to penicillin and erythromycin, but susceptible to the other drugs requested.

Completely susceptible isolates appear at the top of the listing, followed by isolates non-susceptible to one drug, then to two drugs, *etc.* Multi-resistant strains appear at the bottom of the listing. This analysis thus categorizes all of the observed isolates according to resistance phenotype. If a patient has multiple isolates, you can see whether the isolates have the same resistance phenotype, or whether the phenotype changes over time, for example accumulating mutations, resistance genes, and plasmids. By examining the dates and the room numbers, you may also detect possible outbreaks of certain strains of *S. aureus* as defined by their multi-resistance pattern.

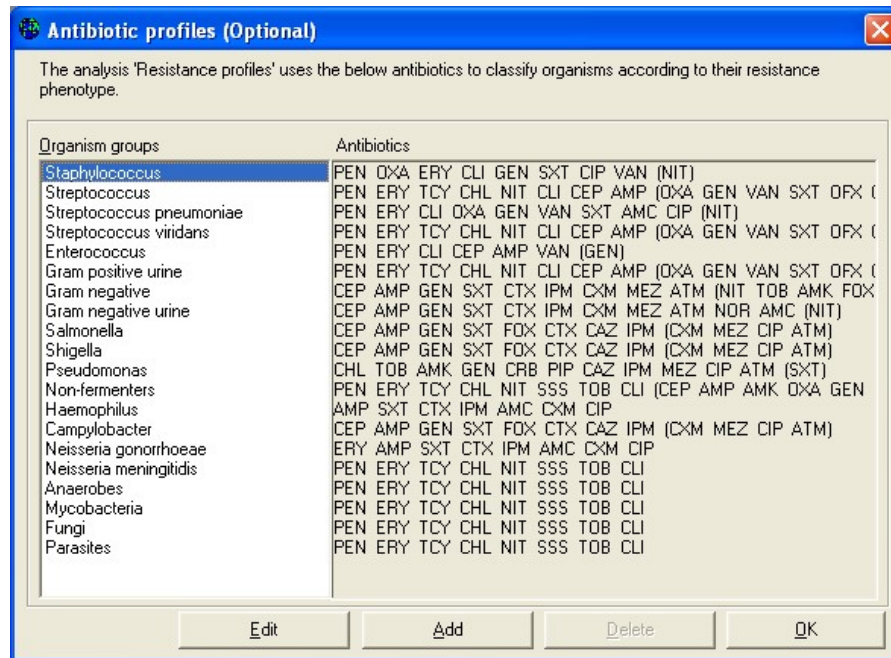
Click on “Continue” to see a summary of this list.



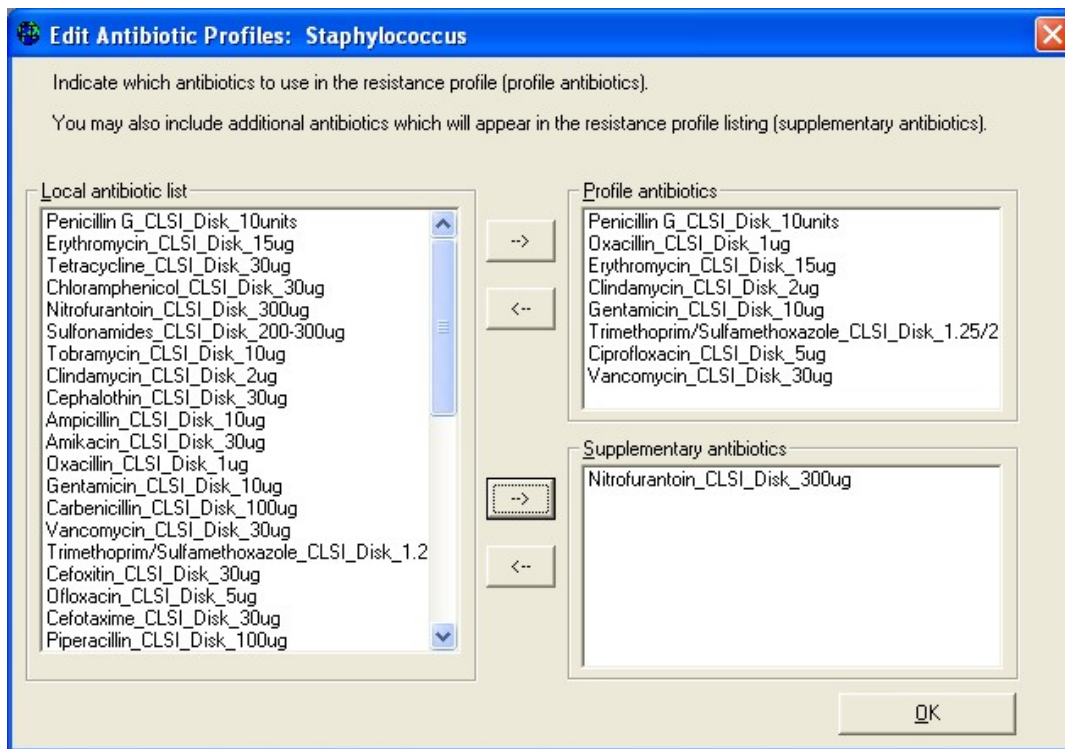
This output summarizes findings for all resistance phenotypes observed in the database. In this sample database, you will see that the most common phenotype is pan-susceptible (susceptible to all drugs requested) followed by “PEN” (non-susceptible to penicillin). Among MRSA isolates, the most common phenotype is “PEN OXA ERY CLI GEN SXT CIP”, in other words non-susceptible to all drugs requested with the exception of vancomycin. The day-by-day distribution of this MRSA phenotype is shown in the graph.

Click on “Continue” to return to the main menu.

In this analysis, WHONET did not analyze results for all of the drugs tested against *S. aureus*. Instead, WHONET used the drugs indicated in the laboratory configuration. To see and modify the drugs used for this analysis, click on “Analysis type” and “Edit profiles”.



This list indicates which antimicrobials will be used, by default, for each organism group. For the item “Staphylococcus”, click on “Edit”.



On this screen, you can select the drugs to use for the resistance profiles. Drugs in the box “Profile antibiotics” will be used to define the resistance phenotype. Drugs appearing under “Supplementary antibiotics” will appear in the line-listing, but are not used to define the resistance phenotype.

Note: Any modifications that you make to the profile antibiotics while you are in the Data analysis area of WHONET will be forgotten as soon as you leave WHONET. Any edits that you do here are temporary. To save the changes permanently for future use, you will need to use the WHONET configuration program. To do this, you will need to go back to the main WHONET screen and select “Modify laboratory” and “Antibiotics”. You will see a button called “Profiles”. After you change the profile antibiotics, click on “Save” to save them into your laboratory configuration.