

Screenshots of the on-line Result form Serotyping EURL-Salmonella Proficiency Test Typing 2023

EURL-Salmonella EURL-Salmonella Proficiency Test Typing 2023 Result form Serotyping							
LABORATORY INFORMATION							
Laboratory code PT 2023							
Name contact person							
E-mail address contact person							
Optional second E-mail address							
Name Institute (laboratory)							
City							
Country	Country:						
Date of arrival of the parcel in your laboratory Was your parcel damaged at arrival?	dd/mm/yyyy						
Medium used for sub-culturing the strains							
QUESTIONS SEROTYPING							
What was the frequency of serotyping of Salmonella at your laboratory in 2023? How many Salmonella strains did your laboratory (approximately) serotype in	Once a week Twice a week Thrice a week Monthly Other						



Is your laboratory accredited for Salmonella serotyping? The strains in this study were serotyped by:	Yes No Own laboratory Other laboratory, namely (strain numbers):			
The determination of the <i>Salmonella</i> serovars in this study was done by (multiple answers possible):	✓ classical serology ✓ molecular method(s), please specify the tool(s) used:			
What kind of sera do you use?	Biorad Own preparation Pro-Lab Sifin Statens Serum Institut (SSI) Other, namely:			
How was this molecular method validated?				
QUESTIONS ON CONFIRMATION BY BIOCHEMICAL TESTS/PCR				
Did you use any biochemical tests or PCR to confirm serotyping results?	✓ Yes No			
Biochemical and/or PCR tests used for strain S1:	None PCR to confirm S. Typhimurium/monophasic variant Dulcitol Malonate ONPG Salicine Sorbitol Other(s), namely:			

(same question for strains S1 - S21)

Biochemical and/or PCR tests used fo strain S21:	PCR to confirm S. Typhimurium/monophasic variant Dulcitol Malonate ONPG Salicine Sorbitol Other(s), namely:	
Reference(s) for the PCR method(s)	Barco et al., 2011 EFSA Journal, 2010 Maurischat et al., 2015 Prendergast et al., 2013 Tennant et al., 2010 Other (e.g. commercial test kit), namely:	



Starting date of serotyping did/mm/yyyy did/mwyyyy did/mm/yyyy did/mm/yyyy did/mwyyyy did/mwyyyyyyy did/mwyyyyyyyyyyyyyyyyyyyyyyyyyyyyyyyyyyyy			TEST RESULT	S SEROTYPII	NG
Please report only those results, on which the identification of the serovar names is based. Do not take over the notation "as it is" from the WKLM scheme, e.g. 1,4,[5],12:g.m,s.[1,2] for Hato, but only report what was tested and detected (see example 8). Be sure to give under "serovar name" the complete serovar name you assign to the strain, even if this means that you have to repeat the O/H antigens as found (see examples 3-6). Examples of expected reporting: Oantigens Hantigens (phase 1) Hantigens (phase 2) Serovar name* 9,12 g.m.	Starting d	late of serotyping	dd/mm/y	yyy 🛗 dd/mm/yyy	у
he notation "as it is" from the WKLM scheme, e.g. 1,4,[5],12 g.m.s.;[1,2] for Hato, but only report what was ested and detected (see example 8). de sure to give under "serovar name" the complete serovar name you assign to the strain, even if this means that you have to repeat the OH antigens as found (see examples 3-6). Examples of expected reporting: D-antigens H-antigens (phase 1) H-antigens (phase 2). Serovar name* 2,12 g.m. 2 Entertidis 1,12 g.m. 4,5,12 l. 1,15 g.m. 4,5,12 l. 1,15 g.m. 4,5,12 l. 1,15 g.m. 4,2 g.m. 1,12,27 l.y. e.n.215 Brandenburg H-antigens (rebase 1) H-antigens (phase 2). Serovar name* Please report the serovar name without indicating "S." or "Salmonellar", to facilitate the overall evaluation of all participants' results. Nas once that for technical reasons it regretably is not possible to use special fonts like "italic" or subscript" in the submission of your data. Extrain O-Antigens H-antigens (phase 1) H-antigens (phase 2) Serovar name 11 G. Antigens H-antigens (phase 1) H-antigens (phase 2) Serovar name 13 G. Antigens H-antigens (phase 1) H-antigens (phase 2) Serovar name 14 g.m. 15	inishing	date of serotyping	dd/mm/y	yyy	у
D-antigens H-antigens (phase 1) H-antigens (phase 2) Serovar name* 1/2 g,m - Ententidis 1/2 i 2 Typhimunum 1/3 i - 4.5 1.5 1 i - 4.5 1.5 1/2 g,t - 42 1.5 1/2 g,t - 42 1.5 1/2 g,ms - 4.5 1/2 g,ms - 4.5	he notation ested and Be sure to	on "as it is" from the d detected (see exa o give under "serov	e WKLM scheme, e.g. 1, ample 8). ar name" the complete s	4,[5],12:g,m,s,:[1,2] for serovar name you assig	Hato, but only report what was in to the strain, even if this
1,1	Example	es of expected r	eporting:		
4,512 i			se 1) H-antigens (phase		
4,5,12 i			2		
1,5 6,71,5 42,gt. 1,2 gt 42,gt. 1,1,2 r. v. e,n,z15 Brandenburg Hato PPlease report the serovar name without indicating "S," or "Salmonella", to facilitate the overall evaluation of all participants' results. Also note that for technical reasons it regrettably is not possible to use special fonts like "Italic" or 'subscript" in the submission of your data. Strain O-Antigens H-antigens (phase 1) H-antigens (phase 2) Serovar name 1		i	-		
42 g.t. Brandenburg 44 g.m.s - Hato Please report the serovar name without indicating "S." or "Salmonella", to facilitate the overall evaluation of all participants' results. Also note that for technical reasons it regrettably is not possible to use special fonts like "italic" or subscript" in the submission of your data. Strain O-Antigens H-antigens (phase 1) H-antigens (phase 2) Serovar name 51		i	-		
4,12,27 Iv e,n,z15 Brandenburg Hato Please report the serovar name without indicating "S." or "Salmonella", to facilitate the overall evaluation of all participants' results. Also note that for technical reasons it regrettably is not possible to use special fonts like "italic" or "subscript" in the submission of your data. Strain O-Antigens H-antigens (phase 1) H-antigens (phase 2) Serovar name 11		- a t	1,5		
Please report the serovar name without indicating "S." or "Salmonella", to facilitate the overall evaluation of all participants' results. Also note that for technical reasons it regrettably is not possible to use special fonts like "italic" or subscript" in the submission of your data. Strain O-Antigens H-antigens (phase 1) H-antigens (phase 2) Serovar name 11			e.n. <i>z</i> 15		
Please report the serovar name without indicating "S." or "Salmonella", to facilitate the overall evaluation of all participants' results. Also note that for technical reasons it regrettably is not possible to use special fonts like "italic" or 'svabscript" in the submission of your data. Strain O-Antigens H-antigens (phase 1) H-antigens (phase 2) Serovar name 11			-		
	train		,	1) H-antigens (phase	e 2) Serovar name
33	51				
54	52				
55 56 57 58 59 510 511 512 513 514 515	53				
56	54				
57	55				
58 59 511 512 513 514	56				
59	57				
510	58				
511	59				
512	510				
513					
514					
515					



S17				
518				
S19				
S20				
521				
OPTIONAL T	EST RESULTS MLVA			
Would you like to submit MLVA results?	✓ Yes○ No			
Strain code:				
Allelic profile:				
Format used:	SENTR7-SENTR5-SENTR6-SENTR4-SE-3 STTR9-STTR5-STTR6-STTR10-STTR3			
	O Other			
(5	strains)			
Strain code:				
Allelic profile:				
Format used:	SENTR7-SENTR5-SENTR6-SENTR4-SE-3 STTR9-STTR5-STTR6-STTR10-STTR3			
	Other Other			
F	FINALLY			
Any comments				
Any comments				
Name of person(s) carrying out the typing				
Name of the person in charge				
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The EURL-Salmonella handles your personal data with the utmost care. Personal data is protected under the General Data Protection Regulation (GDPR). Your data will be encrypted and treated anonymously. Original data is only accessible for EURL-Salmonella staff involved in this project.				
SAVE YOUR PROGRESS RESUM				
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