

# **GASDN: Resulting and Archiving Group A Strep Results**

## **PURPOSE**

This procedure provides instructions for resulting and filing GASDN group A strep results in LIS.

# **POLICY STATEMENT**

Results will be available within 24 h; refer to the <u>Laboratory Services</u> web page

#### **ABBREVIATIONS**

- ASR: analyte-specific reagent
- Ct: crossing threshold
- FDA: US Food and Drug Administration
- IC: internal control
- GASD: group A strep worksheet
- GASDN: Group A Strep Detection (LIS code)
- GASCT: group A strep Ct

## **DOCUMENTATION/RECORDS**

GASR: group A strep resultLIS: laboratory information system

GASIC: group A strep internal control

- PCR: polymerase chain reaction
- SAS: Specimen description
- STYP: specimen type code
- WO: worksheet

- GAS Segment Report
- LIS Incomplete worksheet, SmarTerm Function WO, options #1

PROCEDURE A: Follow the activities below for reporting and archiving patient results

# **Reporting Patient Results**

Activity	Step	Action	Action								
Export results	1	Export patient test results from the Simplexa instrument to LIS; see MB 8.05 procedure H									
	2	Reporting	activit	ies are perf	ormed in the Molecular Office						
Computer		Enter results in Sunquest Result Entry									
Entry			Step	Prompt	Action/Entry						
Molecular	2		а		Click on the Sunquest icon to log on						
Office	3		b	User	СЕхххххх						
s			С	Password	XXXXXXX						
			d	Location	R						
			е		Click OK						
	4	Select Result Entry from Menu options									
		Figure 1: Int	terface	Configuratio	on: Select <b>SIM</b> from drop down box						
SIM	5			Resultin Inter Configurat							
	6	Click on the button located in the lower left corner to populate the transmitted results									



Activity	Step	Action									
	7		If the page says "Waiting for cups", the results were not successfully transmitted or the results page was accessed too quickly before the transmission was completed								
STYP code or missing code	8	STYP codes: This specimen was grayed out and not checked for release because there was a problem with the specimen type code.									
Review		Figure 2: Review messages located on the top and results									
Messages		Apply Defaults	Speci	imen messages Failed o	up criteria	Tes	t messages	INVALID RES A calculation allowed.		r this test and r	nodification is not
Result Entry		Status         Specim           *,L         T80951           T81141         T81141	SIM2	4 2 30.5 5	8PPCT 0.0	внст	BICCT 27.5	BPDNA POS-ASRS	BPPDN NEG-ASRS		Display     Failure     Verify
		*,L T80913 *,L T81296 *,L T81557 *,L T80851	SIM2	6         28.9           7         29.3           8         29.3           9         30.5	0.0		27.5 27.5 27.7 27.5	POS-ASRS POS-ASRS POS-ASRS	NEG-ASRS NEG-ASRS NEG-ASRS		Techn Delta IRange
	9	Solution: STYF SmarTerm MEN		s can be correc	ted or mis	sing co	des can	be add	ed in <i>Sui</i>	nquest O	rder Entry or
		5	Step	Prompt	Action/Ent	ry					
SmarTerm			а	FUNCTION	MEM						
-			b	WORKSHEET	GASD						
			С	TEST	GASDN Add	missing	codo: E	56021			
			d	ACC. NO.	<ul> <li>Rep</li> </ul>	lacing ST	YP code	: M-F669		r modify)	_
			e	SAS	Entering a SAS1	missing	code or	replacin	g an exist	ing code	
Duplicate results		Figure 3: If a rur results	-	ported more th			the dup	olicate r	esults ar	nd releas	e the checked
واللو	10		r	Apply Defaults	Specimen m	essages	Preproces Specimen				
Status         Specimen         Method         Cup         Release         BPCT         I           F66931         SIM3         1         I         0.0         0           F66931         SIM3         2         I         0.0         0											
	11	Click Save	but	ton located on	the lower	left cori	ner				
		Figure 4: Dupli	cate S	pecimens mess	sage						
		_			-						
Duplicate Specimens       ×         You have selected to release one or more specimens with results in multiple cups on the instrument.         Cups selected for release are marked with an asterisk:         Accession											
	12	F6693	F66931 1*, 2								
		Do you	want to r	elease the selected cu	ps with duplicate	speciment	?				
				Do not r	elease duplicates		elease All		cel		
13 Click Release All											



Activity	Step	Action						
	14	Figure 5: Verify Release to Lab; click Accept	Prelim					
	15	lf	Then					
No Problems	16	<ul> <li>Specimen box reads <i>Preprocessing passed</i> with no further messages</li> <li>Test box has no messages</li> <li>Sample results are acceptable</li> </ul>	Click <b>Save</b> and then <b>Accept</b> (Fig. 5)					
<ul> <li>Record Control results from GAS Segment Report on GASD worksheet</li> <li>Record Ct values in GASCT and GASIC columns</li> <li>Interpretation: Valid or Invalid</li> </ul>								
Archive	18	used during testing and GAS Segment Report						
	19	Place report in the GAS result log book in Molecular Office						

# **PROCEDURE C:** Follow the activities below for result manual entry in SmarTerm Function MEM **Inhibitory / Abnormal Results**

Activity	Step	Action				
		Entering re	ring result codes manually; refer to table 1			
	1		Step	Prompt	Action/Entry	
Manual			а	FUNCTION	MEM	
Entry			b	WORKSHEET	skip for manual entry without calculations	
Abnormal			С	TEST	GASDN	
results			d	ACC. NO.	F66931	
lesants			е	SAS1	THR	
SmarTerm			f	GASCT	skip	
			g	GASIC	skip	
			h	GASR	UNR-ASRSG	
			i		Click Phoned Result	
				j	NAME	Enter name and last initial
			k		Click OK, date and time stamp will default	
			Ι	ACCEPT	A	



# **Table 1: Sunquest Result Codes**

Result	Sunquest code	Interpretation
Pos Interface	POS-ASRS Interface POS-L20-ASRS-L20	<ol> <li>Positive</li> <li>Analyte Specific Reagent: This test was developed and its performance determined by Children's Hospitals and Clinics of MN. It has not been cleared or approved by the FDA</li> </ol>
Neg Interface	NEG-ASRS Interface NEG-L20-ASRS-L20	<ol> <li>Negative</li> <li>Analyte Specific Reagent: This test was developed and its performance determined by Children's Hospitals and Clinics of MN. It has not been cleared or approved by the FDA</li> </ol>
Unresolved Results	<b>UNR - ASRSG</b> Function MEM	<ol> <li>Unresolved: This sample is inhibitory to amplification and the results are inconclusive. Consider repeat collection if clinically indicated.</li> <li>ASR / FDA statement</li> <li>Phoned Result Phone Troub [free text name]</li> </ol>
Amplification curve abnormal IC not inhibited	<b>EQ – ASRSG</b> Function MEM	<ol> <li>Equivocal result, suggest repeat specimen</li> <li>ASR / FDA statement</li> <li>Phoned Result [free text name]</li> </ol>
Corrected report	Function MEM	<ol> <li>Modify results in function MEM by tying M hyphen accession number: M-T54966</li> <li>Enter correct numerical result in GASCT AND GASIC fields. Computer will automatically put in a correction statement</li> <li>In field GASR, click Phoned Result Phoned Result [free text name]</li> <li>Refer to Procedure D</li> </ol>

# **PROCEDURE D:** Follow the activities below for modifying results in SmarTerm Function MEM **Corrected Report**

Activity	Step	Action			
	1	Entering I	niscella	ineous codes; r	efer to table 1
Manual			Step	Prompt	Action/Entry
Entry			b	FUNCTION	MEM
Modifying			С	WORKSHEET	Enter GASD (required for worksheet calculations)
results			d	TEST	GASDN
			е	ACC. NO.	M – F66931
SmarTerm			f	SAS1	THR
			g	GASCT	<ul><li>0 – enter correct Ct value below previous value</li><li>34 <i>Correction statement will automatically default</i></li></ul>
			h	GASIC	28 27.5 Correction statement will automatically default
			j	GASR	POS-L20-ASRS-L20; automatic worksheet calculation
					Click Phoned Result
			k	NAME	SmarTerm Called to (Er/ter name) Ruth H, RN OK Cancel
			Ι		Click OK, date and time stamp will default
			0	ACCEPT	А



# REFERENCES

- 1. Laboratory for Windows User's Guide, UD-1599/00-12-GL, Sunquest Information Systems
- 2. CAP Microbiology Checklist, Section Department: Molecular Diagnostics, MIC.64988, 04.21.2014, College of American pathologists, 325 Waukegan Rd, Northfield, IL 60093

#### **Historical Record**

Version Written/Revised by:		Effective Date:	Summary of Revisions		
1	Pat Ackerman	07.20.16	Initial Version		
2	P. Ackerman	07.30.2016	LIS interface protocols added; reformatted for CMS upload		