Pregnancy Evaluation Report **RPL Standard Panel 1.3**

PATIENT NAME: Example, Patient DATE OF REPORT:

CLINICIAN: Clinician Example 11/04/2018

PRIVATE AND CONFIDENTIAL



300 Trade Center, Suite 6540 Woburn, MA 01801 USA tel: 800.667.8893 USA only tel: 781.937.8893 fax: 781.935.3068 CLIA#: 22D0884531 CLIA#: 22D0884531



							Rest	ULTS O	VERVIEW ^{**}
Category		Теятя		Units	Range	In Range	Border- Line	Out of Range	Note
	Endocrine	Thyroid stimulating hor	mone (TSH)	mlU/L	0.40-4.50	1.52			
22	Coagulation	Lupus anticoagulant (LA)	N/A	negative	negative			
	Immunology	Anticardiolipin (aCL)	lgG	GLP	0-14	7			
∇		Anti-β ₂ Glycoprotein I (aBGPI)	IgM	MLP	0-14	4			
			lgG	SGU	0-20			50	positive
			IgM	SMU	0-20			31	positive
0	Consting	Maternal Karyotype (blood)	N/A	46, XX			Х	46,XX,t[3;4;14][q21;p14;q22.3] see comments
ň	Genetics	Chromosomal Microarray, POC ClariSure [®] Oligo-SNP		N/A	46,XX or 46,XY	46,XX			no maternal cell contamination

COMMENTS:

THIS RESULTS OVERVIEW PAGE IS NOT INTENDED TO PROVIDE COMPLETE INFORMATION. PLEASE REVIEW THE INDIVIDUAL TEST RESULTS THAT FOLLOW.

- 1. Out of range results obtained for anti-B2 glycoprotein I IgG and IgM. Please review full results in detail.
- 2. An abnormal female chromosome complement with an apparently balanced complex rearrangement involving chromosomes 3, 4, and 14 is seen. Please review full results in detail.
- 3. POC: normal female microarray result. No maternal cell contamination. Please review full results in detail.

Please contact ReproSource Client Services with any questions tel: 800 667 8893 option 1 email: clientservices@reprosource.com web: www.reprosource.com

CURRENT TESTING GUIDELINES FOR RECURRENT PREGNANCY LOSS (RPL)

REFERENCE SOURCES

ACOG: American College of Obstetrics & Gynecology ASRM: American Society of Reproductive Medicine ACG: Antiphospholipid Consensus Group

Opinion: Expert Opinion Papers & Reviews

Recurrent Pregnancy Loss Testing		b	Σ	(5	RE	nion
Criteria Tests			ASR	ACG	ESH	Opi
	Thyroid Stimulating Hormone		Yes	Yes	Yes	Yes
2 or 3 or more	Lupus anticoagulant (LA)	Yes	Yes	Yes	Yes	Yes
intrauterine	Anticardiolipin (aCL) IgG IgM	Yes	Yes	Yes	Yes	Yes
pregnancy losses	Anti- β_2 GPI IgG IgM	Yes	Yes	Yes	Yes	Yes
	Karyotype X2 (both parents)	Yes	Yes		Yes	Yes

References

- 1. ACOG Practice Bulletin 150 (2015) Early Pregnancy Loss
- 2. ACOG Practice Bulletin 132 (2012) Antiphospholipid Syndrome
- 3. ASRM Committee Opinion (2013) Evaluation and Treatment of RPL
- 4. ACG International Consensus Statement (2006) Antiphospholipid Syndrome (APS)Recommendations
- 5. ESHRE Practice Guidelines (2017) Recurrent Pregnancy Loss

ESHRE: European Society of Human Reproduction & Embryology

- Stephenson M. Clinical Obstetrics & Gynecology (2007) Evaluation and Management of Recurrent Early Pregnancy Loss
- 7. Kutteh W. Seminars in Reproductive Medicine (2015) Novel Strategies for the Management of Recurrent Pregnancy Loss

ReproSource®

nt:Example, Patient	DOB: 01/03	3/1990	Collection Date:	11/02/2018	Specimen Numbe	r: T0001390	
TEST	U	NITS	NORMAL RANGE	RESU	JLT	COMMENT	
RPL Standard Panel 1.3 Q8A Lupus Anti Coagulant			Negative	Nega	tive	Negative	
	The determin Thromboplas (KCT). If any plasma are pe after the mixi to confirm the plasma contai	ation of l tin Time of the the rformed ng studie e presenc ining anti	upus anticoagula (APTT), Dilute F aree tests is prolor (to correct for the es any of the three e of a `lupus inhi coagulants invali	nt in plasma incl Russell Viper Ve nged over a norn e presence of any e tests is still prol bitor` (antiphosp idates testing wh	udes the followin nom Test (dRVV nal control, `mixii 7 factor deficiency longed, a confirm holipid antibodie ich may yield fals	g tests: Activated Par T) and Kaolin Clottin ng studies` using norm y, inhibitors, etc.) If atory test is performed s). Additionally, patie se positive results.	tia g [na] d
TSH	m	IU/L	0.40 - 4.50	1.5	2	Normal	
	Reference Ra	nge abov	e is for adults age	ed 20 years or gr	eater.		
	Pregnancy Ra First trimester Second trimest Third trimester	anges: r 0.26-2. ster 0.55- er 0.43-2	66 2.73 2.91				
Anticardiolipin IgG	(GPL	<10	7		Negative	
Anticardiolipin IgM	Ν	MPL	<10	4		Negative	
Anti-Beta 2 glycoprotein	I-IgG S	SGU	<=20	50)	Positive	
Anti-Beta 2 glycoprotein	I-IgM S	SMU	<=20	31		Positive	
Chromosome Analysis Q8A							
	1	N/A	2	46,XX,t(3;4;14)(q21;p14;q22.3)	See Interpretati	or
	Karytoype: 40	6,XX,t(3;	;4;14)(q21;p14;q2	22.3)			
	Cytogenetic I	Reference	e : CB-18-000073	\$			
	Test Setup Da	ate: 10/05	5/2018				
	Test Complet	urce: Dar	: 10/05/2018				
	Clinical Histo	ory:Not p	rovided				
	Metaphases C	Counted:2	20 Analyzed:5 $d_{0} > -650$	Karyotyped:3			

Pregnancy Evaluation Report – RPL Standard Panel 1.3 cont.

i tient : Example, Patient	DOB: 01/03/1990	Collection Date: 11/02/	2018 Specimen Nu	mber: T0001390	Page: 4 of 6
TEST	UNITS	NORMAL RANGE	RESULT	COMME	NT
	Interpretation and Co Abnormal female kar lement with an appar seen. The rearrangen translocated to 4p, th translocated to 3q. Ba reproductive history of That risk is considera translocations as seer for which this study v abnormal liveborn ha in Madan K. Am J M Genetic counseling is to determine their car couple in which a car unbalanced outcomes	omments yotype, high resolution of ently balanced complex nent is a three way recipr ne 4p terminal region translation (e.g., pregnancy loss, inf ably increased with regar n here. This rearrangeme was indicated. A general is been estimated. A revi led Genet. 2012;58A:947 s recommended as is cyt rrier status. Prenatal diag rrier individual is determ s would likely be clear i	G-banding. An abnorma rearrangement involving rocal translocation with nslocated to 14q and the ocations confer a relativ fertility, chromosomally rd to complex rearrangen nt is very likely the etio risk estimate of 48.3% few of complex balanced 7-963. togenetic analysis of the gnosis is recommended i ined. While the distinct n a prenatal G-banded c	al female chromoso g chromosomes 3, 4 the 3q terminal regio e 14q terminal regio rely high risk for ad unbalanced newbo ments, including the logic factor in the i for pregnancy loss d rearrangements ca patient's first-degree n future pregnancie ion between balanc hromosome analysi	me comp- 4, and 14 is on n verse rn). ree way nfertility and 18.4% for in be found ee relatives es for any ed and is,
	it is recommended th such studies.	at oligo-SNP microarray	analysis be considered	in conjunction with	ı any
	Signature Electronic Signature	on File			
	Steven A. Schonberg Technical Director, C	, Ph.D., FACMG Cytogenetics and Genom	ics, 703-802-7156		
Chromosomal Microar	тау,POC ^{Q8M}		6 D I	0.14	
			See Below	See Interp	retation
	CY IUGENETIC RE	30L1S			
	Test Setup Date: 10/)5/2018			
	Test Completion Date	e: 10/16/2018			
	Specimen Source: M	embrane / Villi Fetal Ti	\$\$11A		
	Clinical History:IUE	D at 22 weeks history o	forior SAB v2		
		SULT.	I phot SAD X2		
	NORMAL FEMALE	SOLT. SMICROARRAY RESI	ШТ		
	$ISCN$ arr $(1-22 X)x^2$				

Pregnancy Evaluation Report – RPL Standard Panel 1.3 cont.

		Pregnancy Evalu	ation Report – RPL Stan	dard Panel 1.3 cont.		
P	atient: Example, Patient	DOB: 01/03/1990	Collection Date: 11/02/2	2018 Specimen N	umber: T0001390	Page: 5 of 6
	TEST	UNITS	NORMAL RANGE	RESULT	COMMEN	JT

INTERPRETATION and COMMENTS:

This array assay did not reveal a genomic imbalance. Maternal cell contamination is common in products of conception (POC). Therefore, a normal female result should be interpreted with caution, as it may not represent the fetus. Couples experiencing recurrent spontaneous abortions are advised to have chromosome analyses to rule out the possibility of a balanced rearrangement carrier (Test code 14596X). For more information, please call Genetics Client Services at 1-866-GENE-INFO.

ASSAY INFORMATION: Thresholds for genome-wide screening are set at >2 Mb for gains, >1 Mb for losses, and 10 Mb for segments of homozygosity. These may be lower in cytogenetic relevant regions. This assay will not detect genomic gains or losses that are smaller than the resolution of the method. The additional ability to detect significant regions of allelic homozygosity (ROH) enhances the diagnostic value of this assay. This assay will detect dosage abnormalities only for genomic sequences represented in the array and will not detect balanced rearrangements (e.g., translocations, inversions). Copy number variations (CNV) with no phenotypic consequences are relatively common; CNVs identified as known variants in the general population and reported in the publicly available database of genomic variants will not be reported[http://projects.tcag.ca/variation/].

CNVs within or involving non-coding regions of the genome may not be reported. The performance of this test for detection of mosaicism has not been established. While this assay is validated and should have the capacity to detect a broad spectrum of specific disorders/syndromes, including aneuploidy and subtelomeric deletions, not all of these disorders have been detected in our laboratory due to their rarity. Other pertinent standard genetic tests should be considered in conjunction with a clinical evaluation of the patient. Please expect the results of any other concurrent test in a separate report. The oligo-SNP (oligonucleotide, single nucleotide polymorphism, Affymetrix CytoScan HD) assay uses a microarray containing over 2.67 million probes, including 1.9 million copy number probes and 750 thousand SNP probes. The overall average inter-probe distance is 1,150 base pairs. This test was developed and its analytical performance characteristics have been determined by Quest Diagnostics Nichols Institute, Chantilly, VA. It has not been cleared or approved by FDA. This assay has been validated pursuant to the CLIA regulations and is used for clinical purposes.

Electronic Signature on File

Haiying Meng, M.D., Ph.D., FACMG

nt: Example, Patient	DOB: 01/03/1990	Collection Date: 11/0	2/2018 Specimen N	umber: T0001390	Page: (
TEST	UNITS	NORMAL RANGE	RESULT	COMME	NT
	Technical Director, C	Cytogenetics and Geno	mics, 703-802-7156		
	Test Performed by Q	uest, Chantilly,			
	Quest Diagnostics Ni	chols Institute,			
	14225 Newbrook Dri	ve, Chantilly, VA 201	51		
	Patrick W Mason, M	.D., Ph.D., Director of	Laboratories		
	(703) 802-6900, CLL	A 49D0221801			
Maternal Cell Contan	nination Q8M				
			See Below	See Interp	retation
	Contamination Study	Maternal Cell:			
	NO MATERNAL CE	ELL CONTAMINATI	ON WAS DETECTED		
	Comparative analysis	s of maternal and fetal	DNA for fifteen short ta	ndem repeats	
	(STRs-CSF1PO, D2S	51338, D3S1358, D5S	818, D7S820, D8S1179,	D13S317, D16S539	2
	D18851, D198433, D	D21S11, FGA, THO1,	TPOX, vWA) was perfo	rmed by multiplex P	CR
	and capillary electrop	bhoresis. This assay ca	n detect a minor DNA sp	pecies	
	of 5% or higher in a r	nixture of two DNA s	amples. This test was pe	rformed using	
	a kit that has not beer	n cleared or approved	by the FDA. The analytic	cal performance	
	characteristics of this	s test have been determ	nined by Quest Diagnost	ics	
	Nichols Institute, Cha	antilly, VA. This test s	hould not be used for dia	agnosis	
	without confirmation	by other medically es	tablished means.		
Tests Derformed At:	084 Quart Diagnostics				
Tests renomed At.	QoA Quest Diagnostics (rive			
	Chantilly VA 2015	51			
	O8M Quest Diagnostics (CLIA·49D0221801			
	14225 Newbrook D	rive.			
	Chantilly, VA 2015	51			
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