

EXAMPLE, REPORT W

WX000003827 M 07/08/1978 45 Y

		Referral	resting				
		Colle	cted: 08/30/2023	3 14:07	Received:	08/30/2023	14:07
Test Name	1	Result	Flag	Ref-Ranges	<u> </u>	<u>Jnits</u>	<u>Site</u>
Celiac [Disease HLA-DQ Genoty	ping					
HLA-DQA1	I, Allele 1	01:02					ARRI
	Performed By: UUH Histoco 417 Wakara Way Suite 3220 Salt Lake City, UT 84108 CLIA Number: 46D0523979	ompatibility and In	nmunogenetic				
HLA-DQA1	I, Allele 2	03:01					ARR
	Performed By: UUH Histoco 417 Wakara Way Suite 3220 Salt Lake City, UT 84108 CLIA Number: 46D0523979	ompatibility and In	nmunogenetic				
HLA-DQB1	I, Allele 1	03:02					ARRI
	Performed By: UUH Histoco 417 Wakara Way Suite 3220 Salt Lake City, UT 84108 CLIA Number: 46D0523979	ompatibility and In	nmunogenetic				
HLA-DQB1	I, Allele 2	06:04					ARRI
	Performed By: UUH Histoco 417 Wakara Way Suite 3220 Salt Lake City, UT 84108 CLIA Number: 46D0523979	ompatibility and I	nmunogenetic				
Celiac HLA	Interpretation	See Note					ARRI
	Positive for HLA-DQ8 (DQF Interpretation: HLA-DQ8 w this allele is observed a individuals with celiac of general population. This diagnosis of celiac disea establish a diagnosis. It asymptomatic relative of disease-associated antibo	vas detected. At 10 in approximately 5 disease and 15-20 result is support ase, but does not 1 f this individual an affected indiv	east one copy -10 percent of percent of the ive of a clin oy itself is an idual, celiad	of ne nical			

Ordered By: KAJAL SITWALA, MD, PhD WX0000000002365



EXAMPLE, REPORT W WX000003827 M 07/08/1978 45 Y

		Referral Colle	cted: 08/30/2023	14.07	Received:	08/30/2023	14.0.	
t Name		Result	00,00,2020	Ref-Ranges		Jnits	Sit	
<u>struanc</u>	three to five year intervals. management of this individual findings. Performed By: UUH Histocompat 417 Wakara Way Suite 3220 Salt Lake City, UT 84108 CLIA Number: 46D0523979 BACKGROUND INFORMATION: Celia	Medical screated screated should rely defined and the second seco	ening and on clinical mmunogenetic		<u> </u>	<u>511163</u>	01	
	CHARACTERISTICS: Celiac disease is a systemic autoimmune disease of the gastrointestinal system caused by exposure to cereal gluten in genetically susceptible individuals.							
	INCIDENCE: On average, 1 in 133 individuals in the United States is affected.							
	INHERITANCE: Multifactorial.							
	CAUSE: The presence of either alleles in combination with d							
	CLINICAL SENSITIVITY: greater than 99 percent.							
	METHODOLOGY: Polymerase Chain Sequencing, or Polymerase Cha Oligonucleotide Probe Hybridi	in Reaction/S						
	ANALYTICAL SENSITIVITY AND SP percent.	ECIFICITY: gr	eater than 99					
	LIMITATIONS: Rare diagnostic site mutations. Other genetic influence celiac disease are an HLA allele cannot be resol assignment will be reported a allele frequencies from the c well-documented alleles catal et al, 2020).	and nongenet not evaluated ved unambiguo s the most con common, interme	ic factors th . In cases wh usly, the all mmon, based o ediate and	at ere ele n				
	ALLELES TESTED: HLA-DQA1 and	HLA-DQB1 alle	les.					
	Most celiac disease patients carry HLA-DQ2.5 heterodimers HLA-DQB1*02 alleles. The rema patients carry HLA-DQ8, encod most commonly in combination minority of patients negative	encoded by HL ining 5-10 pe led by HLA-DQB with HLA-DQA1	A-DQA1*05 and rcent of the 1*03:02 allel *03 alleles.	e, A				

Ordered By: KAJAL SITWALA, MD, PhD WX000000002365



EXAMPLE, REPORT W WX0000003827 M 07/08/1978 45 Y

	Referral Testing	
	Collected: 08/30/2023 14:07 Received: 08/30	/2023 14:07
<u>Test Name</u>	ResultFlagRef-RangesUnitscarry HLA-DQB1*02 but without the DQA1*05 alpha chain, mostcommonly with DQA1*02. The presence of the DQB1*02 allelein combination with either DQ2.5 or DQ8 may furtherincrease celiac disease risk.	Site
	Stratified overall genetic risk for patients carrying the celiac disease-associated HLA-DQ genotypes:	
	GenotypeRisk* DQ2.5 homozygousVery High (greater than 1:10) DQ2.5 + DQB1*02Very High (greater than 1:20) DQ2 homozygousHigh (greater than 1:20) DQ8 homozygousHigh (greater than 1:20) DQ8 + DQB1*02 (without DQA1*05)Intermediate (greater than 1:50) DQ2.5 heterozygousAt risk (greater than 1:50) DQ8 heterozygousAt risk (greater than 1:100) Population risk for unknown genotypeIow DQ81*02 (without DQA1*05)Low DQA1*05 (without DQA1*05)Low DQA1*05 (without DQB1*02)Minimal Negative for DQ2 and DQ8Not at risk * Risk is provided from the references below, and defined according to HLA allele combinations, considering a disease prevalence of 1:100. However, these alleles are common in the general population and the majority of individuals positive for celiac-associated alleles do not develop the disease. Detection of these alleles can support a clinical diagnosis but should not be interpreted as diagnostic of celiac disease.	
	<pre>References: 1. Megiorni F, Mora B, Bonamico M, et al. HLA-DQ and risk gradient for celiac disease. Human Immunology. 2009;70:55-59. 2. Pietzak MM, Schofield TC, McGinnis MJ, et al. Stratifying risk for celiac disease in a large at-risk United States population by using HLA alleles. Clinical Gastroenterology and Hepatology. 2009;7:966-971. 3. Almeida LM, Gandolfi L, Pratesi R, et al. Presence of DQ2.2 associated with DQ2.5 increases the risk for celiac disease. Autoimmune Diseases, 2016. 2016:5409653. 4. Vader W, Stepniak D, Kooy Y, et al. The HLA-DQ2 gene dose effect in celiac disease is directly related to the magnitude and breadth of gluten-specific T cell responses. PNAS. 2003;100:12390-12395.</pre>	
	DISCLAIMER INFORMATION:	

LAB: L - LOW, H - HIGH, AB - ABNORMAL, C - CRITICAL, . - NOT TESTED

Ordered By: KAJAL SITWALA, MD, PhD WX00000000002365



EXAMPLE, REPORT W WX0000003827 M 07/08/1978 45 Y

Referral Testing				
Collected: 08/30/2023	14:07	Received:	08/30/2023	14:07
ResultFlagThis test was developed and its performance characterisdetermined by the Histocompatibility& Immunogeneticslaboratory at the University of Utah Health. It has notbeen cleared or approved by the US Food and DrugAdministration (FDA). The FDA has determined that suchclearance or approval is not necessary. This test is usfor clinical purposes. It should not be regarded asinvestigational or for research. Histocompatibility&Immunogenetics laboratory is certified under the ClinicLaboratory Improvement Amendments of 1988 (CLIA-88) asqualified to perform high complexity clinical laboratortesting.Performed at: Histocompatibility & ImmunogeneticsLaboratory, University of Utah Health, 417 Wakara Way,Suite 3220, Salt Lake City, UT 84108.Counseling and informed consent are recommended for gentesting. Consent forms are available online.	ed al Y	<u>i</u> <u>l</u>	<u>Jnits</u>	Site

 Performing Site:

 ARRL: ARUP REFERENCE LAB 500 Chipeta Way Salt Lake City UT 841081221

 Reported Date:
 2023.08.30
 14:07
 CDHDG

LAB: L - LOW, H - HIGH, AB - ABNORMAL, C - CRITICAL, . - NOT TESTED