## **Interpreting MORL Complement-Mediated Kidney Disease Genetic and Functional Results**

GENETIC TESTING																
Gene		Ch	Chromosomal Location				Interpretation									
beer	ment gene that n reported to be red with TMAs/C	The specia	of a given gene				Pathogenic known: a variant that has been proven to be disease-causing  Likely pathogenic: a variant that is likely to be disease causing based on current data  Unknown significance: a variant for which further interpretation is not possible based on available data  Likely benign: a variant not known to cause disease									
PATHWAYS						AUTOANTIBODIES										
<b>CH50</b> (41-95 Units/mL		<b>APFA</b> (50-130%)	Λεερ		Autoantibody		FB Autoantibody (<200 AU)		Fluid Phase Activity -IFE (<7.5%)		C3Nef - C3CSA (<20%)	C5Nef- C3CSAP (<20%)	C4Nef (<20%)			
Determines whether the CP is overactive or whether a CP protein has been abnormally consumed		Determines whether the AP overactive or whether an AF protein has bee abnormally consumed	abnormal C3		binds H (F inter FH fur comp	hds to Factor binds (FB); can terfere with function and npromise AP binds		body that o Factor B n interfere th C3 vertase ion; often in PIGN	th	ermines if a protein ne blood is causing complement regulation/activatic	converta them f	Antibodies to C3- or C5- convertase, preventing them from naturally falling apart		Similar to C3- or C5- nephritic factors, however they stabilize the classical pathway convertase		
							BIOM	ARKERS								
	<b>C3 level</b> (90-180 mg/dL)	C3c Level (<1.5 mg/L)	<b>C4 Level</b> (15-47 mg/dL)	<b>FB Level</b> (22-50 mg/dL)		Ba Level (<1.2 mg/L)	Bb Level (<2.2 mg/L)	<b>FD Lev</b> (0.78-1 mg/L	.59	<b>C5 Level</b> (13.5-27 mg/L)	Properdin Level (10-33 mg/L)	Soluble C5b-9 (<0.3 mg/L)	FI Level (18-44 mg/L)	FH Level (180-420 mg/L)		
High Result	Represents inflammation or obesity	A breakdown product of C3, suggests overactivity of the AP	Represents inflammation		Represents inflammation of FB; I mean being of excess levels of		High level products h levels sugges at FB is sumed ly; high are also n ESKD High level Factor D ( F		(FD) st ng y on we of nent	Elevated with terminal complement pathway inhibitor		Increased activity of the terminal complement pathway	Represents inflammation	Represents inflammation		
Low Result	Deficient because of a gene abnormality or inappropriately consumed		Deficient because of a gene abnormality or inappropriately consumed	because gene abnormal consumed	Deficient ecause of a gene normality or nsumed due overactive AP			253776	,	Suggests terminal pathway hyperactivity	Suggests terminal pathway hyperactivity	Low if on terminal complement blockade	Deficiency typically reflects a gene abnormality	Deficiency typically reflects a gene abnormality or inappropriate consumption		

<sup>\*</sup> AP = Alternate Pathway; CP = Classical Pathway; Nef = Nephritic Factor, ESKD= End Stage Kidney Disease; laboratory results may be significantly altered by inappropriate specimen handling; due to the extreme complexity of the complement cascade, assessing complement activity and regulation is best performed by pathway analysis, together with autoantibody testing and biomarker profiling as opposed to doing tests in isolation