PHN-02

Form S331: Genetics Core Laboratory Results Form

Section A: KEY IDENTIFYING INFORMATION					
A1. Study Identif	fication Number				
Replaced by blinded ID					
blind_id	Blinded ID				
A2. Name of Core Laboratory Technician		PRINT FULL NAME			
Removed to protect privacy					
A3. Specimen collection date		$\frac{1}{M}$ $\frac{1}{M}$ $\frac{1}{D}$ $\frac{1}{D}$ $\frac{1}{Y}$ $\frac{1}{Y}$ $\frac{1}{Y}$ $\frac{1}{Y}$			
Replaced by age	at specimen collection				
gspec_age	A3. <created var="">Age at specimen collection, days</created>				
Section B: TEST RESULTS					
B1. Date sample received		$\frac{1}{M}$ $\frac{1}{M}$ $\frac{1}{D}$ $\frac{1}{D}$ $\frac{1}{Y}$ $\frac{1}{Y}$ $\frac{1}{Y}$ $\frac{1}{Y}$			
Replaced by age	when sample received				
gsampr_age					
B2. Was genoty	ping completed?	YES 1 NO			
GENODONE	B2. Was genotyping comp	leted			
a. Date ge	enotyping completed	$\frac{1}{M} \frac{1}{M} \frac{1}{D} \frac{1}{D} \frac{1}{Y} \frac{1}{Y} \frac{1}{Y} \frac{1}{Y}$			
Replaced by age	Replaced by age when genotyping completed				
geno_age	geno_age B2a. <created var="">Age when genotyping completed, days</created>				

GENOACE	B2b1. ACE gene
hr1	B2b1. <created var="">ACE gene = DD, yes/no</created>

b. Genotype

ACE gene

1.

DD240 ID 340 II 241 Not done ... -1

	Pediatric Heart Netwo	rk: Infant Single Ventricle	Trial	PHN-02	
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2. Ar	ngiotensinogen gene	TT311	CC 210	Not done1	
GENOAG	B2b2. Angiotensinogen ger]			
hr2	B2b2. <created var=""> Angio</created>				
	ngiotensin II type I ceptor gene	CC210 AC 300	AA200	Not done1	
GENOARG	B2b3. Angiotensin II type I				
hr3	B2b3. <created var="">Angiotensin II type I receptor gene = CC, 1:Yes/0:No</created>				
4. Al	dosterone synthase gene	CC210 TC311	TT 230	Not done1	
GENOALD	B2b4. Aldosterone synthas	e gene			
hr4	B2b4. <created var="">Aldost 1:Yes/0:No</created>	erone synthase gene =	GG,		
5. Ca	ardiac chymase A gene	GG220 AG301	AA 200	Not done1	
GENOCCA	BENOCCA B2b5. Cardiac chymase A gene				
hr5	hr5 B2b5. <created var="">Cardiac chymase A gene = AA, 1:Yes/0:No</created>				
HR	sum of hr1-hr5, from 0 to 5				
HighRisk	1:Yes (HR >1), 2:No (HR=0	0 or 1)			
	SKIP TO B4				
B3. Reason gene	otyping was not completed	INSUFFICIENT QUAN HEMOLYSIS		2	
NOGENO	B3. Reason genotyping wa	s not completed			
a. If OTHE	R, specify:				
NOGENO_S	B3a. If other, specify]	
B4. Was DNA sto	ored for future use?	YES	1 NO	O2 (B5)	
DNASTORE	B4. Was DNA stored for fut	ture use]	

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a. Stored for future cardiovascular disease studies related to this research study
DNANOREL [Added version C] B4a. Stored for future disease studies related to this study
b. Stored for future cardiovascular disease studies that are not related to this research YES
DNAREL [Added version C] B4b. Stored for future disease studies not related to this study
B5. Comments
COMMS331 B5. Comments