## Center for Human Genetics, Inc. Date received: \_\_\_ **Riverside Technology Center** Pedigree #: \_\_\_\_\_ 840 Memorial Drive, Suite 101 Family name: Cambridge, MA 02139 Sample type:\_\_\_\_\_ Co-directors: Aubrey Milunsky, M.D., D.Sc., Jeff Milunsky, M.D. Lab #: Phone: 617-492-7083 Fax: 617-492-7092 Website: http://www.chginc.org Patient Name: □ Male ¬ Female □ Unknown Last First Hospital/Patient ID#: Date of Birth: Partner/Parent of: Address: Phone: State Zip Code Citv NPI #: \_\_\_\_\_ Referring Provider: Referring Laboratory (if different): Physician signature:\_\_\_\_ Name: \_\_ Name: Address: Address: Citv Citv State State Zip Code Zip Code Phone: \_\_\_\_\_ Fax: \_\_\_\_\_ Phone: \_\_\_\_\_ Fax: \_\_\_\_\_ Genetic Counselor: Email: Phone: \_\_\_\_\_ Fax: \_\_\_\_\_ Email: \*\* Billing information (Page 4) and copy of insurance card (front and back) MUST accompany sample and requisition form. \*\* \* Consent form (Page 5) is REQUIRED for all samples from New York and all predictive testing BEFORE testing can be initiated.\*\* **CLINICAL INFORMATION:** Ethnicity: □ Ashkenazi Jewish □ French Canadian □ Caucasian □ African American □ Hispanic □ Asian □ Sephardic Jewish □ Armenian □ Turkish □ Mediterranean □ Arabic □ Other: \_\_\_\_\_ ICD-10 Diagnosis (REQUIRED): **Purpose of Study:** □ Diagnosis □ Carrier Screen □ Prenatal Diagnosis □ Predictive/Presymptomatic\*\* □ No family history (Call before sending samples) □ Family history\* □ Ultrasound abnormality\* \*Please include additional information: \_\_\_\_\_\_ Pregnancy information (if applicable): Gestational age \_\_\_\_\_ □ By LMP \_\_\_\_ □ By ultrasound \_\_\_ Date □ Date Name and relationship of family members previously tested at CHG:

Name of mutation to be tested (if known in family):

**MOLECULAR (DNA) TEST REQUISITION** 

FOR CHG LAB USE ONLY:

M	OLECULAR (DNA) TEST REQUISITION  FOR CHG LA Pedigree #:	B USE ONLY: Lab #:	
Pa	atient Name:	Date of Birth:	
	Last First	<u> </u>	
	**SPECIMEN REQUIREMENTS FOR ALL TESTS LISTED BELO	DW: 7-10 cc BLOOD IN EDTA OR ACD ANTICOAGULANT**	
	NA TEST(S) REQUESTED: LEASE NOTE: <u>ANALYSIS = SEQUENCING AND MLPA)</u>	Date Sample Collected:	
П	Aarskog Scott syndrome ( <i>FGD1</i> analysis)	□ Costello syndrome panel	
	ACTG2 sequencing	Sequencing of □ HRAS □ KRAS □ BRAF	
	Acute Myeloid (Myelogenous) Leukemia	<ul> <li>Creatine (transporter) deficiency (SLC6A8 analysis)</li> </ul>	
	□ FLT3 C835 mutation □ NPM1 exon 12 sequencing	□ Cystic fibrosis	
	ADGRG2 sequencing	□ 40+ mutations panel □ 100+ mutations panel	
	Alpha-thalassemia/XLID syndrome (ATRX analysis)	□ <b>CFTR</b> analysis	
	Aneurysm osteoarthritis syndrome ( <b>SMAD3</b> analysis)	□ Deafness	
	Angelman/Angelman-like syndrome panel	<ul> <li>Connexin 26 sequencing (nonsyndromic deafness)</li> </ul>	
	<ul> <li>Methylation-sensitive MLPA</li> </ul>	□ Connexin 30 deletion (nonsyndromic deafness)	
	Sequencing of UBE3A SLC9A6	□ Mitochondrial A1555G	
	Analysis of □ TCF4 □ ZEB2	Developmental Language Disorders	
	Aortic valve disease (NOTCH1 sequencing)	Sequencing of D FOXP1 D FOXP2	
	ARX analysis	□ Dravet syndrome ( <b>SCN1A</b> analysis)	
	Ashkenazi Jewish panel	□ Duchenne or Becker Muscular Dystrophy ( <i>DMD/BMD</i> MLPA)	
	□ Bloom syndrome □ LDD/DLD □ Consum disease □ MSUD Type 1B	□ Ehlers-Danlos syndrome	
	□ Canavan disease □ MSUD Type 1B	Types I/II: □ COL5A1 analysis □ COL5A2 sequencing	
	□ Factor XI deficiency □ Mucolipidosis type IV	Type IV: □ COL3A1 analysis	
	<ul> <li>□ Familial Dysautonomia</li> <li>□ Remaline Myopathy</li> <li>□ Familial Hyperinsulinemia</li> <li>□ Niemann-Pick disease Type A</li> </ul>	Type VII: Analysis of   COL1A1   COL1A2  Ehlers-Danlos variant with periventricular heterotopia (FLNA analysis)	lveie)
	□ Fanconi anemia Group C □ Tay Sachs disease	□ Epilepsy/Intellectual Disability (Female Restricted) ( <i>PCDH19</i> and	
	□ Gaucher disease □ Usher syndrome 1F	□ Familial Mediterranean Fever	ily SiS)
	□ GSD Type 1A □ Usher syndrome 3A	□ Common mutations only □ <i>MEFV</i> analysis	
	□ Joubert disease □ Walker-Warburg syndrome	□ FG syndrome	
	Ataxia panel	□ <i>MED12</i> sequencing □ <i>FLNA</i> analysis	
	□ Spinocerebellar ataxia (check all that apply)	□ Fragile X syndrome	
	Types = 1 = 2 = 3 = 6 = 7 = 8 = 10 = 12 = 17	□ Hemochromatosis	
	□ DRPLA	☐ Huntington disease**	
	Autism/Autism Spectrum Disorder (53 gene panel)	□ Infantile Spasms	
	Autism (with macrocephaly)	Analysis of   ARX   CDKL5   SCN1A	
	(PTEN analysis and promoter sequencing)	<ul> <li>Infertility testing</li> </ul>	
	Banking (circle type) DNA or Lymphoblast	<ul> <li>Ovarian insufficiency/spermatogenic failure (NR5A1 analysis)</li> </ul>	)
	Beals syndrome (FBN2 sequencing)	<ul> <li>Premature ovarian failure (Fragile X testing)</li> </ul>	
	Borjesen-Forssman-Lehmann syndrome ( <i>PHF6</i> sequencing)	□ SYCP3 sequencing	
	Branchio-oculo-facial syndrome ( <i>TFAP2A</i> analysis)	<ul> <li>Y-microdeletion studies</li> </ul>	
	Breast/Ovarian Cancer**	□ Intracranial aneurysm	
	□ BRCA1/2 analysis	Sequencing of □ NTM □ TGFβR3	
	□ BRCA1/2 (Ashkenazi Jewish mutations only)	□ Kabuki syndrome	
	C9orf72-Related Neurodegenerative Disease	Analysis of DMLL2/KMT2D DKDM6A	
	CADASIL ( <b>NOTCH3</b> sequencing)	□ Kennedy disease ( <i>SBMA</i> )	
Ц	Cardiofaciocutaneous syndrome panel Sequencing of  BRAF  MAP2K1  MAP2K2  KRAS	<ul> <li>□ LADD syndrome</li> <li>Analysis of □ FGF10 □ FGFR2</li> </ul>	
	Charcot-Marie-Tooth disease 1A/HNPP ( <i>PMP22</i> MLPA)	Sequencing of   FGFR3	
	Charcot-Marie-Tooth disease 1B, 2I, 2J (MPZ analysis)	□ LEOPARD syndrome panel	
	CHARGE syndrome ( <i>CHD7</i> analysis)	Sequencing of Description Property Research	
	Chronic Intestinal Pseudo-obstruction ( <i>ACTG2</i> sequencing)	□ Loeys-Dietz syndrome	
	Coffin-Lowry syndrome ( <i>RSK2</i> analysis)	Analysis of $\Box$ TGF $\beta$ R1 $\Box$ TGF $\beta$ R2 $\Box$ SMAD3 $\Box$ TGF $\beta$ 2	
	Colon Cancer	Sequencing of □ <i>TGFβ3</i>	
_	□ Familial Adenomatous Polyposis ( <i>APC</i> analysis)	□ Marfan syndrome ( <i>FBN1</i> analysis)	
	□ Lynch/HNPCC Panel	□ Maternal cell contamination studies	

Analysis of  $\square$  *MLH1*  $\square$  *MSH2*  $\square$  *MSH6*  $\square$  *PMS2* □ **EPCAM/TACSTD1** MLPA □ **MYH** Associated Polyposis (**MUTYH** analysis)

□ Congenital bilateral absence of vas deferens (CBAVD)

□ 100+ mutations panel □ CFTR analysis □ ADGRG2 sequencing

□ Congenital Contractural Arachnodactyly (*FBN2* sequencing)

□ CONNECT1 (Connective Tissue Disorders analysis)

□ CONNECT1 Sequencing Only (22 genes)

□ CONNECT1 MLPA (circle tests)

COL1A1, COL1A2, COL2A1, COL3A1, COL5A1, COL11A1, FBN1, FLNA, MYH11, SMAD3, TGF\$2, TGF\$R1, TGF\$R2 □ MED12 related disorders (MED12 sequencing)

□ Mowat Wilson syndrome (**ZEB2** analysis)

□ All 37 gene analysis □ MELAS □ CPEO/KSS □ LHON □ Leigh syndrome/NARP □ MERRF

□ Multiple Endocrine Neoplasia, type 1 (*MEN1* analysis)

□ Multiple Endocrine Neoplasia, type 2 (*RET* analysis)

□ Melanoma (*CDKN2A* analysis)

□ Mitochondrial diseases panel

FOR <b>CHG</b> LAB USE ONLY:	
Pedigree #:	Lab #:

Patient Name:			Date of Birth:
	Last	First	

DNA TEST(S) REQUESTED:	
(PLEASE NOTE: <u>ANALYSIS = SEQUENCING AND MLPA)</u>	Date Sample Collected:
□ Myeloproliferative Disease	□ Thrombophilia panel
□ <b>JAK2</b> (V617F mutation [reflex to exon 12 sequencing])	· · · ·
□ CALR exon 9 sequencing	□ Prothrombin (G20210A)
□ <i>MPL</i> sequencing	□ Tuberous Sclerosis
□ Neurexin 1 ( <i>NRXN1</i> analysis)	Analysis of $\Box$ <b>TSC1</b> $\Box$ <b>TSC2</b>
<ul> <li>Neurofibromatosis</li> </ul>	□ UPD: (circle test) chromosome 7 14 15
Analysis of   NF1   NF2	<ul> <li>Visceral Myopathy (ACTG2 sequencing)</li> </ul>
<ul> <li>Neurofibromatosis type 1-like (Legius) syndrome</li> </ul>	<ul><li>von-Hippel-Lindau disease (VHL analysis)</li></ul>
(SPRED1 analysis)	□ Waardenburg syndrome
□ Neuroligin	Types 1 and 3:   PAX3 analysis
Sequencing of D NLGN3 D NLGN4	Type 2: Analysis of DMITF DSOX10 DEDNRB
<ul> <li>Noonan syndrome panel</li> </ul>	Type 4: Analysis of □ SOX10 □ EDN3 □ EDNRB
Sequencing of prescription PTPN11 SOS1 SOS2 RAF1	<ul><li>Wilson disease (ATP7B analysis)</li></ul>
□ KRAS □ NRAS □ SHOC2 S2G mutation	<ul> <li>X-inactivation studies</li> </ul>
□ BRAF □ CBL □ RIT1 □ LZTR1	<ul> <li>X-linked lymphoproliferative disease (SH2D1A analysis)</li> </ul>
□ OpitzG/BBB syndrome ( <i>MID1</i> analysis)	<ul> <li>X-linked Intellectual Disability (XLID)</li> </ul>
<ul> <li>Osteogenesis imperfecta Type I, II, III, IV</li> </ul>	(Order whole panel, individual tiers, or single gene)
Analysis of □ COL1A1 □ COL1A2	□ Tier A □ <b>NLGN3</b> sequencing
Pancreatitis (hereditary)	<ul><li>NLGN4 sequencing</li></ul>
□ Pancreatitis panel: □ CTRC sequencing,	<ul><li>Rett syndrome (<i>MECP2</i> analysis)</li></ul>
Analysis of □ PRSS1 □ SPINK1	<ul> <li>Rett syndrome - atypical (CDKL5 analysis)</li> </ul>
□ <b>CFTR</b> analysis	□ Tier 1 □ <b>DLG3</b> sequencing
□ Paraganglioma-Pheochromocytoma syndromes	□ <i>FTSJ1</i> sequencing
Analysis of SDHB SDHC SDHD	□ JARID1C sequencing
<ul> <li>Paternity testing (Call before sending samples)</li> </ul>	□ Borjesen-Forssman-Lehmann syndrome
□ Pelizaeus-Merzbacher disease ( <i>PLP1</i> analysis)	( <b>PHF6</b> sequencing)
□ Pendred syndrome ( <b>SLC26A4</b> sequencing)	□ <b>ZNF41</b> sequencing
<ul><li>Phosphomannomutase 2 (<i>PMM2</i> sequencing)</li></ul>	☐ Tier 2 ☐ Asperger syndrome ( <i>GDI1</i> analysis)
□ Phenylketonuria ( <i>PAH</i> analysis)	□ <b>FACL4</b> analysis
□ Pitt Hopkins syndrome ( <i>TCF4</i> analysis)	□ <b>OPHN1</b> analysis
□ Pitt Hopkins-like syndrome	Renpenning syndrome ( <i>PQBP1</i> analysis)
□ CNTNAP2 sequencing □ Neurexin 1 (NRXN1 analysis	
<ul><li>Prader-Willi syndrome (Methylation-sensitive MLPA)</li><li>PTCHD1 analysis</li></ul>	□ Tier 3 □ Alpha-thalassemia/XLID syndrome ( <i>ATRX</i> analysis)
□ <i>PTEN</i> Hamartoma Tumor syndromes	□ Aarskog Scott syndrome ( <i>FGD1</i> analysis)
( <i>PTEN</i> analysis and promoter sequencing)	□ OpitzG/BBB syndrome ( <i>MID1</i> analysis)
□ Renpenning syndrome ( <i>PQBP1</i> analysis)	□ Pelizaeus-Merzbacher disease ( <i>PLP1</i> analysis)
□ Rett/Rett-like syndrome panel	□ Coffin-Lowry syndrome ( <i>RSK2</i> analysis)
Analysis of   MECP2   CDKL5   FOXG1   TCF4	□ Creatine (transporter) deficiency
□ SCN1A analysis	(SLC6A8 analysis)
□ Sickle cell anemia (HbSS, HbSC)	□ Tier 4 □ <b>AGTR2</b> analysis
□ Small fiber neuropathy (SCN9A sequencing)	□ <i>ARHGEF6</i> analysis
□ Smith-Lemli-Opitz syndrome ( <i>DHCR7</i> sequencing)	□ <i>MED12</i> sequencing
□ SNP microarray	□ <b>PAK3</b> analysis
□ Sotos syndrome ( <b>NSD1</b> analysis)	□ SLC16A2 sequencing
<ul> <li>Spinal Muscular Atrophy (SMN1 exons 7-8 MLPA)</li> </ul>	<ul> <li>Other X-linked intellectual disability genes</li> </ul>
□ Haplotype CNV sequencing	□ <i>IL1RAPL1</i> analysis
□ Stickler syndrome	□ <b>RAB39B</b> sequencing
Type I:   COL2A1 analysis	<ul> <li>X-linked Intellectual Disability/Epilepsy Panels</li> </ul>
Type II: □ COL11A1 analysis	□ Panel 1 □ Angelman-like syndrome (X-linked
Type III:   COL11A2 sequencing	Christianson type) (SLC9A6 sequencing)
□ SYNGAP1 sequencing	<ul> <li>PCDH19 sequencing (females only)</li> </ul>
(Intellectual disability, dominant, nonsyndromic)	□ Rett syndrome ( <i>MECP2</i> analysis)
□ Tay-Sachs disease	□ Rett syndrome - atypical (CDKL5 analysis)
□ Thoracic Aortic Aneurysms/Dissections panel	□ Panel 2 □ <i>ATP6AP2</i> sequencing
Analysis of □ FBN1 □ MYH11 □ SMAD3	□ Creatine (transporter) deficiency (SLC6A8 analys
□ TGFβ2 □ TGFβR1 □ TGFβR2	□ <b>OPHN1</b> analysis
Sequencing of _ ACTA2 _ BGN _ FOXE3 _ LOX _ M.	· · · · · · · · · · · · · · · · · · ·
□ MFAP5 □ MYLK □ NTM □ PRKG1 □ S	KI   XY Disorders of sex development (NR5A1 analysis)

 $\square$  SLC2A10  $\square$  SMAD2  $\square$  TGF $\beta$ 3

 $\hfill\Box$  Y-microdeletion studies/Y-chromosome detection (SRY)

MOLECULAR (DNA) TEST REQUISITION Center for Human Genetics, Inc. Riverside Technology Center 840 Memorial Drive, Suite 101 Cambridge, MA 02139

Co-directors: Aubrey Milunsky, M.D., D.Sc., Jeff Milunsky, M.D.

Phone: (617) 492-7083 Fax: (617) 492-7092

Web: http://www.CHGINC.org

Patient Name:				Date of Birth:	
	Last	First	MI		

Last	First	MI	
INF	ORMED CONS	ENT FOR DI	NA TESTING
signed consent form is strongly re	ecommended for <u>all</u> gene	tic testing. If a sigr	ictive testing BEFORE testing can be initiated. ned consent is not submitted, the Center for Human obtained the patient's informed consent.
isolated from to assess the probability that I (m	y/our fetus/child) am (is)	(sample type) ob affected with or care	Human Genetics, Inc. to analyze a sample of DNA tained on (date) ry the gene for the genetic disease
The test procedure has been exp	lained to me/us and I/we	understand that:	
carrier for, the ab 2. The test results m	nay indicate that it is likely ove disease. nay be indeterminate bec	ause of my (my/our	ny/our fetus/child) am (is) affected with, or a fetus'/child's) genetic patterns or the genetic tations of the current technology.
II. DNA tests are performed with p (when applicable). Turn-around		-	and specific degrees of quoted accuracy
III. One possible result of DNA tes when comparing my (my/our fe	_		dence of previously undisclosed non-paternity family members.
IV. Genetic counseling, further testing process.	sting, or additional physic	cian consults may be	e warranted after testing in order to complete
for a minimum of three months	s. We do not guarantee or that time, any remaining	the future availab g material will be dis	A will be stored at the Center for Human Genetics, Inc. ility of DNA unless specific arrangements have bee sposed of at the discretion of the Laboratory Director, is maintained.
I can request that r	remaining DNA <u>not</u> be u	sed for research p	ourposes by initialing here:
VI. The results of this test are to be regulations.	pe released only to the or	dering physician an	d referral laboratory (if applicable) per HIPAA
· · ·	en satisfactorily explained	d to me/us by my/ou	proposed DNA test(s) and its/their limitations for ir physician or genetic counselor; and (2) I/we
Patient/Guardian Signa	ature		Witness Signature

Date

## \*\*\*BILLING INFORMATION AND COPY OF INSURANCE CARD FRONT AND BACK MUST ACCOMPANY SAMPLE AND REQUISITION FORM\*\*\*

SVC PROVIDER: CENTER FOR HUMAN GENETICS INC CLIA #22D0650242 NPI #1821153156

PATIENT INFOR	MATION:		***************************************
LAST NAME:		DER:	DATE OF BIRTH
FIRST NAME:	(CIRC	CLE)	1 1
MIDDLE:	M	F	MM/DD/YYYY
STREET ADDRESS:			APARTMENT# / FLOOR
CITY:	ST	ATE	ZIP
PHONE: HOME( )	CELL(	)	
PAYMENT INFO: (SELECT ONE) (CIRCLE): LAB/HOS	P/FAC/INST INSURANCE F	PATIENT CR	EDIT CARD
BILLING INFOR	RMATION (MUST BE COMP	LETED)	
LABORATORY/ HOSPITAL/ FA			RESS:
	FFIX LABEL HERE:		
ADDRESS:			
CITY, STATE, ZIP ATTENTION:			
PHONE: ( )			
FAX: ( )			
PURCHASE ORDER#			
PATIENT MEDICAL RECORD#			
INSUR	ANCE INFORMATION:		
INSURANCE COMPANY NAME:			
INSURANCE IDENTIFICATION #			
INSURANCE GROUP #			
SUBSCRIBER NAME:	SUBSCRIBER	DATE OF BIR	гн:
LAST:	/	1	
	2020	DD / YYYY	
FIRST.	(CIRCLE):	1111	
RELATIONSHIP TO PATIENT:	SELF	PARENT	SPOUSE
	SELF	PARENT	SPOUSE
INSURANCE ADDRESS:	INSURANCE T	ELEPHONE A	ND EXTENTION:
STREET:	( )		
	FAX#( )		
	CONTACT NAM		
CITY, STATE, ZIP:	AUTHORIZATI	ON#	
	VALID FROM:	/ / TO	1 1
*SECONDARY INS NAME:	SUB NAME:	R	ELATIONSHIP:
	JB DOB: / /		DER: M F
PATIENT ACKNOWLEDGEMENT: I AUTHORIZE ANY HOLDER OF MEDICAL INFORMATION ABOUT ME TO RELEASE TO ANY INSURANCE PLACE OF THE ORIGINAL AND REQUEST THAT THE PAYMENT OF MEDICAL INSURANCE BE PAID TO INSURANCE COMPANY DOES NOT PAY.	E CARRIER ANY INFORMATION NEEDED FOR THIS CI CHG, INC. I ALSO UNDERSTAND THAT I WILL BE HEL	LAIM. I PERMIT A COPY OF RESPONSIBLE FOR AL	OF THIS AUTHORIZATION TO BE USED IN MY PORTION OF THE CLAIM THAT THE
REQUIRED SIGNATURE:		DATE:	1 1
	FICIARY NAME:		
BENEFICIARY AGREEMENT: I HAVE BEEN NOTIFIED BY THE CENTER FOR HUMAN GENETICS THAT, IN MY CASE, MEDICARE IS LII		TED BELOW, FOR THE R	EASON STATED. IF MEDICARE DENIES PAYMENT. I
AGREE TO BE PERSONALLY AND FULLY RESPONSIBLE FOR PAYMENT.			
REQUIRED BENEFICIARY SIGNATURE:			DATE: / /

MEDICARE WILL ONLY PAY FOR SERVICES THAT IT DETERMINES TO BE "REASONABLE AND NECESSARY" UNDER SECTION 1862(a) (1) OF THE MEDICARE LAW.

IF MEDICARE DETERMINES THAT A PARTICULAR SERVICE, ALTHOUGH IT WOULD OTHERWISE BE COVERED, IS NOT "REASONABLE AND NECESSARY" UNDER MEDICARE PAYMENT STANDARDS, MEDICARE WILL DENY PAYMENT FOR MOLECULAR
THAT SERVICE. THE CENTER FOR HUMAN GENETICS BELIEVES THAT MEDICARE IS LIKELY TO DENY PAYMENT FOR MOLECULAR
DNA TESTING.

A. Notifier:  B. Patient Name:	C. Identification Number:	
Advance Bene	eficiary Notice of Noncoverage (	ABN)
NOTE: If Medicare doesn't pay for	or <b>D.</b> below, you may have to	pay.
	ng, even some care that you or your health ca	
good reason to think you need. We	e expect Medicare may not pay for the <b>D.</b>	below.
D.	E. Reason Medicare May Not Pay:	F. Estimated Cost
<ul> <li>Ask us any questions that</li> <li>Choose an option below all</li> <li>Note: If you choose Option</li> <li>that you might have</li> </ul>	an make an informed decision about your care you may have after you finish reading. bout whether to receive the <b>D.</b> on 1 or 2, we may help you to use any other in e, but Medicare cannot require us to do this.	₋ listed above.
G. OPTIONS: Check only on	e box. We cannot choose a box for you.	
also want Medicare billed for an of Summary Notice (MSN). I understand payment, but I can appeal to Medoes pay, you will refund any pay DPTION 2. I want the D. ask to be paid now as I am respo DPTION 3. I don't want the D.	listed above. You may ask to be pofficial decision on payment, which is sent to not stand that if Medicare doesn't pay, I am respondedicare by following the directions on the MSN ments I made to you, less co-pays or deduction listed above, but do not bill Medionsible for payment. I cannot appeal if Medicare would be anot appeal to see if Medicare would be seen if Medicare would b	ne on a Medicare insible for I. If Medicare bles. care. You may are is not billed. th this choice I
H. Additional Information:  This notice gives our opinion, nothis notice or Medicare billing, call of the control of the contr	ot an official Medicare decision. If you have 1-800-MEDICARE (1-800-633-4227/TTY: 1-8 e received and understand this notice. You als	e other questions o
I. Signature:	J. Date: ersons are required to respond to a collection of information unless it displa	

According to the Paperwork Reduction Act of 1995, no persons are required to respond to a collection of information unless it displays a valid OMB control number. The valid OMB control number for this information collection is 0938-0566. The time required to complete this information collection is estimated to average 7 minutes per response, including the time to review instructions, search existing data resources, gather the data needed, and complete and review the information collection. If you have comments concerning the accuracy of the time estimate or suggestions for improving this form, please write to: CMS, 7500 Security Boulevard, Attn: PRA Reports Clearance Officer, Baltimore, Maryland 21244-1850.