

**PATIENT INFORMATION**

Name (Last, First, MI)	Biological Sex <input type="checkbox"/> F <input type="checkbox"/> M	Date of Birth (MM/DD/YY)	MRN
Ethnicity: <input type="checkbox"/> African American <input type="checkbox"/> Asian <input type="checkbox"/> Caucasian <input type="checkbox"/> Hispanic <input type="checkbox"/> Portuguese <input type="checkbox"/> Other:			Ashkenazi Jewish <input type="checkbox"/> Yes <input type="checkbox"/> No
Address	City	State	Zip
Phone	Email		

**SPECIMEN INFORMATION (For phlebotomy service, select all services you are requesting)**

<input type="checkbox"/> Blood (EDTA preferred) <input type="checkbox"/> Saliva <input type="checkbox"/> Buccal Swab** <input type="checkbox"/> DNA <input type="checkbox"/> Cord blood* <input type="checkbox"/> Other*:	<input type="checkbox"/> Personal history of allogenic bone marrow or peripheral stem cell transplant*
---	--

**Collection Date**   **Specimen ID**   **Medical Record #**

\*Blood/saliva from patients with a history of allogenic bone marrow or stem cell transplant cannot be used for genetic testing. Blood/saliva from patients with active hematological disease is not recommended. An alternative specimen may be needed. Please see [ambrygen.com/specimen-requirements](http://ambrygen.com/specimen-requirements) for details  
\*\*Only for Fragile X syndrome and chromosomal microarray.  
\*Fetal specimens, cord blood and POC will have maternal cell contamination studies added for a charge. Maternal and fetal specimen required. Please see page 2 for Maternal Cell Contamination sample submission test codes.

Phlebotomy Services Request:  Phlebotomy draw    Insurance preverification first    Send blood kit to patient\*\*    Send saliva kit to patient  
\*\*As the patient's clinician, I am unaware of any potential for complication or difficulty in drawing blood for the listed patient(s). I understand that the phlebotomist has full authority to refuse to draw any patient if the safety of the phlebotomist and/or patient(s) are in question.

**INDICATION(S) FOR TESTING**

ICD-10 code(s):	Will the course of treatment change depending upon the results of the test? <input type="checkbox"/> Yes <input type="checkbox"/> No
-----------------	--

**PATIENT HISTORY**    No personal history of neurological disease

**PLEASE SUPPLY CLINIC NOTES AND PEDIGREE**   If pregnant, due date: \_\_\_\_\_   Upcoming procedure date: \_\_\_\_\_

Reasons for Testing: \_\_\_\_\_

<b>Birth and Neonatal History</b> <input type="checkbox"/> N/A Gestational age at birth: _____ Birth weight: _____ Head circumference at birth (if available): _____ <b>Developmental History</b> <input type="checkbox"/> N/A Developmental delay: <input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown Type of delay (choose all that apply): <input type="checkbox"/> Motor <input type="checkbox"/> Language <input type="checkbox"/> Global Intellectual disability: <input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown Regression or plateau: <input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown Does patient meet DSM-V diagnostic criteria for an autism spectrum disorder? <input type="checkbox"/> Yes <input type="checkbox"/> No <b>Seizure History</b> <input type="checkbox"/> N/A   Age at first unprovoked seizure: _____ Has this patient been diagnosed with an epilepsy syndrome? <input type="checkbox"/> Yes <input type="checkbox"/> No <input type="checkbox"/> Unknown If yes, please specify: _____	<b>Other History</b> <input type="checkbox"/> N/A Hypo-/hyperpigmentation: <input type="checkbox"/> Yes <input type="checkbox"/> No   Telangiectasias: <input type="checkbox"/> Yes <input type="checkbox"/> No Other skin abnormality, type: _____ Brain tumor, type: _____ Nerve tumor, type: _____ Other tumor, type: _____ <b>Other Clinical Findings (choose all that apply)</b> <input type="checkbox"/> Ataxia <input type="checkbox"/> Macrocephaly <input type="checkbox"/> Psychiatric disorder <input type="checkbox"/> Dysmorphic features <input type="checkbox"/> Microcephaly <input type="checkbox"/> Spasticity <input type="checkbox"/> Hearing disorder <input type="checkbox"/> Migraine <input type="checkbox"/> Vision disorder <input type="checkbox"/> Hypotonia <input type="checkbox"/> Movement disorder
--	--

Prior Testing: \_\_\_\_\_

**ORDERING PHYSICIAN/SENDING FACILITY (Each listed person will receive a copy of the report)**

Facility Name (Facility Code)	Address	City	State /Country	Zip	Phone
Ordering Licensed Provider Name (Last, First)(Code)	NPI#	Phone	Fax/Email		
Genetic Counselor or Other Medical Provider Name (Last, First) (Code)		Phone/Fax/Email			

**CONFIRMATION OF INFORMED CONSENT, PRE-TEST GENETIC COUNSELING, AND MEDICAL NECESSITY FOR GENETIC TESTING**

The undersigned person (or representative thereof) ensures he/she is a licensed medical professional authorized to order genetic testing and confirms that the patient has given appropriate consent. I confirm that testing is medically necessary and that test results may impact medical management for the patient. I agree to allow Ambry Genetics to facilitate the provision of pre-test genetic counseling services by a third-party service, as required by the patient's insurance provider. Furthermore, all information on this TRF is true to the best of my knowledge. My signature applies to the attached letter of medical necessity.

**Signature Required for Processing   Medical Professional Signature:** \_\_\_\_\_   **Date:** \_\_\_\_\_

**INSURANCE BILLING (Include copy of both sides of insurance card)**

Patient Relation to Policy Holder? <input type="checkbox"/> Self <input type="checkbox"/> Spouse <input type="checkbox"/> Child	Name and DOB of Policy Holder (if not self)	Facility Name <input type="checkbox"/> Send invoice to facility address above
Insurance Company	Policy #   HMO Auth #	Address

Out Of Pocket: We will start testing immediately and will attempt to contact the patient if their estimated out-of-pocket costs are > USD \$100.	Contact Name
Special Billing Notes:	Phone Number   E-mail/Fax

<input type="checkbox"/> <b>PATIENT PAYMENT</b>	<input type="checkbox"/> Check (Payable to Ambry Genetics) <input type="checkbox"/> Credit Card (Call 949-900-5795)
---	--

**Patient Acknowledgement:** I acknowledge that the information provided by me is true and correct. For direct insurance billing: I authorize my insurance benefits to be paid directly to Ambry Genetics Corporation (Ambry), authorize Ambry to release medical information concerning my testing to my insurer, to be my designated representative for purposes of appealing any denial of benefits as needed and to request additional medical records for this purpose. I understand that I am financially responsible for any amounts not covered by my insurer and responsible for sending Ambry money received from my health insurance company. For patient payment by credit card: I hereby authorize Ambry Genetics Corporation to bill my credit card as indicated above. In order to expedite consideration for eligibility for Ambry's Patient Assistance Program, please provide the total annual gross household income: \$ \_\_\_\_\_ and the number of family members in the household supported by the listed income: \_\_\_\_\_. I authorize Ambry Genetics Corporation to verify the above information for the sole purpose of assessing financial need, including the right to seek supporting documentation.

**FOR NY RESIDENTS:**  I am a New York resident and I give Ambry Genetics permission to store my sample for longer than 60 days. **NOTE:** If left blank, consent is interpreted as "NO".

**Signature Required For Insurance/Self-Pay Patients and NY Sample Storage Consent:** Patient Signature \_\_\_\_\_   **Date:** \_\_\_\_\_

# Neurology Test Requisition Form

<input type="checkbox"/> <b>Opt-in to Reporting of Variants of Unknown Significance (VUS)</b> <i>Checking this box indicates that VUS identified on the test(s) ordered below will be reported for this patient. If you do not check this box, VUS will not be reported.</i>	<input type="checkbox"/> <b>Opt-out of Patient For Life</b> <i>Check this box to opt-out of the Ambry Genetics Patient for Life program. If you check this box, Ambry will NOT continually review data for this patient to screen for potential pathogenic or likely pathogenic variants in newly added genes and will NOT proactively issue gene-level reclassification reports.</i>																																																																																																																																
<b>For Reflex or Concurrent Testing:</b> Test 1: _____ <input type="checkbox"/> Reflex to _____ Test 2: _____ <input type="checkbox"/> Concurrent with _____	<table border="1" style="width:100%; border-collapse: collapse;"> <thead> <tr> <th style="width:5%;">Check</th> <th style="width:65%;">Test Name</th> <th style="width:10%;">Test Code</th> <th style="width:20%;">Description</th> </tr> </thead> <tbody> <tr style="background-color: #00a0e3; color: white;"> <td colspan="4"><b>Migraine</b></td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Familial hemiplegic migraine</td> <td style="text-align: center;">6866</td> <td>ATP1A2, CACNA1A, PRRT2, SCN1A <input type="checkbox"/> Check if parental samples are included</td> </tr> <tr style="background-color: #00a0e3; color: white;"> <td colspan="4"><b>Neurodevelopmental Disorders</b></td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>AutismNext®</td> <td style="text-align: center;">6863</td> <td>72 genes for non-syndromic autism spectrum disorders and/or intellectual disability <input type="checkbox"/> Check if parental samples are included</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Autism, macrocephaly</td> <td style="text-align: center;">2106</td> <td>PTEN</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Fragile X syndrome</td> <td style="text-align: center;">4544</td> <td>FMR1 repeat expansion analysis and methylation studies</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>NeurodevelopmentNext™</td> <td style="text-align: center;">6861</td> <td>202 genes known to cause developmental delay, intellectual disability, and/or autism spectrum disorders <input type="checkbox"/> Check if parental samples are included</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>NeurodevelopmentNext-Expanded™</td> <td style="text-align: center;">6860</td> <td>&gt;1,400 genes associated with neonatal to childhood onset developmental delay, seizures, intellectual disability, developmental regression, autism spectrum disorders <input type="checkbox"/> Check if parental samples are included</td> </tr> <tr style="background-color: #00a0e3; color: white;"> <td colspan="4"><b>Hereditary Neuropathy</b></td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Familial transthyretin amyloidosis</td> <td style="text-align: center;">1560</td> <td>TTR</td> </tr> <tr style="background-color: #00a0e3; color: white;"> <td colspan="4"><b>Neurocutaneous/Neuro-Oncology Disorders</b></td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Ataxia-telangiectasia</td> <td style="text-align: center;">9014</td> <td>ATM</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>BrainTumorNext®</td> <td style="text-align: center;">8847</td> <td>27 genes for brain tumors</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>HHTNext®</td> <td style="text-align: center;">8672</td> <td>ACVRL1, ENG, SMAD4, GDF2, RASA1</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Legius syndrome</td> <td style="text-align: center;">5724</td> <td>SPRED1</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Li-Fraumeni syndrome</td> <td style="text-align: center;">2866</td> <td>TP53</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Neurofibromatosis 1</td> <td style="text-align: center;">5704</td> <td>NF1</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Neurofibromatosis 2</td> <td style="text-align: center;">9024</td> <td>NF2</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Nevoid basal cell carcinoma syndrome/Gorlin syndrome</td> <td style="text-align: center;">5684</td> <td>PTCH1</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Schwannomatosis</td> <td style="text-align: center;">7180</td> <td>SMARCB1</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Tuberous sclerosis</td> <td style="text-align: center;">5904</td> <td>TSC1, TSC2</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>von Hippel-Lindau</td> <td style="text-align: center;">2606</td> <td>VHL</td> </tr> <tr style="background-color: #00a0e3; color: white;"> <td colspan="4"><b>SINGLE SITE ANALYSIS (Please include a copy of relative's report)</b></td> </tr> <tr> <td colspan="2">Gene(s): _____</td> <td colspan="2">Mutation(s): _____</td> </tr> <tr> <td colspan="4">Relative Name: _____</td> </tr> <tr> <td colspan="4">Relationship to Relative: _____ Accession # (If tested at Ambry): _____</td> </tr> <tr> <td colspan="4">Positive control sample: <input type="checkbox"/> Will be provided <input type="checkbox"/> Already at Ambry <input type="checkbox"/> Not available</td> </tr> <tr style="background-color: #00a0e3; color: white;"> <td colspan="4"><b>FOR PRENATAL SPECIMENS, POC OR CORD BLOOD: MATERNAL CELL CONTAMINATION ANALYSIS REQUIRED</b></td> </tr> <tr style="background-color: #00a0e3; color: white;"> <td colspan="4"><b>Both test codes required for fetal specimens.</b></td> </tr> <tr> <td colspan="4"><input type="checkbox"/> 1260 MCC for fetal specimen or cord blood</td> </tr> <tr> <td colspan="4"><input type="checkbox"/> 1262 MCC Reference for maternal blood sample (No Charge)</td> </tr> </tbody> </table>	Check	Test Name	Test Code	Description	<b>Migraine</b>				<input type="checkbox"/>	Familial hemiplegic migraine	6866	ATP1A2, CACNA1A, PRRT2, SCN1A <input type="checkbox"/> Check if parental samples are included	<b>Neurodevelopmental Disorders</b>				<input type="checkbox"/>	AutismNext®	6863	72 genes for non-syndromic autism spectrum disorders and/or intellectual disability <input type="checkbox"/> Check if parental samples are included	<input type="checkbox"/>	Autism, macrocephaly	2106	PTEN	<input type="checkbox"/>	Fragile X syndrome	4544	FMR1 repeat expansion analysis and methylation studies	<input type="checkbox"/>	NeurodevelopmentNext™	6861	202 genes known to cause developmental delay, intellectual disability, and/or autism spectrum disorders <input type="checkbox"/> Check if parental samples are included	<input type="checkbox"/>	NeurodevelopmentNext-Expanded™	6860	>1,400 genes associated with neonatal to childhood onset developmental delay, seizures, intellectual disability, developmental regression, autism spectrum disorders <input type="checkbox"/> Check if parental samples are included	<b>Hereditary Neuropathy</b>				<input type="checkbox"/>	Familial transthyretin amyloidosis	1560	TTR	<b>Neurocutaneous/Neuro-Oncology Disorders</b>				<input type="checkbox"/>	Ataxia-telangiectasia	9014	ATM	<input type="checkbox"/>	BrainTumorNext®	8847	27 genes for brain tumors	<input type="checkbox"/>	HHTNext®	8672	ACVRL1, ENG, SMAD4, GDF2, RASA1	<input type="checkbox"/>	Legius syndrome	5724	SPRED1	<input type="checkbox"/>	Li-Fraumeni syndrome	2866	TP53	<input type="checkbox"/>	Neurofibromatosis 1	5704	NF1	<input type="checkbox"/>	Neurofibromatosis 2	9024	NF2	<input type="checkbox"/>	Nevoid basal cell carcinoma syndrome/Gorlin syndrome	5684	PTCH1	<input type="checkbox"/>	Schwannomatosis	7180	SMARCB1	<input type="checkbox"/>	Tuberous sclerosis	5904	TSC1, TSC2	<input type="checkbox"/>	von Hippel-Lindau	2606	VHL	<b>SINGLE SITE ANALYSIS (Please include a copy of relative's report)</b>				Gene(s): _____		Mutation(s): _____		Relative Name: _____				Relationship to Relative: _____ Accession # (If tested at Ambry): _____				Positive control sample: <input type="checkbox"/> Will be provided <input type="checkbox"/> Already at Ambry <input type="checkbox"/> Not available				<b>FOR PRENATAL SPECIMENS, POC OR CORD BLOOD: MATERNAL CELL CONTAMINATION ANALYSIS REQUIRED</b>				<b>Both test codes required for fetal specimens.</b>				<input type="checkbox"/> 1260 MCC for fetal specimen or cord blood				<input type="checkbox"/> 1262 MCC Reference for maternal blood sample (No Charge)			
Check	Test Name	Test Code	Description																																																																																																																														
<b>Migraine</b>																																																																																																																																	
<input type="checkbox"/>	Familial hemiplegic migraine	6866	ATP1A2, CACNA1A, PRRT2, SCN1A <input type="checkbox"/> Check if parental samples are included																																																																																																																														
<b>Neurodevelopmental Disorders</b>																																																																																																																																	
<input type="checkbox"/>	AutismNext®	6863	72 genes for non-syndromic autism spectrum disorders and/or intellectual disability <input type="checkbox"/> Check if parental samples are included																																																																																																																														
<input type="checkbox"/>	Autism, macrocephaly	2106	PTEN																																																																																																																														
<input type="checkbox"/>	Fragile X syndrome	4544	FMR1 repeat expansion analysis and methylation studies																																																																																																																														
<input type="checkbox"/>	NeurodevelopmentNext™	6861	202 genes known to cause developmental delay, intellectual disability, and/or autism spectrum disorders <input type="checkbox"/> Check if parental samples are included																																																																																																																														
<input type="checkbox"/>	NeurodevelopmentNext-Expanded™	6860	>1,400 genes associated with neonatal to childhood onset developmental delay, seizures, intellectual disability, developmental regression, autism spectrum disorders <input type="checkbox"/> Check if parental samples are included																																																																																																																														
<b>Hereditary Neuropathy</b>																																																																																																																																	
<input type="checkbox"/>	Familial transthyretin amyloidosis	1560	TTR																																																																																																																														
<b>Neurocutaneous/Neuro-Oncology Disorders</b>																																																																																																																																	
<input type="checkbox"/>	Ataxia-telangiectasia	9014	ATM																																																																																																																														
<input type="checkbox"/>	BrainTumorNext®	8847	27 genes for brain tumors																																																																																																																														
<input type="checkbox"/>	HHTNext®	8672	ACVRL1, ENG, SMAD4, GDF2, RASA1																																																																																																																														
<input type="checkbox"/>	Legius syndrome	5724	SPRED1																																																																																																																														
<input type="checkbox"/>	Li-Fraumeni syndrome	2866	TP53																																																																																																																														
<input type="checkbox"/>	Neurofibromatosis 1	5704	NF1																																																																																																																														
<input type="checkbox"/>	Neurofibromatosis 2	9024	NF2																																																																																																																														
<input type="checkbox"/>	Nevoid basal cell carcinoma syndrome/Gorlin syndrome	5684	PTCH1																																																																																																																														
<input type="checkbox"/>	Schwannomatosis	7180	SMARCB1																																																																																																																														
<input type="checkbox"/>	Tuberous sclerosis	5904	TSC1, TSC2																																																																																																																														
<input type="checkbox"/>	von Hippel-Lindau	2606	VHL																																																																																																																														
<b>SINGLE SITE ANALYSIS (Please include a copy of relative's report)</b>																																																																																																																																	
Gene(s): _____		Mutation(s): _____																																																																																																																															
Relative Name: _____																																																																																																																																	
Relationship to Relative: _____ Accession # (If tested at Ambry): _____																																																																																																																																	
Positive control sample: <input type="checkbox"/> Will be provided <input type="checkbox"/> Already at Ambry <input type="checkbox"/> Not available																																																																																																																																	
<b>FOR PRENATAL SPECIMENS, POC OR CORD BLOOD: MATERNAL CELL CONTAMINATION ANALYSIS REQUIRED</b>																																																																																																																																	
<b>Both test codes required for fetal specimens.</b>																																																																																																																																	
<input type="checkbox"/> 1260 MCC for fetal specimen or cord blood																																																																																																																																	
<input type="checkbox"/> 1262 MCC Reference for maternal blood sample (No Charge)																																																																																																																																	
<table border="1" style="width:100%; border-collapse: collapse;"> <thead> <tr> <th style="width:5%;">Check</th> <th style="width:65%;">Test Name</th> <th style="width:10%;">Test Code</th> <th style="width:20%;">Description</th> </tr> </thead> <tbody> <tr style="background-color: #00a0e3; color: white;"> <td colspan="4"><b>Comprehensive Testing</b></td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>CustomNext-Neuro™</td> <td style="text-align: center;">9545</td> <td>Customizable test of up to 500 genes from neuro menu</td> </tr> <tr style="background-color: #00a0e3; color: white;"> <td colspan="4"><b>Epilepsy</b></td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>EpiRapid®</td> <td style="text-align: center;">6862</td> <td>22 epilepsy genes with treatment associations <input type="checkbox"/> Check if parental samples are included</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>EpilepsyNext®</td> <td style="text-align: center;">6864</td> <td>124 genes for epilepsy <input type="checkbox"/> Check if parental samples are included</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>EpilepsyNext-Expanded™</td> <td style="text-align: center;">6865</td> <td>&gt;890 genes associated with seizures, primarily with neonatal to childhood onset <input type="checkbox"/> Check if parental samples are included</td> </tr> <tr style="background-color: #00a0e3; color: white;"> <td colspan="4"><b>Clinical Genomics</b></td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>SNP Array <i>Buccal swab accepted</i></td> <td style="text-align: center;">5490</td> <td>Chromosomal microarray (&gt;2.6 million copy number probes and 750,000 SNP probes)</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Familial targeted microarray <i>Buccal swab accepted</i></td> <td style="text-align: center;">5495</td> <td>Paid option. Only available following SNP Array (5490) completed at Ambry. Incidental findings unrelated to the variant(s) detected in the proband, will NOT be reported. Name of proband tested at Ambry: _____</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>ExomeNext®-Proband</td> <td style="text-align: center;">9993</td> <td>Proband only exome sequencing Secondary Findings<sup>†</sup>: <input type="checkbox"/> Opt-out</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>ExomeNext®-Proband plus mtDNA</td> <td style="text-align: center;">9994</td> <td>Proband only exome sequencing plus mtDNA sequencing Secondary Findings<sup>†</sup>: <input type="checkbox"/> Opt-out</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>ExomeNext®-Duo</td> <td style="text-align: center;">9991</td> <td>Duo exome sequencing Secondary Findings<sup>†</sup>: <input type="checkbox"/> Opt-out</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>ExomeNext®-Duo plus mtDNA</td> <td style="text-align: center;">9992</td> <td>Duo exome sequencing plus mtDNA sequencing Secondary Findings<sup>†</sup>: <input type="checkbox"/> Opt-out</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>ExomeNext®-Trio</td> <td style="text-align: center;">9995</td> <td>Trio exome sequencing plus mtDNA sequencing Candidate (Novel) Genetic Etiologies: <input type="checkbox"/> Opt-out Secondary Findings<sup>†</sup>: <input type="checkbox"/> Opt-out</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>ExomeNext®-Trio plus mtDNA</td> <td style="text-align: center;">9996</td> <td>Trio exome sequencing plus mtDNA sequencing plus mtDNA sequencing Candidate (Novel) Genetic Etiologies: <input type="checkbox"/> Opt-out Secondary Findings<sup>†</sup>: <input type="checkbox"/> Opt-out</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>ExomeNext®-Rapid® <i>(Institutional billing or patient payment only)</i></td> <td style="text-align: center;">9999R</td> <td>Rapid Trio exome sequencing plus mtDNA sequencing Candidate (Novel) Genetic Etiologies: <input type="checkbox"/> Opt-out Secondary Findings<sup>†</sup>: <input type="checkbox"/> Opt-out</td> </tr> <tr> <td style="text-align: center;"><input type="checkbox"/></td> <td>Order through Ambry-Port ExomeNext®-Select</td> <td style="text-align: center;">9500</td> <td>Up to 500 gene custom exome sequencing test</td> </tr> </tbody> </table>	Check	Test Name	Test Code	Description	<b>Comprehensive Testing</b>				<input type="checkbox"/>	CustomNext-Neuro™	9545	Customizable test of up to 500 genes from neuro menu	<b>Epilepsy</b>				<input type="checkbox"/>	EpiRapid®	6862	22 epilepsy genes with treatment associations <input type="checkbox"/> Check if parental samples are included	<input type="checkbox"/>	EpilepsyNext®	6864	124 genes for epilepsy <input type="checkbox"/> Check if parental samples are included	<input type="checkbox"/>	EpilepsyNext-Expanded™	6865	>890 genes associated with seizures, primarily with neonatal to childhood onset <input type="checkbox"/> Check if parental samples are included	<b>Clinical Genomics</b>				<input type="checkbox"/>	SNP Array <i>Buccal swab accepted</i>	5490	Chromosomal microarray (>2.6 million copy number probes and 750,000 SNP probes)	<input type="checkbox"/>	Familial targeted microarray <i>Buccal swab accepted</i>	5495	Paid option. Only available following SNP Array (5490) completed at Ambry. Incidental findings unrelated to the variant(s) detected in the proband, will NOT be reported. Name of proband tested at Ambry: _____	<input type="checkbox"/>	ExomeNext®-Proband	9993	Proband only exome sequencing Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out	<input type="checkbox"/>	ExomeNext®-Proband plus mtDNA	9994	Proband only exome sequencing plus mtDNA sequencing Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out	<input type="checkbox"/>	ExomeNext®-Duo	9991	Duo exome sequencing Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out	<input type="checkbox"/>	ExomeNext®-Duo plus mtDNA	9992	Duo exome sequencing plus mtDNA sequencing Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out	<input type="checkbox"/>	ExomeNext®-Trio	9995	Trio exome sequencing plus mtDNA sequencing Candidate (Novel) Genetic Etiologies: <input type="checkbox"/> Opt-out Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out	<input type="checkbox"/>	ExomeNext®-Trio plus mtDNA	9996	Trio exome sequencing plus mtDNA sequencing plus mtDNA sequencing Candidate (Novel) Genetic Etiologies: <input type="checkbox"/> Opt-out Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out	<input type="checkbox"/>	ExomeNext®-Rapid® <i>(Institutional billing or patient payment only)</i>	9999R	Rapid Trio exome sequencing plus mtDNA sequencing Candidate (Novel) Genetic Etiologies: <input type="checkbox"/> Opt-out Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out	<input type="checkbox"/>	Order through Ambry-Port ExomeNext®-Select	9500	Up to 500 gene custom exome sequencing test	<p>† Secondary Findings: If box is left unchecked, the ACMG recommended list of Secondary Findings will be reported. Secondary Findings are not available for ExomeNext-Select orders.</p>																																																								
Check	Test Name	Test Code	Description																																																																																																																														
<b>Comprehensive Testing</b>																																																																																																																																	
<input type="checkbox"/>	CustomNext-Neuro™	9545	Customizable test of up to 500 genes from neuro menu																																																																																																																														
<b>Epilepsy</b>																																																																																																																																	
<input type="checkbox"/>	EpiRapid®	6862	22 epilepsy genes with treatment associations <input type="checkbox"/> Check if parental samples are included																																																																																																																														
<input type="checkbox"/>	EpilepsyNext®	6864	124 genes for epilepsy <input type="checkbox"/> Check if parental samples are included																																																																																																																														
<input type="checkbox"/>	EpilepsyNext-Expanded™	6865	>890 genes associated with seizures, primarily with neonatal to childhood onset <input type="checkbox"/> Check if parental samples are included																																																																																																																														
<b>Clinical Genomics</b>																																																																																																																																	
<input type="checkbox"/>	SNP Array <i>Buccal swab accepted</i>	5490	Chromosomal microarray (>2.6 million copy number probes and 750,000 SNP probes)																																																																																																																														
<input type="checkbox"/>	Familial targeted microarray <i>Buccal swab accepted</i>	5495	Paid option. Only available following SNP Array (5490) completed at Ambry. Incidental findings unrelated to the variant(s) detected in the proband, will NOT be reported. Name of proband tested at Ambry: _____																																																																																																																														
<input type="checkbox"/>	ExomeNext®-Proband	9993	Proband only exome sequencing Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out																																																																																																																														
<input type="checkbox"/>	ExomeNext®-Proband plus mtDNA	9994	Proband only exome sequencing plus mtDNA sequencing Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out																																																																																																																														
<input type="checkbox"/>	ExomeNext®-Duo	9991	Duo exome sequencing Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out																																																																																																																														
<input type="checkbox"/>	ExomeNext®-Duo plus mtDNA	9992	Duo exome sequencing plus mtDNA sequencing Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out																																																																																																																														
<input type="checkbox"/>	ExomeNext®-Trio	9995	Trio exome sequencing plus mtDNA sequencing Candidate (Novel) Genetic Etiologies: <input type="checkbox"/> Opt-out Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out																																																																																																																														
<input type="checkbox"/>	ExomeNext®-Trio plus mtDNA	9996	Trio exome sequencing plus mtDNA sequencing plus mtDNA sequencing Candidate (Novel) Genetic Etiologies: <input type="checkbox"/> Opt-out Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out																																																																																																																														
<input type="checkbox"/>	ExomeNext®-Rapid® <i>(Institutional billing or patient payment only)</i>	9999R	Rapid Trio exome sequencing plus mtDNA sequencing Candidate (Novel) Genetic Etiologies: <input type="checkbox"/> Opt-out Secondary Findings <sup>†</sup> : <input type="checkbox"/> Opt-out																																																																																																																														
<input type="checkbox"/>	Order through Ambry-Port ExomeNext®-Select	9500	Up to 500 gene custom exome sequencing test																																																																																																																														
<b>FAMILY HISTORY (Completion of this section is required for order including parental samples. If available, please also submit a 3-generation pedigree)</b> <table border="1" style="width:100%; border-collapse: collapse;"> <thead> <tr> <th style="width:20%;">Relative</th> <th style="width:30%;">Name</th> <th style="width:15%;">DOB</th> <th style="width:20%;">Affected status<sup>††</sup></th> <th style="width:15%;">Samples included?</th> </tr> </thead> <tbody> <tr> <td> </td> <td> </td> <td> </td> <td style="text-align: center;"><input type="checkbox"/> Yes <input type="checkbox"/> No</td> <td style="text-align: center;"><input type="checkbox"/></td> </tr> <tr> <td> </td> <td> </td> <td> </td> <td style="text-align: center;"><input type="checkbox"/> Yes <input type="checkbox"/> No</td> <td style="text-align: center;"><input type="checkbox"/></td> </tr> </tbody> </table> <p>†† If affected, please list symptoms and age at diagnosis: _____</p>		Relative	Name	DOB	Affected status <sup>††</sup>	Samples included?				<input type="checkbox"/> Yes <input type="checkbox"/> No	<input type="checkbox"/>				<input type="checkbox"/> Yes <input type="checkbox"/> No	<input type="checkbox"/>	<p><b>Note:</b> Gene lists for NeurodevelopmentNext-Expanded and EpilepsyNext-Expanded are regularly updated due to proactive review of current literature using an internal, peer-reviewed clinical validity scheme (Smith ED, Radtke K, Rossi M, et al. 2017 Human mutation 38(5):600-608). The patient's test report will include a list of genes evaluated. For up-to-date gene lists, visit ambrygen.com</p>																																																																																																																
Relative	Name	DOB	Affected status <sup>††</sup>	Samples included?																																																																																																																													
			<input type="checkbox"/> Yes <input type="checkbox"/> No	<input type="checkbox"/>																																																																																																																													
			<input type="checkbox"/> Yes <input type="checkbox"/> No	<input type="checkbox"/>																																																																																																																													

## ExomeNext Medical Necessity Form

### REQUIRED FOR INSURANCE ORDERS ONLY (NOT REQUIRED FOR CIGNA MEMBERS)

This form is ONLY required if you are requesting reflex to Exome sequencing and wish to have the patient's insurance billed. Please complete and submit with the TRF and a copy of clinical notes. This form replaces the Letter of Medical Necessity.

1. Has the patient had previous Whole Exome Sequencing (WES) performed?

- Yes, date performed: \_\_\_\_\_  
 No

2. Does this patient have a clinical presentation consistent with the following (select all that apply):

- Multiple abnormalities affecting unrelated organ systems (please specify): \_\_\_\_\_  
**OR two of the following:**  
 Abnormality affecting a single organ system(specify): \_\_\_\_\_  
 Significant intellectual disability, symptoms of a complex neurodevelopmental disorder (i.e. self-injurious behavior, reverse sleep-wake cycle, or seizure/epilepsy), or severe neuropsychiatric condition (e.g. schizophrenia, bipolar, Tourette syndrome)  
 Family history strongly implicating a genetic etiology (please specify findings and relationships)  
 Period of unexplained developmental regression (unrelated to autism or epilepsy)

3. Are the results of this WES test expected to directly influence this patient's medical management recommendations and clinical outcome?

- Yes (please describe): \_\_\_\_\_  
 No

4. Please describe the genetic tests that would be indicated if WES were NOT performed (i.e., single gene tests, gene panels, etc.):

- Chromosomal microarray  
 Single gene test(s): \_\_\_\_\_  
 Multigene panel(s): \_\_\_\_\_  
 Other genetic test(s): \_\_\_\_\_

5. Please describe follow-up procedures & frequency that would be needed if WES were NOT performed (i.e., lumbar puncture, imaging studies, brain MRI, etc.):

- Imaging study: \_\_\_\_\_  
 Surgery: \_\_\_\_\_  
 Biopsy: \_\_\_\_\_  
 Other: \_\_\_\_\_