

COMPLETE ENTIRE FORM AND SUBMIT CLINIC NOTES/PEDIGREE TO AVOID DELAYS

PATIENT INFORMATION						
Name (Last, First, MI)		Date of Birth (MM/DD/YY)	Date of Death (if applicable)	Phone Number/Email		
Address	City	State	Zip	Biological Sex <input type="checkbox"/> F <input type="checkbox"/> M	Ethnicity: <input type="checkbox"/> African American <input type="checkbox"/> Asian <input type="checkbox"/> Caucasian <input type="checkbox"/> Hispanic <input type="checkbox"/> Jewish (Ashkenazi) <input type="checkbox"/> Portuguese <input type="checkbox"/> Other:	
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 #		
<small>*Blood or saliva from patients with active/recent hematological disease will undergo additional review and may not be accepted in some cases. For these, cultured fibroblasts or fresh/fresh frozen normal tissue are preferred. See ambrygen.com/specimen-requirements for details. **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. *Only for Fragile X syndrome and chromosomal microarray.</small>						
Phlebotomy Services Request: <input type="checkbox"/> Phlebotomy draw <input type="checkbox"/> Insurance preverification first <input type="checkbox"/> Send kit to patient**						
<small>**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.</small>						
INDICATION(S) FOR TESTING						
ICD-10 code(s):			Will patient management be changed depending on the test results? <input type="checkbox"/> Yes <input type="checkbox"/> No			
PATIENT HISTORY <input type="checkbox"/> No personal history of neurological disease						
PLEASE SUPPLY CLINIC NOTES AND PEDIGREE If pregnant, due date:			Upcoming procedure date:			
Reasons for Testing:						
Birth and Neonatal History <input type="checkbox"/> N/A Gestational age at birth: _____ Birth weight: _____ Head circumference at birth (if available): _____ Developmental History <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 Seizure History <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: _____ Other History <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: _____			Other Features (choose all that apply) <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						
<small>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.</small>						
Signature Required for Processing Medical Professional Signature:				Date:		
INSURANCE BILLING (Include copy of both sides of insurance card)			INSTITUTIONAL BILLING			
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, unless you check the box below. We will attempt to contact you if your estimated out-of-pocket costs are > USD \$100. <input type="checkbox"/> Do not start testing until the patient approves payment terms regarding estimated out-of-pocket costs. By checking this box, I understand that there will be a delay in starting this test until Ambry is able to reach the patient to communicate OOP costs.			Contact Name			
Special Billing Notes: 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.			Phone Number		E-mail/Fax	
			<input type="checkbox"/> PATIENT PAYMENT		<input type="checkbox"/> Check (Payable to Ambry Genetics) <input type="checkbox"/> Credit Card (Call 949-900-5795)	
FOR NY RESIDENTS: <input type="checkbox"/> 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 - Page 2 of 4

PLEASE SUBMIT THE FOLLOWING WITH THE TRF:

1. Clinic Notes 2. Pedigree 3. Insurance Card

Please check the box next to the test(s) being ordered below. If this TRF is sent to Ambyr without or ahead of the sample, it will be treated as a preverification. If test ordered is different than the test preverified, we will honor what is on the TRF order form with the sample.

For multiple test orders, testing will be run concurrently (multiple tests initiated at the same time) unless otherwise specified. To order reflexive testing (second test starts pending first test outcome), please clearly indicate the order of reflexive tests in the notes section or next to the test check box. For reflex test orders, any positive findings (pathogenic/likely pathogenic) in the first test will be reported out to the clinician, and the requested second test will be canceled; all other findings will automatically reflex (including VUS).

Check to order	Test Name	Test Code	Description	Check to order	Test Name	Test Code	Description
Comprehensive Testing (Required: completed CustomNext-Neuro ordering form on page 3)				Migraine			
<input type="checkbox"/>	CustomNext-Neuro	9540	Up to 196 gene custom neurology test	<input type="checkbox"/>	Familial hemiplegic migraine	7035	ATP1A2, CACNA1A, PRRT2, SCN1A <input type="checkbox"/> Check if parental samples are included
Epilepsy				Neurodevelopmental Disorders			
<input type="checkbox"/>	EpiRapid® <i>blood only, no saliva</i>	7033	16 epilepsy genes with treatment associations	<input type="checkbox"/>	Rett syndrome	2026	MECP2
<input type="checkbox"/>	EpiRapid reflex to EpilepsyNext <i>blood only, no saliva</i>	7034	16 epilepsy genes with treatment associations, reflex to 100 genes for epilepsy	<input type="checkbox"/>	AutismNext®	7024	48 genes for syndromic and non-syndromic ASDs <input type="checkbox"/> Check if parental samples are included
<input type="checkbox"/>	EpiFirst-Fever®	7011	13 genes for febrile seizures <input type="checkbox"/> Check if parental samples are included	<input type="checkbox"/>	Autism, macrocephaly	2106	PTEN
<input type="checkbox"/>	EpiFirst-Focal®	7017	11 genes for non-lesional focal epilepsy <input type="checkbox"/> Check if parental samples are included	<input type="checkbox"/>	Fragile X syndrome	4544	FMRI repeat expansion analysis and methylation studies
<input type="checkbox"/>	EpiFirst-IS®	7013	17 genes for infantile spasms <input type="checkbox"/> Check if parental samples are included	<input type="checkbox"/>	IDNext®	7027	140 genes for syndromic and non-syndromic ID <input type="checkbox"/> Check if parental samples are included
<input type="checkbox"/>	EpilepsyNext®	7019	100 genes for epilepsy <input type="checkbox"/> Check if parental samples are included	<input type="checkbox"/>	Neurodevelopment-Expanded	7028	196 genes for ID, ASDs, and epilepsy <input type="checkbox"/> Check if parental samples are included
<input type="checkbox"/>	Neurodevelopment-Expanded®	7028	196 genes for epilepsy, ID, ASDs <input type="checkbox"/> Check if parental samples are included	Hereditary Neuropathy			
Clinical Genomics				<input type="checkbox"/>	Familial transthyretin amyloidosis	1560	TTR
<input type="checkbox"/>	SNP Array	5490	Chromosomal microarray (>2.6 million copy number probes and 750,000 SNP probes)	Neurocutaneous/Neuro-Oncology Disorders			
<input type="checkbox"/>	Familial targeted microarray	5495	Paid option. Only available following SNP Array (5490) completed at Ambyr. Incidental findings unrelated to the variant(s) detected in the proband, will NOT be reported. Name of proband tested at Ambyr: _____	<input type="checkbox"/>	Ataxia-telangiectasia	9014	ATM
<input type="checkbox"/>	ExomeNext-Proband	9993	Proband only exome sequencing Secondary Findings [^] : <input type="checkbox"/> Opt-out	<input type="checkbox"/>	BrainTumorNext®	8847	27 genes for brain tumors
<input type="checkbox"/>	ExomeNext-Proband plus mtDNA	9994	Proband only exome sequencing plus mtDNA sequencing Secondary Findings [^] : <input type="checkbox"/> Opt-out	<input type="checkbox"/>	HHTNext®	8672	ACVRL1, ENG, SMAD4, GDF2, RASA1
<input type="checkbox"/>	ExomeNext-Duo	9991	Duo exome sequencing <input type="checkbox"/> Opt-out of Candidate (novel) Genetic Etiologies	<input type="checkbox"/>	Legius syndrome	5724	SPRED1
<input type="checkbox"/>	ExomeNext-Duo plus mtDNA	9992	Duo exome sequencing <input type="checkbox"/> Opt-out of Candidate (novel) Genetic Etiologies	<input type="checkbox"/>	Li-Fraumeni syndrome	2866	TP53
<input type="checkbox"/>	ExomeNext-Trio	9995	Trio exome sequencing Candidate (Novel) Genetic Etiologies: <input type="checkbox"/> Opt-out Secondary Findings [^] : <input type="checkbox"/> Opt-out	<input type="checkbox"/>	Neurofibromatosis 1	5704	NF1
<input type="checkbox"/>	ExomeNext-Trio plus mtDNA	9996	Trio exome sequencing plus mtDNA sequencing Candidate (Novel) Genetic Etiologies: <input type="checkbox"/> Opt-out Secondary Findings [^] : <input type="checkbox"/> Opt-out	<input type="checkbox"/>	Neurofibromatosis 2	9024	NF2
<input type="checkbox"/>	Order through AP*	9500	Up to 500 gene custom exome sequencing test	<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
				SINGLE SITE ANALYSIS (Please include a copy of relative's report)			
				Gene(s): _____ Mutation(s): _____			
				Relative Name: _____			
				Relationship to Relative: _____ Accession # (If tested at Ambyr): _____			
				Positive control sample: <input type="checkbox"/> will be provided <input type="checkbox"/> already at Ambyr <input type="checkbox"/> not available			
				FOR PRENATAL SPECIMENS, POC OR CORD BLOOD: MATERNAL CELL CONTAMINATION ANALYSIS REQUIRED			
				Both test codes required for fetal specimens.			
				<input type="checkbox"/> 1260 MCC for fetal specimen or cord blood			
				<input type="checkbox"/> 1262 MCC Reference for maternal blood sample (No Charge)			

[^]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.

*AP is AmbyrPort, our online portal ambyr.com/ap

FAMILY HISTORY (Completion of this section is required for order including parental samples. If available, please also submit a 3-generation pedigree)

Relative	Name	DOB	Affected status*	Samples included?	Secondary findings**
			<input type="checkbox"/> yes <input type="checkbox"/> no	<input type="checkbox"/>	<input type="checkbox"/> opt in <input type="checkbox"/> opt out
			<input type="checkbox"/> yes <input type="checkbox"/> no	<input type="checkbox"/>	<input type="checkbox"/> opt in <input type="checkbox"/> opt out

*If affected, please list symptoms and age at diagnosis:

** If reflexing to whole exome sequencing, please indicate opt-in or opt-out of the ACMG Secondary Findings minimum list

CustomNext-Neuro (9540) Ordering Form - Page 3 of 4

(This form is ONLY required if ordering CustomNext-Neuro, test code 9540)

PATIENT INFORMATION			
Last Name	First Name	Middle Initial	DOB (MM/DD/YY)

CUSTOMIZE YOUR PANEL

To create your own panel, add all desired genes below **OR** select one of the following multi-gene tests and add or remove all desired genes below. All genes include sequencing and deletion/duplication analysis.

- | | | |
|---|--|---|
| <input type="checkbox"/> EpiFirst-Fever: 13 genes for febrile seizures | <input type="checkbox"/> EpilepsyNext: 100 genes for epilepsy | <input type="checkbox"/> IDNext: 140 genes for intellectual disability |
| <input type="checkbox"/> EpiFirst-IS: 17 genes for infantile spasms | <input type="checkbox"/> AutismNext: 48 genes for autism spectrum disorder | <input type="checkbox"/> Neurodevelopment-Expanded: All 196 genes in CustomNext-Neuro |
| <input type="checkbox"/> EpiFirst-Focal: 11 genes for non-lesional focal epilepsy | | |

A list of genes for each multi-gene test is available on ambrygen.com

- Rett/Angelman syndrome: 22 genes associated with Rett syndrome, Angelman syndrome, and phenocopies (ARX, ATRX, CDKL5, CNTNAP2, DYRK1A, EHMT1, FOXG1, IQSEC2, MBD5, MECP2, MEF2C, NRXN1, PCDH19, PNKP, SATB2, SHANK3, SLC2A1, SLC9A6, STXBP1, TCF4, UBE3A, ZEB2)
- Batten disease: 13 genes associated with NCLs, or Batten disease (ATP13A2, CLN3, CLN5, CLN6, CLN8, CTSD, CTSF, DNAJC5, GRN, KCTD7, MFSD8, PPT1, TPP1)

CUSTOM GENE SELECTION (Check "+" to add and "-" to remove)													
--	--	--	--	--	--	--	--	--	--	--	--	--	--

+	GENE	-	+	GENE	-	+	GENE	-	+	GENE	-	+	GENE	-
<input type="checkbox"/>	ABCD1	<input type="checkbox"/>	<input type="checkbox"/>	CREBBP	<input type="checkbox"/>	<input type="checkbox"/>	GNAO1	<input type="checkbox"/>	<input type="checkbox"/>	MAOA	<input type="checkbox"/>	<input type="checkbox"/>	PNPO	<input type="checkbox"/>
<input type="checkbox"/>	ACSL4	<input type="checkbox"/>	<input type="checkbox"/>	CRH	<input type="checkbox"/>	<input type="checkbox"/>	GOSR2	<input type="checkbox"/>	<input type="checkbox"/>	MBD5	<input type="checkbox"/>	<input type="checkbox"/>	POGZ	<input type="checkbox"/>
<input type="checkbox"/>	ADNP	<input type="checkbox"/>	<input type="checkbox"/>	CSTB	<input type="checkbox"/>	<input type="checkbox"/>	GPC3	<input type="checkbox"/>	<input type="checkbox"/>	MECP2	<input type="checkbox"/>	<input type="checkbox"/>	POLG	<input type="checkbox"/>
<input type="checkbox"/>	ALDH7A1	<input type="checkbox"/>	<input type="checkbox"/>	CTCF	<input type="checkbox"/>	<input type="checkbox"/>	GRIA3	<input type="checkbox"/>	<input type="checkbox"/>	MED12	<input type="checkbox"/>	<input type="checkbox"/>	PORCN	<input type="checkbox"/>
<input type="checkbox"/>	ALG13	<input type="checkbox"/>	<input type="checkbox"/>	CTSD	<input type="checkbox"/>	<input type="checkbox"/>	GRIN1	<input type="checkbox"/>	<input type="checkbox"/>	MED23	<input type="checkbox"/>	<input type="checkbox"/>	PPT1	<input type="checkbox"/>
<input type="checkbox"/>	ANKRD11	<input type="checkbox"/>	<input type="checkbox"/>	CTSF	<input type="checkbox"/>	<input type="checkbox"/>	GRIN2A	<input type="checkbox"/>	<input type="checkbox"/>	MEF2C	<input type="checkbox"/>	<input type="checkbox"/>	PQBP1	<input type="checkbox"/>
<input type="checkbox"/>	AP1S2	<input type="checkbox"/>	<input type="checkbox"/>	CUL4B	<input type="checkbox"/>	<input type="checkbox"/>	GRIN2B	<input type="checkbox"/>	<input type="checkbox"/>	MFSD8	<input type="checkbox"/>	<input type="checkbox"/>	PRICKLE1	<input type="checkbox"/>
<input type="checkbox"/>	AP4B1	<input type="checkbox"/>	<input type="checkbox"/>	DCX	<input type="checkbox"/>	<input type="checkbox"/>	GRN	<input type="checkbox"/>	<input type="checkbox"/>	MID1	<input type="checkbox"/>	<input type="checkbox"/>	PRRT2	<input type="checkbox"/>
<input type="checkbox"/>	ARHGEF9	<input type="checkbox"/>	<input type="checkbox"/>	DDX3X	<input type="checkbox"/>	<input type="checkbox"/>	HCN1	<input type="checkbox"/>	<input type="checkbox"/>	NDP	<input type="checkbox"/>	<input type="checkbox"/>	PTCHD1	<input type="checkbox"/>
<input type="checkbox"/>	ARID1B	<input type="checkbox"/>	<input type="checkbox"/>	DEPDC5	<input type="checkbox"/>	<input type="checkbox"/>	HDAC8	<input type="checkbox"/>	<input type="checkbox"/>	NDUFA1	<input type="checkbox"/>	<input type="checkbox"/>	PTEN	<input type="checkbox"/>
<input type="checkbox"/>	ARX	<input type="checkbox"/>	<input type="checkbox"/>	DHCR7	<input type="checkbox"/>	<input type="checkbox"/>	HNRNPU	<input type="checkbox"/>	<input type="checkbox"/>	NHLRC1	<input type="checkbox"/>	<input type="checkbox"/>	PTPN11	<input type="checkbox"/>
<input type="checkbox"/>	ATP13A2	<input type="checkbox"/>	<input type="checkbox"/>	DLG3	<input type="checkbox"/>	<input type="checkbox"/>	HOXA1	<input type="checkbox"/>	<input type="checkbox"/>	NHS	<input type="checkbox"/>	<input type="checkbox"/>	PURA	<input type="checkbox"/>
<input type="checkbox"/>	ATPIA2	<input type="checkbox"/>	<input type="checkbox"/>	DNAJC5	<input type="checkbox"/>	<input type="checkbox"/>	HPRT1	<input type="checkbox"/>	<input type="checkbox"/>	NIPBL	<input type="checkbox"/>	<input type="checkbox"/>	RAB39B	<input type="checkbox"/>
<input type="checkbox"/>	ATP7A	<input type="checkbox"/>	<input type="checkbox"/>	DNM1	<input type="checkbox"/>	<input type="checkbox"/>	HUWE1	<input type="checkbox"/>	<input type="checkbox"/>	NLGN3	<input type="checkbox"/>	<input type="checkbox"/>	RAD21	<input type="checkbox"/>
<input type="checkbox"/>	ATRX	<input type="checkbox"/>	<input type="checkbox"/>	DYNC1H1	<input type="checkbox"/>	<input type="checkbox"/>	IDS	<input type="checkbox"/>	<input type="checkbox"/>	NLGN4X	<input type="checkbox"/>	<input type="checkbox"/>	RAI1	<input type="checkbox"/>
<input type="checkbox"/>	BRWD3	<input type="checkbox"/>	<input type="checkbox"/>	DYRK1A	<input type="checkbox"/>	<input type="checkbox"/>	IQSEC2	<input type="checkbox"/>	<input type="checkbox"/>	NRXN1	<input type="checkbox"/>	<input type="checkbox"/>	RPL10	<input type="checkbox"/>
<input type="checkbox"/>	CA8	<input type="checkbox"/>	<input type="checkbox"/>	EEF1A2	<input type="checkbox"/>	<input type="checkbox"/>	KAT6A	<input type="checkbox"/>	<input type="checkbox"/>	NSD1	<input type="checkbox"/>	<input type="checkbox"/>	RPS6KA3	<input type="checkbox"/>
<input type="checkbox"/>	CACNA1A	<input type="checkbox"/>	<input type="checkbox"/>	EHMT1	<input type="checkbox"/>	<input type="checkbox"/>	KATNAL2	<input type="checkbox"/>	<input type="checkbox"/>	NSUN2	<input type="checkbox"/>	<input type="checkbox"/>	SATB2	<input type="checkbox"/>
<input type="checkbox"/>	CACNA1C	<input type="checkbox"/>	<input type="checkbox"/>	EPM2A	<input type="checkbox"/>	<input type="checkbox"/>	KCNA2	<input type="checkbox"/>	<input type="checkbox"/>	OCRL	<input type="checkbox"/>	<input type="checkbox"/>	SCARB2	<input type="checkbox"/>
<input type="checkbox"/>	CASK	<input type="checkbox"/>	<input type="checkbox"/>	FGD1	<input type="checkbox"/>	<input type="checkbox"/>	KCNC1	<input type="checkbox"/>	<input type="checkbox"/>	OFD1	<input type="checkbox"/>	<input type="checkbox"/>	SCN1A	<input type="checkbox"/>
<input type="checkbox"/>	CC2D1A	<input type="checkbox"/>	<input type="checkbox"/>	FLNA	<input type="checkbox"/>	<input type="checkbox"/>	KCNJ10	<input type="checkbox"/>	<input type="checkbox"/>	OPHN1	<input type="checkbox"/>	<input type="checkbox"/>	SCN1B	<input type="checkbox"/>
<input type="checkbox"/>	CDKL5	<input type="checkbox"/>	<input type="checkbox"/>	FMR1 *	<input type="checkbox"/>	<input type="checkbox"/>	KCNQ2	<input type="checkbox"/>	<input type="checkbox"/>	OTC	<input type="checkbox"/>	<input type="checkbox"/>	SCN2A	<input type="checkbox"/>
<input type="checkbox"/>	CHD2	<input type="checkbox"/>	<input type="checkbox"/>	FOLR1	<input type="checkbox"/>	<input type="checkbox"/>	KCNQ3	<input type="checkbox"/>	<input type="checkbox"/>	PACS1	<input type="checkbox"/>	<input type="checkbox"/>	SCN8A	<input type="checkbox"/>
<input type="checkbox"/>	CHD7	<input type="checkbox"/>	<input type="checkbox"/>	FOXG1	<input type="checkbox"/>	<input type="checkbox"/>	KCNT1	<input type="checkbox"/>	<input type="checkbox"/>	PAK3	<input type="checkbox"/>	<input type="checkbox"/>	SHANK3	<input type="checkbox"/>
<input type="checkbox"/>	CHD8	<input type="checkbox"/>	<input type="checkbox"/>	FOXP1	<input type="checkbox"/>	<input type="checkbox"/>	KCTD7	<input type="checkbox"/>	<input type="checkbox"/>	PCDH19	<input type="checkbox"/>	<input type="checkbox"/>	SIK1	<input type="checkbox"/>
<input type="checkbox"/>	CHRNA2	<input type="checkbox"/>	<input type="checkbox"/>	FOXP2	<input type="checkbox"/>	<input type="checkbox"/>	KDM5C	<input type="checkbox"/>	<input type="checkbox"/>	PDHA1	<input type="checkbox"/>	<input type="checkbox"/>	SLC13A5	<input type="checkbox"/>
<input type="checkbox"/>	CHRNA4	<input type="checkbox"/>	<input type="checkbox"/>	FTSJ1	<input type="checkbox"/>	<input type="checkbox"/>	KIAA2022	<input type="checkbox"/>	<input type="checkbox"/>	PHF6	<input type="checkbox"/>	<input type="checkbox"/>	SLC16A2	<input type="checkbox"/>
<input type="checkbox"/>	CHRN2	<input type="checkbox"/>	<input type="checkbox"/>	GABRA1	<input type="checkbox"/>	<input type="checkbox"/>	KIF1A	<input type="checkbox"/>	<input type="checkbox"/>	PHF8	<input type="checkbox"/>	<input type="checkbox"/>	SLC25A22	<input type="checkbox"/>
<input type="checkbox"/>	CLN3	<input type="checkbox"/>	<input type="checkbox"/>	GABRB3	<input type="checkbox"/>	<input type="checkbox"/>	L1CAM	<input type="checkbox"/>	<input type="checkbox"/>	PIGA	<input type="checkbox"/>	<input type="checkbox"/>	SLC2A1	<input type="checkbox"/>
<input type="checkbox"/>	CLN5	<input type="checkbox"/>	<input type="checkbox"/>	GABRG2	<input type="checkbox"/>	<input type="checkbox"/>	LAMP2	<input type="checkbox"/>	<input type="checkbox"/>	PIGN	<input type="checkbox"/>	<input type="checkbox"/>	SLC35A2	<input type="checkbox"/>
<input type="checkbox"/>	CLN6	<input type="checkbox"/>	<input type="checkbox"/>	GAMT	<input type="checkbox"/>	<input type="checkbox"/>	LG1	<input type="checkbox"/>	<input type="checkbox"/>	PLCB1	<input type="checkbox"/>	<input type="checkbox"/>	SLC6A1	<input type="checkbox"/>
<input type="checkbox"/>	CLN8	<input type="checkbox"/>	<input type="checkbox"/>	GATM	<input type="checkbox"/>	<input type="checkbox"/>	LINS	<input type="checkbox"/>	<input type="checkbox"/>	PLP1	<input type="checkbox"/>	<input type="checkbox"/>	SLC6A8	<input type="checkbox"/>
<input type="checkbox"/>	CNTNAP2	<input type="checkbox"/>	<input type="checkbox"/>	GDI1	<input type="checkbox"/>	<input type="checkbox"/>	MAN1B1	<input type="checkbox"/>	<input type="checkbox"/>	PNKP	<input type="checkbox"/>	<input type="checkbox"/>	SLC9A6	<input type="checkbox"/>

*FMR1 CGG repeat analysis as an additional test bill separately (test code 4544) **ADD**

Total Gene Count (REQUIRED: include total number of genes on your CustomNext-Neuro panel): _____

Notes:

ExomeNext Medical Necessity Form - Page 4 of 4

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: _____