

NULISaseq™ Neuro 220 Panel

Table 3: Performance Validation Data by Target. Targets added to Neuro 220 Panel, respective to the CNS 120 Panel, are highlighted in bold text.

Target	UniProt ID	Protein Name	Detectability (%)		CV (%)				LOD (NPQ) Median (IQR)	Cross Reactivity
			Plasma normal n=37	Plasma disease n=135	Plasma		CSF			
					Intra	Inter	Intra	Inter		
Abeta38	P05067	Amyloid-beta precursor protein	41.86	100	n/a	n/a	8.2	10.6	7.3 (1.5, 8.4)	< 1%
Abeta40	P05067	Amyloid-beta precursor protein	67.44	97.87	16.8	10.9	6.8	11.1	5.7 (0.0, 7.0)	< 1%
Abeta42	P05067	Amyloid-beta precursor protein	58.41	97.87	12.4	19.4	7.9	11.7	11.0 (10.7, 11.3)	< 1%
ACHE	P22303	Acetylcholinesterase	100.0	100.0	6.9	2.2	11.2	2.1	4.2 (3.1, 4.6)	n/a
AGRN	O00468	Agtrin	100.0	100.0	7.3	5.3	4.8	6.0	9.4 (9.2, 9.7)	n/a
ANXA5	P08758	Annexin A5	100.0	100.0	4.6	4.1	9.1	10.5	1.8 (1.2, 2.3)	n/a
APOA1	P02647	Apolipoprotein A-I	n/a	n/a	4.5	1.2	6.5	3.5	n/a	n/a
APOA2	P02652	Apolipoprotein A-II	n/a	n/a	5.1	5.2	4.9	3.8	n/a	< 1%
APOE	P02649	Apolipoprotein E	n/a	n/a	6.3	0.8	3.4	0.8	n/a	< 1%
APOE4	P02649	Apolipoprotein E	24.3	43.0	5.8	1.0	7.1	5.8	0.0 (0.0, 0.6)	< 1%
APOH	P02749	Beta-2-glycoprotein 1	n/a	n/a	12.0	15.7	4.8	7.0	n/a	n/a
AQP4	P55087	Aquaporin-4	100.0	100.0	8.1	14.0	4.7	8.0	7.8 (7.6, 8.1)	n/a
ARSA	P15289	Arylsulfatase A	97.30	100.0	4.2	5.0	8.1	3.5	9.5 (9.3, 9.6)	< 1%
ATXN2	Q99700	Ataxin 2	100.0	100.0	5.8	6.3	10.8	3.2	10.2 (9.9, 10.7)	n/a
ATXN3	P54252	Ataxin 3	100.0	100.0	4.6	9.3	8.8	8.9	7.0 (6.8, 7.8)	< 1%
AXL	P30530	AXL receptor tyrosine kinase	100.0	100.0	5.3	3.4	9.3	1.6	0.0 (0.0, 0.0)	n/a
BACE1	P56817	Beta-secretase 1	100.0	100.0	6.2	3.5	4.7	5.2	9.3 (9.0, 9.5)	n/a
BASP1	P80723	Brain abundant membrane attached signal protein 1	100.0	100.0	32.5	45.9	69.4	85.6	7.3 (6.7, 8.3)	n/a
BCAN	Q96GW7	Brevican core protein	100.0	100.0	11.8	8.9	5.7	10.3	0.0 (0.0, 0.0)	n/a
BD-MAPT	P10636	Microtubule-associated protein tau (Brain-Derived)	81.1	77.8	7.9	4.2	10.7	3.8	12.5 (12.4, 12.6)	< 1%
BD-pTau-181	P10636	Microtubule-associated protein tau (Brain-Derived)	100.0	100.0	8.0	9.4	9.8	4.7	11.2 (10.6, 11.6)	< 1%
BD-pTau-205	P10636	Microtubule-associated protein tau (Brain-Derived)	86.5	83.7	7.1	5.5	5.4	4.3	12.6 (12.4, 12.7)	n/a
BD-pTau-212	P10636	Microtubule-associated protein tau (Brain-Derived)	100.0	100.0	6.5	4.0	5.5	2.2	11.7 (10.8, 11.8)	n/a
BD-pTau-217	P10636	Microtubule-associated protein tau (Brain-Derived)	83.8	88.9	12.2	8.6	9.6	1.1	8.4 (8.1, 8.6)	< 1%
BD-pTau-231	P10636	Microtubule-associated protein tau (Brain-Derived)	62.2	68.1	5.8	6.4	4.3	5.7	13.2 (13.1, 13.2)	n/a
BDNF	P23560	Brain-derived neurotrophic factor	100.0	100.0	6.2	6.8	n/a	n/a	3.2 (0.0, 4.9)	< 1%
C1q	P02745, P02746, P02747	Complement C1q and subunits A, B, C	n/a	n/a	11.4	9.5	6.6	10.1	n/a	< 1%
CALB2	P22676	Calretinin	75.7	78.5	20.2	17.0	8.8	2.1	7.2 (6.8, 7.5)	n/a
CCL11	P51671	Eotaxin	100.0	100.0	13.4	8.1	n/a	n/a	6.6 (6.1, 6.9)	n/a
CCL13	Q99616	C-C motif chemokine 13	100.0	100.0	5.0	3.2	15.0	3.0	0.0 (0.0, 3.0)	< 1%
CCL17	Q92583	C-C motif chemokine 17	100.0	100.0	7.0	1.4	14.8	0.0	0.0 (0.0, 4.5)	n/a
CCL18	P55774	C-C motif chemokine 18	100.0	100.0	4.9	2.3	n/a	n/a	0.0 (0.0, 0.0)	n/a
CCL2	P13500	C-C motif chemokine 2	100.0	100.0	6.6	3.7	5.2	3.5	3.2 (0.0, 4.3)	n/a
CCL22	O00626	C-C motif chemokine 22	100.0	100.0	5.6	2.9	13.8	6.5	0.0 (0.0, 1.7)	n/a
CCL26	Q9Y258	C-C motif chemokine 26	100.0	100.0	6.2	2.0	14.1	3.4	6.0 (5.7, 6.3)	n/a
CCL3	P10147	C-C motif chemokine 3	100.0	100.0	4.4	2.5	4.1	2.4	3.4 (3.0, 3.8)	294.3% ^a
CCL4	P13236	C-C motif chemokine ligand 4	100.0	100.0	9.9	3.9	19.3	0.0	0.0 (0.0, 4.5)	1438.1% ^b
CCL5	P13501	C-C motif chemokine 18	100.0	100.0	9.0	5.3	n/a	n/a	3.8 (3.2, 4.2)	n/a
CD33	P20138	Myeloid cell surface antigen CD33	100.0	100.0	7.7	3.2	13.3	4.2	5.1 (4.2, 5.6)	n/a
CD40LG	P29965	CD40 ligand	100.0	100.0	6.3	3.9	n/a	n/a	0.0 (0.0, 0.0)	n/a
CD63	P08962	CD63 antigen	100.0	100.0	4.4	1.8	6.9	1.5	6.8 (6.7, 6.9)	< 1%
CGRP	P06881	Calcitonin gene-related peptide 1	18.9	25.2	25.4	40.6	28.5	24.8	9.2 (8.8, 9.7)	n/a
CHI3L1	P36222	Chitinase-3-like protein 1	100.0	100.0	6.4	2.3	4.6	2.4	6.1 (5.2, 6.7)	< 1%
CHIT1	Q13231	Chitotriosidase-1	95.35	100.0	8.9	3.9	9.5	3.8	0.0 (0.0, 0.0)	< 1%
CNDP1	Q96KN2	Beta-Ala-His dipeptidase	90.7	89.36	5.9	6.5	7.3	7.2	0.0 (0.0, 0.0)	< 1%
CNTN2	Q02246	Contactin-2	100.0	98.5	n/a	n/a	5.8	6.9	0.0 (0.0, 0.0)	n/a

NULISaseq™ Neuro 220 Panel

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Target	UniProt ID	Protein Name	Detectability (%)		CV (%)				LOD (NPQ) Median (IQR)	Cross Reactivity
			Plasma n=43	CSF n=47	Plasma		CSF			
					Intra	Inter	Intra	Inter		
CRH	P06850	Corticoliberin	97.3	99.3	13.3	7.2	7.8	10.7	8.9 (8.7, 9.4)	n/a
CRP	P02741	C-reactive protein	n/a	n/a	8.7	3.0	3.2	15.8	n/a	< 1%
CSF1R	P07333	Macrophase colony-stimulating factor 1 receptor	100.0	100.0	6.4	3.1	8.5	4.0	0.0 (0.0, 0.0)	n/a
CSF2	P04141	Granulocyte-macrophage colony-stimulating factor	97.3	100.0	6.5	4.7	7.7	2.7	9.8 (9.2, 10.1)	n/a
CST3	P01034	Cystatin-C	100.0	100.0	5.2	3.6	4.3	5.3	6.5 (6.2, 6.9)	n/a
CST5	P28325	Cystatin-D	100	99.3	6.9	32.7	n/a	n/a	5.9 (4.6, 6.8)	n/a
CTNNB1	P35222	Catenin beta-1	73.0	84.4	6.9	8.0	8.7	3.5	12.4 (12.2, 12.6)	n/a
CX3CL1	P78423	Fractalkine	100.0	100.0	5.2	4.3	6.7	4.7	6.0 (5.7, 6.5)	n/a
CXCL1	P09341	Growth-regulated alpha protein	100.0	95.74	6.5	14.9	10.7	15.4	8.4 (8.1, 8.6)	< 1%
CXCL10	P02778	C-X-C motif chemokine 10	100.0	100.0	8.0	2.9	5.8	5.9	0.0 (0.0, 0.0)	n/a
CXCL13	O43927	C-X-C motif chemokine 13	100.0	100.0	6.0	2.9	14.2	4.2	0.0 (0.0, 0.0)	n/a
CXCL8	P10145	Interleukin-8, IL8	100.0	100.0	5.0	11.3	3.8	10.7	7.5 (7.2, 7.9)	< 1%
DDC	P20711	Aromatic-L-amino-acid decarboxylase	100.0	100.0	3.6	2.3	8.0	1.6	2.9 (2.6, 3.1)	n/a
DKK1	O94907	Dickkopf-related protein 1	100.0	100.0	6.6	8.0	16.1	10.0	7.2 (6.6, 7.7)	< 1%
DLG4	P78352	Disks large homolog4	73.0	86.7	9.5	9.2	10.9	6.3	11.3 (10.1, 11.5)	n/a
DNM1L	O00429	Dynamin 1 like	81.1	91.9	9.1	7.7	12.3	7.5	7.7 (7.3, 8.0)	n/a
EDA2R	Q9HAV5	Tumor necrosis factor receptor superfamily member 27	86.5	91.9	14.3	5.3	14.3	8.8	8.3 (7.9, 8.6)	n/a
EFEMP1	Q12805	EGF-containing fibulin-like extracellular matrix protein 1	100.0	100.0	6.8	0.2	10.3	0.4	0.0 (0.0, 0.0)	n/a
EGFR	P00533	Epidermal growth factor receptor	100.0	100.0	6.7	4.3	16.6	6.6	0.0 (0.0, 0.0)	n/a
EIF4EBP1	Q13541	Eukaryotic translation initiation factor 4E-binding protein 1	100.0	100.0	6.6	5.7	16.4	10.3	8.9 (8.3, 9.3)	n/a
ENO2	P09104	Gamma-enolase	100.0	100.0	4.5	5.2	5.0	4.3	4.0 (3.1, 5.3)	n/a
F2R	P25116	Coagulation factor II thrombin receptor	100.0	100.0	7.5	5.9	15.3	3.9	7.6 (1.5, 8.5)	n/a
FABP3	P05413	Fatty acid-binding protein, heart	100.0	100.0	5.9	2.4	5.3	3.3	3.2 (2.4, 3.8)	n/a
FAM3B	P58499	Protein FAM3B	83.8	85.2	10.9	5.0	10.9	6.2	7.3 (6.7, 7.7)	< 1%
FCGR2A	P12318	Low affinity immunoglobulin gamma Fc region receptor II-a	100.0	98.5	10.5	4.2	27.9	7.5	0.0 (0.0, 0.0)	n/a
FCGR2B	P31994	Low affinity immunoglobulin gamma Fc region receptor II-b	21.6	19.3	17.1	15.5	8.8	21.0	9.4 (8.9, 9.7)	n/a
FCN2	Q15485	Ficolin-2	100.0	100.0	8.1	9.1	n/a	n/a	0.0 (0.0, 0.0)	< 1%
FGF2	P09038	Fibroblast growth factor 2	94.6	95.6	8.0	4.3	8.9	5.7	8.8 (8.7, 8.9)	n/a
FGF21	Q9NSA1	Fibroblast growth factor 21	100.0	100.0	4.6	3.2	15.4	0.0	4.5 (4.1, 5.1)	n/a
FLT1	P17948	Vascular endothelial growth factor receptor 1	100.0	100.0	6.4	3.4	8.7	6.0	8.3 (8.0, 8.4)	n/a
FOLR1	P15328	Folate receptor alpha	100.0	100.0	8.1	2.4	6.6	2.9	2.1 (0.0, 3.1)	n/a
GAP43	P17677	Neuromodulin	100.0	100.0	7.9	4.7	11.0	12.1	10.8 (10.4, 11.5)	n/a
GAS6	Q14393	Growth arrest-specific protein 6	100.0	100.0	9.8	6.4	12.7	7.6	0.0 (0.0, 0.0)	n/a
GBA	P04062	Glucosylceramidase beta	100.0	99.3	7.0	8.0	11.2	7.3	7.7 (7.4, 8.0)	< 1%
GDF15	Q99988	Growth/differentiation factor 15	100.0	100.0	5.2	4.6	5.5	1.9	8.9 (8.5, 12.4)	n/a
GDI1	P31150	Rab GDP dissociation inhibitor alpha	100.0	97.87	7.0	5.2	8.6	4.9	8.9 (8.4, 9.3)	n/a
GDNF	P39905	Glial cell line-derived neurotrophic factor	78.4	88.1	6.0	1.4	8.5	1.6	9.4 (9.2, 9.5)	< 1%
GFAP	P14136	Glial fibrillary acidic protein	100.0	100.0	7.8	2.5	4.6	3.1	6.7 (5.9, 7.3)	n/a
GLP1R	P43220	Glucagon-like peptide 1 receptor	100.0	100.0	8.8	5.0	7.7	4.5	8.5 (8.1, 8.8)	n/a
GOT1	P17174	Aspartate aminotransferase, cytoplasmic	100.0	100.0	6.4	1.0	7.3	2.7	8.0 (7.7, 8.3)	< 1%
GNPMB	Q14956	Transmembrane glycoprotein NMB	100.0	100.0	7.8	6.1	18.2	12.2	6.5 (5.9, 7.3)	n/a
GRN	P28799	Progranulin	100.0	100.0	6.3	3.0	19.1	10.5	7.3 (6.8, 7.5)	n/a
GSDMD	P57764	Gasdermin D	35.1	40.0	8.5	10.0	6.3	16.0	11.6 (11.4, 11.8)	n/a
GSDME	O60443	Gasdermin E	100.0	98.5	13.1	15.2	13.5	7.4	9.8 (9.3, 10.4)	n/a

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			Plasma n=43	CSF n=47	Plasma		CSF			
					Intra	Inter	Intra	Inter		
HBA1	P69905	Hemoglobin subunit alpha	100.0	100.0	5.3	2.9	12.3	4.9	1.6 (1.1, 2.0)	< 1%
HMOX1	P09601	Heme oxygenase 1	100.0	100.0	7.6	4.2	n/a	n/a	2.7 (0.0, 5.7)	n/a
HPGDS	O60760	Hematopoietic prostaglandin D synthase	100.0	100.0	8.9	7.3	14.3	1.9	9.9 (9.4, 10.3)	n/a
HSPB1	P04792	Heat Shock protein beta-1	100.0	100.0	9.9	3.4	20.3	0.0	0.0 (0.0, 0.7)	< 1%
HTT	P42858	Huntingtin	100.0	100.0	5.4	3.3	8.1	5.1	8.8 (8.5, 9.1)	< 1%
ICAM1	P05362	Intercellular adhesion molecule 1	100.0	100.0	8.4	3.9	n/a	n/a	0.0 (0.0, 0.0)	n/a
IFNG	P01579	Interferon gamma	100.0	100.0	6.2	3.3	11.6	30.6	8.9 (8.4, 9.4)	n/a
IGF1R	P08069	Insulin-like growth factor 1 receptor	100.0	100.0	5.1	1.4	4.9	1.0	11.4 (11.4, 11.5)	n/a
IGFBP7	Q16270	Insulin-like growth factor-binding protein 7	100.0	100.0	7.2	3.7	5.9	5.1	8.1 (7.8, 8.7)	n/a
IL10	P22301	Interleukin-10	100.0	100.0	6.9	2.2	11.5	1.5	6.2 (6.0, 6.6)	n/a
IL12p70	P29459_ P29460	Interleukin-12 subunit beta_Interleukin-12 subunit alpha	100.0	99.3	7.3	2.7	15.3	6.2	8.4 (8.0, 8.8)	n/a
IL13	P35225	Interleukin-13	100.0	100.0	6.4	10.4	7.0	8.7	10.0 (9.9, 10.2)	n/a
IL15	P40933	Interleukin-15	100.0	100.0	6.3	4.1	5.4	4.2	8.3(7.9, 8.7)	n/a
IL16	Q14005	Pro-interleukin-16	100.0	100.0	4.5	2.2	8.7	2.0	4.1 (3.7, 5.6)	n/a
IL17A	Q16552	Interleukin-17A	97.3	90.4	7.0	3.8	7.4	5.5	10.7 (10.5, 10.8)	n/a
IL18	Q14116	Interleukin-18	100.0	100.0	3.9	3.3	15.4	4.0	2.9 (1.4, 3.5)	n/a
IL1B	P01584	Interleukin-1 beta	51.4	64.4	10.0	3.0	16.7	0.0	5.3 (4.9, 5.8)	n/a
IL2	P60568	Interleukin-2	100.0	100.0	8.8	9.0	11.2	9.7	6.7 (6.4, 7.1)	n/a
IL33	O95760	Interleukin 33	91.9	89.6	10.0	8.0	15.0	8.2	7.2 (6.9, 7.8)	n/a
IL4	P05112	Interleukin-4	86.5	89.6	13.6	13.9	18.3	7.9	7.0 (6.7, 7.3)	n/a
IL5	P05113	Interleukin-5	100.0	100.0	9.7	13.2	6.7	13.3	6.3 (5.5, 6.7)	n/a
IL6	P05231	Interleukin-6	100.0	100.0	6.5	2.9	5.4	1.9	8.9 (8.5, 9.3)	n/a
IL6R	P08887	Interleukin-6 receptor subunit alpha	100.0	100.0	7.2	7.6	n/a	n/a	0.0 (0.0, 4.8)	n/a
IL7	P13232	Interleukin-7	100.0	100.0	6.0	6.6	11.2	4.6	8.7 (8.2, 8.8)	n/a
IL9	P15248	Interleukin-9	100.0	100.0	7.5	3.8	6.7	1.7	7.9 (0.0, 8.2)	n/a
ITGAV	P06756	Integrin subunit alpha V	100.0	100.0	6.3	6.2	10.3	18.0	9.8 (9.5, 10.2)	n/a
KDR	P35968	Vascular endothelial growth factor receptor 2	100.0	100.0	9.1	0.4	15.6	2.2	0.0 (0.0, 0.0)	< 1%
KLK6	Q92876	Kallikrein-6	100.0	100.0	11.2	6.1	4.8	8.9	7.1 (6.3, 7.9)	< 1%
L1CAM	P32004	Neural cell adhesion molecule L1	100.0	100.0	5.5	2.5	14.1	4.1	3.8 (2.4, 4.8)	n/a
LDLR	P01130	Low-density lipoprotein receptor	100.0	100.0	8.5	4.0	20.5	2.1	5.8 (0.0, 6.5)	n/a
LGALS3	P17931	Galectin-3	100.0	100.0	6.9	3.5	n/a	n/a	0.0 (0.0, 0.0)	< 1%
LRRK2	Q5S007	Leucine-rich repeat serine/threonine-protein kinase 2	100.0	100.0	6.0	10.4	14.2	9.5	9.1 (8.6, 9.9)	< 1%
MAG	P20916	Myelin associated glycoprotein	94.6	90.4	4.2	4.1	3.8	5.0	13.1 (13.0, 13.2)	n/a
MDH1	P40925	Malate dehydrogenase, cytoplasmic	100.0	100.0	4.8	2.6	8.9	6.7	3.1 (1.9, 3.7)	n/a
MDK	P21741	Midkine	97.3	95.6	6.5	5.0	7.4	7.4	12.8 (12.5, 12.9)	n/a
mHTT-exon1	P42858	Huntingtin	2.7	12.6	n/a	n/a	n/a	n/a	10.5 (10.1, 11.1)	< 1%
MME	P08473	Membrane metalloendopeptidase	97.3	100.0	8.5	5.3	n/a	n/a	2.2 (0.0, 4.0)	n/a
MMP10	P09238	Stromelysin-2	100.0	100.0	6.7	5.9	14.9	7.1	11.7 (8.5, 11.8)	n/a
MMP9	P14780	Matrix metalloproteinase-9	100.0	100.0	5.5	1.4	7.9	5.1	4.0 (3.3, 4.5)	< 1%
MOG	Q16653	Myelin-oligodendrocyte glycoprotein	100.0	100.0	7.5	4.7	4.0	6.8	0.0 (0.0, 5.3)	n/a
MSLN	Q13421	Mesothelin	100.0	100.0	4.6	3.8	11.9	3.0	5.8 (4.5, 6.5)	n/a
NCAM1	P13591	Neural cell adhesion molecule 1	100.0	100.0	5.6	0.6	6.4	0.9	0.0 (0.0, 0.0)	n/a
NEFH	P12036	Neurofilament heavy polypeptide	100.0	99.3	6.3	3.1	5.1	3.2	10.6 (10.5, 10.7)	< 1%
NEFL	P07196	Neurofilament light polypeptide	100.0	100.0	6.1	3.9	3.7	4.5	10.6 (10.5, 10.8)	n/a
NELL1	Q92832	Protein kinase C-binding protein NELL1	100.0	100.0	10.3	8.6	16.9	8.6	8.9 (8.7, 9.2)	< 1%
NFKB1	P19838	Nuclear factor NF-kappa-B p105 subunit	100.0	100.0	6.0	6.3	18.7	0.0	8.3 (7.5, 8.7)	n/a

Table 3: Performance Validation Data by Target. Targets added to Neuro 220 Panel, respective to the CNS 120 Panel, are highlighted in bold text.

Target	UniProt ID	Protein Name	Detectability (%)		CV (%)				LOD (NPQ) Median (IQR)	Cross Reactivity
			Plasma n=43	CSF n=47	Plasma		CSF			
					Intra	Inter	Intra	Inter		
NFKB2	Q00653	Nuclear factor kappa B subunit2	100.0	100.0	8.0	9.0	14.5	13.4	8.7 (8.1, 10.0)	n/a
NGF	P01138	Beta-nerve growth factor	100.0	100.0	5.3	1.8	8.7	1.4	10.2 (10.1, 10.4)	n/a
NLRP3	Q96P20	NLR family pyrin domain containing 3	51.4	61.5	5.9	4.5	5.7	5.2	10.0 (9.9, 10.2)	n/a
NPTX1	Q15818	Neuronal pentraxin-1	100.0	100.0	6.4	1.7	3.6	3.8	11.3 (11.2, 11.5)	n/a
NPTX2	P47972	Neuronal pentraxin-2	100.0	100.0	5.8	2.3	4.7	2.3	10.6 (10.3, 10.8)	n/a
NPTXR	O95502	Neuronal pentraxin receptor	100.0	100.0	7.9	5.2	6.1	6.3	8.8 (8.5, 9.0)	n/a
NPY	P01303	Neuropeptide Y	100.0	100.0	11.3	8.6	11.4	7.4	6.0 (5.8, 6.2)	n/a
NRGN	Q92686	Neurogranin	100.0	100.0	6.2	6.9	17.3	10.0	7.7 (7.4, 7.8)	n/a
NT-proBNP	P16860	N-terminal prohormone of bain natriuretic peptide	100.0	99.3	10.5	4.8	18.7	10.0	6.2 (5.9, 6.6)	< 1%
NTRK2	Q16620	BDNF/NT-3 growth factors receptor	100.0	100.0	6.7	5.2	10.2	12.2	7.9 (7.7, 8.2)	n/a
NTRK3	Q16288	NT-3 growth factor receptor	100.0	100.0	7.1	3.4	8.4	6.4	9.5 (9.2, 10.5)	n/a
Oligo-SNCA	P37840	Alpha-synuclein	67.6	76.3	12.8	10.7	27.3	19.0	2.9 (2.0, 3.4)	99.3% ^c
PAFAH1B3	Q15102	Platelet-activating factor acetylhydrolase IB subunit gamma	100.0	100.0	4.2	4.0	16.1	2.6	7.8 (7.6, 8.3)	n/a
PARK7	Q99497	Protein/nucleic acid deglycase DJ-1	94.6	98.5	8.5	21.5	14.4	7.7	9.3 (8.9, 9.8)	< 1%
PARP1	P09874	Poly [ADP-ribose] polymerase 1	37.8	45.9	5.2	5.9	5.0	4.1	9.1 (8.9, 11.0)	n/a
PDGFC	Q9NRA1	Platelet-derived growth factor C	100.0	100.0	5.2	4.6	12.0	3.4	6.5 (6.0, 6.9)	n/a
PDGFRB	P09619	Platelet-derived growth factor receptor beta	100.0	100.0	6.3	3.0	14.7	3.2	0.0 (0.0, 1.8)	n/a
PDLIM5	Q96HC4	PDZ and LIM domain 5	81.1	68.1	8.2	5.3	6.2	0.5	8.4 (8.3, 8.6)	n/a
PGF	P49763	Placenta growth factor	100.0	100.0	5.2	2.4	5.3	2.0	9.7 (9.6, 9.9)	n/a
PGK1	P00558	Phosphoglycerate kinase 1	100.0	100.0	4.6	4.3	11.4	12.7	10.2 (9.7, 10.4)	22.7% ^d
PLAUR	Q03405	Urokinase plaminogen activator surface receptor	100.0	100.0	5.1	2.8	7.3	3.0	5.8 (5.3, 6.2)	n/a
pLRRK2-S1292	Q5S007	Leucine-rich repeat serine/threonine-protein kinase 2	56.8	68.1	22.9	29.2	26.7	21.7	10.8 (10.5, 11.7)	n/a
POSTN	Q15063	Periostin	100.0	100.0	8.4	4.0	n/a	n/a	0.0 (0.0, 0.0)	n/a
PPBP	P02775	Platelet basic protein	n/a	n/a	6.4	1.0	5.6	5.2	n/a	n/a
pPRKN-S65	O60260	E3 ubiquitin-protein ligase parkin (Parkin)	27.0	28.9	21.3	130.2	12.3	22.0	9.3 (8.7, 9.6)	10.1%^e
PPY	P01298	Pancreatic prohormone	100.0	100.0	7.9	4.0	n/a	n/a	6.7 (5.5, 7.2)	n/a
pQ-ATXN3	P54252	Ataxin-3	78.4	79.3	9.3	23.0	30.4	31.9	8.6 (8.2, 9.0)	5.1%^f
pQ-HTT	P42858	Huntingtin	5.4	20.7	17.9	54.3	19.9	57.0	10.5 (10.0, 11.2)	1.8%^g
pRAB10-T73	P61026	Ras-related protein Rab-10	24.3	41.5	21.4	27.5	23.6	22.8	10.0 (9.7, 10.5)	< 1%
pRAB12-S106	Q61Q22	Ras-related protein Rab-12	21.6	37.0	52.2	17.4	18.4	21.2	11.6 (9.3, 11.8)	< 1%
pRAB29-T71	O14966	Ras-related protein Rab-7L1	37.8	39.3	18.9	116.6	23.9	26.8	6.1 (5.6, 6.40)	< 1%
PRDX5	P30044	Peroxiredoxin-5, mitochondrial	81.1	85.9	8.9	9.8	21.3	18.7	10.0 (9.7, 11.5)	n/a
PRDX6	P30041	Peroxiredoxin-6	100.0	100.0	4.8	6.8	10.1	4.2	5.9 (5.6, 6.0)	n/a
PRKN	O60260	E3 ubiquitin-protein ligase parkin	100.0	100.0	4.9	3.8	5.6	4.7	10.3 (5.0, 10.4)	< 1%
PROS1	P07225	Vitamin K-dependent protein S	n/a	n/a	6.0	1.5	4.7	1.8	n/a	n/a
PSEN1	P49768	Presenilin 1	100.0	100.0	9.5	10.8	15.9	11.5	8.4 (7.7, 8.7)	n/a
PSME1	Q06323	Proteasome activator complex subunit1	97.3	100.0	7.2	3.0	14.2	5.0	9.4 (8.0, 10.0)	n/a
pSNCA-S129	P37840	Alpha-synuclein	100.0	100.0	6.1	3.5	11.1	10.8	9.5 (9.4, 9.7)	< 1%
pTARDBP-S409	Q13148	TAR DNA-binding protein 43	100.0	100.0	3.4	2.8	4.1	2.6	8.6 (8.4, 8.8)	< 1%
pTau-181	P10636	Microtubule-associated protein tau	100.0	100.0	6.2	4.9	7.4	2.4	10.2 (9.8, 10.5)	< 1%
pTau-205	P10636	Microtubule-associated protein tau	100.0	100.0	6.5	7.9	5.8	5.7	9.4 (9.0, 10.5)	< 1%
pTau-212	P10636	Microtubule-associated protein tau	100.0	100.0	8.6	5.7	5.6	3.6	8.5 (8.0, 9.1)	< 1%
pTau-217	P10636	Microtubule-associated protein tau	100.0	97.0	13.6	8.1	8.2	0.2	7.7 (6.9, 8.2)	< 1%
pTau-231	P10636	Microtubule-associated protein tau	100.0	100.0	5.2	4.9	3.3	5.9	9.7 (9.5, 10.6)	< 1%
pTDP43-409	Q13148	TAR DNA-binding protein 43	73.0	80.7	7.7	13.5	10.9	8.7	13.0 (12.7, 13.2)	< 1%

Table 3: Performance Validation Data by Target. Targets added to Neuro 220 Panel, respective to the CNS 120 Panel, are highlighted in bold text.

Target	UniProt ID	Protein Name	Detectability (%)		CV (%)				LOD (NPQ) Median (IQR)	Cross Reactivity
			Plasma n=43	CSF n=47	Plasma		CSF			
					Intra	Inter	Intra	Inter		
PTN	P21246	Pleiotrophin	5.4	9.6	12.1	8.1	20.5	11.5	5.1 (4.8, 5.4)	n/a
PTPRS	Q13332	Receptor-type tyrosine-protein phosphate S	100.0	100.0	4.6	2.9	9.3	2.7	4.1 (3.7, 4.6)	n/a
RAB10	P161026	RAB10, member RAS oncogene family	100	100.0	6.1	4.2	8.4	6.9	9.4 (9.2, 9.6)	< 1%
RAB12	Q6IQ22	RAB12, member RAS oncogene family	62.2	83.7	14.4	17.9	28.9	16.9	10.0 (9.4, 10.5)	< 1%
RAB29	O14966	RAB29, member RAS oncogene family	100.0	100.0	5.4	2.4	10.3	2.4	6.7 (6.3, 7.1)	< 1%
REST	Q13127	RE1 silencing transcription factor	100.0	100.0	6.0	3.6	13.4	6.1	10.1 (9.8, 10.2)	n/a
RUVBL2	Q9Y230	RuvB like AAA ATPase 2	78.4	86.7	3.8	3.2	4.2	1.4	10.2 (10.1, 11.1)	n/a
S100A12	P80511	Protein S100-A12	100.0	100.0	6.6	9.2	10.4	12.4	5.1 (4.5, 5.5)	< 1%
S100B	P04271	S100 calcium binding protein B	100.0	100.0	6.3	4.8	5.7	2.6	7.5 (7.2, 7.6)	n/a
SAA1	P0DJH8	Serum amyloid A-1 protein	100.0	100.0	3.4	1.0	18.2	9.0	0.0 (0.0, 0.0)	< 1%
SERPINA3	P01011	Alpha-1-antichymotrypsin	n/a	n/a	6.8	3.4	2.7	3.8	n/a	< 1%
SFRP1	Q8N474	Secreted frizzled-related protein 1	100.0	96.3	7.1	10.4	7.4	8.3	12.6 (12.5, 12.8)	n/a
SFTPD	P35247	Pulmonary surfactant-associated protein D	100.0	100.0	7.0	3.5	11.7	2.5	6.4 (6.2, 6.7)	n/a
SLIT2	O94813	Slit homolog 2 protein	100.0	100.0	6.7	3.8	12.0	6.1	9.9 (9.6, 10.0)	< 1%
SMOC1	Q9H4F8	SPARC-related modular calcium-binding protein 1	100.0	100.0	9.5	4.2	20.9	11.1	7.9 (7.5, 8.2)	n/a
SNAP25	P60880	Synaptosomal-associated protein 25	10.8	5.2	3.5	0.7	3.8	1.5	13.2 (13.1, 13.2)	n/a
SNCA	P37840	Alpha-synuclein	100.0	100.0	4.7	3.8	12.6	6.2	4.4 (3.6, 5.0)	< 1%
SNCB	Q16143	Synuclein beta	5.4	16.3	16.2	33.7	11.1	3.6	10.0 (9.8, 10.1)	n/a
SOD1	P00441	Superoxide dismutase [Cu-Zn]	100.0	100.0	6.4	3.0	4.2	3.2	5.9 (5.3, 6.2)	< 1%
SPP1	P10451	Osteopontin	100.0	100.0	8.2	6.3	5.8	6.7	5.5 (4.8, 6.0)	< 1%
SQSTM1	Q13501	Sequestosome-1	100.0	100.0	7.6	3.2	9.3	3.1	5.5 (5.1, 6.1)	n/a
STMN2	Q93045	Stathmin-2	100.0	100.0	6.4	5.4	3.9	4.5	9.2 (9.0, 9.4)	< 1%
STX4	Q12846	Syntaxin-4	100.0	100.0	5.5	3.5	16.8	0.2	4.7 (3.4, 5.1)	n/a
TARDBP	Q13148	TAR DNA-binding protein 43	100.0	100.0	3.4	2.8	4.1	2.6	8.6 (8.4, 8.8)	n/a
TEK	Q02763	Angiopoietin-1 receptor	100.0	100.0	5.0	2.3	15.3	5.7	2.0 (0.0, 3.2)	n/a
TIMP3	P35625	Metalloproteinase inhibitor 3	100.0	100.0	5.7	6.1	16.3	5.4	0.0 (0.0, 5.0)	n/a
TNF	P01375	Tumor necrosis factor	100.0	100.0	6.0	3.6	15.7	3.6	8.8 (8.2, 9.1)	n/a
TNFSF14	O43557	Tumor necrosis factor ligand superfamily member 14	100.0	100.0	4.3	3.3	22.0	0.0	4.1 (3.5, 4.9)	n/a
TREM1	Q9NP99	Triggering receptor expressed on myeloid cells 1	100.0	100.0	5.5	3.7	14.7	5.2	6.5 (5.9, 7.3)	n/a
TREM2	Q9NZC2	Triggering receptor expressed on myeloid cells 2	100.0	100.0	6.0	1.9	6.8	3.0	0.0 (0.0, 0.0)	n/a
tTau (MAPT)	P10636	Microtubule-associated protein tau	100.0	100.0	7.9	4.7	8.4	1.5	8.9 (8.4, 9.7)	< 1%
UBB	P0CG47	Polyubiquitin-B	45.9	54.8	5.5	3.3	4.1	2.9	11.4 (11.3, 11.4)	n/a
UCHL1	P09936	Ubiquitin carboxyl-terminal hydrolase isozyme L1	23.3	31.9	1.1	0.0	6.2	2.9	12.4 (12.3, 12.6)	n/a
VCAM1	P19320	Vascular cell adhesion protein 1	100.0	100.0	5.4	4.1	16.7	5.0	0.0 (0.0, 0.0)	n/a
VEGFA	P15692	Vascular endothelial growth factor A	100.0	100.0	4.8	3.1	4.3	2.5	8.1 (7.9, 8.3)	< 1%
VEGFD	O43915	Vascular endothelial growth factor D	100.0	100.0	8.7	2.8	9.8	3.8	10.4 (10.3, 10.6)	n/a
VEGF	O15240	VEGF nerve growth factor inducible	100.0	100.0	6.8	5.8	5.2	4.4	8.3 (7.9, 9.0)	n/a
VSNL1	P62760	Visinin-like protein 1	100.0	100.0	8.0	5.6	4.8	5.1	9.4 (9.0, 9.6)	9.9% ^h
YWHAG	P61981	Tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein gamma	100.0	100.0	7.2	3.8	5.8	2.3	11.7 (11.4, 11.9)	< 1%
YWHAZ	P63104	14-3-3 protein zeta/delta	2.7	2.2	8.2	6.1	10.9	0.0	8.9 (8.7, 9.0)	n/a

a. Cross-reactivity with CCL3L1, which has 95% homology with CCL3. CCL3L1 is not in the Neuro 220 panel.
b. Cross-reactivity with CCL4L1, which has 97% homology with CCL4. CCL4L1 is not in the Neuro 220 panel.
c. Cross-reactivity with pSNCA, < 1% with unphosphorylated SNCA Ag. Potential aggregation of the pSNCA antigen during testing has not been assessed.
d. Cross-reactivity with PGK2, which has 88% homology with PGK1. PGK2 is not in the Neuro 220 panel.
e. Cross-reactivity with PRKN, but affinity against p-PRKN-S65 much higher.
f. Cross-reactivity with ATXN3. Due to uniqueness of this target (polyQ-length mutation), 1% criteria may not apply in this situation.
g. Cross-reactivity with HTT. Due to uniqueness of this target (polyQ-length mutation), 1% criteria may not apply in this situation.
h. Cross-reactivity with HPCAL4, which has 89% homology with VSNL1. HPCAL4 is not in the Neuro 220 panel.