AALBORG UNIVERSITETSHOSPITAL
– i gode hænder

Aalborg Universitetshospital
Nordjyllands Gensekventeringscenter
Reberbansgade 15, 9000 Aalborg

Hereditær neuropati basis panel

Panelbeskrivelse

Navn: Hereditær neuropati basis panel **version:** 2.0 **Ibrugtagningsdato:** 01/06-2023

Metodebeskrivelse: Analysen udføres ved helgenomsekventering (WGS) med Illumina PCR free library prep (tagmentation) og Illumina sekventering (Novaseq). Efterfølgende fortages in silico filtrering af data til relevante genregioner baseret på nedenstående tabel. Middelsekventeringsdybden er minimum 30X, og minimumsandelen af de kodede regioner der dækket minimum 10X er 90 %.

Genliste

Gener hvor middelsekventeringsdybden er < 30X er markeret med[1]. Gener, hvor andelen af de kodede regioner der dækket minimum 10X er < 90 % er markeret med[2]. Værdier er angivet med +/- standardafvigelse.

Gen	Middelsekventeringsdybde [x]	Andel >= 10x [%]	Andel >= 20x [%]	Andel >= 30x [%]
AARS1	52.2 +/- 8.16	100.0 +/- 0.0	99.99 +/- 0.09	98.83 +/- 2.63
ABCA1	53.85 +/- 7.91	100.0 +/- 0.0	99.99 +/- 0.05	99.26 +/- 2.45
ABHD12	50.52 +/- 8.43	100.0 +/- 0.01	99.74 +/- 0.8	96.96 +/- 6.38
AGTPBP1	55.11 +/- 7.36	100.0 +/- 0.0	100.0 +/- 0.0	99.45 +/- 1.0
AGXT	52.73 +/- 8.05	100.0 +/- 0.0	99.94 +/- 0.21	98.69 +/- 4.4
AIFM1	40.87 +/- 15.75	99.86 +/- 0.59	91.64 +/- 13.41	65.89 +/- 38.83
AP1S1	48.22 +/- 8.25	100.0 +/- 0.0	99.81 +/- 0.9	95.16 +/- 8.63
APOA1	54.64 +/- 8.92	100.0 +/- 0.0	99.98 +/- 0.12	99.12 +/- 4.13
APTX	53.48 +/- 7.91	100.0 +/- 0.0	100.0 +/- 0.01	99.62 +/- 0.82
ARHGEF10	54.2 +/- 7.9	100.0 +/- 0.02	99.78 +/- 0.61	98.09 +/- 3.29
ARSA	49.07 +/- 7.79	100.0 +/- 0.01	99.77 +/- 0.84	96.75 +/- 5.82
ATL1	54.35 +/- 7.55	100.0 +/- 0.0	100.0 +/- 0.0	99.42 +/- 1.26
ATL3	55.1 +/- 7.43	100.0 +/- 0.0	99.99 +/- 0.03	99.52 +/- 1.03
ATM	55.39 +/- 7.08	100.0 +/- 0.0	99.99 +/- 0.03	99.5 +/- 0.98
ATP1A1	52.95 +/- 7.56	100.0 +/- 0.0	99.94 +/- 0.25	98.33 +/- 3.65
ATP7A	43.34 +/- 15.44	99.96 +/- 0.18	95.8 +/- 8.34	72.24 +/- 33.0
B4GALNT1	49.56 +/- 8.18	100.0 +/- 0.0	99.92 +/- 0.46	97.45 +/- 7.28
BAG3	49.59 +/- 8.1	99.95 +/- 0.05	99.68 +/- 1.08	96.74 +/- 7.05
BCKDHB	54.92 +/- 7.0	99.94 +/- 0.05	99.92 +/- 0.04	99.18 +/- 1.9
BICD2	54.9 +/- 8.01	100.0 +/- 0.0	99.92 +/- 0.36	98.74 +/- 3.23
BSCL2	52.37 +/- 8.4	100.0 +/- 0.0	99.99 +/- 0.06	98.79 +/- 6.57
CD59	52.63 +/- 8.34	100.0 +/- 0.0	99.82 +/- 0.3	98.44 +/- 2.21
CFAP276	51.84 +/- 8.81	100.0 +/- 0.0	100.0 +/- 0.0	98.57 +/- 4.92
CHCHD10	45.02 +/- 7.99	100.0 +/- 0.0	99.88 +/- 0.44	92.29 +/- 13.11
CNTNAP1	49.7 +/- 8.08	100.0 +/- 0.0	99.83 +/- 0.52	96.55 +/- 6.11
COA7	53.39 +/- 7.6	100.0 +/- 0.0	100.0 +/- 0.0	99.15 +/- 2.42
COX6A1	54.37 +/- 7.93	100.0 +/- 0.0	100.0 +/- 0.0	99.63 +/- 1.08
CPOX	54.28 +/- 7.9	100.0 +/- 0.0	100.0 +/- 0.0	99.45 +/- 1.68
CPOX	54.01 +/- 6.99	100.0 +/- 0.0	99.77 +/- 1.08	97.26 +/- 4.85
CTDP1	56.34 +/- 8.42	100.0 +/- 0.03	99.78 +/- 0.7	98.42 +/- 3.27
CYP27A1	51.96 +/- 8.62	100.0 +/- 0.0	99.86 +/- 0.92	98.15 +/- 6.72

Gen	Middelsekventeringsdybde [x]	Andel >= 10x [%]	Andel >= 20x [%]	Andel >= 30x [%]
DARS2	52.97 +/- 7.53	100.0 +/- 0.0	99.98 +/- 0.12	99.09 +/- 2.37
DCAF8	52.16 +/- 7.74	100.0 +/- 0.0	100.0 +/- 0.03	98.73 +/- 2.59
DCTN1	51.49 +/- 8.28	100.0 +/- 0.0	99.94 +/- 0.39	98.03 +/- 6.63
DEGS1	57.17 +/- 7.68	100.0 +/- 0.0	99.97 +/- 0.19	99.38 +/- 1.78
DNAJB2	50.9 +/- 8.84	100.0 +/- 0.0	99.84 +/- 0.47	97.6 +/- 3.74
DNAJC3	54.53 +/- 7.34	99.93 +/- 0.15	99.9 +/- 0.17	99.24 +/- 2.07
DNM2	50.48 +/- 8.2	100.0 +/- 0.0	99.92 +/- 0.46	98.07 +/- 5.13
DNMT1	54.48 +/- 8.09	100.0 +/- 0.0	99.98 +/- 0.11	99.37 +/- 1.94
DRP2	40.01 +/- 15.38	99.79 +/- 0.7	90.98 +/- 15.25	65.78 +/- 39.11
DST	54.62 +/- 7.32	100.0 +/- 0.0	99.99 +/- 0.04	99.46 +/- 1.27
DYNC1H1	53.25 +/- 7.78	100.0 +/- 0.0	99.97 +/- 0.18	98.9 +/- 3.38
EGR2	51.81 +/- 8.39	100.0 +/- 0.0	99.96 +/- 0.3	97.17 +/- 7.48
ELP1	54.83 +/- 7.55	100.0 +/- 0.0	99.99 +/- 0.08	99.33 +/- 1.55
ERCC6	54.52 +/- 7.2	100.0 +/- 0.0	99.99 +/- 0.06	99.41 +/- 1.57
ERCC8	54.54 +/- 7.47	100.0 +/- 0.0	100.0 +/- 0.03	99.43 +/- 1.25
ETFDH	55.85 +/- 7.78	99.95 +/- 0.11	99.79 +/- 0.25	99.09 +/- 1.77
FAH	51.24 +/- 8.32	100.0 +/- 0.0	99.76 +/- 0.84	97.67 +/- 6.81
FBLN5	50.92 +/- 8.2	100.0 +/- 0.0	99.99 +/- 0.09	97.89 +/- 6.0
FBXO38	54.97 +/- 7.35	100.0 +/- 0.0	100.0 +/- 0.03	99.21 +/- 1.93
FGD4	54.36 +/- 7.4	100.0 +/- 0.0	100.0 +/- 0.02	99.24 +/- 1.7
FIG4	55.98 +/- 7.24	100.0 +/- 0.0	99.96 +/- 0.21	99.19 +/- 3.24
FLVCR1	54.93 +/- 7.3	99.83 +/- 0.23	99.66 +/- 0.39	98.61 +/- 2.83
FXN	53.74 +/- 7.71	99.99 +/- 0.03	99.96 +/- 0.14	99.15 +/- 2.07
GALC	56.72 +/- 7.98	99.97 +/- 0.05	99.96 +/- 0.05	99.54 +/- 1.38
GAN	54.67 +/- 7.4	100.0 +/- 0.0	100.0 +/- 0.01	99.36 +/- 1.66
GARS1	55.24 +/- 7.5	100.0 +/- 0.0	100.0 +/- 0.0	99.26 +/- 1.64
GBA2	50.05 +/- 8.63	100.0 +/- 0.0	99.77 +/- 1.07	96.91 +/- 7.5
GDAP1	55.29 +/- 7.74	100.0 +/- 0.0	100.0 +/- 0.0	99.52 +/- 1.58
GJB1	38.69 +/- 14.75	99.93 +/- 0.41	88.54 +/- 19.21	64.23 +/- 40.08
GJC2	51.68 +/- 10.23	100.0 +/- 0.0	99.86 +/- 0.31	95.11 +/- 7.27
GLA	43.48 +/- 15.86	99.99 +/- 0.04	95.69 +/- 8.38	73.06 +/- 32.79
GNB4	54.36 +/- 7.2	99.96 +/- 0.05	99.93 +/- 0.16	99.16 +/- 1.48
PRNP	51.12 +/- 8.53	100.0 +/- 0.0	99.96 +/- 0.28	98.29 +/- 5.37
HADHA	53.12 +/- 7.75	100.0 +/- 0.0	99.91 +/- 0.35	98.23 +/- 5.5
HADHB	55.78 +/- 7.5	100.0 +/- 0.0	100.0 +/- 0.0	99.73 +/- 0.95
HARS1	52.53 +/- 8.08	100.0 +/- 0.0	99.96 +/- 0.28	98.46 +/- 6.7
HINT1	54.88 +/- 8.13	100.0 +/- 0.0	100.0 +/- 0.0	99.14 +/- 3.44
HK1	52.73 +/- 8.56	100.0 +/- 0.0	99.93 +/- 0.39	98.5 +/- 5.51
HMBS	52.21 +/- 8.84	100.0 +/- 0.0	99.99 +/- 0.04	98.55 +/- 5.6
HSPB1	47.77 +/- 9.55	100.0 +/- 0.0	99.5 +/- 2.68	94.75 +/- 14.0
HSPB8	49.7 +/- 7.5	99.99 +/- 0.06	99.94 +/- 0.22	97.57 +/- 6.13
HYCC1	54.33 +/- 7.45	100.0 +/- 0.0	99.98 +/- 0.08	99.3 +/- 1.36
IARS2	55.74 +/- 7.11	100.0 +/- 0.0	99.98 +/- 0.07	99.38 +/- 1.42
IGHMBP2	53.51 +/- 8.33	100.0 +/- 0.0	99.99 +/- 0.05	98.59 +/- 4.46
INF2	49.37 +/- 8.46	99.51 +/- 0.58	97.2 +/- 2.36	90.6 +/- 8.15
JAG1	54.22 +/- 7.67	100.0 +/- 0.0	99.94 +/- 0.37	98.84 +/- 2.65

Gen	Middelsekventeringsdybde [x]	Andel >= 10x [%]	Andel >= 20x [%]	Andel >= 30x [%]
KCNA2	52.71 +/- 7.71	100.0 +/- 0.0	99.99 +/- 0.05	99.16 +/- 2.6
KIF1A	51.47 +/- 8.41	100.0 +/- 0.0	99.96 +/- 0.19	98.05 +/- 5.33
KIF1B	53.73 +/- 7.74	100.0 +/- 0.0	99.95 +/- 0.23	98.96 +/- 2.69
KIF5A	50.98 +/- 7.66	100.0 +/- 0.0	99.98 +/- 0.11	98.48 +/- 3.39
LITAF	54.0 +/- 8.01	100.0 +/- 0.0	99.94 +/- 0.29	99.19 +/- 2.21
LMNA	49.81 +/- 8.27	100.0 +/- 0.0	99.96 +/- 0.16	97.29 +/- 6.96
LRSAM1	51.12 +/- 7.79	100.0 +/- 0.0	99.95 +/- 0.22	97.72 +/- 5.2
LYST	55.53 +/- 7.3	100.0 +/- 0.0	100.0 +/- 0.0	99.59 +/- 1.01
MARS1	52.56 +/- 7.73	100.0 +/- 0.0	100.0 +/- 0.03	99.09 +/- 3.13
MCM3AP	54.28 +/- 7.88	100.0 +/- 0.0	99.98 +/- 0.12	99.26 +/- 2.55
MED25	47.8 +/- 8.31	100.0 +/- 0.0	99.95 +/- 0.2	95.72 +/- 10.84
MFN2	52.55 +/- 8.18	100.0 +/- 0.0	99.96 +/- 0.22	98.5 +/- 3.53
MMACHC	49.2 +/- 8.64	100.0 +/- 0.0	99.82 +/- 0.84	96.32 +/- 8.15
MME	55.24 +/- 7.7	100.0 +/- 0.0	99.99 +/- 0.06	99.55 +/- 1.42
MORC2	50.99 +/- 7.63	100.0 +/- 0.0	99.98 +/- 0.09	98.45 +/- 3.48
MPV17	53.45 +/- 8.53	100.0 +/- 0.0	99.98 +/- 0.12	98.83 +/- 4.25
MPZ	45.17 +/- 7.9	100.0 +/- 0.0	99.28 +/- 2.83	91.11 +/- 12.78
MTMR2	56.03 +/- 7.65	100.0 +/- 0.0	100.0 +/- 0.0	99.67 +/- 0.87
MTRFR	54.45 +/- 8.5	100.0 +/- 0.0	99.98 +/- 0.12	99.31 +/- 1.67
MYH14	49.18 +/- 7.97	100.0 +/- 0.0	99.54 +/- 1.14	96.13 +/- 5.99
NAGA	50.92 +/- 8.27	100.0 +/- 0.0	99.99 +/- 0.08	98.48 +/- 5.23
NAGLU	53.54 +/- 8.38	100.0 +/- 0.0	99.96 +/- 0.3	98.55 +/- 4.21
NDRG1	50.38 +/- 7.56	100.0 +/- 0.0	99.9 +/- 0.47	97.97 +/- 3.29
NEFH	52.81 +/- 8.39	100.0 +/- 0.0	99.98 +/- 0.09	98.0 +/- 5.66
NEFL	53.52 +/- 7.38	100.0 +/- 0.0	99.94 +/- 0.4	98.91 +/- 2.98
NGF	54.93 +/- 8.75	100.0 +/- 0.0	99.99 +/- 0.07	99.17 +/- 3.72
NTRK1	48.9 +/- 8.21	100.0 +/- 0.0	99.99 +/- 0.08	97.0 +/- 7.44
OPA1	56.32 +/- 7.2	100.0 +/- 0.0	100.0 +/- 0.0	99.76 +/- 0.6
OPA3	51.76 +/- 8.25	99.98 +/- 0.04	99.96 +/- 0.06	98.68 +/- 3.1
PDHA1	43.74 +/- 16.86	99.88 +/- 0.67	94.1 +/- 10.7	70.21 +/- 35.65
PDK3	44.61 +/- 16.86	99.97 +/- 0.19	95.7 +/- 8.38	70.98 +/- 34.89
PDYN	51.43 +/- 8.62	100.0 +/- 0.0	100.0 +/- 0.0	98.61 +/- 4.33
PEX10	56.38 +/- 8.62	100.0 +/- 0.0	99.93 +/- 0.45	99.24 +/- 2.69
PEX7	56.2 +/- 7.37	100.0 +/- 0.0	100.0 +/- 0.0	99.46 +/- 1.84
PHYH	55.21 +/- 7.7	99.96 +/- 0.05	99.95 +/- 0.05	99.45 +/- 1.74
PLEKHG5	52.65 +/- 8.97	100.0 +/- 0.0	99.95 +/- 0.25	97.87 +/- 6.59
PLP1	43.2 +/- 16.12	99.9 +/- 0.54	94.53 +/- 8.67	70.22 +/- 34.85
PMM2	53.74 +/- 8.4	100.0 +/- 0.0	99.87 +/- 0.85	98.37 +/- 4.97
PMP2	56.22 +/- 7.98	100.0 +/- 0.0	99.99 +/- 0.03	99.65 +/- 0.91
PMP22	54.3 +/- 9.42	99.99 +/- 0.03	99.74 +/- 1.61	97.34 +/- 13.99
PNKP	51.37 +/- 8.13	100.0 +/- 0.0	100.0 +/- 0.0	98.05 +/- 3.57
PNPLA6	52.18 +/- 8.57	100.0 +/- 0.0	99.94 +/- 0.24	97.88 +/- 5.9
POLG	52.85 +/- 8.09	100.0 +/- 0.0	99.94 +/- 0.37	98.88 +/- 3.87
POLR3A	54.1 +/- 7.83	100.0 +/- 0.0	99.97 +/- 0.19	99.29 +/- 2.62
PPOX	52.02 +/- 8.17	100.0 +/- 0.0	99.98 +/- 0.13	98.22 +/- 5.32
PRDM12	46.65 +/- 7.8	99.99 +/- 0.07	99.54 +/- 1.29	93.36 +/- 11.24

Gen	Middelsekventeringsdybde [x]	Andel >= 10x [%]	Andel >= 20x [%]	Andel >= 30x [%]
PRKCG	45.14 +/- 7.84	100.0 +/- 0.01	99.46 +/- 2.64	93.32 +/- 12.08
PRNP	52.49 +/- 8.2	100.0 +/- 0.0	99.96 +/- 0.19	98.74 +/- 2.14
PRPS1	39.07 +/- 15.13	99.41 +/- 1.94	89.05 +/- 16.93	63.46 +/- 37.47
PRX	51.86 +/- 8.79	100.0 +/- 0.0	99.82 +/- 1.19	98.09 +/- 7.42
PTEN	54.68 +/- 6.95	100.0 +/- 0.0	99.92 +/- 0.28	98.97 +/- 2.41
PTPN11	55.1 +/- 7.8	100.0 +/- 0.0	99.96 +/- 0.18	99.29 +/- 1.99
PTRH2	53.96 +/- 9.25	100.0 +/- 0.0	100.0 +/- 0.0	99.14 +/- 3.67
RAB7A	54.2 +/- 8.16	100.0 +/- 0.0	100.0 +/- 0.0	99.42 +/- 2.03
REEP1	54.21 +/- 7.58	100.0 +/- 0.0	100.0 +/- 0.02	99.09 +/- 2.55
RETREG1	54.44 +/- 8.24	100.0 +/- 0.0	99.88 +/- 0.45	98.36 +/- 3.1
SACS	54.68 +/- 7.46	100.0 +/- 0.0	99.98 +/- 0.12	99.5 +/- 1.27
SBF1	52.02 +/- 8.61	99.98 +/- 0.1	99.74 +/- 0.65	96.98 +/- 6.56
SBF2	54.54 +/- 7.36	100.0 +/- 0.0	99.99 +/- 0.05	99.16 +/- 1.87
SCARB2	54.75 +/- 7.95	100.0 +/- 0.0	99.97 +/- 0.18	99.32 +/- 2.51
SCN10A	52.42 +/- 7.89	100.0 +/- 0.0	100.0 +/- 0.0	99.17 +/- 2.12
SCN11A	53.73 +/- 7.25	100.0 +/- 0.0	99.72 +/- 0.43	98.27 +/- 2.53
SCN9A	53.78 +/- 7.42	99.98 +/- 0.04	99.97 +/- 0.05	98.87 +/- 1.69
SCYL1	50.49 +/- 8.7	100.0 +/- 0.0	99.49 +/- 2.22	95.46 +/- 9.31
SEPTIN9	48.36 +/- 7.61	99.98 +/- 0.09	98.72 +/- 1.59	93.09 +/- 7.5
SETX	54.63 +/- 7.53	100.0 +/- 0.0	99.98 +/- 0.11	99.36 +/- 1.53
SH3TC2	53.81 +/- 7.8	100.0 +/- 0.0	99.97 +/- 0.11	99.27 +/- 1.87
SIGMAR1	52.09 +/- 8.66	100.0 +/- 0.0	100.0 +/- 0.03	98.76 +/- 3.44
SLC12A6	53.29 +/- 7.44	100.0 +/- 0.03	99.99 +/- 0.04	98.86 +/- 2.22
SLC25A19	53.14 +/- 8.09	100.0 +/- 0.0	99.76 +/- 1.28	98.79 +/- 5.73
SLC25A46	54.4 +/- 7.28	100.0 +/- 0.0	99.99 +/- 0.07	99.43 +/- 1.22
SLC52A2	53.88 +/- 9.82	99.99 +/- 0.04	99.72 +/- 1.85	97.97 +/- 7.75
SLC52A3	49.53 +/- 7.72	99.99 +/- 0.04	99.94 +/- 0.15	98.21 +/- 4.2
SLC5A7	55.8 +/- 7.58	100.0 +/- 0.0	99.96 +/- 0.23	99.0 +/- 2.04
SMN1	52.69 +/- 11.54	100.0 +/- 0.0	99.81 +/- 0.88	95.69 +/- 7.64
SORD	53.79 +/- 8.39	99.7 +/- 0.57	98.52 +/- 1.06	96.22 +/- 4.51
SOX10	49.57 +/- 8.72	100.0 +/- 0.0	99.73 +/- 0.95	95.55 +/- 7.42
SPAST	53.38 +/- 6.55	100.0 +/- 0.0	99.87 +/- 0.39	98.12 +/- 2.89
SPG11	54.82 +/- 7.58	100.0 +/- 0.0	99.99 +/- 0.05	99.49 +/- 1.56
SPG7	52.87 +/- 7.91	100.0 +/- 0.0	99.94 +/- 0.3	98.32 +/- 3.91
SPTBN4	51.0 +/- 8.07	100.0 +/- 0.0	99.88 +/- 0.45	97.11 +/- 6.92
SPTLC1	55.71 +/- 7.72	100.0 +/- 0.0	100.0 +/- 0.0	99.5 +/- 1.16
SPTLC2	54.72 +/- 7.69	100.0 +/- 0.0	99.99 +/- 0.04	99.62 +/- 0.99
SUCLA2	57.39 +/- 7.73	100.0 +/- 0.0	100.0 +/- 0.0	99.97 +/- 0.14
SURF1	51.22 +/- 8.29	100.0 +/- 0.0	99.78 +/- 1.21	97.57 +/- 4.8
SYT2	50.46 +/- 8.19	100.0 +/- 0.0	99.88 +/- 0.38	97.28 +/- 6.12
TFG	52.97 +/- 7.45	100.0 +/- 0.0	99.92 +/- 0.44	97.85 +/- 3.81
TRIM2	55.02 +/- 7.41	100.0 +/- 0.0	100.0 +/- 0.01	99.37 +/- 1.59
TRPA1	55.29 +/- 7.37	100.0 +/- 0.0	100.0 +/- 0.03	99.36 +/- 1.0
TRPV4	49.83 +/- 8.64	100.0 +/- 0.0	99.94 +/- 0.31	97.87 +/- 5.65
TTPA	54.8 +/- 7.2	99.98 +/- 0.04	99.97 +/- 0.05	99.0 +/- 3.24
TTR	55.48 +/- 8.34	100.0 +/- 0.0	100.0 +/- 0.0	99.52 +/- 1.77

Gen	Middelsekventeringsdybde [x]	Andel >= 10x [%]	Andel >= 20x [%]	Andel >= 30x [%]
TUBB3	56.14 +/- 8.69	100.0 +/- 0.0	99.91 +/- 0.52	98.61 +/- 3.96
TWNK	52.44 +/- 8.87	100.0 +/- 0.0	99.91 +/- 0.44	98.51 +/- 5.9
TYMP	53.69 +/- 9.95	100.0 +/- 0.0	99.96 +/- 0.2	97.63 +/- 6.4
VCP	53.01 +/- 7.83	100.0 +/- 0.0	100.0 +/- 0.01	99.09 +/- 2.71
VPS13A	55.32 +/- 7.42	100.0 +/- 0.0	99.99 +/- 0.04	99.49 +/- 1.07
VRK1	55.99 +/- 7.79	100.0 +/- 0.0	99.99 +/- 0.04	99.47 +/- 1.23
VWA1	52.66 +/- 9.13	100.0 +/- 0.0	99.72 +/- 1.01	96.8 +/- 7.62
WARS1	52.98 +/- 8.35	100.0 +/- 0.0	100.0 +/- 0.0	98.98 +/- 3.14
WNK1	51.7 +/- 7.44	99.99 +/- 0.04	99.78 +/- 0.51	97.83 +/- 3.27
XK	43.22 +/- 16.07	99.96 +/- 0.25	94.56 +/- 8.99	70.3 +/- 34.89
XPA	55.18 +/- 7.69	100.0 +/- 0.0	99.99 +/- 0.06	99.28 +/- 2.4
XRCC1	46.93 +/- 7.79	100.0 +/- 0.03	99.68 +/- 1.3	95.1 +/- 10.3
YARS1	50.56 +/- 8.06	100.0 +/- 0.0	99.99 +/- 0.05	98.46 +/- 5.15
ZFYVE26	52.87 +/- 7.83	100.0 +/- 0.0	99.94 +/- 0.28	98.84 +/- 3.35

For supplerende oplysninger vedrørende analysen kan afdelingen kontaktes på mail: mol-dia@rn.dk. Rapport genereret: 09/09-2024