

PN	Description	CZK	EUR	PLN
Library preparation (Includes sample QC, library preparation and pooling)				
Illumina				
LP-050	DNA library (PCR-free on request)	5 500	232	1 100
LP-080	Shotgun genomic DNA library (8 at least)	2 600	110	520
LP-060	Total RNA or mRNA (polyA-selected) library	6 900	290	1 380
LP-070	RNA library including rRNA depletion (H/M/R/Bacteria/custom)	7 800	328	1 560
LP-170	miRNA library	7 200	303	1 440
LP-200	3' mRNA library (QuantSeq/UMI)	5 300	223	1 060

For projects requiring numerous library preps please contact us for a quotation!

TA-024	Tailed Amplicon library, up to 24 samples	29 100	1 223	5 820
TA-048	Tailed Amplicon library, up to 48 samples	35 100	1 475	7 020
TA-096	Tailed Amplicon library, up to 94 samples	47 100	1 979	9 420
TA-192	Tailed Amplicon library, up to 190 samples	69 200	2 908	13 840

Pacific Biosciences and Oxford Nanopore Technologies				
LP-110	DNA library - PacBio	Please contact us for a quotation		
LP-150	DNA library - Oxford Nanopore	Please contact us for a quotation		

Quality control of libraries prepared by customers				
Illumina				
QC-080	Library QC, customer-prepared (Qubit, Bioanalyzer, qPCR)	4 200	177	840

If you send more libraries at a time or need other QC procedures (e.g. quantification, fragment analysis, qPCR, size selection etc.) please contact us for a quotation!

Sequencing - Short and Long reads				
Illumina - 1 million reads data package				
DP004-PE250	ShareSeq Data Package, PE250, 1 million reads / 0,25 Gb	7 100	299	1 420
Illumina - 20 million reads data packages				
DP020-PE150	ShareSeq Data Package, PE150, 20 million reads / 3 Gb	7 100	299	1 420
DP020-PE250	ShareSeq Data Package, PE250, 20 million reads / 5 Gb	17 600	740	3 520
DP020-SE50	ShareSeq Data Package, SE50, 20 million reads / 1 Gb	12 400	522	2 480
Illumina - 100 million reads data packages				
DP100-PE150	ShareSeq Data Package, PE150, 100 million reads / 15 Gb	18 300	769	3 660
DP100-PE250	ShareSeq Data Package, PE250, 100 million reads / 25 Gb	55 100	2 316	11 020
DP100-SE50	ShareSeq Data Package, SE50, 100 million reads / 5 Gb	47 000	1 975	9 400
Illumina - other data packages				
DP600-PE150	ShareSeq Data Package, PE150, 600 million reads / 90 Gb	56 300	2 366	11 260
DP800-PE150	ShareSeq Data Package, PE150, 800 million reads / 120 Gb	66 300	2 786	13 260
Illumina - NovaSeq6000, price per lane or flowcell				
DS-240	NovaSeq6000 S4 lane, PE150, 5000-6000 million reads	195 000	8 194	39 000
DS-300	NovaSeq6000 S4 flowcell, PE150, 20000-24000 million reads	583 400	24 513	116 680
DS-150	NovaSeq6000 SP lane, PE150, 650-800 million reads	Please contact us for a quotation		
DS-290	NovaSeq6000 SP flowcell, PE150, 1300-1600 million reads	161 900	6 803	32 380
Pacific Biosciences and Oxford Nanopore Technologies				
DS-200	PacBio Sequel II, 10-30 hrs movie, upto 5 million reads	Please contact us for a quotation		
DS-210	Oxford Nanopore MinION FC, single-end, 1-200 kb reads, 5-10 Gb	Please contact us for a quotation		
DS-280	Oxford Nanopore Flongle FC, single-end, 1-200 kb reads, 1-2 Gb	Please contact us for a quotation		
Bioinformatic services				
BS-SB	Bioinformatic services, price per order	2 900	122	580

For other sequencing settings or large scale projects please contact us for a quotation!

PN	Description	CZK	EUR	PLN
Metagenome analysis (Illumina)				
Sample QC, amplicon generation using a single primer set, double indexing (unique-dual strategy), library prep, 250 b paired-end read sequencing (NovaSeq6000), demultiplexing, analysis report. For primer sets and other details please see our website.				
Output >20K reads/sample				
SO-024	Metagenomic profiling, up to 24 samples, >20K reads/sample	45 600	1 916	9 120
SO-048	Metagenomic profiling, up to 48 samples, >20K reads/sample	67 200	2 824	13 440
SO-096	Metagenomic profiling, up to 94 samples, >20K reads/sample	96 000	4 034	19 200
SO-192	Metagenomic profiling, up to 190 samples, >20K reads/sample	124 800	5 244	24 960
Output >60K reads/sample				
SO-024C	Metagenomic profiling, up to 24 samples, >60K reads/sample	52 800	2 219	10 560
SO-048C	Metagenomic profiling, up to 48 samples, >60K reads/sample	77 900	3 274	15 580
SO-096C	Metagenomic profiling, up to 94 samples, >60K reads/sample	111 300	4 677	22 260
SO-192C	Metagenomic profiling, up to 190 samples, >60K reads/sample	144 700	6 080	28 940
Output >120K reads/sample				
SO-024D	Metagenomic profiling, up to 24 samples, >120K reads/sample	60 000	2 522	12 000
SO-048D	Metagenomic profiling, up to 48 samples, >120K reads/sample	88 500	3 719	17 700
SO-096D	Metagenomic profiling, up to 94 samples, >120K reads/sample	126 500	5 316	25 300
SO-192D	Metagenomic profiling, up to 190 samples, >120K reads/sample	164 500	6 912	32 900

For large scale or custom metagenome projects please contact us for a quotation!

Whole-exome and whole-genome human / mouse sequencing (Illumina)				
Sample QC, library prep, 150 b paired-end read sequencing (Illumina), analysis report				
SO-006	WES - Human, Agilent V6 (58Mb), 6 Gb	9 800	412	1 960
SO-012	WES - Human, Agilent V6 (58Mb), 12 Gb	13 300	559	2 660
SO-006-M	WES - Mouse, Agilent SureSelectXT, 6 Gb	11 600	488	2 320
SO-012-M	WES - Mouse, Agilent SureSelectXT, 12 Gb	15 100	635	3 020
SO-090	WGS - human, 90 Gb	23 800	1 000	4 760
Bioinformatic services				
BS-SB	Bioinformatic services, price per order	2 900	122	580

For large scale or custom WES / WGS projects please contact us for a quotation!

Data analysis services				
Differential expression analysis, metagenome data analysis or any other bioinformatic analysis ...				
Please contact us to receive a quote of our data analysis services!				

Notice

This is an abbreviated pricelist of our services. If you require additional information, special sequencing services and/or data analysis services, please contact us.

All prices quoted without VAT and costs of sample shipment. This pricelist is valid since February 1, 2023. Subject to change at any time.
General terms and conditions and additional information available at www.seqme.eu apply.

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