



CF Genomics price list

Service	Unit	Full service	Service only (user supplied consumables)
Library preparation			
DNA library preparation (1 sample)	1 sample	5 128	NA
DNA library preparation (4 samples)	4 samples	10 107	NA
DNA library preparation (8 samples)	8 samples	16 345	NA
RNA library preparation (polyA)	1 sample	7 167	NA
RNA library preparation (polyA)	4 samples	14 665	NA
RNA library preparation (polyA)	8 samples	24 264	NA
RNA library preparation (rRNA depletion)	1 sample	7 377	NA
RNA library preparation (rRNA depletion)	4 samples	15 505	NA
RNA library preparation (rRNA depletion)	8 samples	25 944	NA
MinION Library preparation	1 - 12 samples	12 268	NA
Covaris fragmentation	1 sample	275	NA
PippinPrep size selection	sample	1 120	NA
Library QC	sample	277	NA
Sequencing			
MiSeq v2 50 cycles	run	29 821	3 532
MiSeq v2 300 cycles	run	41 149	7 985
MiSeq v2 500 cycles	run	49 100	11 695
MiSeq v2 Nano 500 cycles	run	30 124	7 985
MiSeq v3 150 cycles	run	36 198	7 243
MiSeq v3 600 cycles	run	65 427	15 901
MiSeq v3 600 cycles shared run (1M reads)	1M reads	6 543	NA
NextSeq high-output 75 cycles	run	55 322	9 541
NextSeq high-output 75 cycles shared run (10M reads)	10M reads	2 213	NA
NextSeq high-output 150 cycles	run	107 530	20 156
NextSeq high-output 300 cycles	run	160 662	20 781
NextSeq mid-output 150 cycles	run	46 298	12 039
NextSeq mid-output 300 cycles	run	73 718	18 908
Sequencing packages			
RNA-Seq (polyA, 30M+ reads, 1x75bp)	sample	10 661	NA
Human exome (5Gb raw data 2x75bp, 80x+ mean coverage)	sample	18 763	NA
Human exome (4Gb raw data 2x75bp, 50x+ mean coverage)	sample	15 919	NA
Flow cytometry			
Cell sorting – setup (first hour)	hour	3 825	NA
Cell sorting aseptic mode – setup (first hour)	hour	4 087	NA
Cell sorting – hour	hour	3 444	NA
Cell sorting aseptic mode -hour	hour	3 497	NA
Flow cytometry – initial training	hour	2 375	NA
Flow cytometry (self service)	hour	1 723	NA
Flow cytometry (full service)	hour	2 322	NA
Microdissection (self service)	hour	1 023	NA

All prices in CZK excl. VAT (21%)

Discounts for academic users – contact us for details.