

BEA Service Fees (updated Dec 2025)

- BEA offers services for the **Illumina, 10X Genomics, Nanopore, Affymetrix, ABI and Agilent** platforms.
- Service fees listed below generally include all reagents, labor and analysis costs, if not indicated.
- **Prices can be subjected with changes without prior notice due to currency fluctuations.**
- All prices include the INDI (Indirect costs) Surcharge at KI.
- Observe that prices often apply to a fix number of samples and that orders sometimes only can be made in multiples of that number.
- External customers outside Swedish academia pay a surcharge on indicated prices.

Illumina

- BEA is offering sequencing services on the Illumina Nextseq and Miseq platforms. Services are complete and include sample QC, library preparation, sequencing and data analysis. Prices are dependent on the level of multiplexing and sequencing depth.
- Minimum number of samples, special formats and different conditions apply to sequencing services. Please enquire for more information.
- Illumina beadarrays are sold in different sample formats as described below. Some arrays are restricted with a minimum order. Prices include all reagents, arrays and analysis costs.

Illumina library preparation services

Library Prep for Illumina sequencing (small<12 samples<large)		Price / sample
Illumina mRNA stranded ligation	(small-large)	1600-1400
Illumina stranded ligation Total RNA seq with rRNA depletion	(small-large)	2400-2200
NEBNext DNA ultra II Library Prep (DNA and ChIPseq)	(small-large)	1600-1400
NEBNext UltraExpress RNA Library Prep	(small-large)	1400-1600
Takara SMART-seq ultra low Single Cell and Pico samples	(small-large)	1400-1200
Illumina Nextera XT/Flex (DNA)	(small-large)	1200-1000
Illumina Nextera Exome (24rxn)		2600
RRBS DNA Methylation Profiling	(small-large)	1800-1600
16S metagenomic Profiling		500
Pooling, Fragmentation, depletion and other requests		Please enquire

Illumina sequencing services

Illumina sequencing on Novaseq Plus Platform NGI Stockholm		Price / run
10B 300 (Total 338 cycles) ~1200 Mreads (SR/PE) price per lane	(lane)	Please enquire
25B 300 (Total 338 cycles) ~3200 Mreads (SR/PE) price per lane	(lane)	Please enquire
Illumina XLeap sequencing on Nextseq 2000 Platform (new version)		Price / run
P1 XLeap 100 (Total 138 cycles) ~100Mreads (SR/PE) price per flowcell	(full flowcell)	13000
P1 XLeap 300 (Total 338 cycles) ~100Mreads (SR/PE) price per flowcell	(full flowcell)	18000
P1 XLeap 600 (Total 638 cycles) ~100Mreads (SR/PE) price per flowcell	(full flowcell)	24000
P2 XLeap 100 (Total 138 cycles) ~400Mreads (SR/PE) price per flowcell	(full flowcell)	18000
P2 XLeap 300 (Total 338 cycles) ~400Mreads (SR/PE) price per flowcell	(full flowcell)	39000
P3 XLeap 100 (Total 138 cycles) ~1200Mreads (SR/PE) price per flowcell	(full flowcell)	35000
P4 XLeap 100 (Total 138 cycles) ~1800Mreads (SR/PE) price per flowcell	(full flowcell)	47000
Illumina sequencing on MiSeq Platform		Price / flowcell
300 cycles v.2 nano ~1Mreads (SR/PE) price per lane/flowcell	(full flowcell)	6000
300 cycles v.2 micro ~4Mreads (SR/PE) price per lane/flowcell	(full flowcell)	8000
150 cycles v.3 ~25Mreads (SR/PE) price per lane/flowcell	(full flowcell)	17000
600 cycles v.3 ~25Mreads (SR/PE) price per lane/flowcell	(full flowcell)	24000



Illumina Infinium DNA Microarray Analysis Assays		
Infinium methylation EPIC 2.0 (8 sample format)	Price / sample	Arrays Included
Minimum order (16 samples)	4600	Yes
>32 samples	4400	Yes
Infinium methylation Mouse 265K (12 sample fixed format)	Price / sample	Arrays Included
Minimum order (24 samples)	3600	Yes
>48 samples	3400	Yes
Infinium Human Genotyping arrays	Price / sample	Arrays Included
4,8,16,24 samples (different formats and versions)	Please enquire	Yes

Oxford Nanopore Services

- BEA provides sequencing services in different formats with Promethion, Minion and Flongle flowcells.
- ONT platform and flowcells can be used for many applications including high-accuracy, direct sequencing of native DNA and RNA.
- BEA will help users to design experiments, provide reagents and service including library preparation, sequencing and Bioinformatic service.

ONT Promethion, Minion, Flongle flowcells available	Price / run	
Promethion (Minimum order 4 flowcells) 6000 pores ~20Gb data	(full flowcell)	15000
Library preps available for different purposes and formats	Please enquire	Please enquire
Flongle flowcell (96 pores) minimum order apply	Please enquire	12 flowcells
Minion flowcell (512 pores)	Please enquire	1 flowcells

10X Chromium

- BEA core facility offers service with the 10X Genomics Chromium platform allowing solutions for single cell genomics, characterization and gene expression profiling.
- The Chromium single cell assays provide solutions for cell characterization and gene expression profiling of hundreds to of thousands of cells.
- BEA will help users to design experiments, provide reagents and service from the initial cell separation step. Bioinformatic service with the Cell Ranger software is included.
- Service includes different options and versions of GEM kits and library preparation methods for gene expression and cell profiling.

10X Chromium GEM-X Single Cell Gene Expression v4	Price / sample	Format
Separation and library construction (sequencing not included)	Please enquire	1rxn
Separation and library construction (sequencing not included)	Please enquire	4rxn
Cell profiling and additional library preparation	2000	1rxn

10X Chromium Flex GEM_X v2	Price / sample	Format
Separation and library construction (sequencing not included)	Please enquire	1rxn
Separation and library construction (sequencing not included)	Please enquire	4rxn
Cell profiling and additional library preparation	2000	1rxn

Affymetrix

- Affymetrix provide two types of expression arrays: **(WT) whole transcript** arrays and **3' (IVT) *in vitro* transcribed** arrays.
- Affymetrix cartridge arrays are sometimes, when indicated, not included in the price and will be added to the bill separately. Please enquire for specific array prices.

Affymetrix Whole Transcript (WT) Assay		
GCAS Cartridge format for Clariom D, HTA 2.0 (mouse, human, rat)	Price / sample	Arrays Included
Individual samples in cartridge array format		
1-12 samples	4700	Yes
> 12 samples	4500	Yes
> 48 samples	4300	Yes
GCAS Cartridge format for Clariom S, (mouse, human, rat)	Price / sample	Arrays Included
Individual samples in cartridge array format		
1-12 samples	3200	Yes
> 12 samples	3000	Yes
> 48 samples	2800	Yes
Small and/or fragmented or degraded samples. Additional costs:	Price / sample	Arrays Included
Affymetrix Pico WT assay (Small sample)	500	NA

Hybridization only	Price per sample
Hybridisation, washing, staining and scanning of individual arrays	800

Additional Services

- BEA provides RNA and DNA quality control services with the Agilent Bioanalyzer 2100 and TapeStation 2200. We also provide services for RNA and DNA extraction from various tissues and cell samples.
- BEA has standardized qPCR services using the ABI 7900 platform. Validation and qPCR projects can also be customized to fulfill other demands. Apart from this we also have access to pyrosequencing and digital PCR platforms.
- Extraction services are limited to projects with downstream genomic analysis at BEA.

Custom services	Price per sample
RNA isolation and purification from blood PAXgene or FFPE samples	500
RNA/DNA isolation and purification from tissue	400
RNA/DNA isolation and purification from cells	300
Allprep RNA and DNA isolation and purifications from tissue	500
Extra cost for RNA/DNA isolation and purification with QIAzol or TRIzol	100
Agilent Bioanalyzer RNA/DNA quality control– Nano, Pico, smallRNA, DNA (min 10 samples)	100
AgilentTapeStation RNA/DNA quality control– D1000, HS1000, HS5000 (min 2 samples)	100
Qubit RNA/DNA concentration measurement (min 10 samples)	10

qPCR analysis

96 well Taqman/SYBR Assays	Price
Project Setup fee (includes initial planning and dispensing excluding reagent cost)	1000
Additional hourly cost (if needed)	500
Reverse Transcription including DNase treatment/sample	200
Taqman Low Density Arrays (TLDA)	Price
Running cost excluding reagents/card	1000
Reverse Transcription including DNase treatment/sample	200

Bioinformatic services

Basic bioinformatics included in the analysis cost and free of charge.	
Bioinformatic services handling external data not analysed at BEA. Charged by the hour.	500



FAQ

- Sequencing data obtained from samples at BEA may only be used for academic and scientific purposes.
- Users are responsible for data analysis and interpretation. The vast majority of the samples going through the BEA Core Facility yield good results. However, while every care will be taken, the core facility cannot take any responsibility for any particular result, lost samples and data.
- BEA disclaims any liability regarding quality, reliability and reproducibility. Users are required to contribute financially to project costs and take full responsibility for selected methods, processing procedures and the selected instrument and mode of sequencing.
- In case of publication contributions have to be acknowledged to BEA.
- BEA provides backup of generated data and generated samples for >12 months. After this period BEA takes no responsibility. Customers are strongly advised to collect data and leftover samples before this period.