### Sequencing

**NovaSeq 6000**

<table>
<thead>
<tr>
<th>Cycles</th>
<th>Reads</th>
<th>Cost</th>
</tr>
</thead>
<tbody>
<tr>
<td>35</td>
<td>10'000 mio</td>
<td>12'690 CHF</td>
</tr>
<tr>
<td>100</td>
<td>400 mio</td>
<td>1'670 CHF</td>
</tr>
<tr>
<td>100</td>
<td>800 mio</td>
<td>2'915 CHF</td>
</tr>
<tr>
<td>100</td>
<td>1'600 mio</td>
<td>5'190 CHF</td>
</tr>
<tr>
<td>100</td>
<td>4'100 mio</td>
<td>9'575 CHF</td>
</tr>
<tr>
<td>200</td>
<td>800 mio</td>
<td>3'755 CHF</td>
</tr>
<tr>
<td>200</td>
<td>1'600 mio</td>
<td>6'495 CHF</td>
</tr>
<tr>
<td>200</td>
<td>4'100 mio</td>
<td>11'860 CHF</td>
</tr>
<tr>
<td>200</td>
<td>10'000 mio</td>
<td>16'570 CHF</td>
</tr>
<tr>
<td>300</td>
<td>800 mio</td>
<td>4'105 CHF</td>
</tr>
<tr>
<td>300</td>
<td>1'600 mio</td>
<td>7'060 CHF</td>
</tr>
<tr>
<td>300</td>
<td>4'100 mio</td>
<td>12'720 CHF</td>
</tr>
<tr>
<td>300</td>
<td>10'000 mio</td>
<td>18'590 CHF</td>
</tr>
<tr>
<td>500</td>
<td>800 mio</td>
<td>5'670 CHF</td>
</tr>
</tbody>
</table>

*For 10XG scRNA-seq or BRB-seq, use 100 cycles (or 200 cycles for 10'000 mio reads).*

**NextSeq 500**

<table>
<thead>
<tr>
<th>Cycles</th>
<th>Reads</th>
<th>Cost</th>
</tr>
</thead>
<tbody>
<tr>
<td>75</td>
<td>400 mio</td>
<td>1'779 CHF</td>
</tr>
<tr>
<td>150</td>
<td>130 mio</td>
<td>1'347 CHF</td>
</tr>
<tr>
<td>150</td>
<td>400 mio</td>
<td>3'380 CHF</td>
</tr>
<tr>
<td>300</td>
<td>130 mio</td>
<td>2'250 CHF</td>
</tr>
<tr>
<td>300</td>
<td>400 mio</td>
<td>5'295 CHF</td>
</tr>
</tbody>
</table>

**MiSeq**

<table>
<thead>
<tr>
<th>Cycles</th>
<th>Reads</th>
<th>Cost</th>
</tr>
</thead>
<tbody>
<tr>
<td>50</td>
<td>15 mio</td>
<td>1'197 CHF</td>
</tr>
<tr>
<td>150</td>
<td>25 mio</td>
<td>1'294 CHF</td>
</tr>
<tr>
<td>300</td>
<td>1 mio</td>
<td>564 CHF</td>
</tr>
<tr>
<td>300</td>
<td>4 mio</td>
<td>759 CHF</td>
</tr>
<tr>
<td>300</td>
<td>15 mio</td>
<td>1'472 CHF</td>
</tr>
<tr>
<td>500</td>
<td>1 mio</td>
<td>637 CHF</td>
</tr>
<tr>
<td>500</td>
<td>15 mio</td>
<td>1'618 CHF</td>
</tr>
<tr>
<td>600</td>
<td>25 mio</td>
<td>2'046 CHF</td>
</tr>
</tbody>
</table>

### Library preparation

**ISML (Illumina Stranded mRNA Ligation)**

<table>
<thead>
<tr>
<th>Samples</th>
<th>Reagents Fee</th>
<th>Labour Fee, 1-16 samples</th>
<th>Labour Fee, 17-32 samples</th>
<th>Labour Fee, 33-48 samples</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>95 CHF</td>
<td>280 CHF</td>
<td>350 CHF</td>
<td>420 CHF</td>
</tr>
</tbody>
</table>

**NebNext Ultra II stranded mRNA**

<table>
<thead>
<tr>
<th>Samples</th>
<th>Reagents Fee</th>
<th>Labour Fee, 1-16 samples</th>
<th>Labour Fee, 17-32 samples</th>
<th>Labour Fee, 33-48 samples</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>115 CHF</td>
<td>280 CHF</td>
<td>350 CHF</td>
<td>420 CHF</td>
</tr>
</tbody>
</table>
### NebNext Ultra II stranded total RNA

<table>
<thead>
<tr>
<th>Item</th>
<th>Cost (CHF)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reagents fee, per sample</td>
<td>175</td>
</tr>
<tr>
<td>Labour fee, 1-16 samples</td>
<td>280</td>
</tr>
<tr>
<td>Labour fee, 17-32 samples</td>
<td>350</td>
</tr>
<tr>
<td>Labour fee, 33-48 samples</td>
<td>420</td>
</tr>
</tbody>
</table>

### SMART-seq mRNA (ultra low input mRNA)

<table>
<thead>
<tr>
<th>Item</th>
<th>Cost (CHF)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reagents fee, per sample</td>
<td>150</td>
</tr>
<tr>
<td>Labour fee, 1-16 samples</td>
<td>280</td>
</tr>
<tr>
<td>Labour fee, 17-32 samples</td>
<td>350</td>
</tr>
<tr>
<td>Labour fee, 33-48 samples</td>
<td>420</td>
</tr>
</tbody>
</table>

### SMARTer mammalian pico v3 (ultra low input total RNA)

<table>
<thead>
<tr>
<th>Item</th>
<th>Cost (CHF)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reagents fee, per sample</td>
<td>235</td>
</tr>
<tr>
<td>Labour fee, 1-16 samples</td>
<td>210</td>
</tr>
<tr>
<td>Labour fee, 17-32 samples</td>
<td>280</td>
</tr>
<tr>
<td>Labour fee, 33-48 samples</td>
<td>350</td>
</tr>
</tbody>
</table>

### BRB-seq (Mercurius, 3’ end)

<table>
<thead>
<tr>
<th>Item</th>
<th>Cost (CHF)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reagents fee, per sample</td>
<td>26</td>
</tr>
<tr>
<td>Additional reagents fee, per plate</td>
<td>177</td>
</tr>
<tr>
<td>Labour fee, 1-32 samples</td>
<td>210</td>
</tr>
<tr>
<td>Labour fee, 33-64 samples</td>
<td>280</td>
</tr>
<tr>
<td>Labour fee, 65-96 samples</td>
<td>350</td>
</tr>
</tbody>
</table>

### NebNext Ultra II DNA

<table>
<thead>
<tr>
<th>Item</th>
<th>Cost (CHF)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reagents fee, per sample</td>
<td>78</td>
</tr>
<tr>
<td>Labour fee, 1-16 samples</td>
<td>210</td>
</tr>
<tr>
<td>Labour fee, 17-32 samples</td>
<td>280</td>
</tr>
<tr>
<td>Labour fee, 33-48 samples</td>
<td>350</td>
</tr>
</tbody>
</table>

### HiC (Arima)

<table>
<thead>
<tr>
<th>Item</th>
<th>Cost (CHF)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Reagents fee, per sample</td>
<td>345</td>
</tr>
<tr>
<td>Labour fee, 1-16 samples</td>
<td>350</td>
</tr>
</tbody>
</table>

*Other methods available on demand: QuantSeq 3’, Qiagen miRNA, Illumina DNA Prep, Nextera XT, TruSeq PCR-free, exome sequencing, 16S metagenomics, HiC...*

### Single cells

#### 10X Genomics 3’ scRNA-seq

<table>
<thead>
<tr>
<th>Item</th>
<th>Cost (CHF)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Per sample</td>
<td>2'300</td>
</tr>
<tr>
<td>Assistance (per chip)</td>
<td>315</td>
</tr>
</tbody>
</table>

#### 10X Genomics 3’ scRNA-seq CellPlex

<table>
<thead>
<tr>
<th>Item</th>
<th>Cost (CHF)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Per samples pool</td>
<td>2'685</td>
</tr>
<tr>
<td>Assistance (per chip)</td>
<td>315</td>
</tr>
<tr>
<td>Each tag</td>
<td>85</td>
</tr>
</tbody>
</table>

#### 10X Genomics FLEX fixed cells scRNA-seq

<table>
<thead>
<tr>
<th>Item</th>
<th>Cost (CHF)</th>
</tr>
</thead>
<tbody>
<tr>
<td>4 samples</td>
<td>5'500</td>
</tr>
<tr>
<td>8 samples</td>
<td>9'500</td>
</tr>
<tr>
<td>16 samples</td>
<td>13'430</td>
</tr>
<tr>
<td>32 samples</td>
<td>24'065</td>
</tr>
<tr>
<td>64 samples</td>
<td>38'910</td>
</tr>
<tr>
<td>Assistance (per chip)</td>
<td>315</td>
</tr>
</tbody>
</table>

#### 10X Genomics VDJ + 5’ scRNA-seq

<table>
<thead>
<tr>
<th>Item</th>
<th>Cost (CHF)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Per sample</td>
<td>2'680</td>
</tr>
<tr>
<td>Assistance (per chip)</td>
<td>420</td>
</tr>
</tbody>
</table>

#### 10X Genomics multiome (scRNA-seq + scATAC-seq)

<table>
<thead>
<tr>
<th>Item</th>
<th>Cost (CHF)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Per sample</td>
<td>4'015</td>
</tr>
<tr>
<td>Assistance (per chip)</td>
<td>490</td>
</tr>
</tbody>
</table>
### 10X Chromium independent use

Per run 96 CHF

*Prices don’t include sequencing.*

*Other methods on demand: 10XG scATAC-seq, CRISPR screen, protein detection...; PARSE scRNA-seq; Plate-based single-cell (f.sight.omics/i.dot/c.wash)*

### Spatial Transcriptomics

<table>
<thead>
<tr>
<th>Service Description</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Visium Cytassist 6.5mm x 6.5mm</strong></td>
<td>3'982 CHF</td>
</tr>
<tr>
<td>Per slide of 2 samples</td>
<td></td>
</tr>
<tr>
<td>Assistance (per slide)</td>
<td>490 CHF</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Service Description</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Visium Cytassist 11mm x 11mm</strong></td>
<td>7'834 CHF</td>
</tr>
<tr>
<td>Per slide of 2 samples</td>
<td></td>
</tr>
<tr>
<td>Assistance (per slide)</td>
<td>490 CHF</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Service Description</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Visium non-Cytassist 6.5mm x 6.5mm</strong></td>
<td>6'840 CHF</td>
</tr>
<tr>
<td>Per slide of 4 samples</td>
<td></td>
</tr>
<tr>
<td>Assistance (per slide)</td>
<td>350 CHF</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Service Description</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Visium non-Cytassist optimization</strong></td>
<td>1'275 CHF</td>
</tr>
<tr>
<td>Per optimization slide</td>
<td></td>
</tr>
<tr>
<td>Assistance</td>
<td>175 CHF</td>
</tr>
</tbody>
</table>

*Prices don’t include sequencing.*

*Other methods on demand (Curio Slide-seq).*

### Quantification and QC of nucleic acids and libraries

<table>
<thead>
<tr>
<th>Service Description</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Fragment Analyzer profiling (libraries, dsDNA, ssDNA...)</strong></td>
<td></td>
</tr>
<tr>
<td>1 lane (11 samples + ladder), reagents incl., user-operated</td>
<td>41 CHF</td>
</tr>
<tr>
<td>Assistance, per lane</td>
<td>35 CHF</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Service Description</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>TapeStation 4200 profiling (RNA)</strong></td>
<td></td>
</tr>
<tr>
<td>Per sample (ladder + pos. ctrl free), reagents incl., user-operated</td>
<td>7 CHF</td>
</tr>
<tr>
<td>Assistance, per run</td>
<td>35 CHF</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Service Description</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>qPCR-based library quantification (GECF-operated)</strong></td>
<td></td>
</tr>
<tr>
<td>Basal fee (up to 48 samples)</td>
<td>72 CHF</td>
</tr>
<tr>
<td>Each sample</td>
<td>4.20 CHF</td>
</tr>
</tbody>
</table>

<table>
<thead>
<tr>
<th>Service Description</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Qubit</strong></td>
<td></td>
</tr>
<tr>
<td>Qubit HS dsDNA, per sample</td>
<td>2 CHF</td>
</tr>
<tr>
<td>Qubit HS RNA, per sample</td>
<td>2.50 CHF</td>
</tr>
</tbody>
</table>

*Picogreen and ribogreen also available on demand for large projects*

### qPCR & BioMark

<table>
<thead>
<tr>
<th>Service Description</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>qPCR</strong></td>
<td></td>
</tr>
<tr>
<td>qPCR instrument booking, per hour (no reagent included)</td>
<td>7.10 CHF</td>
</tr>
<tr>
<td>Robotized 384w qPCR plate assembly, plasticware incl., per plate</td>
<td>35 CHF</td>
</tr>
</tbody>
</table>

*Fluidigm BioMark high-throughput qPCR on-demand*

*All prices without assistance. Assistance available on demand.*

### Digital PCR

<table>
<thead>
<tr>
<th>Service Description</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>QIAcuity One</strong></td>
<td></td>
</tr>
<tr>
<td>Nanoplate 8.5K 24-well</td>
<td>42 CHF</td>
</tr>
<tr>
<td>Nanoplate 8.5K 96-well</td>
<td>159 CHF</td>
</tr>
<tr>
<td>Nanoplate 26K 24-well</td>
<td>68 CHF</td>
</tr>
</tbody>
</table>

*Plate + run included. No mastermix provided.*

*All prices without assistance. Assistance available on demand.*
## Oxford Nanopore

### Sequencing runs (labour included)

<table>
<thead>
<tr>
<th>Service</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td>PromethION Solo2 run</td>
<td>1'625 CHF</td>
</tr>
<tr>
<td>MiniION run</td>
<td>1'285 CHF</td>
</tr>
<tr>
<td>Flongle run</td>
<td>290 CHF</td>
</tr>
</tbody>
</table>

### Ligation prep (Minion/Promethion)

<table>
<thead>
<tr>
<th>Service</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td>Per sample</td>
<td>235 CHF</td>
</tr>
<tr>
<td>Labour (total)</td>
<td>140 CHF</td>
</tr>
</tbody>
</table>

### Ligation prep (Flongle)

<table>
<thead>
<tr>
<th>Service</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td>Per sample</td>
<td>130 CHF</td>
</tr>
<tr>
<td>Labour (total)</td>
<td>140 CHF</td>
</tr>
</tbody>
</table>

### Native barcoding prep (Flongle/Minion/Promethion)

<table>
<thead>
<tr>
<th>Service</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td>Per multiplexed prep of 1-8 samples</td>
<td>310 CHF</td>
</tr>
<tr>
<td>Per multiplexed prep of 9-16 samples</td>
<td>510 CHF</td>
</tr>
<tr>
<td>Per multiplexed prep of 17-24 samples</td>
<td>710 CHF</td>
</tr>
<tr>
<td>Labour (total)</td>
<td>210 CHF</td>
</tr>
</tbody>
</table>

*Prices of other preps on demand.*

### ORF clones

<table>
<thead>
<tr>
<th>Service</th>
<th>Price</th>
</tr>
</thead>
<tbody>
<tr>
<td>Per clone</td>
<td>20 CHF</td>
</tr>
</tbody>
</table>

### Versions log

- April 2018: minor changes to match rounding system from FNS
- July 2023: major update.