

## Nanopore Sequencing

### Preparing sample:

- Mix 9  $\mu\text{l}$  of each sample with 1  $\mu\text{l}$  of index.
- Incubate in thermocycler with the following settings:
  - 1 minute at 30°C
  - 1 minute at 80°C
- Pipet all samples in a reaction tube.
- Add 40  $\mu\text{l}$  AM pure XP (beads) to buffer.
- Put in magnet separator and remove the remaining suspension.
- Carefully wash with alcohol two times, removing the remaining suspension quickly each time.
- Centrifuge briefly and remove the rest of the remaining alcohol.
- Dissolve in 10  $\mu\text{l}$  water and mix by snapping the reaction tube gently. Do not let the beads dry out.
- Put in magnet separator.
- Add 1  $\mu\text{l}$  rapid adapter (RAP) in a new reaction tube.
- Add 10  $\mu\text{l}$  of the beads and water mix to the RAP-mix.
- Incubate for 10 minutes

### Preparing the flow cell:

- Add 30  $\mu\text{l}$  FLT in a tube of Flush Buffer (FB)
- Open the priming port and try to remove air from the flow cell by pipetting.
- Add 750  $\mu\text{l}$  Flush Buffer inside.

### Loading the samples:

- Mix 35  $\mu\text{l}$  sequencing buffer with 25  $\mu\text{l}$  loading beads in a reaction tube.
- Load 150  $\mu\text{l}$  flush buffer in the priming port.
- Load the samples at the spot on.

From: iGEM Bielefeld-CeBiTec