Impore Field Effect Transistor (FET) for Protein Analysis **KU LEUVEN**

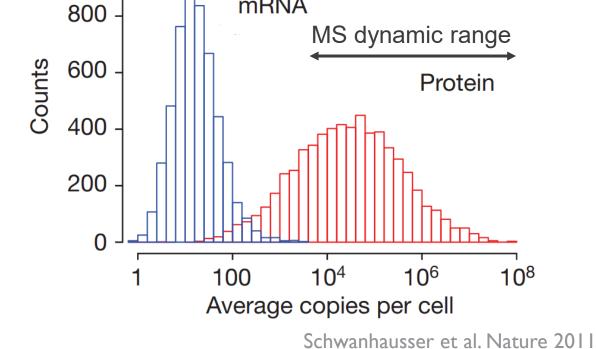
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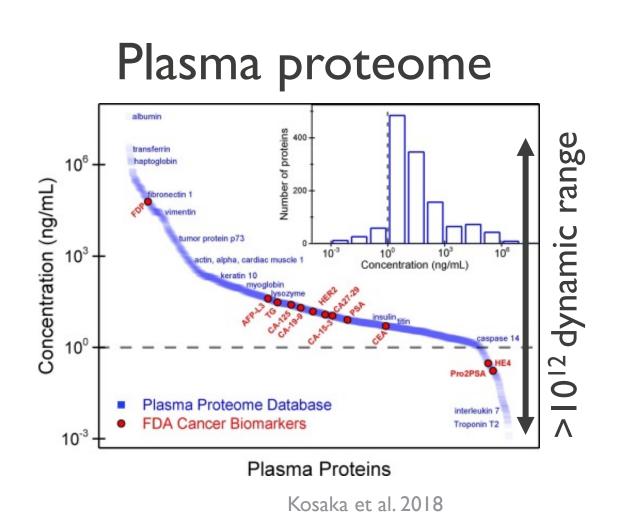
Key technology challenge for proteomics-on-chip

Nanopore FET single-molecule proteomics-on-chip vision

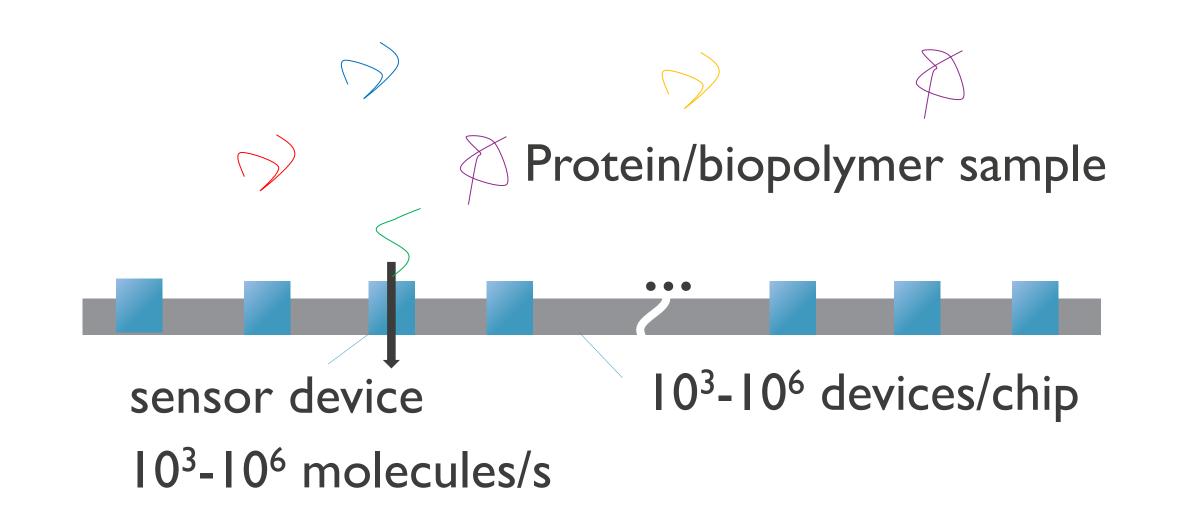
Cell proteome

• Proteomics-on-chip favors





- single-molecule approach
- Enormous concentration differences (= large dynamic range) in proteomes
- To exceed MS dynamic range (10³-10⁵) by multiple orders of magnitude one needs to analyze billions of single molecules
- For 10¹² dynamic range in an hour, billions of molecules per second need to be analyzed

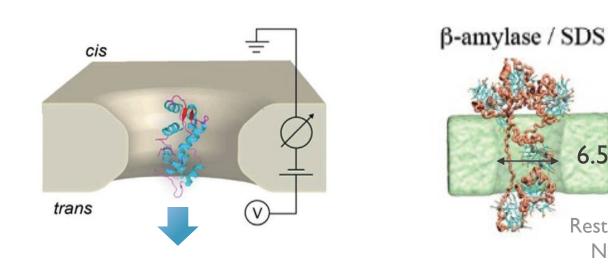


NPFET potentially breaks the translocation speed limit currently limiting nanopore technology

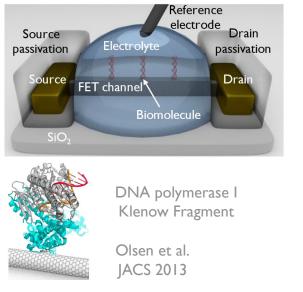
Solid-state nanopore

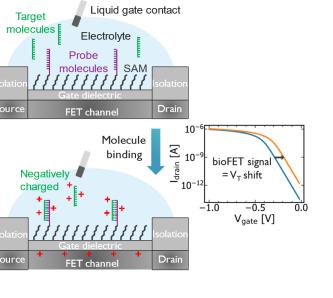




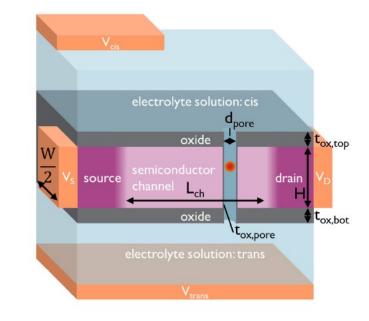




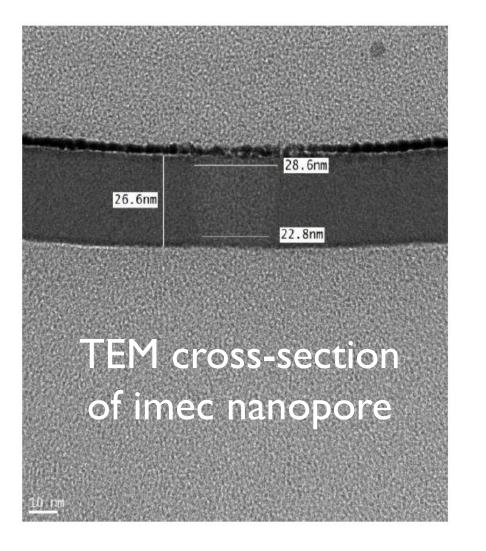


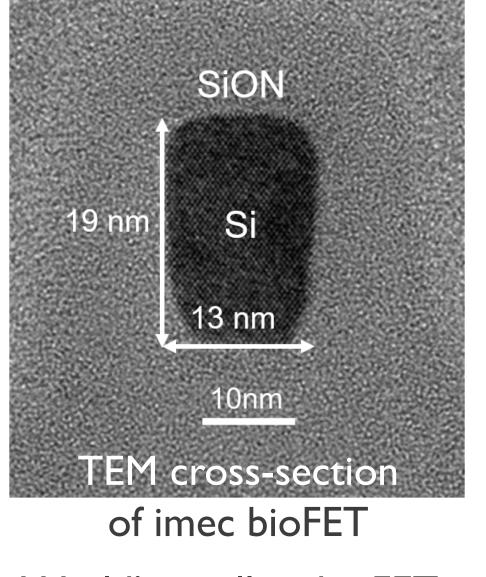


- Translocating molecule modulates the recorded ionic current passing through pore
- Chemically robust (vs. biopore)
 - Concentration dynamic range
 - misses large part of translocating proteins
 - senses long DNA
- Difficult to integrate on chip
- Molecular resolution toward sequencing undemonstrated
- Molecule docks close (I-I0nm) to FET surface. Molecule's charge modulates channel charge and FET current
- Chemically robust
 - Concentration dynamic range
 - surface attachment
- Gigascale integration on chip
 - Up to billions per chip
 - Molecular resolution determined by biochemistry



- A nanopore wrapped by a nanoscale fieldeffect transistor (FET). The FET senses single molecules translocating through the pore
- Chemically robust (vs. biopore)
- High concentration dynamic range
 - Potentially orders of magnitude more than MS
 - Chip potentially reads up to billions of molecules/s
- Megascale integration on chip
 - Up to millions per chip
- Molecular resolution toward sequencing undemonstrated





World's smallest bioFETs

