

# Introduction to “Next-Generation” Sequencing and its Applications

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**11AM** Introduction to NGS (M. Sammons)

**11:10AM** Practical advice for getting started in NGS (M. Sammons)

**11:40AM** Center for Functional Genomics NGS Workflow/Recommendations (S. Chittur)

**12:10PM** Applications: ATAC-seq to define cell lineages (M. Sammons)

**12:25PM** Q&A

**12:35PM** Short break

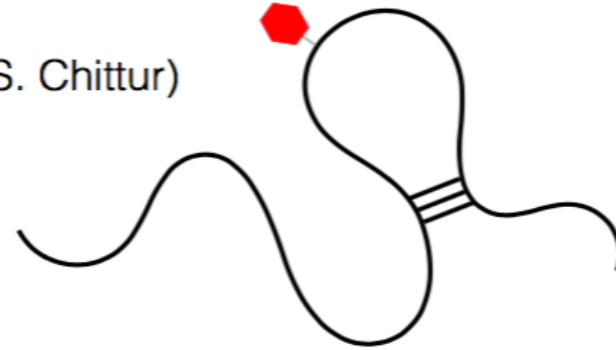
**12:45PM** Brief Primer: Using Galaxy for NGS data analysis (O. Novikova)

**1PM** Applications: RNA methylome analysis by NGS (J. Herschkowitz)

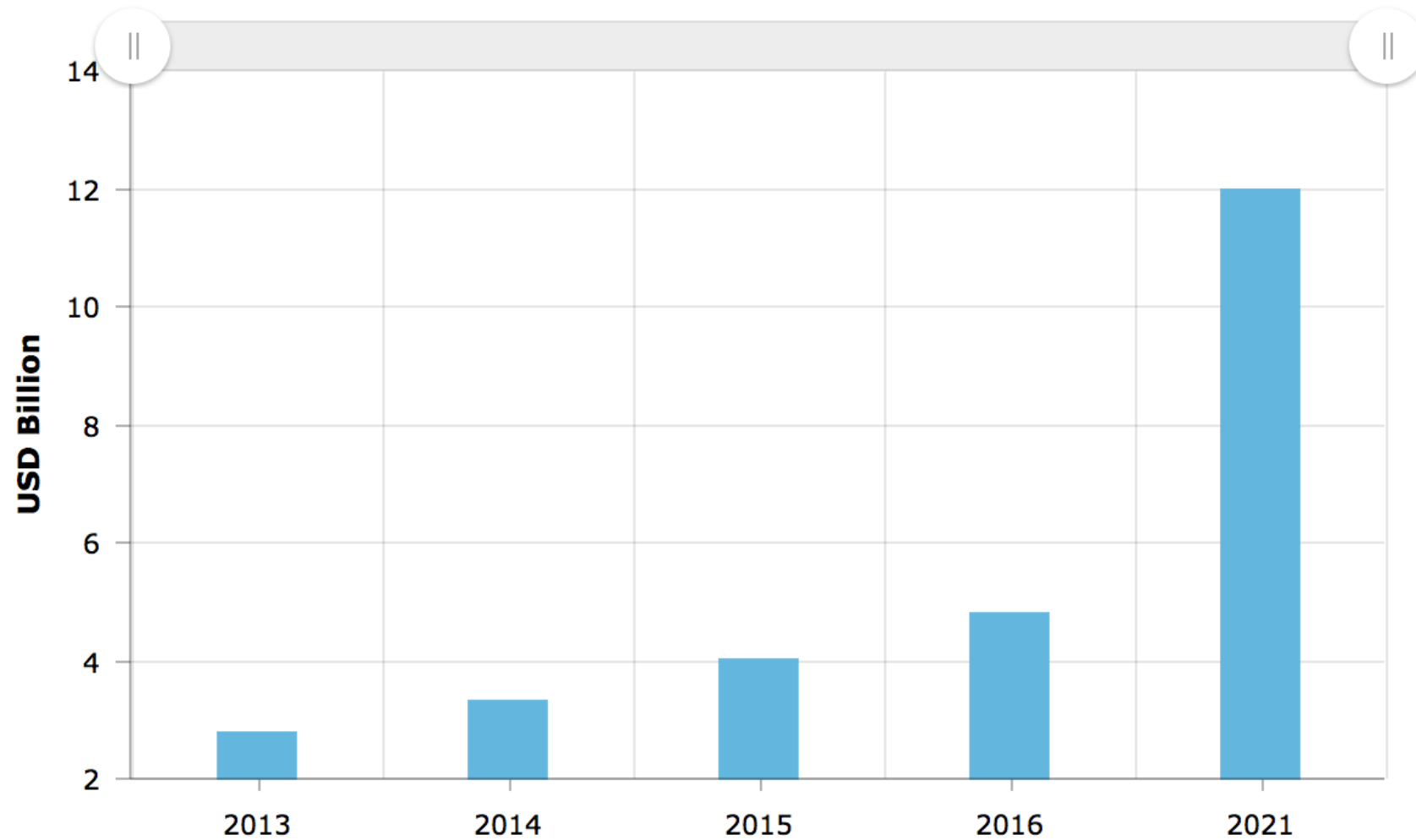
**1:15PM** Center for Functional Genomics Core Services (S. Chittur)

**1:45PM** Final Q&A

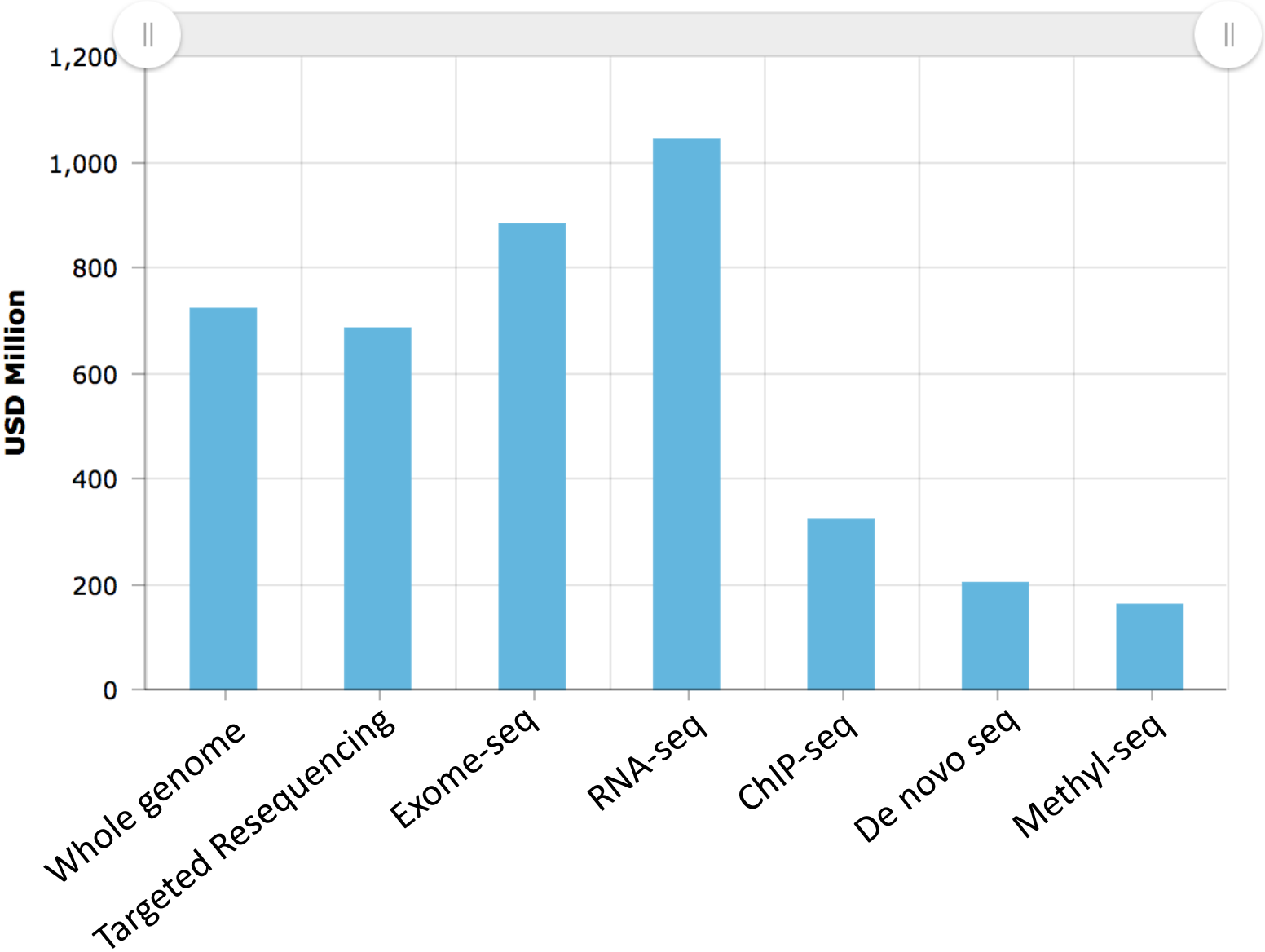
**2PM** End



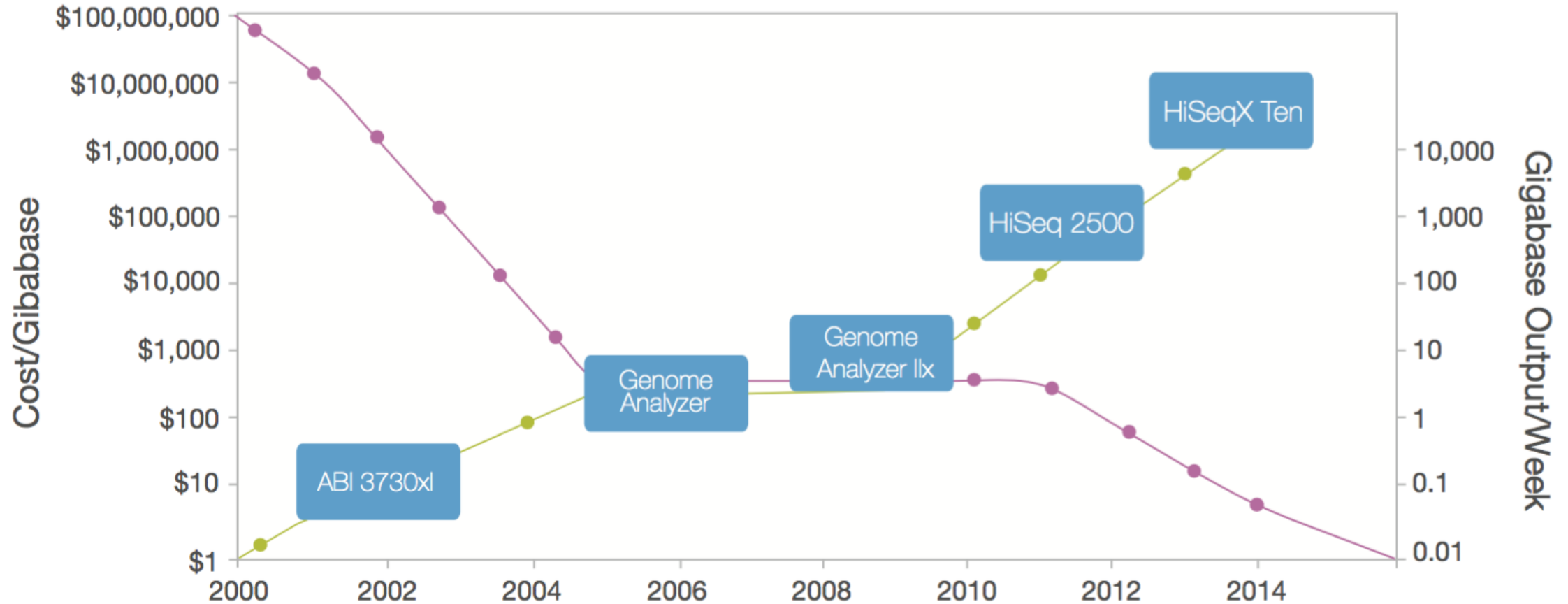
# Global Next-Generation Sequencing Expenditures



# Sequencing Applications by Dollars Spent



# Rapid rise in sequencing ability (and drop in cost)



## **RNA-seq**

Differential gene expression analysis → replacing/complimenting microarray

Discovery/quantification of novel genes/RNA species

Differential promoter, UTR, polyA signal analysis

Splicing analysis

Transcriptome assembly

*Ribosome footprinting*

Analysis of RBP binding to RNA

## **Whole genome sequencing/resequencing/exome sequencing**

Genome assembly (reference or sample/patient-specific)

Identification of mutations/rearrangements in disease

*Rapid identification of strain/viruses/metagenomic populations*

## **ChIP-seq/Chromatin analysis**

Identification of DNA binding protein genomewide localization

Epigenetics analysis (location of histones/modifications)

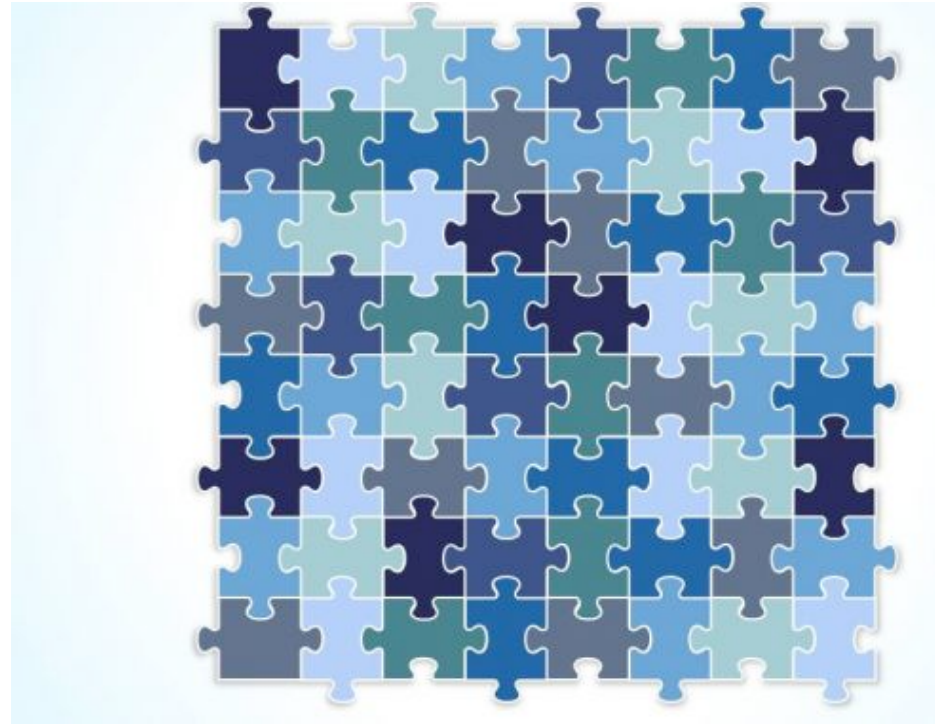
Identification of accessible/inaccessible chromatin regions

# What is next-generation sequencing?

- Catch-all term for high-throughput DNA sequencing
- Different companies, different technologies, *different uses*
- **Illumina** dominates short read “sequencing by synthesis”
- **Thermo** has short read options with different chemistry
  - 454/Roche and ABI are basically dead...Thermo is hanging in there
- **PacBio** excels at long reads (genome assembly/metagenomics)
- **Oxford Nanopore** does long and short reads with unique set of advantages over others (and disadvantages)

# PacBio

- Long reads help assembly of new genomes
- Complete 16S sequencing



Short reads

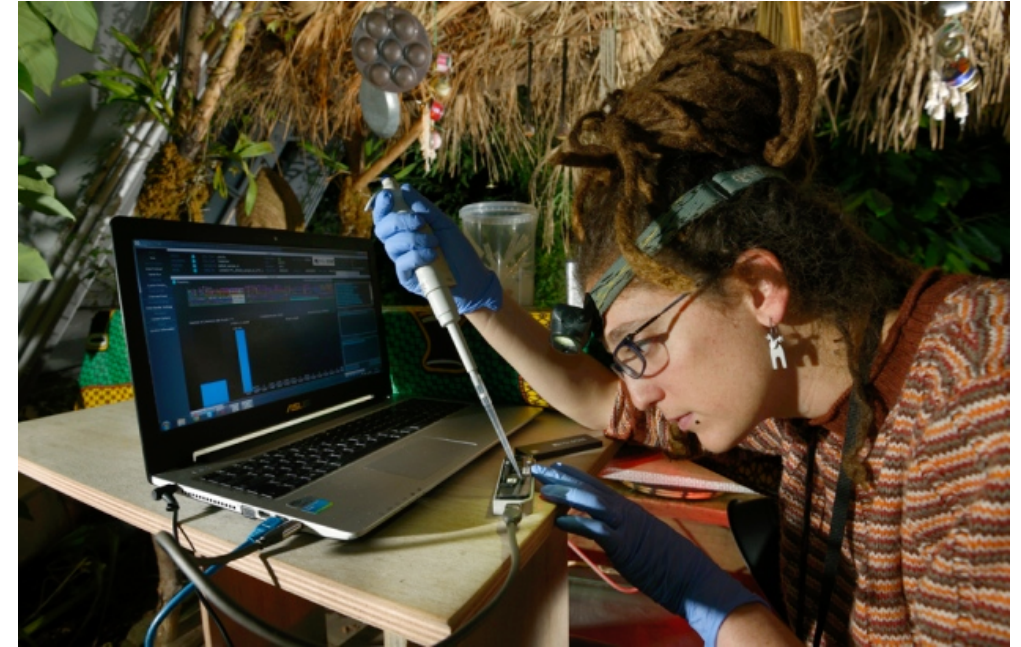


Long reads



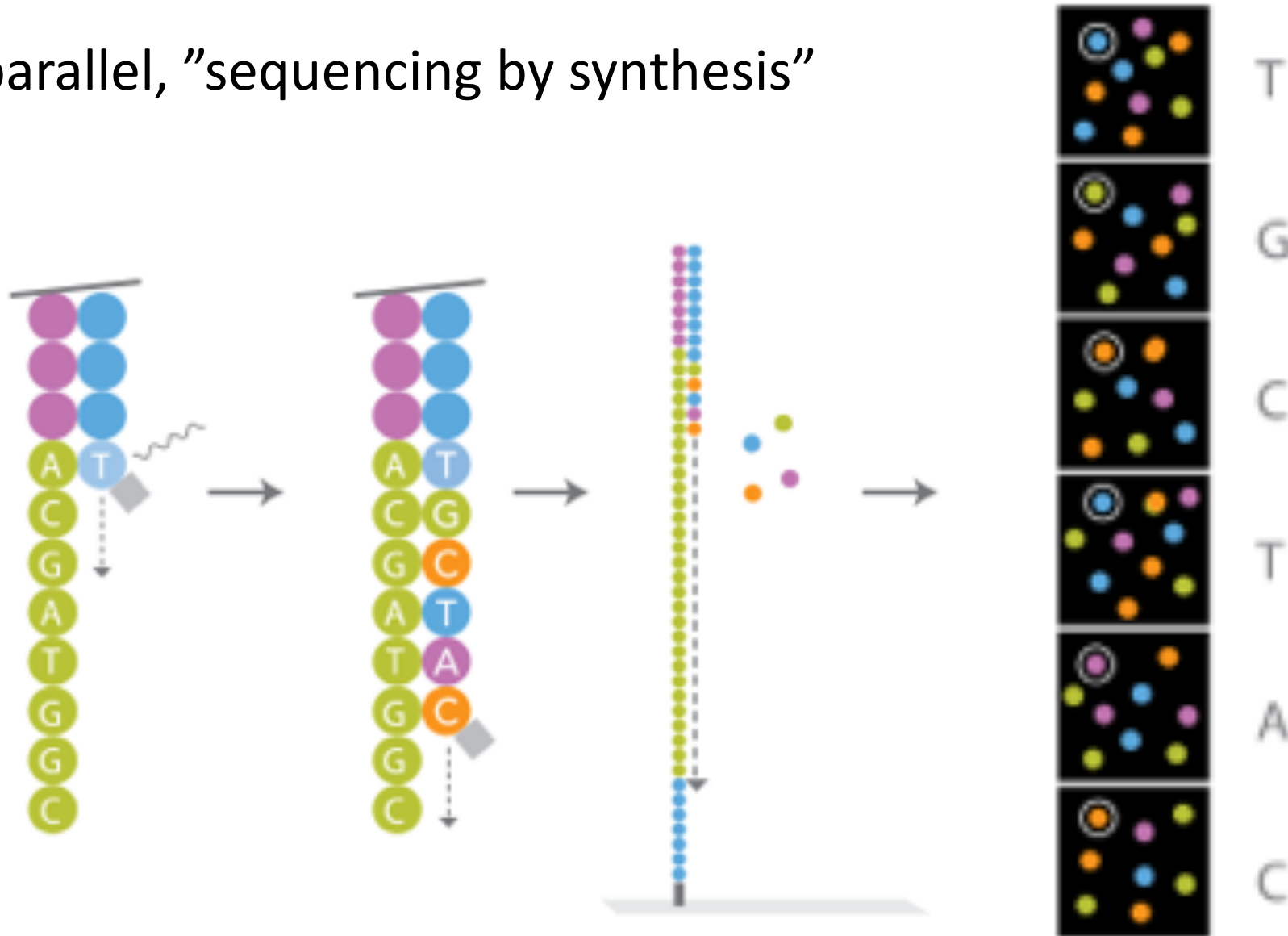
# Oxford Nanopore

- "Unique" nanopore sequencing chemistry
- Mobile, small, short and long reads possible



# Illumina

- Highly parallel, "sequencing by synthesis"



# For more information

[Introduction to Next-Generation Sequencing Technology](http://www.illumina.com/technology/next-generation-sequencing.html)

[www.illumina.com/technology/next-generation-sequencing.html](http://www.illumina.com/technology/next-generation-sequencing.html)

[EMBL-EBI Next-Gen Sequencing Practical Course Online](https://www.ebi.ac.uk/training/online/course/ebi-next-generation-sequencing-practical-course) (*Highly recommended!*)

<https://www.ebi.ac.uk/training/online/course/ebi-next-generation-sequencing-practical-course>



An Introduction to Next-Generation Sequencing Technology