## Three Generations of DNA Sequencing



## DNA Sequencing

Polymerase Chain Reaction (PCR) revolutionized the fields of biology and biochemistry


What we need for PCR Reaction:


Primers and DNA Polymerase

## PCR Termination



2',3'-dideoxyATP

## DNA Sequencing



## DNA Sequencing



## DNA Sequencing

Primer:
5 , $\qquad$ GG $\mathbf{3}^{\prime}$


## Electropherograms



## Next Generation Sequencing

Recent advances in sequencing methods has made genome sequencing very fast and cheap!

Human Genome Project (used chain termination method) took 10 years to complete and over $\$ 300$ million!

Next-Gen techniques allowed the genome of James Watson (of Watson and Crick fame) - the $3^{\text {rd }}$ sequenced human genome) to be sequenced in 2 months for less than $\$ 1 \mathrm{M}$.

Genome sequencing cost as estimated by NHGRI


## Next Generation Sequencing - Pyrosequencing

DNA is randomly sheared by sonication 300-500 bp fragments produced

dsDNA is melted apart to form single strands.
ssDNA are bound to DNA "capture" Beads SPRI Beads = solid phase reversible immobilization


This capture reaction happens under REALLY dilute conditions (< 1 DNA molecule per bead)

Under high salt conditions (2.5 M NaCl) and PEG, DNA binds to the carboxylate. Believe it, it


DNA is amplified on the bead by PCR until ~10 million copies are present

## Next Generation Sequencing - Pyrosequencing

Beads are transferred to a fiber optic slide with one bead per well.


Each well is 75 picoliters and one slide contains 1.6
million wells


> Pyrosequencing reactions occur in these wells.

1. One nucleotide (let's say dGTP) is added to the slide.
2. A 'burst' of light is seen (sensed through the fiber optic cable) if the dNTP is complementary.
3. Wash off residual dGTP
4. Add dATP - burst - wash
5. Add dCTP - burst - wash
6. Add dTTP - burst - wash
7. Back to $1^{\text {st }}$ step

## Next Generation Sequencing - Pyrosequencing

$$
\text { DNA }_{n \text { residues }}+\text { dNTP } \frac{\text { DNA polymerase }}{1} \text { DNA }_{n+1 \text { residues }}+\underset{\text { Pyrophosphate }}{\mathrm{P}_{2} \mathrm{O}_{7}^{4-}}
$$

$$
\mathrm{P}_{2} \mathrm{O}_{7}^{4-}+\quad \text { Adenosine } \underset{\mathrm{O}^{-}}{\stackrel{\mathrm{O}}{\mathrm{P}}-\mathrm{OSO}_{3}^{2-}} \xrightarrow[2]{\text { ATP sulfurylase }} \text { ATP }+\mathrm{SO}_{4}^{2-}
$$

Adenosine-5 '-phosphosulfate


Preparation for next step (d) $\mathrm{NTP}+2 \mathrm{H}_{2} \mathrm{O} \xrightarrow[4]{\text { apyrase }}$ (d) $\mathrm{NMP}+2 \mathrm{PO}_{4}^{3-}$

Once sequencing reactions are done, genome sequence is pieced back together (shotgun approach)

