

## X-RAY CRYSTALLOGRAPHY

- xray beam thru protein crystal → collection plate → fast fourier transform → electron density map
- density map
  - +in 3D
  - +needs to be high resolution to see alpha carbon backbone and side chains
    - >this obtained using fourier transformation software

## DOING A MOLECULAR DYNAMIC SIMULATION

- designed around Newtons laws of motion
- atoms test degrees of freedom
- enables protein to 'find' dif transition states
  - +for the mechanism of function

## LEC 8 – POLYMERASE CHAIN REACTION

### PCR

- amplify specific DNA sequences
  - +using oligonucleotide primers
    - >that're complementary to the 2 strands
    - >designed to allow extension towards eachother by a DNA Pol
- in vitro enzymatic reaction
  - +primer directed
  - +produces specific DNA sequence
- primers
  - +delineate DNA region to be amplified
- thermostable DNA Pol required
- cycles
  - +20-40 cycles
  - +denaturation
    - >one DNA target
    - >heated to 90-96 degrees
    - > 15-60 secs (longer for longer fragments)
    - >separated into single strands
  - +primer annealing
    - >50-70 degrees (depends on primer length and the sequence)
    - >two primers anneal (to the complementary sequences)
    - >primers define region for amplification
  - +primer extension (DNA Pol)
    - > 68-72 degrees (the optimal temp of enzyme)
    - >DNA synthesis
    - >DNA Pol (grabs the nucleotides and joins em) makes a copy of the template DNA
      - adds nucleotides to ends of primers
      - extends primer 5' → 3' on both strands
- needs
  - +target/template DNA
    - >single or double stranded
    - > 25 ng to 1 ug
  - +primers (forwards and reverse)

- >single stranded
- >18-30 bases
- >made in DNA synthesiser
- >distance between them defines length of PCR product
- +DNA Pol
  - >adds nucleotides to 3' end
  - >needs to be thermostable
  - >isolated from thermophilic bacteria
  - >Taq Pol from *Thermus aquaticus* (half life of 1.6 hours at 95 degrees)
  - >Pfy Pol from *Pyrococcus furiosus* (high fidelity)
  - >Tth Pol from *Thermus thermophilus* (long range PCR)
- +nucleotides (aka dNTPs)
  - >A (dATP)
  - >T (TTP)
  - >G (dGTP)
  - >C (dCTP)
- +Mg<sup>2+</sup> ions (via MgCl<sub>2</sub> usually)
  - >amount has effect on primer annealing
  - >too little = low product
  - >too much = nonspecific products

## PCR PRODUCTS

- aka amplicons
- mispriming
  - +band above product
  - +due to nonspecific binding of primers (to non-target sequences)
- primer dimers
  - +band below product

## PCR AUTOMATION

- thermal cyclers
- repeated temp change
- can sometimes have 1536 slots so large amounts can be run

## VARIATIONS OF PCR

- multiplex
  - +add multiple pairs of primers
  - +amplifies more than one target
  - +can distinguish b/t products due to dif sizes
  - +used to test for microorganisms in contaminated food/water
    - >test for a number of organisms at once
- long range
  - +normal pcr is 2-3kb
  - +can do fragments as big at 50kb
  - +extension step can be 30mins
- reverse transcription (RT-PCR)

