Tools for 3D molecule interactive visualization (115 mins)

Hugo Samano - Marta Strumillo

- 1.) Marta let's play
- 2.) Hugo let's work

Intro - hands on!

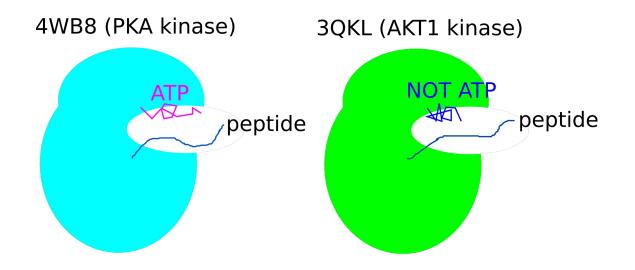
".ent", ".pdb", ".mol", ".xplor", ".mmod", ".ccp4", ".r3d", ".trj" and ".pse".

Intro - hands on!

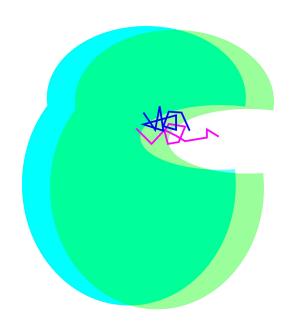
".ent", ".pdb", ".mol", ".xplor", ".mmod", ".ccp4", ".r3d", ".trj" and ".pse".

https://sites.google.com/site/afonsomduarte/pymol_tut

What do we have?



What are we going to do?



What do we want?





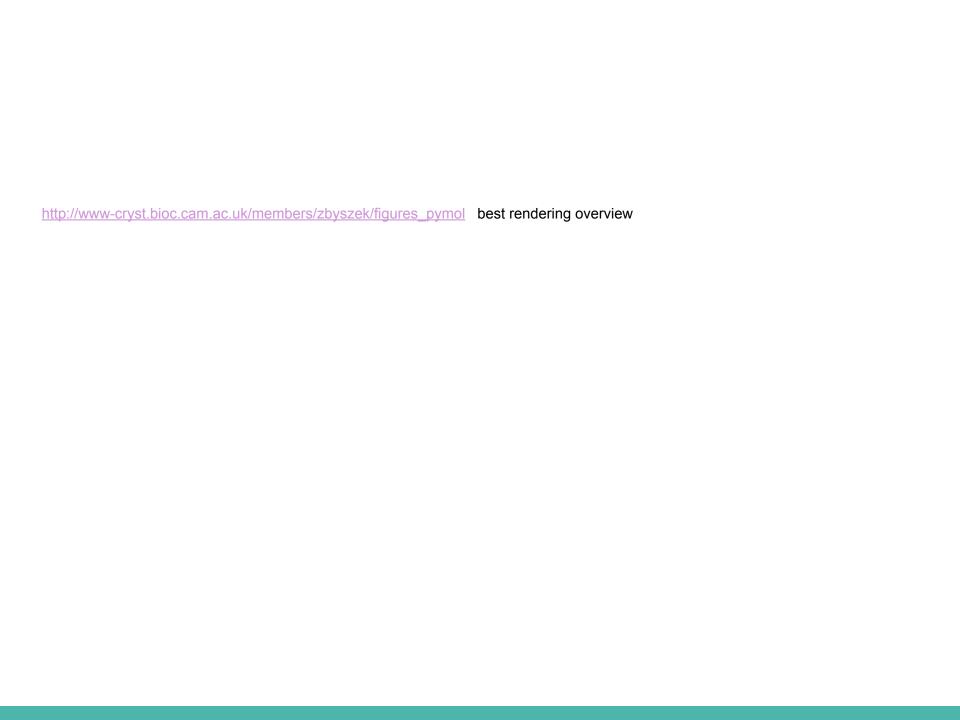
Let's go!

Take 4WB8 kinase, and move ATP to 3QKL

4WB8 - how many chains, what's in it

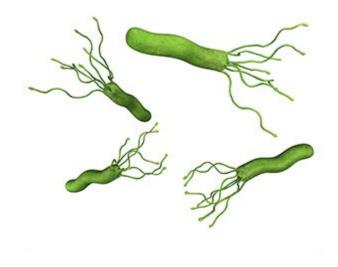
3QKL - how many chains, what's in it

Align, delete 4wb8, but not atp, delete inhibitor, but not 4wb8



Helicobacter pylori

Helicobacter pylori infects more than 50% of the human population Clinical isolates are categorized as Western or East Asian strains They differ by polymorphisms in the CagA gene

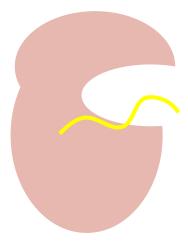


CagA inhibits a protein kinase



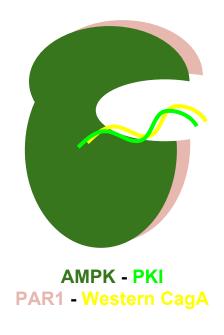
AMPK and PKI

both are endogenous proteins



Human PAR1 is inhibited by *Helicobacter pylori*CagA

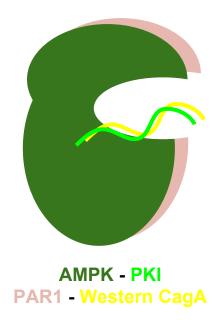
CagA inhibits a protein kinase



AMPK and PAR1 are structurally very similar

Western CagA mimics PKI in the way it binds to PAR1

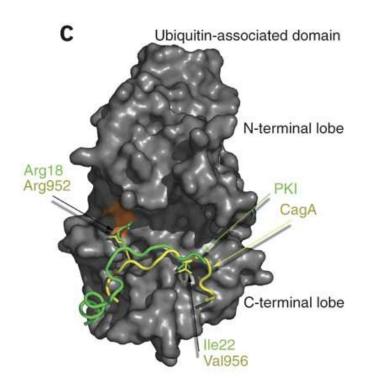
CagA inhibits a protein kinase



Is this mimicry property exclusive of the Western strain?



Modeling by homology



http://www.nature.com/nsmb/journal/v17/n1/full/nsmb.1705.html