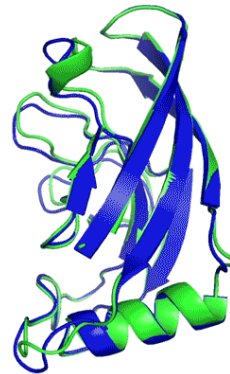


Augustin Žídek (Deepmind)

Protein Structure Prediction with Alphafold



T1037 / 6vr4
90.7 GDT
(RNA polymerase domain)



T1049 / 6y4f
93.3 GDT
(adhesin tip)

● Experimental result
● Computational prediction

12. 10. 2021. 3 p.m.

Fotochemie lecture hall, Viničná 7

OR

<https://cuni-cz.zoom.us/j/92629475400?pwd=dWhaOGR4VEc4SUFIVEgwZGZKQTJLdz09>

<https://bioinformatika.mff.cuni.cz/seminar/>