

DESIGN AND SYNTHESIS OF SMALL MOLECULES AS PROTEIN-PROTEIN INTERACTION STABILIZERS.

Markella Konstantinidou,⁽¹⁾ Zlata Boiarska,⁽¹⁾ Dario Antonacci,⁽¹⁾ Ave Kuusk,^(2,3) Hongming Chen,⁽³⁾ Christian Ottmann^(2,4) and Alexander Dömling⁽¹⁾

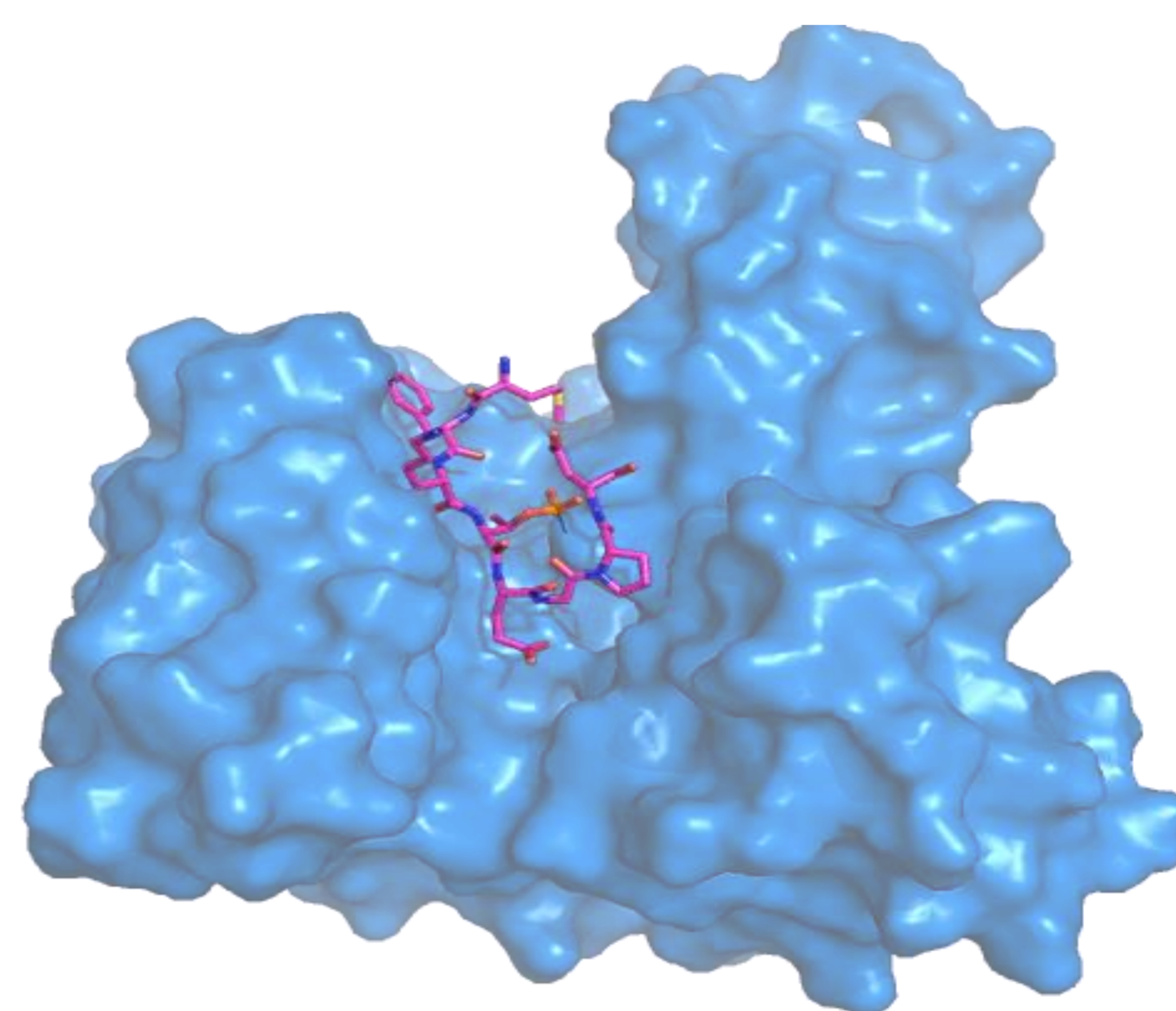
(1) Group of Drug Design, Groningen Research Institute of Pharmacy, University of Groningen, A. Deusinglaan 1, 9713 AV, Groningen the Netherlands, (2) Laboratory of Chemical Biology, Department of Biomedical Engineering and Institute for Complex Molecular Systems, Eindhoven University of Technology, The Netherlands, (3) Discovery Sciences, Innovative Medicines and Early Development Biotech Unit, AstraZeneca R&D Gothenburg, Mölndal, Sweden, (4) Department of Chemistry, University of Duisburg-Essen, Germany.

1 Introduction

Targeting protein – protein interactions (PPI) has emerged as a significant strategy in drug discovery.^[1] In this project we are focusing on the 14-3-3 protein family and more specifically on the interaction between 14-3-3 and p53.^[2] Since 14-3-3 is a positive regulator of p53 and p53 is mutated in 50% of human cancers, stabilization of the PPI is expected to lead to tumor suppression.^[3]

2 Design

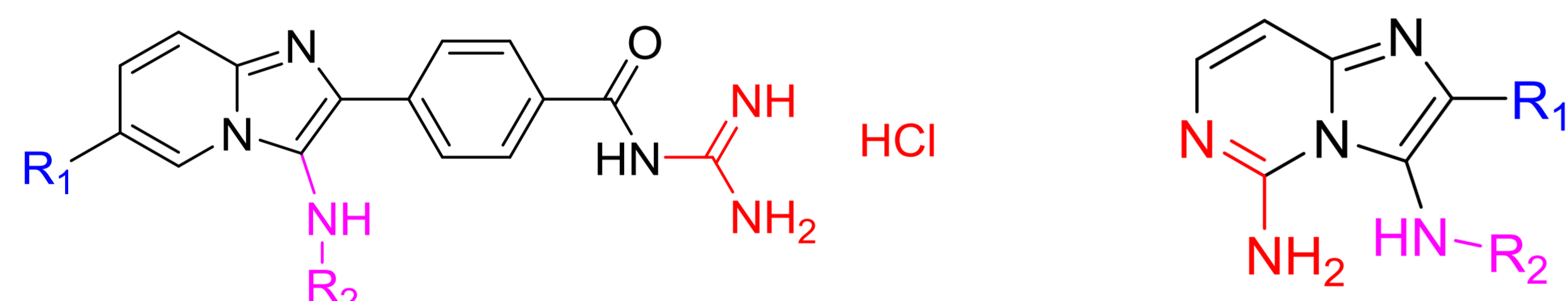
- The 14-3-3/p53 PPI is large, hydrophobic, with no deep pockets
- A polar fragment was co-crystallized with the PPI and showed a stabilization effect
- Having the fragment as starting point, we chose two scaffolds based on multi-component reaction chemistry (MCR) for the rapid generation of diverse libraries
- The scaffolds derive from the Groebke-Blackburn-Bienaymé reaction (GBB), which leads to flat, aromatic structures, with promising shape complementarity



The PPI of 14-3-3 (blue surface) with p53 (magenta sticks).

3 Synthesis

- Derivatives from 2 scaffolds were synthesized after docking studies



9 derivatives (Scaffold 1), 10 derivatives (Scaffold 2) were synthesized

- The two scaffolds can be accessed through a two step synthesis
- Commercially available starting materials are used
- Diversity is achieved through the two points of variation in each scaffold (blue and magenta substituents)
- The moieties in red are participating in the main interactions with the PPI

4 Biological evaluation

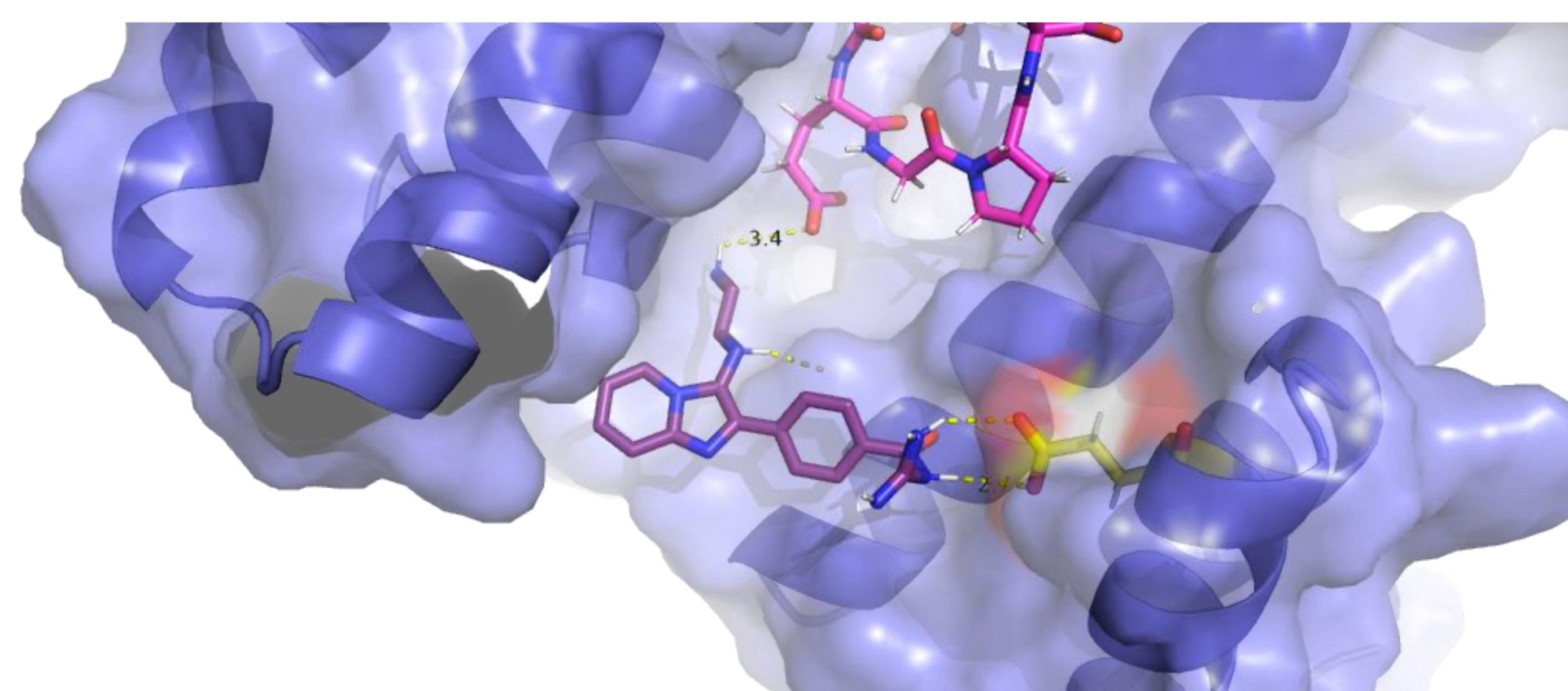
- Isothermal titration calorimetry (ITC)
- Surface plasmon resonance (SPR)
- Fluorescence polarization assay (FP)
- Crystallography

References

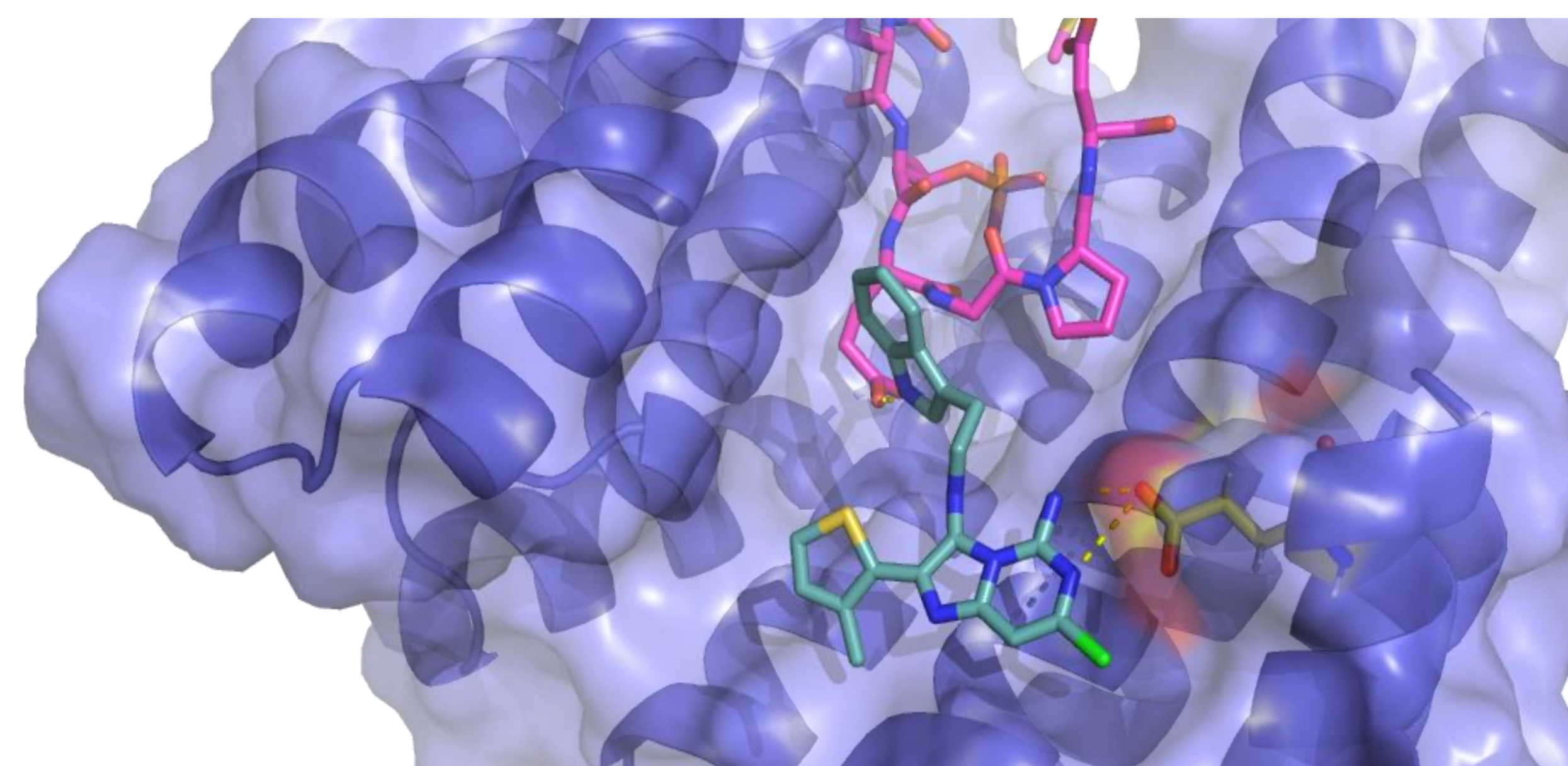
- [1] Jin L, Wang W, Fang G, Targeting protein-protein interaction by small molecules. *Annu. Rev. Pharmacol. Toxicol.* **2014**, *54*, 435-56
- [2] Hartman AM, Hirsch AKH. Molecular insight into specific 14-3-3 modulators: Inhibitors and stabilizers of protein-protein interactions of 14-3-3. *Eur. J. Med. Chem.* **2017**, *136*, 573-584
- [3] Doveston RG, Kuusk A, Andrei SA, Levsen S, Cao Q, Castaldi MP, Hendricks A, Brunsveld L, Chen H, Boyd H, Ottmann C. Small-molecule stabilization of the p53 - 14-3-3 protein-protein interaction. *FEBS Lett.* **2017**, *591*, 2449-2457

Acknowledgement

This project has received funding from the European Union's Framework Programme for Research and Innovation Horizon 2020 (2014-2020) under the Marie Skłodowska-Curie Grant Agreement No. 675555, Accelerated Early staGe drug discovery (AEGIS).



Docking pose for derivative of Scaffold 1



Docking pose for derivative of Scaffold 2