

## Posters Session Monday

- 1 Esam Abualrous **Natural polymorphism modulates the susceptibility of MHC class II proteins to HLA-DM**
- 2 Reid Alderson **Structural basis for the dysregulation of HSP27 in Charcot-Marie-Tooth disease**
- 3 Robert Arbon **Assessing the generalizability of adaptive sampling policies**
- 4 Ernest Awoonor-Williams **Modelling the binding free energy of peptidomimetic inhibitors to SARS-CoV-2 M<sup>pro</sup>**
- 5 Cory Ayres **Force pull quasiequilibrium: investigating the impact of force on tcr/peptide-mhc stability and immunological recognition**
- 6 Hossein Batebi **Formation of a  $\beta 2AR^*$ -Gs<sup>GDP</sup> intermediate complex**
- 7 Miriam Bertazzon **Investigation of the functional role of CD2BP2**
- 8 Johann Biedermann **Investigation of cation permeation through AMPA receptors by molecular dynamics simulations**
- 9 Yasemin Bozkurt Varolgünes **Allostery in proteins as point-to-point telecommunication in a network: frequency decomposed signal-to-noise-ratio and channel capacity analysis**
- 10 Matteo Castelli **Exploring the effects of post-translational modifications through computational approaches**
- 11 Liwei Chang **Inferring protein folding pathways from accelerated molecular dynamics**

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| 12 | Yaoyi Chen        | <b>Machine learning implicit solvent method for molecular dynamics</b>   |
| 13 | Moumita Dasgupta  | <b>Biophysical characterization of interactions of T cell receptor with viral antigens derived from lymphocytic choriomeningitis virus</b> |
| 14 | Andrea Di Luca    | <b>How global motions are involved in the deactivation of complex I</b>  |
| 15 | Kathrin Funck     | <b>Structural and functional explorations of the MICOS complex</b>   |
| 16 | Toni Giorgino     | <b>Nanobody interaction unveils structure, dynamics and proteotoxicity of the Finnish-type amyloidogenic gelsolin variant</b>              |
| 17 | Giulia Glorani    | <b>An unusual aspartic acid cluster in the reovirus attachment fiber <math>\sigma</math>1 mediates stability at low pH</b>                 |
| 18 | Sebastian Günther | <b>Temperature jump X-ray crystallography to analyze protein dynamics in macromolecular crystals</b>                                       |
| 19 | Shozeb Haider     | <b>Allosteric communication in Class A <math>\beta</math>-lactamases</b>   |
| 20 | Nandan Haloi      | <b>Role of the internal loops in gating of outer membrane porins</b>   |
| 21 | Tim Hempel        | <b>Molecular mechanism of inhibiting the SARS-CoV-2 cell entry facilitator TMPRSS2 with camostat and nafamostat</b>                        |
| 22 | Miriam Jäger      | <b>Predicting ion channel conductance from dissipation-corrected TMD and LE simulations</b>  |

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| <b>23</b> | Eoin Kearney        | <b>Molecular dynamics simulations of uranyl adsorption on montmorillonite clay with desferrioxamine-B</b>              |
| <b>24</b> | Grant Keller        | <b>Determinants of antigenicity in tumor neoepitopes for the development of personalized/multiple peptide vaccines</b> |
| <b>25</b> | Christopher Kolloff | <b>NOE spin diffusion in large biomolecules</b>  |