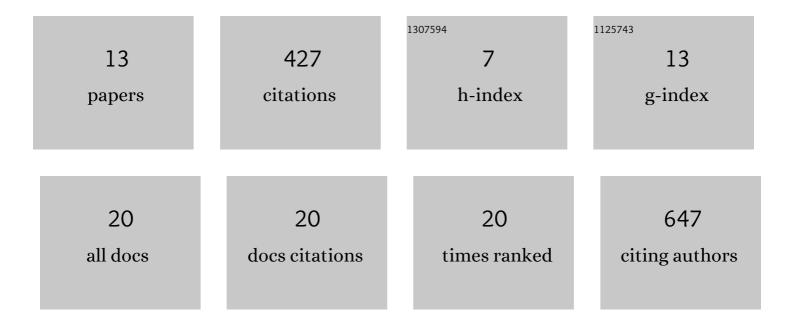
## Marton Olbei

List of Publications by Year in descending order

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MADTON OLDEL

| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | SignaLink3: a multi-layered resource to uncover tissue-specific signaling networks. Nucleic Acids Research, 2022, 50, D701-D709.  | 14.5 | 19        |
| 2  | Analysing miRNA-Target Gene Networks in Inflammatory Bowel Disease and Other Complex Diseases<br>Using Transcriptomic Data. Genes, 2022, 13, 370.   | 2.4  | 4         |
| 3  | Mapping the epithelial–immune cell interactome upon infection in the gut and the upper airways. Npj<br>Systems Biology and Applications, 2022, 8, 15.   | 3.0  | 3         |
| 4  | Evolution of regulatory networks associated with traits under selection in cichlids. Genome Biology, 2021, 22, 25.  | 8.8  | 17        |
| 5  | ViralLink: An integrated workflow to investigate the effect of SARS-CoV-2 on intracellular signalling and regulatory pathways. PLoS Computational Biology, 2021, 17, e1008685.                          | 3.2  | 11        |
| 6  | SARS-CoV-2 Causes a Different Cytokine Response Compared to Other Cytokine Storm-Causing Respiratory Viruses in Severely III Patients. Frontiers in Immunology, 2021, 12, 629193.                       | 4.8  | 91        |
| 7  | Integrated intra―and intercellular signaling knowledge for multicellular omics analysis. Molecular<br>Systems Biology, 2021, 17, e9923.   | 7.2  | 152       |
| 8  | CytokineLink: A Cytokine Communication Map to Analyse Immune Responses—Case Studies in<br>Inflammatory Bowel Disease and COVID-19. Cells, 2021, 10, 2242.   | 4.1  | 5         |
| 9  | COVID19 Disease Map, a computational knowledge repository of virus–host interaction mechanisms.<br>Molecular Systems Biology, 2021, 17, e10387.   | 7.2  | 53        |
| 10 | ULK1 and ULK2 are less redundant than previously thought: computational analysis uncovers distinct regulation and functions of these autophagy induction proteins. Scientific Reports, 2020, 10, 10940. | 3.3  | 23        |
| 11 | Network Biology Approaches to Identify Molecular and Systems-Level Differences Between Salmonella<br>Pathovars. Methods in Molecular Biology, 2019, 1918, 265-273.                                      | 0.9  | 2         |
| 12 | SalmoNet, an integrated network of ten Salmonella enterica strains reveals common and distinct pathways to host adaptation. Npj Systems Biology and Applications, 2017, 3, 31.                          | 3.0  | 28        |
| 13 | Sherlock: an open-source data platform to store, analyze and integrate Big Data for biology.<br>F1000Research, 0, 10, 409.  | 1.6  | 1         |