Veljko Veljkovic

List of Publications by Year in descending order

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687363 642732 30 617 13 23 citations h-index g-index papers 32 32 32 701 docs citations times ranked citing authors all docs

| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Identification of SARSâ€CoVâ€2 Papainâ€like Protease (PLpro) Inhibitors Using Combined Computational Approach**. ChemistryOpen, 2022, 11, e202100248. | 1.9 | 8 |
| 2 | <i>Ehrlichia</i> SLiM Ligand Mimetic Activates Notch Signaling in Human Monocytes. MBio, 2022, 13, e0007622. | 4.1 | 11 |
| 3 | Biological Rationale for the Repurposing of BCG Vaccine against SARS-CoV-2. Journal of Proteome Research, 2020, 19, 4649-4654. | 3.7 | 11 |
| 4 | Use of the informational spectrum methodology for rapid biological analysis of the novel coronavirus 2019-nCoV: prediction of potential receptor, natural reservoir, tropism and therapeutic/vaccine target. F1000Research, 2020, 9, 52. | 1.6 | 20 |
| 5 | Use of the informational spectrum methodology for rapid biological analysis of the novel coronavirus 2019-nCoV: prediction of potential receptor, natural reservoir, tropism and therapeutic/vaccine target. F1000Research, 2020, 9, 52. | 1.6 | 19 |
| 6 | Zika virus infection elicits auto-antibodies to C1q. Scientific Reports, 2018, 8, 1882. | 3.3 | 21 |
| 7 | Using electronic biology based platform to predict flu vaccine efficacy for 2018/2019. F1000Research, 2018, 7, 298. | 1.6 | 10 |
| 8 | Using electronic biology based platform to predict flu vaccine efficacy for 2018/2019. F1000Research, 2018, 7, 298. | 1.6 | 5 |
| 9 | A simple method for calculation of basic molecular properties of nutrients and their use as a criterion for a healthy diet. F1000Research, 2017, 6, 13. | 1.6 | 2 |
| 10 | Prediction of influenza vaccine effectiveness for the influenza season 2017/18 in the US. F1000Research, 2017, 6, 2067. | 1.6 | 12 |
| 11 | Possible repurposing of seasonal influenza vaccine for prevention of Zika virus infection. F1000Research, 2016, 5, 190. | 1.6 | 8 |
| 12 | Possible repurposing of seasonal influenza vaccine for prevention of Zika virus infection. F1000Research, 2016, 5, 190. | 1.6 | 9 |
| 13 | Predicted Enhanced Human Propensity of Current Avian-Like H1N1 Swine Influenza Virus from China. PLoS ONE, 2016, 11, e0165451. | 2.5 | 2 |
| 14 | In Silico Prediction and Experimental Confirmation of HA Residues Conferring Enhanced Human Receptor Specificity of H5N1 Influenza A Viruses. Scientific Reports, 2015, 5, 11434. | 3.3 | 53 |
| 15 | Evolution of 2014/15 H3N2 Influenza Viruses Circulating in US: Consequences for Vaccine Effectiveness and Possible New Pandemic. Frontiers in Microbiology, 2015, 6, 1456. | 3.5 | 13 |
| 16 | In silico analysis suggests interaction between Ebola virus and the extracellular matrix. Frontiers in Microbiology, 2015, 6, 135. | 3.5 | 24 |
| 17 | Virtual screen for repurposing approved and experimental drugs for candidate inhibitors of EBOLA virus infection. F1000Research, 2015, 4, 34. | 1.6 | 41 |
| 18 | In silico analysis suggests repurposing of ibuprofen for prevention and treatment of EBOLA virus disease. F1000Research, 2015, 4, 104. | 1.6 | 23 |

| # | Article | lF | CITATIONS |
|----|---|-----|-----------|
| 19 | Influenza vaccine as prevention for cardiovascular diseases: Possible molecular mechanism. Vaccine, 2014, 32, 6569-6575. | 3.8 | 51 |
| 20 | Novel Phylogenetic Algorithm to Monitor Human Tropism in Egyptian H5N1-HPAIV Reveals Evolution toward Efficient Human-to-Human Transmission. PLoS ONE, 2013, 8, e61572. | 2.5 | 33 |
| 21 | The role of long-range intermolecular interactions in discovery of new drugs. Expert Opinion on Drug Discovery, 2011, 6, 1263-1270. | 5.0 | 26 |
| 22 | Characterization of conserved properties of hemagglutinin of H5N1 and human influenza viruses: possible consequences for therapy and infection control. BMC Structural Biology, 2009, 9, 21. | 2.3 | 67 |
| 23 | Identification of hemagglutinin structural domain and polymorphisms which may modulate swine H1N1 interactions with human receptor. BMC Structural Biology, 2009, 9, 62. | 2.3 | 62 |
| 24 | Discovery of New Therapeutic Targets by the Informational Spectrum Method. Current Protein and Peptide Science, 2008, 9, 493-506. | 1.4 | 34 |
| 25 | Evolution of SARS-CoV-2 virus and assessment of the effectiveness of COVID-19 vaccine. F1000Research, 0, 10, 28. | 1.6 | 2 |
| 26 | Should safety of the flu vaccine for cancer patients be reexamined?. F1000Research, 0, 7, 1. | 1.6 | 12 |
| 27 | (Re)emergence of A(H1N1)pdm09Â influenza viruses with pandemic markers in the 2018/2019 flu season in the USA. F1000Research, 0, 8, 141. | 1.6 | 1 |
| 28 | Use of the informational spectrum methodology for rapid biological analysis of the novel coronavirus 2019-nCoV: prediction of potential receptor, natural reservoir, tropism and therapeutic/vaccine target. F1000Research, 0, 9, 52. | 1.6 | 4 |
| 29 | Use of the informational spectrum methodology for rapid biological analysis of the novel coronavirus 2019-nCoV: prediction of potential receptor, natural reservoir, tropism and therapeutic/vaccine target. F1000Research, 0, 9, 52. | 1.6 | 23 |
| 30 | Prediction of the effectiveness of COVID-19 vaccine candidates. F1000Research, 0, 9, 365. | 1.6 | 5 |