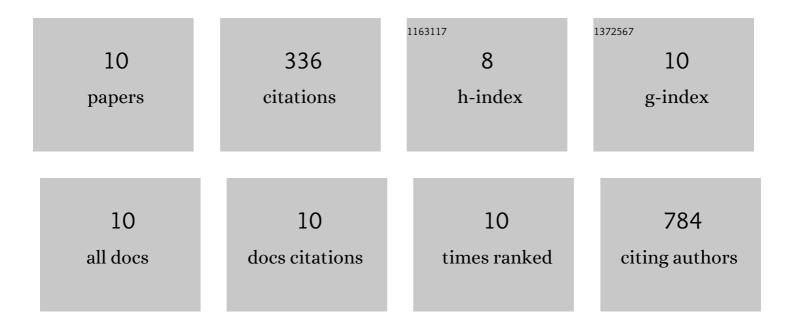
Bhaskar Dutta

List of Publications by Year in descending order

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RHASKAD DUITTA

| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | A small sustained increase in NOD1 abundance promotes ligand-independent inflammatory and oncogene transcriptional responses. Science Signaling, 2020, 13, . | 3.6 | 6 |
| 2 | IFN-mediated negative feedback supports bacteria class-specific macrophage inflammatory responses. ELife, 2019, 8, . | 6.0 | 16 |
| 3 | Dual Roles for Ikaros in Regulation of Macrophage Chromatin State and Inflammatory Gene Expression. Journal of Immunology, 2018, 201, 757-771. | 0.8 | 43 |
| 4 | Proteome and Secretome Analysis Reveals Differential Post-transcriptional Regulation of Toll-like Receptor Responses. Molecular and Cellular Proteomics, 2017, 16, S172-S186. | 3.8 | 29 |
| 5 | Genome-wide siRNA screen of genes regulating the LPS-induced NF-κB and TNF-α responses in mouse macrophages. Scientific Data, 2017, 4, 170008. | 5.3 | 7 |
| 6 | Genome-wide siRNA screen of genes regulating the LPS-induced TNF-α response in human macrophages. Scientific Data, 2017, 4, 170007. | 5.3 | 11 |
| 7 | Systematic Investigation of Multi-TLR Sensing Identifies Regulators of Sustained Gene Activation in Macrophages. Cell Systems, 2017, 5, 25-37.e3. | 6.2 | 48 |
| 8 | Distinct NF-κB and MAPK Activation Thresholds Uncouple Steady-State Microbe Sensing from Anti-pathogen Inflammatory Responses. Cell Systems, 2016, 2, 378-390. | 6.2 | 97 |
| 9 | An interactive web-based application for Comprehensive Analysis of RNAi-screen Data. Nature Communications, 2016, 7, 10578. | 12.8 | 13 |
| 10 | Comprehensive RNAi-based screening of human and mouse TLR pathways identifies species-specific preferences in signaling protein use. Science Signaling, 2016, 9, ra3. | 3.6 | 66 |