## Hagit T Porath

## List of Publications by Year

 in descending orderSource: https:/|exaly.com/author-pdf/5901099/publications.pdf
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| 1 | Analysis of Intron Sequences Reveals Hallmarks of Circular RNA Biogenesis in Animals. Cell Reports, 2015, 10, 170-177. | 6.4 | 918 |
| :---: | :---: | :---: | :---: |
| 2 | Trade-off between Transcriptome Plasticity and Genome Evolution in Cephalopods. Cell, 2017, 169, 191-202.ell. | 28.9 | 268 |
| 3 | Elevated RNA Editing Activity Is a Major Contributor to Transcriptomic Diversity in Tumors. Cell Reports, 2015, 13, 267-276. | 6.4 | 262 |
| 4 | A genome-wide map of hyper-edited RNA reveals numerous new sites. Nature Communications, 2014, 5, 4726. | 12.8 | 193 |
| 5 | Massive A-to-I RNA editing is common across the Metazoa and correlates with dsRNA abundance. Genome Biology, 2017, 18, 185. | 8.8 | 118 |
| 6 | RNA editing by ADAR1 leads to context-dependent transcriptome-wide changes in RNA secondary structure. Nature Communications, 2017, 8, 1440. | 12.8 | 77 |
| 7 | Fmrp Interacts with Adar and Regulates RNA Editing, Synaptic Density and Locomotor Activity in Zebrafish. PLoS Genetics, 2015, 11, el005702. | 3.5 | 76 |
| 8 | RNA editing is abundant and correlates with task performance in a social bumblebee. Nature Communications, 2019, 10, 1605. | 12.8 | 57 |
| 9 | Dynamic hyper-editing underlies temperature adaptation in Drosophila. PLoS Genetics, 2017, 13, e1006931. | 3.5 | 51 |
| 10 | A-to-I RNA Editing in the Earliest-Diverging Eumetazoan Phyla. Molecular Biology and Evolution, 2017, 34, 1890-1901. | 8.9 | 45 |
| 11 | ROP: dumpster diving in RNA-sequencing to find the source of 1 trillion reads across diverse adult human tissues. Genome Biology, 2018, 19, 36. | 8.8 | 42 |
| 12 | Purifying selection of long dsRNA is the first line of defense against false activation of innate immunity. Genome Biology, 2020, 21, 26. | 8.8 | 31 |
| 13 | A-to-I RNA editing in honeybees shows signals of adaptation and convergent evolution. IScience, 2021, 24, 101983. | 4.1 | 18 | 24, 101983.

