## Lovorka Stojic

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/809276/publications.pdf

Version: 2024-02-01

21 papers 1,924 citations

430874 18 h-index 713466 21 g-index

23 all docs

23 docs citations

 $\begin{array}{c} 23 \\ times \ ranked \end{array}$ 

3283 citing authors

| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | CCT3- <i>LINC00326</i> axis regulates hepatocarcinogenic lipid metabolism. Gut, 2022, 71, 2081-2092.   | 12.1 | 32        |
| 2  | Long Noncoding RNAs at the Crossroads of Cell Cycle and Genome Integrity. Trends in Genetics, 2021, 37, 528-546.   | 6.7  | 23        |
| 3  | A high-content RNAi screen reveals multiple roles for long noncoding RNAs in cell division. Nature Communications, 2020, 11, 1851.   | 12.8 | 43        |
| 4  | Tuning the Expression of Long Noncoding RNA Loci with CRISPR Interference. Methods in Molecular Biology, 2020, 2161, 1-16.   | 0.9  | 2         |
| 5  | SAM68 is required for regulation of Pumilio by the NORAD long noncoding RNA. Genes and Development, 2018, 32, 70-78.   | 5.9  | 61        |
| 6  | Specificity of RNAi, LNA and CRISPRi as loss-of-function methods in transcriptional analysis. Nucleic Acids Research, 2018, 46, 5950-5966.   | 14.5 | 101       |
| 7  | Neurodevelopmental protein Musashi-1 interacts with the Zika genome and promotes viral replication. Science, 2017, 357, 83-88.   | 12.6 | 152       |
| 8  | Aging increases cell-to-cell transcriptional variability upon immune stimulation. Science, 2017, 355, 1433-1436.   | 12.6 | 265       |
| 9  | Transcriptional silencing of long noncoding RNA GNG12-AS1 uncouples its transcriptional and product-related functions. Nature Communications, 2016, 7, 10406.  | 12.8 | 77        |
| 10 | 5-hydroxymethylcytosine marks promoters in colon that resist DNA hypermethylation in cancer. Genome Biology, 2015, 16, 69.   | 8.8  | 60        |
| 11 | Imprinted Chromatin around DIRAS3 Regulates Alternative Splicing of GNG12-AS1, a Long Noncoding RNA. American Journal of Human Genetics, 2013, 93, 224-235.  | 6.2  | 41        |
| 12 | Molecular mechanisms of genomic imprinting and clinical implications for cancer. Expert Reviews in Molecular Medicine, 2011, 13, e2.   | 3.9  | 66        |
| 13 | Chromatin regulated interchange between polycomb repressive complex 2 (PRC2)-Ezh2 and PRC2-Ezh1 complexes controls myogenin activation in skeletal muscle cells. Epigenetics and Chromatin, 2011, 4, 16. | 3.9  | 113       |
| 14 | Enhancer of Zeste Homolog 2 Overexpression Has a Role in the Development of Anaplastic Thyroid Carcinomas. Journal of Clinical Endocrinology and Metabolism, 2011, 96, 1029-1038.                        | 3.6  | 62        |
| 15 | Mismatch Repair Status and the Response of Human Cells to Cisplatin. Cell Cycle, 2007, 6, 1796-1802.   | 2.6  | 40        |
| 16 | High Doses of SN1 Type Methylating Agents Activate DNA Damage Signaling Cascades that are Largely Independent of Mismatch Repair. Cell Cycle, 2005, 4, 473-477.  | 2.6  | 40        |
| 17 | Mismatch repair-dependent G2 checkpoint induced by low doses of SN1 type methylating agents requires the ATR kinase. Genes and Development, 2004, 18, 1331-1344.   | 5.9  | 206       |
| 18 | Is mismatch repair really required for ionizing radiation–induced DNA damage signaling?. Nature Genetics, 2004, 36, 432-433.   | 21.4 | 18        |

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|----|---|-----|-----------|
| 19 | Mismatch repair and DNA damage signalling. DNA Repair, 2004, 3, 1091-1101.  | 2.8 | 340       |
| 20 | Methylation-induced G2/M arrest requires a full complement of the mismatch repair protein hMLH1. EMBO Journal, 2003, 22, 2245-2254.   | 7.8 | 160       |
| 21 | Mismatch repair-dependent transcriptome changes in human cells treated with the methylating agent N-methyl-n'-nitro-N-nitrosoguanidine. Cancer Research, 2003, 63, 8158-66. | 0.9 | 18        |