## Lissette Delgado-Cruzata

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3776455/publications.pdf

Version: 2024-02-01

18 papers

1,101 citations

567281 15 h-index 18 g-index

18 all docs 18 docs citations

18 times ranked

2092 citing authors

| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Mismatch Repair Polymorphisms as Markers of Breast Cancer Prevalence in the Breast Cancer Family Registry. Anticancer Research, 2016, 36, 4437-4442.   | 1.1  | 24        |
| 2  | Dietary Modifications, Weight Loss, and Changes in Metabolic Markers Affect Global DNA Methylation in Hispanic, African American, and Afro-Caribbean Breast Cancer Survivors,. Journal of Nutrition, 2015, 145, 783-790. | 2.9  | 59        |
| 3  | Genome-Wide Methylation Analyses in Glioblastoma Multiforme. PLoS ONE, 2014, 9, e89376.  | 2.5  | 45        |
| 4  | Differences in DNA methylation by extent of breast cancer family history in unaffected women. Epigenetics, 2014, 9, 243-248.   | 2.7  | 23        |
| 5  | Correlations in global DNA methylation measures in peripheral blood mononuclear cells and granulocytes. Epigenetics, 2014, 9, 1504-1510.   | 2.7  | 15        |
| 6  | DNA double-strand break repair genotype and phenotype and breast cancer risk within sisters from the New York site of the Breast Cancer Family Registry (BCFR). Cancer Causes and Control, 2013, 24, 2157-2168.          | 1.8  | 14        |
| 7  | Genomic Methylation Changes Over Time in Peripheral Blood Mononuclear Cell DNA: Differences by Assay Type and Baseline Values. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1314-1318.                       | 2.5  | 24        |
| 8  | Global DNA methylation levels in white blood cell DNA from sisters discordant for breast cancer from the New York site of the Breast Cancer Family Registry. Epigenetics, 2012, 7, 868-874.                              | 2.7  | 40        |
| 9  | DNA Methylation Changes Correlate with Gleason Score and Tumor Stage in Prostate Cancer. DNA and Cell Biology, 2012, 31, 187-192.  | 1.9  | 39        |
| 10 | Adult global DNA methylation in relation to pre-natal nutrition. International Journal of Epidemiology, 2012, 41, 116-123.   | 1.9  | 64        |
| 11 | Repetitive element DNA methylation levels in white blood cell DNA from sisters discordant for breast cancer from the New York site of the Breast Cancer Family Registry. Carcinogenesis, 2012, 33, 1946-1952.            | 2.8  | 66        |
| 12 | Smoking and hepatocellular carcinoma mortality. Experimental and Therapeutic Medicine, 2012, 3, 124-128.   | 1.8  | 11        |
| 13 | DNA methylation in white blood cells. Epigenetics, 2011, 6, 828-837.   | 2.7  | 304       |
| 14 | Global DNA methylation levels in girls with and without a family history of breast cancer. Epigenetics, 2011, 6, 29-33.  | 2.7  | 31        |
| 15 | Global methylation profiles in DNA from different blood cell types. Epigenetics, 2011, 6, 76-85.   | 2.7  | 128       |
| 16 | Mus81-Eme1-Dependent and -Independent Crossovers Form in Mitotic Cells during Double-Strand Break Repair in Schizosaccharomyces pombe. Molecular and Cellular Biology, 2007, 27, 3828-3838.                              | 2.3  | 24        |
| 17 | RMI1/NCE4, a suppressor of genome instability, encodes a member of the RecQ helicase/Topo III complex. EMBO Journal, 2005, 24, 2024-2033.  | 7.8  | 150       |
| 18 | The severe slow growth of Deltasrs2Deltarqh1 in Schizosaccharomyces pombe is suppressed by loss of recombination and checkpoint genes. Nucleic Acids Research, 2002, 30, 4781-4792.                                      | 14.5 | 40        |