Lissette Delgado-Cruzata

List of Publications by Year
in descending order

18 papers	1,101 citations	567281 15 h-index	839539 18 g-index
18 all docs	18 docs citations	18 times ranked	2092 citing authors

#	Article	IF	Citations
1	DNA methylation in white blood cells. Epigenetics, 2011, 6, 828-837.	2.7	304
2	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
3	Global methylation profiles in DNA from different blood cell types. Epigenetics, 2011, 6, 76-85.	2.7	128
4	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
5	Adult global DNA methylation in relation to pre-natal nutrition. International Journal of Epidemiology, 2012, 41, 116-123.	1.9	64
6	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
7	Genome-Wide Methylation Analyses in Glioblastoma Multiforme. PLoS ONE, 2014, 9, e89376.	2.5	45
8	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
9	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
10	DNA Methylation Changes Correlate with Gleason Score and Tumor Stage in Prostate Cancer. DNA and Cell Biology, 2012, 31, 187-192.	1.9	39
11	Global DNA methylation levels in girls with and without a family history of breast cancer. Epigenetics, 2011, 6, 29-33.	2.7	31
12	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
13	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
14	Mismatch Repair Polymorphisms as Markers of Breast Cancer Prevalence in the Breast Cancer Family Registry. Anticancer Research, 2016, 36, 4437-4442.	1.1	24
15	Differences in DNA methylation by extent of breast cancer family history in unaffected women. Epigenetics, 2014, 9, 243-248.	2.7	23
16	Correlations in global DNA methylation measures in peripheral blood mononuclear cells and granulocytes. Epigenetics, 2014, 9, 1504-1510.	2.7	15
17	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
18	Smoking and hepatocellular carcinoma mortality. Experimental and Therapeutic Medicine, 2012, 3, 124-128.	1.8	11