Sabine A Langie

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A pooled analysis of molecular epidemiological studies on modulation of DNA repair by host factors. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2022, 876-877, 503447.	1.7	2
2	Methylome-wide analysis of IVF neonates that underwent embryo culture in different media revealed no significant differences. Npj Genomic Medicine, 2022, 7, .	3.8	4
3	Diagnostic characterization of respiratory allergies by means of a multiplex immunoassay. Clinical and Experimental Immunology, 2021, 203, 183-193.	2.6	2
4	The hCOMET project: International database comparison of results with the comet assay in human biomonitoring. Baseline frequency of DNA damage and effect of main confounders. Mutation Research - Reviews in Mutation Research, 2021, 787, 108371.	5.5	45
5	Collection and storage of human white blood cells for analysis of DNA damage and repair activity using the comet assay in molecular epidemiology studies. Mutagenesis, 2021, 36, 193-212.	2.6	20
6	Different epigenetic signatures of newborn telomere length and telomere attrition rate in early life. Aging, 2021, 13, 14630-14650.	3.1	13
7	DNA damage in circulating leukocytes measured with the comet assay may predict the risk of death. Scientific Reports, 2021, 11, 16793.	3.3	36
8	Upregulation of mNEIL3 in Ogg1-null cells is a potential backup mechanism for 8-oxoG repair. Mutagenesis, 2021, 36, 437-444.	2.6	1
9	An optimized comet-based in vitro DNA repair assay to assess base and nucleotide excision repair activity. Nature Protocols, 2020, 15, 3844-3878.	12.0	33
10	Minimum Information for Reporting on the Comet Assay (MIRCA): recommendations for describing comet assay procedures and results. Nature Protocols, 2020, 15, 3817-3826.	12.0	189
11	Regenerative responses following DNA damage: β-catenin mediates head regrowth in the planarian Schmidtea mediterranea. Journal of Cell Science, 2020, 133, .	2.0	3
12	Potassium bromate as positive assay control for the Fpg-modified comet assay. Mutagenesis, 2020, 35, 341-348.	2.6	32
13	The Influence of the Duration of Breastfeeding on the Infant's Metabolic Epigenome. Nutrients, 2019, 11, 1408.	4.1	29
14	The enzyme-modified comet assay: Enzyme incubation step in 2 vs 12-gels/slide systems. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2019, 845, 402981.	1.7	14
15	Impact of Weight Loss Strategies on Obesityâ€Induced DNA Damage. Molecular Nutrition and Food Research, 2019, 63, 1900045.	3.3	17
16	Meta-analysis of epigenome-wide association studies in neonates reveals widespread differential DNA methylation associated with birthweight. Nature Communications, 2019, 10, 1893.	12.8	140
17	DNA repair as a human biomonitoring tool: Comet assay approaches. Mutation Research - Reviews in Mutation Research, 2019, 781, 71-87.	5.5	40
18	<i>In vivo</i> Toxicity Assessment of Silver Nanoparticles in Homeostatic versus Regenerating Planarians. Nanotoxicology, 2019, 13, 476-491.	3.0	21

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19	DNA methylation and the hygiene hypothesis: connecting respiratory allergy and childhood acute lymphoblastic leukemia. Epigenomics, 2019, 11, 1519-1537.	2.1	4
20	The comet assay in human biomonitoring: cryopreservation of whole blood and comparison with isolated mononuclear cells. Mutagenesis, 2018, 33, 41-47.	2.6	25
21	GLI2 promoter hypermethylation in saliva of children with a respiratory allergy. Clinical Epigenetics, 2018, 10, 50.	4.1	19
22	Cohort Profile: Pregnancy And Childhood Epigenetics (PACE) Consortium. International Journal of Epidemiology, 2018, 47, 22-23u.	1.9	105
23	Impact of obesity and overweight on DNA stability: Few facts and many hypotheses. Mutation Research - Reviews in Mutation Research, 2018, 777, 64-91.	5.5	61
24	Maternal intake of methyl-group donors affects DNA methylation of metabolic genes in infants. Clinical Epigenetics, 2017, 9, 16.	4.1	129
25	The effect of paternal methyl-group donor intake on offspring DNA methylation and birth weight. Journal of Developmental Origins of Health and Disease, 2017, 8, 311-321.	1.4	21
26	Salivary <scp>DNA</scp> Methylation Profiling: Aspects to Consider for Biomarker Identification. Basic and Clinical Pharmacology and Toxicology, 2017, 121, 93-101.	2.5	56
27	RELIC: a novel dye-bias correction method for Illumina Methylation BeadChip. BMC Genomics, 2017, 18, 4.	2.8	96
28	Dietary and supplemental maternal methyl-group donor intake and cord blood DNA methylation. Epigenetics, 2017, 12, 1-10.	2.7	112
29	The Ageing Brain: Effects on DNA Repair and DNA Methylation in Mice. Genes, 2017, 8, 75.	2.4	28
30	Whole-Genome Saliva and Blood DNA Methylation Profiling in Individuals with a Respiratory Allergy. PLoS ONE, 2016, 11, e0151109.	2.5	44
31	A child's spit epigenome can reveal its respiratory allergy risk. , 2016, , .		0
32	The comet assay: past, present, and future. Frontiers in Genetics, 2015, 6, 266.	2.3	103
33	Assessing the carcinogenic potential of low-dose exposures to chemical mixtures in the environment: the challenge ahead. Carcinogenesis, 2015, 36, S254-S296.	2.8	239
34	Causes of genome instability: the effect of low dose chemical exposures in modern society. Carcinogenesis, 2015, 36, S61-S88.	2.8	149
35	Functional evaluation of DNA repair in human biopsies and their relation to other cellular biomarkers. Frontiers in Genetics, 2014, 5, 116.	2.3	13
36	Comet assay to measure DNA repair: approach and applications. Frontiers in Genetics, 2014, 5, 288.	2.3	130

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37	Metabolic effects of a highâ€fat diet postâ€weaning after low maternal dietary folate during pregnancy and lactation. Molecular Nutrition and Food Research, 2014, 58, 1087-1097.	3.3	24
38	Redox and epigenetic regulation of the APE1 gene in the hippocampus of piglets: The effect of early life exposures. DNA Repair, 2014, 18, 52-62.	2.8	15
39	A Standardized Protocol for the In Vitro Comet-Based DNA Repair Assay. Methods in Pharmacology and Toxicology, 2014, , 377-395.	0.2	3
40	DNA-repair measurements by use of the modified comet assay: An inter-laboratory comparison within the European Comet Assay Validation Group (ECVAG). Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2013, 757, 60-67.	1.7	37
41	Maternal folate depletion and highâ€fat feeding from weaning affects DNA methylation and DNA repair in brain of adult offspring. FASEB Journal, 2013, 27, 3323-3334.	0.5	73
42	Measurement of DNA base and nucleotide excision repair activities in mammalian cells and tissues using the comet assay – A methodological overview. DNA Repair, 2013, 12, 1007-1010.	2.8	40
43	Tissue differences in BER-related incision activity and non-specific nuclease activity as measured by the comet assay. Mutagenesis, 2013, 28, 673-681.	2.6	10
44	Epigenetic regulation of DNA base excision repair during ageing and dietary restriction. Proceedings of the Nutrition Society, 2013, 72, .	1.0	0
45	Comparison of Methods for Quantification of Global DNA Methylation in Human Cells and Tissues. PLoS ONE, 2013, 8, e79044.	2.5	143
46	The effect of ageing and short-term dietary restriction on the epigenetic, transcriptomic and phenotypic profile of base excision repair in mouse brain and liver. Proceedings of the Nutrition Society, 2012, 71, .	1.0	1
47	Early determinants of the ageing trajectory. Best Practice and Research in Clinical Endocrinology and Metabolism, 2012, 26, 613-626.	4.7	39
48	Effects of micronutrients on DNA repair. European Journal of Nutrition, 2012, 51, 261-279.	3.9	63
49	Measuring DNA repair incision activity of mouse tissue extracts towards singlet oxygen-induced DNA damage: a comet-based in vitro repair assay. Mutagenesis, 2011, 26, 461-471.	2.6	39
50	Modulation of nucleotide excision repair in human lymphocytes by genetic and dietary factors. British Journal of Nutrition, 2010, 103, 490-501.	2.3	34
51	The effect of early-life nutrition on DNA repair and DNA methylation in the brain of newborn piglets. Proceedings of the Nutrition Society, 2010, 69, .	1.0	Ο
52	An ECVAG trial on assessment of oxidative damage to DNA measured by the comet assay. Mutagenesis, 2010, 25, 125-132.	2.6	99
53	Variation in the measurement of DNA damage by comet assay measured by the ECVAGÂ inter-laboratory validation trial. Mutagenesis, 2010, 25, 113-123.	2.6	155
54	The effect of oxidative stress on nucleotide-excision repair in colon tissue of newborn piglets. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2010, 695, 75-80.	1.7	39

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55	Adult-onset, short-term dietary restriction reduces cell senescence in mice. Aging, 2010, 2, 555-566.	3.1	116
56	Formation of lysine 63-linked poly-ubiquitin chains protects human lung cells against benzo[a]pyrene-diol-epoxide-induced mutagenicity. DNA Repair, 2007, 6, 852-862.	2.8	11
57	The role of glutathione in the regulation of nucleotide excision repair during oxidative stress. Toxicology Letters, 2007, 168, 302-309.	0.8	63
58	Redox-dependent regulation of nucleotide excision repair. Toxicology Letters, 2006, 164, S264-S265.	0.8	0
59	Development and validation of a modified comet assay to phenotypically assess nucleotide excision repair. Mutagenesis, 2006, 21, 153-158.	2.6	81