

# Guro E Lind

## List of Publications by Year in descending order

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Version: 2024-02-01

38  
papers

2,729  
citations

279487

23  
h-index

301761

39  
g-index

39  
all docs

39  
docs citations

39  
times ranked

5241  
citing authors

#	ARTICLE	IF	CITATIONS
1	Cholangiocarcinoma: current knowledge and future perspectives consensus statement from the European Network for the Study of Cholangiocarcinoma (ENS-CCA). <i>Nature Reviews Gastroenterology and Hepatology</i> , 2016, 13, 261-280.	8.2	964
2	Multi-omics of 34 colorectal cancer cell lines - a resource for biomedical studies. <i>Molecular Cancer</i> , 2017, 16, 116.	7.9	232
3	Circulating biomarkers for early detection and clinical management of colorectal cancer. <i>Molecular Aspects of Medicine</i> , 2019, 69, 107-122.	2.7	214
4	Identification of an epigenetic biomarker panel with high sensitivity and specificity for colorectal cancer and adenomas. <i>Molecular Cancer</i> , 2011, 10, 85.	7.9	126
5	Gene methylation profiles of normal mucosa, and benign and malignant colorectal tumors identify early onset markers. <i>Molecular Cancer</i> , 2008, 7, 94.	7.9	102
6	Four DNA methylation biomarkers in biliary brush samples accurately identify the presence of cholangiocarcinoma. <i>Hepatology</i> , 2015, 61, 1651-1659.	3.6	94
7	MicroRNAs as growth regulators, their function and biomarker status in colorectal cancer. <i>Oncotarget</i> , 2016, 7, 6476-6505.	0.8	93
8	The novel colorectal cancer biomarkers <i>CDO1</i> , <i>ZSCAN18</i> and <i>ZNF331</i> are frequently methylated across gastrointestinal cancers. <i>International Journal of Cancer</i> , 2015, 136, 844-853.	2.3	76
9	Multilevel genomics of colorectal cancers with microsatellite instability clinical impact of JAK1 mutations and consensus molecular subtype 1. <i>Genome Medicine</i> , 2017, 9, 46.	3.6	71
10	Novel target genes and a valid biomarker panel identified for cholangiocarcinoma. <i>Epigenetics</i> , 2012, 7, 1249-1257.	1.3	68
11	The epigenome of testicular germ cell tumors. <i>Apmis</i> , 2007, 115, 1147-1160.	0.9	61
12	Epigenetic biomarkers in gastrointestinal cancers: The current state and clinical perspectives. <i>Seminars in Cancer Biology</i> , 2018, 51, 36-49.	4.3	59
13	The recently suggested intestinal cancer stem cell marker <i>DCLK1</i> is an epigenetic biomarker for colorectal cancer. <i>Epigenetics</i> , 2014, 9, 346-350.	1.3	55
14	Hypermethylated MAL gene – a silent marker of early colon tumorigenesis. <i>Journal of Translational Medicine</i> , 2008, 6, 13.	1.8	48
15	DNA-Methylation-Based Detection of Urological Cancer in Urine: Overview of Biomarkers and Considerations on Biomarker Design, Source of DNA, and Detection Technologies. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2657.	1.8	48
16	CpG island methylator phenotype identifies high risk patients among microsatellite stable <i>BRAF</i> mutated colorectal cancers. <i>International Journal of Cancer</i> , 2017, 141, 967-976.	2.3	40
17	DNA methylation analyses of the connexin gene family reveal silencing of <i>GJC1</i> (Connexin45) by promoter hypermethylation in colorectal cancer. <i>Epigenetics</i> , 2011, 6, 602-609.	1.3	39
18	Regulator of Chromosome Condensation 2 Identifies High-Risk Patients within Both Major Phenotypes of Colorectal Cancer. <i>Clinical Cancer Research</i> , 2015, 21, 3759-3770.	3.2	32

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19	Protein expression of BIRC5, TK1, and TOP2A in malignant peripheral nerve sheath tumours – A prognostic test after surgical resection. <i>Molecular Oncology</i> , 2015, 9, 1129-1139.	2.1	32
20	A Tissue-Based Comparative Effectiveness Analysis of Biomarkers for Early Detection of Colorectal Tumors. <i>Clinical and Translational Gastroenterology</i> , 2012, 3, e27.	1.3	30
21	Colorectal cancer DNA methylation marker panel validated with high performance in Non-Hodgkin lymphoma. <i>Epigenetics</i> , 2014, 9, 428-436.	1.3	29
22	Contribution of <i>MLH1</i> constitutional methylation for Lynch syndrome diagnosis in patients with tumor <i>MLH1</i> downregulation. <i>Cancer Medicine</i> , 2018, 7, 433-444.	1.3	28
23	A robust internal control for high-precision DNA methylation analyses by droplet digital PCR. <i>Clinical Epigenetics</i> , 2018, 10, 24.	1.8	26
24	DNA Hypermethylation of MAL: A Promising Diagnostic Biomarker for Colorectal Tumors. <i>Gastroenterology</i> , 2007, 132, 1631-1632.	0.6	22
25	Methylated RASSF1A in malignant peripheral nerve sheath tumors identifies neurofibromatosis type 1 patients with inferior prognosis. <i>Neuro-Oncology</i> , 2015, 17, 63-69.	0.6	17
26	Identification of Highly Methylated Genes across Various Types of B-Cell Non-Hodgkin Lymphoma. <i>PLoS ONE</i> , 2013, 8, e79602.	1.1	16
27	Experimental factors affecting the robustness of DNA methylation analysis. <i>Scientific Reports</i> , 2016, 6, 33936.	1.6	15
28	Improved prognostication of glioblastoma beyond molecular subtyping by transcriptional profiling of the tumor microenvironment. <i>Molecular Oncology</i> , 2020, 14, 1016-1027.	2.1	15
29	Early and accurate detection of cholangiocarcinoma in patients with primary sclerosing cholangitis by methylation markers in bile. <i>Hepatology</i> , 2022, 75, 59-73.	3.6	15
30	Re-assessing ZNF331 as a DNA methylation biomarker for colorectal cancer. <i>Clinical Epigenetics</i> , 2018, 10, 70.	1.8	14
31	A Gene Panel, Including LRP12, Is Frequently Hypermethylated in Major Types of B-Cell Lymphoma. <i>PLoS ONE</i> , 2014, 9, e104249.	1.1	13
32	miR-486-5p expression is regulated by DNA methylation in osteosarcoma. <i>BMC Genomics</i> , 2022, 23, 142.	1.2	8
33	Detecting cholangiocarcinoma in patients with primary sclerosing cholangitis – The promise of DNA methylation and molecular biomarkers. <i>JHEP Reports</i> , 2020, 2, 100143.	2.6	6
34	Sequencing IDH1/2 glioma mutation hotspots in gliomas and malignant peripheral nerve sheath tumors. <i>Neuro-Oncology</i> , 2014, 16, 320-322.	0.6	5
35	Details matter: the role of genomic location and assay standardization in DNA methylation analyses. <i>Epigenomics</i> , 2017, 9, 933-935.	1.0	5
36	Prognostic relevance of an epigenetic biomarker panel in sentinel lymph nodes from colon cancer patients. <i>Clinical Epigenetics</i> , 2017, 9, 97.	1.8	4

#	ARTICLE	IF	CITATIONS
37	Multiregional assessment of CIMP in primary colorectal cancers: Phenotype concordance but marker variability. <i>International Journal of Cancer</i> , 2021, 148, 1652-1657.	2.3	4
38	Targeted genetic and epigenetic profiling of esophageal adenocarcinomas and non-dysplastic Barrett's esophagus. <i>Clinical Epigenetics</i> , 2022, 14, .	1.8	2