

# Iver Nordentoft

## List of Publications by Year in descending order

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Version: 2024-02-01

34  
papers

4,450  
citations

218677

26  
h-index

434195

31  
g-index

36  
all docs

36  
docs citations

36  
times ranked

7364  
citing authors

| #  | ARTICLE  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Circulating tumor <scp>DNA</scp> for prognosis assessment and postoperative management after curative-intent resection of colorectal liver metastases. International Journal of Cancer, 2022, 150, 1537-1548.                                      | 5.1  | 22        |
| 2  | An integrated multi-omics analysis identifies prognostic molecular subtypes of non-muscle-invasive bladder cancer. Nature Communications, 2021, 12, 2301.  | 12.8 | 159       |
| 3  | Genome-wide circulating tumor DNA monitoring for bladder cancer treatment management and organ preservation.. Journal of Clinical Oncology, 2021, 39, e16527-e16527.   | 1.6  | 0         |
| 4  | SPTAN1, APC, and FGFR3 Mutation Status and APOBEC Mutation Signatures are Predictive of Mitomycin C Response in Non-muscle-invasive Bladder Cancer. European Urology Open Science, 2021, 34, 59-67.  | 0.4  | 1         |
| 5  | Pervasive chromosomal instability and karyotype order in tumour evolution. Nature, 2020, 587, 126-132.   | 27.8 | 221       |
| 6  | Molecular correlates of cisplatin-based chemotherapy response in muscle invasive bladder cancer by integrated multi-omics analysis. Nature Communications, 2020, 11, 4858.   | 12.8 | 124       |
| 7  | Epigenetic and transcriptomic consequences of excess X-chromosome material in 47,XXX syndrome—A comparison with Turner syndrome and 46,XX females. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2020, 184, 279-293. | 1.6  | 21        |
| 8  | Representative Sequencing: Unbiased Sampling of Solid Tumor Tissue. Cell Reports, 2020, 31, 107550.  | 6.4  | 51        |
| 9  | Mutational Analysis of Field Cancerization in Bladder Cancer. Bladder Cancer, 2020, 6, 253-264.  | 0.4  | 12        |
| 10 | Analysis of Plasma Cell-Free DNA by Ultradeep Sequencing in Patients With Stages I to III Colorectal Cancer. JAMA Oncology, 2019, 5, 1124.   | 7.1  | 538       |
| 11 | Early Detection of Metastatic Relapse and Monitoring of Therapeutic Efficacy by Ultra-Deep Sequencing of Plasma Cell-Free DNA in Patients With Urothelial Bladder Carcinoma. Journal of Clinical Oncology, 2019, 37, 1547-1557.                    | 1.6  | 298       |
| 12 | Abstract 913: Early detection of metastatic relapse and monitoring of therapeutic efficacy by ultra-deep sequencing of plasma cell-free DNA in patients with urothelial bladder carcinoma. , 2019, , .   |      | 3         |
| 13 | Optimized targeted sequencing of cell-free plasma DNA from bladder cancer patients. Scientific Reports, 2018, 8, 1917.   | 3.3  | 50        |
| 14 | Monitoring Treatment Response and Metastatic Relapse in Advanced Bladder Cancer by Liquid Biopsy Analysis. European Urology, 2018, 73, 535-540.  | 1.9  | 112       |
| 15 | Characterization of genetic intratumor heterogeneity in colorectal cancer and matching patient-derived spheroid cultures. Molecular Oncology, 2018, 12, 132-147.   | 4.6  | 49        |
| 16 | Discordant molecular subtype classification in the basal-squamous subtype of bladder tumors and matched lymph-node metastases. Modern Pathology, 2018, 31, 1869-1881.  | 5.5  | 47        |
| 17 | Liquid Biopsy Analysis of FGFR3 and PIK3CA Hotspot Mutations for Disease Surveillance in Bladder Cancer. European Urology, 2017, 71, 961-969.  | 1.9  | 154       |
| 18 | Altered gene expression and repressed markers of autophagy in skeletal muscle of insulin resistant patients with type 2 diabetes. Scientific Reports, 2017, 7, 43775.  | 3.3  | 57        |

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|----|--|------|-----------|
| 19 | Clinical Implications of Monitoring Circulating Tumor DNA in Patients with Colorectal Cancer. <i>Clinical Cancer Research</i> , 2017, 23, 5437-5445.   | 7.0  | 232       |
| 20 | Profiling of long non-coding RNAs identifies LINC00958 and LINC01296 as candidate oncogenes in bladder cancer. <i>Scientific Reports</i> , 2017, 7, 395.   | 3.3  | 117       |
| 21 | Comprehensive multiregional analysis of molecular heterogeneity in bladder cancer. <i>Scientific Reports</i> , 2017, 7, 11702.   | 3.3  | 110       |
| 22 | RHCG and TCAF1 promoter hypermethylation predicts biochemical recurrence in prostate cancer patients treated by radical prostatectomy. <i>Oncotarget</i> , 2017, 8, 5774-5788.   | 1.8  | 22        |
| 23 | Paired Exome Analysis Reveals Clonal Evolution and Potential Therapeutic Targets in Urothelial Carcinoma. <i>Cancer Research</i> , 2016, 76, 5894-5906.  | 0.9  | 87        |
| 24 | Spatial and temporal clonal evolution during development of metastatic urothelial carcinoma. <i>Molecular Oncology</i> , 2016, 10, 1450-1460.  | 4.6  | 44        |
| 25 | Widespread DNA hypomethylation and differential gene expression in Turner syndrome. <i>Scientific Reports</i> , 2016, 6, 34220.  | 3.3  | 106       |
| 26 | Comprehensive Transcriptional Analysis of Early-Stage Urothelial Carcinoma. <i>Cancer Cell</i> , 2016, 30, 27-42.  | 16.8 | 486       |
| 27 | Analysis of circulating tumour DNA to monitor disease burden following colorectal cancer surgery. <i>Gut</i> , 2016, 65, 625-634.  | 12.1 | 381       |
| 28 | Genomic Alterations in Liquid Biopsies from Patients with Bladder Cancer. <i>European Urology</i> , 2016, 70, 75-82.   | 1.9  | 174       |
| 29 | Next-Generation Sequencing of RNA and DNA Isolated from Paired Fresh-Frozen and Formalin-Fixed Paraffin-Embedded Samples of Human Cancer and Normal Tissue. <i>PLoS ONE</i> , 2014, 9, e98187.   | 2.5  | 284       |
| 30 | Cellular Disposal of miR23b by RAB27-Dependent Exosome Release Is Linked to Acquisition of Metastatic Properties. <i>Cancer Research</i> , 2014, 74, 5758-5771.  | 0.9  | 237       |
| 31 | Mutational Context and Diverse Clonal Development in Early and Late Bladder Cancer. <i>Cell Reports</i> , 2014, 7, 1649-1663.  | 6.4  | 128       |
| 32 | miRNAs associated with chemo-sensitivity in cell lines and in advanced bladder cancer. <i>BMC Medical Genomics</i> , 2012, 5, 40.  | 1.5  | 86        |
| 33 | Increased expression of transcription factor TFAP2 $\pm$ correlates with chemosensitivity in advanced bladder cancer. <i>BMC Cancer</i> , 2011, 11, 135.   | 2.6  | 35        |
| 34 | Expression Analysis of cPLA2 Alpha Interacting TIP60 in Diabetic KK $\Delta$ y and Non-Diabetic C57BL Wild-Type Mice: No Impact of Transient and Stable TIP60 Overexpression on Glucose-Stimulated Insulin Secretion in Pancreatic Beta-Cells. <i>Review of Diabetic Studies</i> , 2007, 4, 147-158. | 1.3  | 0         |