List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Artificial intelligence and pathology: From principles to practice and future applications in histomorphology and molecular profiling. Seminars in Cancer Biology, 2022, 84, 129-143. | 9.6 | 41 |
| 2 | Trailblazing precision medicine in Europe: A joint view by Genomic Medicine Sweden and the Centers for Personalized Medicine, ZPM, in Germany. Seminars in Cancer Biology, 2022, 84, 242-254. | 9.6 | 22 |
| 3 | Deciphering the immunosuppressive tumor microenvironment in ALK- and EGFR-positive lung adenocarcinoma. Cancer Immunology, Immunotherapy, 2022, 71, 251-265. | 4.2 | 22 |
| 4 | Mutations in TP53 or DNA damage repair genes define poor prognostic subgroups in primary prostate cancer. Urologic Oncology: Seminars and Original Investigations, 2022, 40, 8.e11-8.e18. | 1.6 | 8 |
| 5 | Assigning evidence to actionability: An introduction to variant interpretation in precision cancer medicine. Genes Chromosomes and Cancer, 2022, 61, 303-313. | 2.8 | 15 |
| 6 | Cutaneous epithelioid haemangiomas show somatic mutations in the mitogenâ€activated protein kinase pathway. British Journal of Dermatology, 2022, 186, 553-563. | 1.5 | 3 |
| 7 | The Different Immune Profiles of Normal Colonic Mucosa in Cancer-Free Lynch Syndrome Carriers and Lynch Syndrome Colorectal Cancer Patients. Gastroenterology, 2022, 162, 907-919.e10. | 1.3 | 27 |
| 8 | Fusionâ€positive nonâ€small cell lung carcinoma: Biological principles, clinical practice, and diagnostic implications. Genes Chromosomes and Cancer, 2022, 61, 244-260. | 2.8 | 32 |
| 9 | TP53 co-mutations as an independent prognostic factor in 2nd and further line therapy—EGFR mutated non-small cell lung cancer IV patients treated with osimertinib. Translational Lung Cancer Research, 2022, 11, 4-13. | 2.8 | 13 |
| 10 | Homologous Recombination Deficiency: Concepts, Definitions, and Assays. Oncologist, 2022, 27, 167-174. | 3.7 | 69 |
| 11 | Standards for the classification of pathogenicity of somatic variants in cancer (oncogenicity): Joint recommendations of Clinical Genome Resource (ClinGen), Cancer Genomics Consortium (CGC), and Variant Interpretation for Cancer Consortium (VICC). Genetics in Medicine, 2022, 24, 986-998. | 2.4 | 55 |
| 12 | Histological and Molecular Plasticity of ALK-positive Non-Small-Cell Lung Cancer under Targeted Therapy - a Case Report. Journal of Physical Education and Sports Management, 2022, , mcs.a006156. | 1.2 | 5 |
| 13 | Impact of Surgeon's Experience in Rigid Versus Elastic MRI/TRUS-Fusion Biopsy to Detect Significant Prostate Cancer Using Targeted and Systematic Cores. Cancers, 2022, 14, 886. | 3.7 | 3 |
| 14 | <scp>Homologous recombination deficiency</scp> is inversely correlated with <scp>microsatellite instability</scp> and identifies immunologically cold tumors in most cancer types. Journal of Pathology: Clinical Research, 2022, 8, 371-382. | 3.0 | 10 |
| 15 | p53 partial loss-of-function mutations sensitize to chemotherapy. Oncogene, 2022, 41, 1011-1023. | 5.9 | 28 |
| 16 | Prognostic impact of copy number alterations and tumor mutational burden in carcinoma of unknown primary. Genes Chromosomes and Cancer, 2022, 61, 551-560. | 2.8 | 4 |
| 17 | The impact of TP53 co-mutations and immunologic microenvironment on outcome of lung cancer with EGFR exon 20 insertions. European Journal of Cancer, 2022, 170, 106-118. | 2.8 | 15 |
| 18 | Early Development of Ubiquitous Acanthocytosis and Extravascular Hemolysis in Lung Cancer Patients Receiving Alectinib. Cancers, 2022, 14, 2720. | 3.7 | 5 |

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|----|--|------|-----------|
| 19 | Brief Report: A Blood-Based MicroRNA Complementary Diagnostic Predicts Immunotherapy Efficacy in Advanced-Stage NSCLC WithÂHighÂProgrammed Death-Ligand 1ÂExpression. JTO Clinical and Research Reports, 2022, 3, 100369. | 1.1 | 3 |
| 20 | Pan-cancer analysis of genomic scar patterns caused by homologous repair deficiency (HRD). Npj Precision Oncology, 2022, 6, . | 5.4 | 23 |
| 21 | Standardized Magnetic Resonance Imaging Reporting Using the Prostate Cancer Radiological Estimation of Change in Sequential Evaluation Criteria and Magnetic Resonance Imaging/Transrectal Ultrasound Fusion with Transperineal Saturation Biopsy to Select Men on Active Surveillance. European Urology Focus. 2021, 7, 102-110. | 3.1 | 28 |
| 22 | Biomarker testing in non-small cell lung cancer in routine care: Analysis of the first 3,717 patients in the German prospective, observational, nation-wide CRISP Registry (AIO-TRK-0315). Lung Cancer, 2021, 152, 174-184. | 2.0 | 53 |
| 23 | Simulated clinical deployment of fully automatic deep learning for clinical prostate MRI assessment. European Radiology, 2021, 31, 302-313. | 4.5 | 24 |
| 24 | <scp>PARP</scp> inhibition in prostate cancer. Genes Chromosomes and Cancer, 2021, 60, 344-351. | 2.8 | 2 |
| 25 | Comparison of Prostate MRI Lesion Segmentation Agreement Between Multiple Radiologists and a Fully Automatic Deep Learning System. RoFo Fortschritte Auf Dem Gebiet Der Rontgenstrahlen Und Der Bildgebenden Verfahren, 2021, 193, 559-573. | 1.3 | 18 |
| 26 | Novel GATA6-FOXO1 fusions in a subset of epithelioid hemangioma. Modern Pathology, 2021, 34, 934-941. | 5.5 | 27 |
| 27 | Ruxolitinib is effective in the treatment of a patient with refractory Tâ€ALL. EJHaem, 2021, 2, 139-142. | 1.0 | 4 |
| 28 | Integrating proteomics into precision oncology. International Journal of Cancer, 2021, 148, 1438-1451. | 5.1 | 15 |
| 29 | The Value of Prostate-specific Antigen Density for Prostate Imaging-Reporting and Data System 3 Lesions on Multiparametric Magnetic Resonance Imaging: A Strategy to Avoid Unnecessary Prostate Biopsies. European Urology Focus, 2021, 7, 325-331. | 3.1 | 34 |
| 30 | Real-world implementation of sequential targeted therapies for EGFR-mutated lung cancer. Therapeutic Advances in Medical Oncology, 2021, 13, 175883592199650. | 3.2 | 24 |
| 31 | Accurate and efficient detection of gene fusions from RNA sequencing data. Genome Research, 2021, 31, 448-460. | 5.5 | 215 |
| 32 | A gene expression signature associated with B cells predicts benefit from immune checkpoint blockade in lung adenocarcinoma. Oncolmmunology, 2021, 10, 1860586. | 4.6 | 40 |
| 33 | Strength in numbers: predicting response to checkpoint inhibitors from large clinical datasets. Cell, 2021, 184, 571-573. | 28.9 | 3 |
| 34 | Homologous recombination repair deficiency (HRD): From biology to clinical exploitation. Genes Chromosomes and Cancer, 2021, 60, 299-302. | 2.8 | 16 |
| 35 | Clinical and molecular practice of European thoracic pathology laboratories during the COVID-19 pandemic. The past and the near future. ESMO Open, 2021, 6, 100024. | 4.5 | 13 |
| 36 | Practical considerations for optimising homologous recombination repair mutation testing in patients with metastatic prostate cancer. Journal of Pathology: Clinical Research, 2021, 7, 311-325. | 3.0 | 19 |

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|----|--|------|-----------|
| 37 | Hidden Variables in Deep Learning Digital Pathology and Their Potential to Cause Batch Effects: Prediction Model Study. Journal of Medical Internet Research, 2021, 23, e23436. | 4.3 | 36 |
| 38 | SARS-CoV-2 infects and replicates in cells of the human endocrine and exocrine pancreas. Nature Metabolism, 2021, 3, 149-165. | 11.9 | 378 |
| 39 | Morphological and molecular breast cancer profiling through explainable machine learning. Nature Machine Intelligence, 2021, 3, 355-366. | 16.0 | 72 |
| 40 | KRAS / GNAS â€ŧesting by highly sensitive deep targeted next generation sequencing improves the endoscopic ultrasoundâ€guided workup of suspected mucinous neoplasms of the pancreas. Genes Chromosomes and Cancer, 2021, 60, 489-497. | 2.8 | 13 |
| 41 | Case Report: Abdominal Lymph Node Metastases of Parathyroid Carcinoma: Diagnostic Workup, Molecular Diagnosis, and Clinical Management. Frontiers in Endocrinology, 2021, 12, 643328. | 3.5 | 12 |
| 42 | Primary neoplasms of the parapharyngeal space: diagnostic and therapeutic pearls and pitfalls. European Archives of Oto-Rhino-Laryngology, 2021, 278, 4933-4941. | 1.6 | 10 |
| 43 | Comparison of single-scanner single-protocol quantitative ADC measurements to ADC ratios to detect clinically significant prostate cancer. European Journal of Radiology, 2021, 136, 109538. | 2.6 | 7 |
| 44 | Combination of Crizotinib and Osimertinib in T790M+ EGFR-Mutant Non-Small Cell Lung Cancer with Emerging MET Amplification Post-Osimertinib Progression in a 10-Year Survivor: A Case Report. Case Reports in Oncology, 2021, 14, 477-482. | 0.7 | 8 |
| 45 | De Novo Versus Secondary Metastatic EGFR-Mutated Non-Small-Cell Lung Cancer. Frontiers in Oncology, 2021, 11, 640048. | 2.8 | 4 |
| 46 | CATCH: A Prospective Precision Oncology Trial in Metastatic Breast Cancer. JCO Precision Oncology, 2021, 5, 676-686. | 3.0 | 20 |
| 47 | KRAS G12C-mutated advanced non-small cell lung cancer: A real-world cohort from the German prospective, observational, nation-wide CRISP Registry (AIO-TRK-0315). Lung Cancer, 2021, 154, 51-61. | 2.0 | 43 |
| 48 | <scp><i>RREB1â€MKL2</i></scp> fusion in a spindle cell sinonasal sarcoma: biphenotypic sinonasal sarcoma or ectomesenchymal chondromyxoid tumor in an unusual site?. Genes Chromosomes and Cancer, 2021, 60, 565-570. | 2.8 | 10 |
| 49 | Targeting rare and non-canonical driver variants in NSCLC – An uncharted clinical field. Lung Cancer, 2021, 154, 131-141. | 2.0 | 8 |
| 50 | Recurrent YAP1-TFE3 Gene Fusions in Clear Cell Stromal Tumor of the Lung. American Journal of Surgical Pathology, 2021, 45, 1541-1549. | 3.7 | 16 |
| 51 | Conventional and semi-automatic histopathological analysis of tumor cell content for multigene sequencing of lung adenocarcinoma. Translational Lung Cancer Research, 2021, 10, 1666-1678. | 2.8 | 6 |
| 52 | Feasibility and Challenges for Sequential Treatments in ALK-Rearranged Non-Small-Cell Lung Cancer. Frontiers in Oncology, 2021, 11, 670483. | 2.8 | 10 |
| 53 | Deconvolution of sarcoma methylomes reveals varying degrees of immune cell infiltrates with association to genomic aberrations. Journal of Translational Medicine, 2021, 19, 204. | 4.4 | 5 |
| 54 | Earlier extracranial progression and shorter survival in ALK- rearranged lung cancer with positive liquid rebiopsies. Translational Lung Cancer Research, 2021, 10, 2118-2131. | 2.8 | 16 |

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|----|---|-----|-----------|
| 55 | Knowledge bases and software support for variant interpretation in precision oncology. Briefings in Bioinformatics, 2021, 22, . | 6.5 | 9 |
| 56 | Complete Metabolic Response in FDG-PET-CT Scan before Discontinuation of Immune Checkpoint Inhibitors Correlates with Long Progression-Free Survival. Cancers, 2021, 13, 2616. | 3.7 | 8 |
| 57 | Role of Synaptophysin, Chromogranin and CD56 in adenocarcinoma and squamous cell carcinoma of the lung lacking morphological features of neuroendocrine differentiation: a retrospective large-scale study on 1170 tissue samples. BMC Cancer, 2021, 21, 486. | 2.6 | 21 |
| 58 | Brigatinib versus other second-generation ALK inhibitors as initial treatment of anaplastic lymphoma kinase positive non-small cell lung cancer with deep phenotyping: study protocol of the ABP trial. BMC Cancer, 2021, 21, 743. | 2.6 | 3 |
| 59 | SWI/SNF-deficient undifferentiated/rhabdoid carcinoma of the gallbladder carrying a POLE mutation in a 30-year-old woman: a case report. Diagnostic Pathology, 2021, 16, 52. | 2.0 | 6 |
| 60 | Fully Automatic Deep Learning in Bi-institutional Prostate Magnetic Resonance Imaging. Investigative Radiology, 2021, 56, 799-808. | 6.2 | 27 |
| 61 | Comprehensive Genomic and Transcriptomic Analysis for Guiding Therapeutic Decisions in Patients with Rare Cancers. Cancer Discovery, 2021, 11, 2780-2795. | 9.4 | 125 |
| 62 | Therapeutic and Prognostic Implications of Immune-Related Adverse Events in Advanced Non-Small-Cell Lung Cancer. Frontiers in Oncology, 2021, 11, 703893. | 2.8 | 33 |
| 63 | Effect of timing, technique and molecular features on brain control with local therapies in oncogene-driven lung cancer. ESMO Open, 2021, 6, 100161. | 4.5 | 9 |
| 64 | Distinct Mutational Profile of Lynch Syndrome Colorectal Cancers Diagnosed under Regular Colonoscopy Surveillance. Journal of Clinical Medicine, 2021, 10, 2458. | 2.4 | 3 |
| 65 | Status quo of ALK testing in lung cancer: results of an EQA scheme based on in-situ hybridization, immunohistochemistry, and RNA/DNA sequencing. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2021, 479, 247-255. | 2.8 | 5 |
| 66 | Comprehensive Dissection of Treatment Patterns and Outcome for Patients With Metastatic Large-Cell Neuroendocrine Lung Carcinoma. Frontiers in Oncology, 2021, 11, 673901. | 2.8 | 8 |
| 67 | Detection of PD-L1 in the urine of patients with urothelial carcinoma of the bladder. Scientific Reports, 2021, 11, 14244. | 3.3 | 9 |
| 68 | Efficacy of docetaxel plus ramucirumab as palliative second-line therapy following first-line chemotherapy plus immune-checkpoint-inhibitor combination treatment in patients with non-small cell lung cancer (NSCLC) UICC stage IV. Translational Lung Cancer Research, 2021, 10, 3093-3105. | 2.8 | 23 |
| 69 | Intimal sarcomas and undifferentiated cardiac sarcomas carry mutually exclusive MDM2, MDM4, and CDK6 amplifications and share a common DNA methylation signature. Modern Pathology, 2021, 34, 2122-2129. | 5.5 | 17 |
| 70 | Efficacy of Immune Checkpoint Inhibitors Alone or in Combination With Chemotherapy in NSCLC Harboring ERBB2 Mutations. Journal of Thoracic Oncology, 2021, 16, 1952-1958. | 1.1 | 32 |
| 71 | Abstract 449: A standard operating procedure for the curation of gene fusions. , 2021, , . | | 0 |
| 72 | Validation of a Targeted Next-Generation Sequencing Panel for Tumor Mutation Burden Analysis. Journal of Molecular Diagnostics, 2021, 23, 882-893. | 2.8 | 2 |

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| # | Article | IF | CITATIONS |
|----|---|-------------------|---------------|
| 73 | Precision oncology: a clinical and patient perspective. Future Oncology, 2021, 17, 3995-4009. | 2.4 | 22 |
| 74 | High tumour mutational burden and EGFR/MAPK pathway activation are therapeutic targets in metastatic porocarcinoma. British Journal of Dermatology, 2021, , . | 1.5 | 6 |
| 75 | Hidden Treasures: Macrophage Long Non-Coding RNAs in Lung Cancer Progression. Cancers, 2021, 13, 4127. | 3.7 | 7 |
| 76 | GOPC:ROS1 and other ROS1 fusions represent a rare but recurrent drug target in a variety of glioma types. Acta Neuropathologica, 2021, 142, 1065-1069. | 7.7 | 16 |
| 77 | Association of the advanced lung cancer inflammation index (ALI) with immune checkpoint inhibitor efficacy in patients with advanced non-small-cell lung cancer. ESMO Open, 2021, 6, 100254. | 4.5 | 35 |
| 78 | The immune microenvironment in EGFR- and ERBB2-mutated lung adenocarcinoma. ESMO Open, 2021, 6, 100253. | 4.5 | 17 |
| 79 | Aligning tumor mutational burden (TMB) quantification across diagnostic platforms: phase II of the Friends of Cancer Research TMB Harmonization Project. Annals of Oncology, 2021, 32, 1626-1636. | 1.2 | 86 |
| 80 | Local ablative treatment with surgery and/or radiotherapy in single-site and oligometastatic carcinoma of unknown primary. European Journal of Cancer, 2021, 157, 179-189. | 2.8 | 13 |
| 81 | Sarcoma classification by DNA methylation profiling. Nature Communications, 2021, 12, 498. | 12.8 | 237 |
| 82 | Laboratory-Developed Tests in the New European Union 2017/746 Regulation: Opportunities and Risks. Clinical Chemistry, 2021, 68, 40-42. | 3.2 | 11 |
| 83 | Prolonged Survival of a Patient with Advanced-Stage Combined Hepatocellular-Cholangiocarcinoma. Case Reports in Gastroenterology, 2021, 14, 658-667. | 0.6 | 6 |
| 84 | Rationale and design of the CRAFT (Continuous ReAssessment with Flexible ExTension in Rare) Tj ETQq0 0 0 rgB | T /Qverloc 4.5 | k 10 Tf 50 30 |
| 85 | Early identification of disease progression in ALK-rearranged lung cancer using circulating tumor DNA analysis. Npj Precision Oncology, 2021, 5, 100. | 5.4 | 21 |
| 86 | Combined Clinical Parameters and Multiparametric Magnetic Resonance Imaging for the Prediction of Extraprostatic Disease—A Risk Model for Patient-tailored Risk Stratification When Planning Radical Prostatectomy. European Urology Focus, 2020, 6, 1205-1212. | 3.1 | 39 |
| 87 | Identification and characterization of a BRAF fusion oncoprotein with retained autoinhibitory domains. Oncogene, 2020, 39, 814-832. | 5.9 | 19 |
| 88 | Patients Resistant Against PSMA-Targeting α-Radiation Therapy Often Harbor Mutations in DNA Damage-Repair–Associated Genes. Journal of Nuclear Medicine, 2020, 61, 683-688. | 5.0 | 61 |
| 89 | Testing <i>NTRK</i> testing: Wetâ€lab and in silico comparison of RNAâ€based targeted sequencing assays. Genes Chromosomes and Cancer, 2020, 59, 178-188. | 2.8 | 52 |
| 90 | Genomic Characterization of Cholangiocarcinoma in Primary Sclerosing Cholangitis Reveals Therapeutic Opportunities. Hepatology, 2020, 72, 1253-1266. | 7.3 | 42 |

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|-----|---|------|-----------|
| 91 | Primary pulmonary myxoid sarcoma with an unusual gene fusion between exon 7 of EWSR1 and exon 5 of CREB1. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2020, 476, 787-791. | 2.8 | 7 |
| 92 | Rearranged ERG confers robustness to prostate cancer cells by subverting the function of p53. Urologic Oncology: Seminars and Original Investigations, 2020, 38, 736.e1-736.e10. | 1.6 | 2 |
| 93 | Comprehensive serial biobanking in advanced NSCLC: feasibility, challenges and perspectives. Translational Lung Cancer Research, 2020, 9, 1000-1014. | 2.8 | 9 |
| 94 | Risk stratification of EGFR+ lung cancer diagnosed with panel-based next-generation sequencing. Lung Cancer, 2020, 148, 105-112. | 2.0 | 26 |
| 95 | Recommendations for the use of next-generation sequencing (NGS) for patients with metastatic cancers: a report from the ESMO Precision Medicine Working Group. Annals of Oncology, 2020, 31, 1491-1505. | 1.2 | 658 |
| 96 | Longitudinal therapy monitoring of ALK-positive lung cancer by combined copy number and targeted mutation profiling of cell-free DNA. EBioMedicine, 2020, 62, 103103. | 6.1 | 32 |
| 97 | EWSR1/FUS–CREB fusions define a distinctive malignant epithelioid neoplasm with predilection for mesothelial-lined cavities. Modern Pathology, 2020, 33, 2233-2243. | 5.5 | 49 |
| 98 | Tumor Mutational Burden as a Pan-cancer Biomarker for Immunotherapy: The Limits and Potential for Convergence. Cancer Cell, 2020, 38, 624-625. | 16.8 | 35 |
| 99 | Successful BRAF/MEK inhibition in a patient with <i>BRAF</i> ^{V600E} -mutated extrapancreatic acinar cell carcinoma. Journal of Physical Education and Sports Management, 2020, 6, a005553. | 1.2 | 13 |
| 100 | Mass Spectrometry Imaging for Reliable and Fast Classification of Non-Small Cell Lung Cancer Subtypes. Cancers, 2020, 12, 2704. | 3.7 | 13 |
| 101 | Mass Spectrometry Imaging Differentiates Chromophobe Renal Cell Carcinoma and Renal Oncocytoma with High Accuracy. Journal of Cancer, 2020, 11, 6081-6089. | 2.5 | 8 |
| 102 | Adaptive Immunity and Pathogenesis of Diabetes: Insights Provided by the α4–Integrin Deficient NOD Mouse. Cells, 2020, 9, 2597. | 4.1 | 4 |
| 103 | Tumor Mutational Burden as a Predictive Biomarker in Solid Tumors. Cancer Discovery, 2020, 10, 1808-1825. | 9.4 | 388 |
| 104 | The landscape of chromothripsis across adult cancer types. Nature Communications, 2020, 11, 2320. | 12.8 | 75 |
| 105 | Distinct immune evasion in <scp>APOBEC</scp> â€enriched, <scp>HPV</scp> â€negative <scp>HNSCC</scp> . International Journal of Cancer, 2020, 147, 2293-2302. | 5.1 | 10 |
| 106 | Conceptual framework for precision cancer medicine in Germany: Consensus statement of the Deutsche Krebshilfe working group †Molecular Diagnostics and Therapy'. European Journal of Cancer, 2020, 135, 1-7. | 2.8 | 23 |
| 107 | Deep Learning for the Classification of Small-Cell and Non-Small-Cell Lung Cancer. Cancers, 2020, 12, 1604. | 3.7 | 63 |
| 108 | Immunoâ€oncology gene expression profiling of formalinâ€fixed and paraffinâ€embedded clear cell renal cell carcinoma: Performance comparison of the <scp>NanoString nCounter</scp> technology with targeted <scp>RNA</scp> sequencing. Genes Chromosomes and Cancer, 2020, 59, 406-416. | 2.8 | 10 |

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|-----|--|-----|-----------|
| 109 | Automated sample preparation with <scp>SP</scp> 3 for lowâ€input clinical proteomics. Molecular Systems Biology, 2020, 16, e9111. | 7.2 | 133 |
| 110 | Harmonization and Standardization of Panel-Based Tumor Mutational Burden Measurement: Real-World Results and Recommendations ofÂtheÂQuality in Pathology Study. Journal of Thoracic Oncology, 2020, 15, 1177-1189. | 1.1 | 81 |
| 111 | Immunohistological expression of oestrogen receptor, progesterone receptor, mammaglobin, human epidermal growth factor receptor 2 and GATAâ€binding protein 3 in nonâ€smallâ€cell lung cancer. Histopathology, 2020, 77, 900-914. | 2.9 | 6 |
| 112 | Metastatic adult pancreatoblastoma: Multimodal treatment and molecular characterization of a very rare disease. Pancreatology, 2020, 20, 425-432. | 1.1 | 11 |
| 113 | Quantifying potential confounders of panel-based tumor mutational burden (TMB) measurement. Lung Cancer, 2020, 142, 114-119. | 2.0 | 28 |
| 114 | Associations of Pathogenic Variants in MLH1, MSH2, and MSH6 With Risk of Colorectal Adenomas and Tumors and With Somatic Mutations in Patients With Lynch Syndrome. Gastroenterology, 2020, 158, 1326-1333. | 1.3 | 60 |
| 115 | Integrated clinicomolecular characterization identifies RAS activation and CDKN2A deletion as independent adverse prognostic factors in cancer of unknown primary. International Journal of Cancer, 2020, 146, 3053-3064. | 5.1 | 14 |
| 116 | Endometrial stromal sarcomas with <i>BCOR</i> â€rearrangement harbor <i>MDM2</i> amplifications. Journal of Pathology: Clinical Research, 2020, 6, 178-184. | 3.0 | 32 |
| 117 | Establishing guidelines to harmonize tumor mutational burden (TMB): in silico assessment of variation in TMB quantification across diagnostic platforms: phase I of the Friends of Cancer Research TMB Harmonization Project. , 2020, 8, e000147. | | 329 |
| 118 | <scp>NTRK</scp> testing: First results of the <scp>QuiPâ€EQA</scp> scheme and a comprehensive map of <scp><i>NTRK</i></scp> fusion variants and their diagnostic coverage by targeted <scp>RNA</scp> â€based <scp>NGS</scp> assays. Genes Chromosomes and Cancer, 2020, 59, 445-453. | 2.8 | 27 |
| 119 | Durvalumab in frail and elderly patients with stage four non-small cell lung cancer: Study protocol of the randomized phase II DURATION trial. Trials, 2020, 21, 352. | 1.6 | 7 |
| 120 | High prevalence of DNA damage repair gene defects and TP53 alterations in men with treatment-naÃ⁻ve metastatic prostate cancer –Results from a prospective pilot study using a 37 gene panel. Urologic Oncology: Seminars and Original Investigations, 2020, 38, 637.e17-637.e27. | 1.6 | 12 |
| 121 | Evaluation of a Hybrid Capture–Based Pan-Cancer Panel for Analysis of Treatment Stratifying Oncogenic Aberrations and Processes. Journal of Molecular Diagnostics, 2020, 22, 757-769. | 2.8 | 42 |
| 122 | Recurrent YAP1 and MAML2 Gene Rearrangements in Retiform and Composite Hemangioendothelioma. American Journal of Surgical Pathology, 2020, 44, 1677-1684. | 3.7 | 51 |
| 123 | Targetable ERBB2 mutations identified in neurofibroma/schwannoma hybrid nerve sheath tumors. Journal of Clinical Investigation, 2020, 130, 2488-2495. | 8.2 | 23 |
| 124 | Safety and Preliminary Efficacy Results from a Phase II Study Evaluating Combined BRAF and MEK Inhibition in Relapsed/Refractory Multiple Myeloma (rrMM) Patients with Activating BRAF V600E Mutations: The GMMG-Birma Trial. Blood, 2020, 136, 44-45. | 1.4 | 16 |
| 125 | IMPACT OF DEEP TARGETED NEXT GENERATION SEQUENCING ON THE WORK-UP OF PATIENTS WITH PANCREAS CYSTS OR DILATED DUCT - A PROSPECTIVE STUDY WITH EUS-GUIDED FNA. Endoscopy, 2020, 52, . | 1.8 | 0 |
| 126 | Clinical and molecular profile of de novo vs. secondary EGFR mutated metastatic non-small-cell lung cancer. Pneumologie, 2020, 74, . | 0.1 | 0 |

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|-----|---|------|-----------|
| 127 | Combined Immunohistochemistry after Mass Spectrometry Imaging for Superior Spatial Information. Proteomics - Clinical Applications, 2019, 13, e1800035. | 1.6 | 23 |
| 128 | KIT-Dependent and KIT-Independent Genomic Heterogeneity of Resistance in Gastrointestinal Stromal Tumors — TORC1/2 Inhibition as Salvage Strategy. Molecular Cancer Therapeutics, 2019, 18, 1985-1996. | 4.1 | 22 |
| 129 | Optimizing panel-based tumor mutational burden (TMB) measurement. Annals of Oncology, 2019, 30, 1496-1506. | 1.2 | 123 |
| 130 | Classification of Cancer at Prostate MRI: Deep Learning versus Clinical PI-RADS Assessment. Radiology, 2019, 293, 607-617. | 7.3 | 214 |
| 131 | Prediction of significant prostate cancer in biopsy-naÃ ⁻ ve men: Validation of a novel risk model combining MRI and clinical parameters and comparison to an ERSPC risk calculator and PI-RADS. PLoS ONE, 2019, 14, e0221350. | 2.5 | 13 |
| 132 | Morphomolecular analysis of the immune tumor microenvironment in human head and neck cancer. Cancer Immunology, Immunotherapy, 2019, 68, 1443-1454. | 4.2 | 13 |
| 133 | RNA-Based Detection of Gene Fusions in Formalin-Fixed and Paraffin-Embedded Solid Cancer Samples. Cancers, 2019, 11, 1309. | 3.7 | 32 |
| 134 | In-house Implementation of Tumor Mutational Burden Testing to Predict Durable Clinical Benefit in Non-small Cell Lung Cancer and Melanoma Patients. Cancers, 2019, 11, 1271. | 3.7 | 27 |
| 135 | Spatial and Temporal Heterogeneity of Panel-Based Tumor Mutational Burden in Pulmonary Adenocarcinoma: Separating Biology From Technical Artifacts. Journal of Thoracic Oncology, 2019, 14, 1935-1947. | 1.1 | 69 |
| 136 | The BRCA2 mutation status shapes the immune phenotype of prostate cancer. Cancer Immunology, Immunotherapy, 2019, 68, 1621-1633. | 4.2 | 38 |
| 137 | Tumor mutational burden standardization initiatives: Recommendations for consistent tumor mutational burden assessment in clinical samples to guide immunotherapy treatment decisions. Genes Chromosomes and Cancer, 2019, 58, 578-588. | 2.8 | 173 |
| 138 | Detection of TP53 Mutations in Tissue or Liquid Rebiopsies at Progression Identifies ALK+ Lung Cancer Patients with Poor Survival. Cancers, 2019, 11, 124. | 3.7 | 36 |
| 139 | Variant classification in precision oncology. International Journal of Cancer, 2019, 145, 2996-3010. | 5.1 | 76 |
| 140 | Several genotypes, one phenotype: PIK3CA/AKT1 mutation-negative hidradenoma papilliferum show genetic lesions in other components of the signalling network. Pathology, 2019, 51, 362-368. | 0.6 | 10 |
| 141 | Comparative genetic profiling aids diagnosis and clinical decision making in challenging cases of CUP syndrome. International Journal of Cancer, 2019, 145, 2963-2973. | 5.1 | 24 |
| 142 | Mutational Diversity and Therapy Response in Breast Cancer: A Sequencing Analysis in the Neoadjuvant GeparSepto Trial. Clinical Cancer Research, 2019, 25, 3986-3995. | 7.0 | 32 |
| 143 | Defective homologous recombination DNA repair as therapeutic target in advanced chordoma. Nature Communications, 2019, 10, 1635. | 12.8 | 64 |
| 144 | Label-Free Enrichment and Molecular Characterization of Viable Circulating Tumor Cells from Diagnostic Leukapheresis Products. Clinical Chemistry, 2019, 65, 549-558. | 3.2 | 37 |

| # | Article | IF | CITATIONS |
|-----|--|------|-----------|
| 145 | Serial liquid biopsies for detection of treatment failure and profiling of resistance mechanisms in <i>KLC1–ALK</i> -rearranged lung cancer. Journal of Physical Education and Sports Management, 2019, 5, a004630. | 1.2 | 13 |
| 146 | Digital PCR After MALDI–Mass Spectrometry Imaging to Combine Proteomic Mapping and Identification of Activating Mutations in Pulmonary Adenocarcinoma. Proteomics - Clinical Applications, 2019, 13, e1800034. | 1.6 | 19 |
| 147 | Proteogenomic systems analysis identifies targeted therapy resistance mechanisms in EGFRâ€mutated lung cancer. International Journal of Cancer, 2019, 144, 545-557. | 5.1 | 8 |
| 148 | Development of tumor mutation burden as an immunotherapy biomarker: utility for the oncology clinic. Annals of Oncology, 2019, 30, 44-56. | 1.2 | 1,742 |
| 149 | Measurement of tumor mutational burden (TMB) in routine molecular diagnostics: <i>in silico</i> and realâ€life analysis of three larger gene panels. International Journal of Cancer, 2019, 144, 2303-2312. | 5.1 | 95 |
| 150 | Integrative Analysis Defines Distinct Prognostic Subgroups of Intrahepatic Cholangiocarcinoma. Hepatology, 2019, 69, 2091-2106. | 7.3 | 63 |
| 151 | Combined targeted DNA and RNA sequencing of advanced NSCLC in routine molecular diagnostics: Analysis of the first 3,000 Heidelberg cases. International Journal of Cancer, 2019, 145, 649-661. | 5.1 | 85 |
| 152 | Position of a panel of international lung cancer experts on the approval decision for use of durvalumab in stage III non-small-cell lung cancer (NSCLC) by the Committee for Medicinal Products for Human Use (CHMP). Annals of Oncology, 2019, 30, 161-165. | 1.2 | 60 |
| 153 | Global DNA methylation reflects spatial heterogeneity and molecular evolution of lung adenocarcinomas. International Journal of Cancer, 2019, 144, 1061-1072. | 5.1 | 22 |
| 154 | Histopathological to multiparametric MRI spatial mapping of extended systematic sextant and MR/TRUS-fusion-targeted biopsy of the prostate. European Radiology, 2019, 29, 1820-1830. | 4.5 | 24 |
| 155 | Size matters: Dissecting key parameters for panelâ€based tumor mutational burden analysis. International Journal of Cancer, 2019, 144, 848-858. | 5.1 | 131 |
| 156 | Next generation sequencing of the cellular and liquid fraction of pancreatic cyst fluid supports discrimination of IPMN from pseudocysts and reveals cases with multiple mutated driver clones: First findings from the prospective ZYSTEUS biomarker study. Genes Chromosomes and Cancer, 2019, 58, 3-11. | 2.8 | 14 |
| 157 | Identification of a highly lethal V3 ⁺ TP53 ⁺ subset in ALK ⁺ lung adenocarcinoma. International Journal of Cancer, 2019, 144, 190-199. | 5.1 | 67 |
| 158 | <i>RSPO2</i> gene rearrangement: a powerful driver of β-catenin activation in liver tumours. Gut, 2019, 68, 1287-1296. | 12.1 | 29 |
| 159 | Defining molecular risk in ALK+ NSCLC. Oncotarget, 2019, 10, 3093-3103. | 1.8 | 35 |
| 160 | Comparison of different semi-automated cfDNA extraction methods in combination with UMI-based targeted sequencing. Oncotarget, 2019, 10, 5690-5702. | 1.8 | 15 |
| 161 | MiRNAs involved in development of intraepithelial precursor lesions and progression to cholangiocarcinoma. Zeitschrift Fur Gastroenterologie, 2019, 57, . | 0.5 | 0 |
| 162 | Cholangiocarcinogenesis is a developmental process driven by distinct sequential alterations of the | | 0 |

cellular transcriptome., 2019, 57,.

| # | Article | IF | CITATIONS |
|-----|--|------|-----------|
| 163 | RSPO2 gene rearrangement – a new cancer driver in the liver. Zeitschrift Fur Gastroenterologie, 2019, 57, . | 0.5 | 0 |
| 164 | TP53 status conversion defines an unfavourable patient subset with inferior overall survival in ALK+ lung adenocarcinoma. , 2019, 73, . | | 0 |
| 165 | A framework for risk stratification in EGFR+ lung adenocarcinoma treated with tyrosine kinase inhibitors. , 2019, 73, . | | 0 |
| 166 | Nivolumab maintenance after salvage autologous stem cell transplantation results in longâ€ŧerm remission in multiple relapsed primary <scp>CNS</scp> lymphoma. European Journal of Haematology, 2018, 101, 115-118. | 2.2 | 19 |
| 167 | Oncogene-induced senescence: a potential breakpoint mechanism against malignant transformation in plasma cell disorders. Leukemia and Lymphoma, 2018, 59, 2660-2669. | 1.3 | 3 |
| 168 | Three molecular pathways model colorectal carcinogenesis in <scp>L</scp> ynch syndrome. International Journal of Cancer, 2018, 143, 139-150. | 5.1 | 129 |
| 169 | <i>EML4â€ALK</i> fusion variant V3 is a highâ€risk feature conferring accelerated metastatic spread, early treatment failure and worse overall survival in ALK ⁺ nonâ€small cell lung cancer. International Journal of Cancer, 2018, 142, 2589-2598. | 5.1 | 93 |
| 170 | Analysis of the proliferative activity in lung adenocarcinomas with specific driver mutations. Pathology Research and Practice, 2018, 214, 408-416. | 2.3 | 4 |
| 171 | Targeted molecular profiling reveals genetic heterogeneity of poromas and porocarcinomas. Pathology, 2018, 50, 327-332. | 0.6 | 27 |
| 172 | Appendiceal goblet cell carcinoids and adenocarcinomas ex-goblet cell carcinoid are genetically distinct from primary colorectal-type adenocarcinoma of the appendix. Modern Pathology, 2018, 31, 829-839. | 5.5 | 44 |
| 173 | Integrative genomic and transcriptomic analysis of leiomyosarcoma. Nature Communications, 2018, 9, 144. | 12.8 | 197 |
| 174 | Targeted deep sequencing of effusion cytology samples is feasible, informs spatiotemporal tumor evolution, and has clinical and diagnostic utility. Genes Chromosomes and Cancer, 2018, 57, 70-79. | 2.8 | 19 |
| 175 | Simultaneous whole-body 18F–PSMA-1007-PET/MRI with integrated high-resolution multiparametric imaging of the prostatic fossa for comprehensive oncological staging of patients with prostate cancer: a pilot study. European Journal of Nuclear Medicine and Molecular Imaging, 2018, 45, 340-347. | 6.4 | 32 |
| 176 | Implementation of a novel efficacy score to compare sealing and cutting devices in a porcine model. Surgical Endoscopy and Other Interventional Techniques, 2018, 32, 1002-1011. | 2.4 | 6 |
| 177 | Pancreatic Ductal Adenocarcinoma Subtyping Using the Biomarkers Hepatocyte Nuclear Factor-1A and Cytokeratin-81 Correlates with Outcome and Treatment Response. Clinical Cancer Research, 2018, 24, 351-359. | 7.0 | 81 |
| 178 | Validating Comprehensive Next-Generation Sequencing Results for Precision Oncology: The NCT/DKTK Molecularly Aided Stratification for Tumor Eradication Research Experience. JCO Precision Oncology, 2018, 2, 1-13. | 3.0 | 20 |
| 179 | Implementing tumor mutational burden (TMB) analysis in routine diagnostics—a primer for molecular pathologists and clinicians. Translational Lung Cancer Research, 2018, 7, 703-715. | 2.8 | 152 |
| 180 | EML4-ALK V3, treatment resistance, and survival: refining the diagnosis of ALK+ NSCLC. Journal of Thoracic Disease, 2018, 10, S1989-S1991. | 1.4 | 26 |

| # | Article | IF | CITATIONS |
|-----|---|-----|-----------|
| 181 | Subclonal evolution of pulmonary adenocarcinomas delineated by spatially distributed somatic mitochondrial mutations. Lung Cancer, 2018, 126, 80-88. | 2.0 | 16 |
| 182 | Genetic profiling of melanoma in routine diagnostics: assay performance and molecular characteristics in a consecutive series of 274 cases. Pathology, 2018, 50, 703-710. | 0.6 | 21 |
| 183 | Integrated analysis of the immunological and genetic status in and across cancer types: impact of mutational signatures beyond tumor mutational burden. Oncolmmunology, 2018, 7, e1526613. | 4.6 | 60 |
| 184 | <i>NRG1</i> Fusions in <i>KRAS</i> Wild-Type Pancreatic Cancer. Cancer Discovery, 2018, 8, 1087-1095. | 9.4 | 189 |
| 185 | Molecular dissection of large cell carcinomas of the lung with null immunophenotype. Pathology, 2018, 50, 530-535. | 0.6 | 7 |
| 186 | loncopy: an R Shiny app to call copy number alterations in targeted NGS data. BMC Bioinformatics, 2018, 19, 157. | 2.6 | 4 |
| 187 | Prostatic metastasis from intrahepatic cholangiocarcinoma. Urology Case Reports, 2018, 20, 90-91. | 0.3 | 4 |
| 188 | Integrated Histogenetic Analysis Reveals BAP1 -Mutated Epithelioid Mesothelioma in a Patient With Cancer of Unknown Primary. Journal of the National Comprehensive Cancer Network: JNCCN, 2018, 16, 677-682. | 4.9 | 6 |
| 189 | A field guide for cancer diagnostics using cellâ€free DNA: From principles to practice and clinical applications. Genes Chromosomes and Cancer, 2018, 57, 123-139. | 2.8 | 155 |
| 190 | Profiling of Oncogenic Signaling in Multiple Myeloma — Association with Biology, Disease Progression and Prognosis. Blood, 2018, 132, 3206-3206. | 1.4 | 1 |
| 191 | Colorectal mixed adenoneuroendocrine carcinomas and neuroendocrine carcinomas are genetically closely related to colorectal adenocarcinomas. Modern Pathology, 2017, 30, 610-619. | 5.5 | 131 |
| 192 | PD-L1 (CD274) copy number gain, expression, and immune cell infiltration as candidate predictors for response to immune checkpoint inhibitors in soft-tissue sarcoma. Oncolmmunology, 2017, 6, e1279777. | 4.6 | 50 |
| 193 | Marker chromosomes can arise from chromothripsis and predict adverse prognosis in acute myeloid leukemia. Blood, 2017, 129, 1333-1342. | 1.4 | 57 |
| 194 | Chloroquine enhances the antimycobacterial activity of isoniazid and pyrazinamide by reversing inflammation-induced macrophage efflux. International Journal of Antimicrobial Agents, 2017, 50, 55-62. | 2.5 | 15 |
| 195 | Mutation patterns in genes encoding interferon signaling and antigen presentation: A panâ€cancer survey with implications for the use of immune checkpoint inhibitors. Genes Chromosomes and Cancer, 2017, 56, 651-659. | 2.8 | 35 |
| 196 | Precision oncology based on omics data: The NCT Heidelberg experience. International Journal of Cancer, 2017, 141, 877-886. | 5.1 | 133 |
| 197 | Prevalence of somatic mitochondrial mutations and spatial distribution of mitochondria in non-small cell lung cancer. British Journal of Cancer, 2017, 117, 220-226. | 6.4 | 25 |
| 198 | MiR-200b and miR-155 as predictive biomarkers for the efficacy of chemoradiation in locally advanced head and neck squamous cell carcinoma. European Journal of Cancer, 2017, 77, 3-12. | 2.8 | 51 |

| # | Article | IF | CITATIONS |
|-----|--|-----|-----------|
| 199 | Genomics of Immunotherapy-Associated Hyperprogressors—Letter. Clinical Cancer Research, 2017, 23, 6374-6375. | 7.0 | 11 |
| 200 | EGFR T790M mutation testing of non-small cell lung cancer tissue and blood samples artificially spiked with circulating cell-free tumor DNA: results of a round robin trial. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2017, 471, 509-520. | 2.8 | 29 |
| 201 | Mutational profiles of Brenner tumors show distinctive features uncoupling urothelial carcinomas and ovarian carcinoma with transitional cell histology. Genes Chromosomes and Cancer, 2017, 56, 758-766. | 2.8 | 21 |
| 202 | Spatial distribution of <i>EGFR</i> and <i>KRAS</i> mutation frequencies correlates with histological growth patterns of lung adenocarcinomas. International Journal of Cancer, 2017, 141, 1841-1848. | 5.1 | 21 |
| 203 | Qualitative Comparison Between Carrier-based and Classical Tissue Microarrays. Applied Immunohistochemistry and Molecular Morphology, 2017, 25, e74-e79. | 1.2 | 15 |
| 204 | Targeted nextâ€generation sequencing enables reliable detection of HER2 (ERBB2) status in breast cancer and provides ancillary information of clinical relevance. Genes Chromosomes and Cancer, 2017, 56, 255-265. | 2.8 | 21 |
| 205 | Mutations of cancer-related genes in benign tumors: the example of hidradenoma papilliferum. Human Pathology, 2017, 62, 246-247. | 2.0 | 3 |
| 206 | Synonymous EGFR variant p.Q787Q is neither prognostic nor predictive in patients with lung adenocarcinoma. Genes Chromosomes and Cancer, 2017, 56, 214-220. | 2.8 | 8 |
| 207 | Tubular, lactating, and ductal adenomas are devoid of MED12 Exon2 mutations, and ductal adenomas show recurrent mutations in GNAS and the PI3K–AKT pathway. Genes Chromosomes and Cancer, 2017, 56, 11-17. | 2.8 | 27 |
| 208 | PARP inhibition in BRCA2-mutated prostate cancer. Annals of Oncology, 2017, 28, 189-191. | 1.2 | 12 |
| 209 | Mutant KIT as imatinib-sensitive target in metastatic sinonasal carcinoma. Annals of Oncology, 2017, 28, 142-148. | 1.2 | 30 |
| 210 | A non-controlled, single arm, open label, phase II study of intravenous and intratumoral administration of ParvOryx in patients with metastatic, inoperable pancreatic cancer: ParvOryx02 protocol. BMC Cancer, 2017, 17, 576. | 2.6 | 36 |
| 211 | Patient-specific molecular alterations are associated with metastatic clear cell renal cell cancer progressing under tyrosine kinase inhibitor therapy. Oncotarget, 2017, 8, 74049-74057. | 1.8 | 14 |
| 212 | Oncogenic driver mutations, treatment, and EGFR-TKI resistance in a Caucasian population with non-small cell lung cancer: survival in clinical practice. Oncotarget, 2017, 8, 77897-77914. | 1.8 | 19 |
| 213 | Implementation of a novel efficacy score to compare sealing and cutting devices in a porcine model. , 2017, 77, . | | 0 |
| 214 | Targeting irradiationâ€induced mitogenâ€activated protein kinase activation in vitro and in an ex vivo model for human head and neck cancer. Head and Neck, 2016, 38, E2049-61. | 2.0 | 16 |
| 215 | Mutations in genes encoding <scp>PI3Kâ€AKT</scp> and <scp>MAPK</scp> signaling define anogenital papillary hidradenoma. Genes Chromosomes and Cancer, 2016, 55, 113-119. | 2.8 | 29 |
| 216 | Highâ€ŧhroughput diagnostic profiling of clinically actionable gene fusions in lung cancer. Genes Chromosomes and Cancer, 2016, 55, 30-44. | 2.8 | 65 |

| # | Article | IF | CITATIONS |
|-----|--|------------------|-----------|
| 217 | Panâ€cancer analysis of copy number changes in programmed deathâ€ligand 1 (PDâ€L1, CD274) – association with gene expression, mutational load, and survival. Genes Chromosomes and Cancer, 2016, 55, 626-639. | ^S 2.8 | 80 |
| 218 | Cadherin-6 is a putative tumor suppressor and target of epigenetically dysregulated miR-429 in cholangiocarcinoma. Epigenetics, 2016, 11, 780-790. | 2.7 | 33 |
| 219 | Targeted next-generation sequencing identifies molecular subgroups in squamous cell carcinoma of the head and neck with distinct outcome after concurrent chemoradiation. Annals of Oncology, 2016, 27, 2262-2268. | 1.2 | 38 |
| 220 | Reliable Entity Subtyping in Non-small Cell Lung Cancer by Matrix-assisted Laser Desorption/Ionization Imaging Mass Spectrometry on Formalin-fixed Paraffin-embedded Tissue Specimens. Molecular and Cellular Proteomics, 2016, 15, 3081-3089. | 3.8 | 72 |
| 221 | Health Care Infrastructure for Financially Sustainable Clinical Genomics. Journal of Molecular Diagnostics, 2016, 18, 697-706. | 2.8 | 15 |
| 222 | Next-generation sequencing facilitates detection of the classic E13-A20 EML4-ALK fusion in an ALK-FISH/IHC inconclusive biopsy of a stage IV lung cancer patient: a case report. Diagnostic Pathology, 2016, 11, 133. | 2.0 | 8 |
| 223 | Integration of genomics and histology revises diagnosis and enables effective therapy of refractory cancer of unknown primary with <i>PDL1</i> amplification. Journal of Physical Education and Sports Management, 2016, 2, a001180. | 1.2 | 57 |
| 224 | Genetic heterogeneity in synchronous colorectal cancers impacts genotyping approaches and therapeutic strategies. Genes Chromosomes and Cancer, 2016, 55, 268-277. | 2.8 | 28 |
| 225 | Genotyping of colorectal cancer for cancer precision medicine: Results from the IPH Center for Molecular Pathology. Genes Chromosomes and Cancer, 2016, 55, 505-521. | 2.8 | 34 |
| 226 | Copy number changes of clinically actionable genes in melanoma, nonâ€small cell lung cancer and colorectal cancer—A survey across 822 routine diagnostic cases. Genes Chromosomes and Cancer, 2016, 55, 821-833. | 2.8 | 43 |
| 227 | NGS-based BRCA1/2 mutation testing of high-grade serous ovarian cancer tissue: results and conclusions of the first international round robin trial. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2016, 468, 697-705. | 2.8 | 24 |
| 228 | Individualized medicine and demographic change as determining workload factors in pathology: quo vadis?. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2016, 468, 101-108. | 2.8 | 19 |
| 229 | CYP3A5 mediates basal and acquired therapy resistance in different subtypes of pancreatic ductal adenocarcinoma. Nature Medicine, 2016, 22, 278-287. | 30.7 | 184 |
| 230 | The putative oncogene CEP72 inhibits the mitotic function of BRCA1 and induces chromosomal instability. Oncogene, 2016, 35, 2398-2406. | 5.9 | 22 |
| 231 | Molecular driver alterations and their clinical relevance in cancer of unknown primary site. Oncotarget, 2016, 7, 44322-44329. | 1.8 | 47 |
| 232 | Role of <i>TP53</i> mutations in triple negative and HER2-positive breast cancer treated with neoadjuvant anthracycline/taxane-based chemotherapy. Oncotarget, 2016, 7, 67686-67698. | 1.8 | 50 |
| 233 | loncopy: a novel method for calling copy number alterations in amplicon sequencing data including significance assessment. Oncotarget, 2016, 7, 13236-13247. | 1.8 | 23 |
| 234 | Genetic changes of non-small cell lung cancer under neoadjuvant therapy. Oncotarget, 2016, 7, 29761-29769. | 1.8 | 16 |

| # | Article | IF | CITATIONS |
|-----|---|-----|-----------|
| 235 | Histone deacetylase inhibition sensitizes osteosarcoma to heavy ion radiotherapy. Radiation Oncology, 2015, 10, 146. | 2.7 | 14 |
| 236 | Basket Trials: Just the End of the First Quarter. Journal of Clinical Oncology, 2015, 33, 2823-2824. | 1.6 | 15 |
| 237 | Distribution of <i>MED12</i> mutations in fibroadenomas and phyllodes tumors of the breast—implications for tumor biology and pathological diagnosis. Genes Chromosomes and Cancer, 2015, 54, 444-452. | 2.8 | 55 |
| 238 | Distinctive Spatiotemporal Stability of Somatic Mutations in Metastasized Microsatellite-stable Colorectal Cancer. American Journal of Surgical Pathology, 2015, 39, 1140-1147. | 3.7 | 35 |
| 239 | The landscape of metastatic progression patterns across major human cancers. Oncotarget, 2015, 6, 570-583. | 1.8 | 208 |
| 240 | Establishment of a patient-derived orthotopic osteosarcoma mouse model. Journal of Translational Medicine, 2015, 13, 136. | 4.4 | 35 |
| 241 | Semiconductorâ€Based Sequencing of Formalinâ€Fixed, Paraffinâ€Embedded Colorectal Cancer Samples. Oncologist, 2015, 20, e10-1. | 3.7 | 10 |
| 242 | Contribution of human papilloma virus to the incidence of squamous cell carcinoma of the head and neck in a European population with high smoking prevalence. European Journal of Cancer, 2015, 51, 514-521. | 2.8 | 75 |
| 243 | CCI-779 (Temsirolimus) exhibits increased anti-tumor activity in low EGFR expressing HNSCC cell lines and is effective in cells with acquired resistance to cisplatin or cetuximab. Journal of Translational Medicine, 2015, 13, 106. | 4.4 | 11 |
| 244 | Prognostic Impact and Clinicopathological Correlations of the Cribriform Pattern in Pulmonary Adenocarcinoma. Journal of Thoracic Oncology, 2015, 10, 638-644. | 1.1 | 83 |
| 245 | Allelic Ratio of <i>KRAS</i> Mutations in Pancreatic Cancer. Oncologist, 2015, 20, e8-e9. | 3.7 | 36 |
| 246 | Loss of SOX2 expression induces cell motility via vimentin upâ€regulation and is an unfavorable risk factor for survival ofÂhead and neck squamous cell carcinoma. Molecular Oncology, 2015, 9, 1704-1719. | 4.6 | 60 |
| 247 | Classical pathology and mutational load of breast cancer – integration of two worlds. Journal of Pathology: Clinical Research, 2015, 1, 225-238. | 3.0 | 91 |
| 248 | Major histocompatibility complex class I expression impacts on patient survival and type and density of immune cells in biliary tract cancer. British Journal of Cancer, 2015, 113, 1343-1349. | 6.4 | 54 |
| 249 | Histological tumor typing in the age of molecular profiling. Pathology Research and Practice, 2015, 211, 897-900. | 2.3 | 10 |
| 250 | ALK-FISH borderline cases in non-small cell lung cancer: Implications for diagnostics and clinical decision making. Lung Cancer, 2015, 90, 465-471. | 2.0 | 36 |
| 251 | GOT1/AST1 expression status as a prognostic biomarker in pancreatic ductal adenocarcinoma. Oncotarget, 2015, 6, 4516-4526. | 1.8 | 22 |
| 252 | EGFR, KRAS, BRAF and ALK gene alterations in lung adenocarcinomas: patient outcome, interplay with morphology and immunophenotype. European Respiratory Journal, 2014, 43, 872-883. | 6.7 | 97 |

| # | Article | IF | CITATIONS |
|-----|--|------|-----------|
| 253 | BRAF V600E-specific immunohistochemistry reveals low mutation rates in biliary tract cancer and restriction to intrahepatic cholangiocarcinoma. Modern Pathology, 2014, 27, 1028-1034. | 5.5 | 96 |
| 254 | KIT mutations in primary mediastinal B-cell lymphoma. Blood Cancer Journal, 2014, 4, e241-e241. | 6.2 | 2 |
| 255 | Increased microtubule assembly rates influence chromosomal instability in colorectal cancer cells. Nature Cell Biology, 2014, 16, 779-791. | 10.3 | 174 |
| 256 | Cancer beyond organ and tissue specificity: Nextâ€generationâ€sequencing gene mutation data reveal complex genetic similarities across major cancers. International Journal of Cancer, 2014, 135, 2362-2369. | 5.1 | 36 |
| 257 | Global alterations of DNA methylation in cholangiocarcinoma target the Wnt signaling pathway. Hepatology, 2014, 59, 544-554. | 7.3 | 97 |
| 258 | KRAS Mutations in Codon 12 or 13 Are Associated With Worse Prognosis in Pancreatic Ductal Adenocarcinoma. Pancreas, 2014, 43, 578-583. | 1.1 | 36 |
| 259 | <i><scp>ROS</scp>1</i> expression and translocations in nonâ€smallâ€cell lung cancer: clinicopathological analysis of 1478 cases. Histopathology, 2014, 65, 187-194. | 2.9 | 96 |
| 260 | Comprehensive analysis of clinico-pathological data reveals heterogeneous relations between atherosclerosis and cancer. Journal of Clinical Pathology, 2014, 67, 482-490. | 2.0 | 13 |
| 261 | Tumour cell proliferation (Ki-67) in non-small cell lung cancer: a critical reappraisal of its prognostic role. British Journal of Cancer, 2014, 111, 1222-1229. | 6.4 | 114 |
| 262 | Mutations in POLE and survival of colorectal cancer patients – link to disease stage and treatment. Cancer Medicine, 2014, 3, 1527-1538. | 2.8 | 56 |
| 263 | Clinical and molecular characteristics of HNSCC patients with brain metastases: a retrospective study. European Archives of Oto-Rhino-Laryngology, 2014, 271, 1715-1722. | 1.6 | 16 |
| 264 | The DNA index is a strong predictive marker in intrahepatic cholangiocarcinoma: the results of a five-year prospective study. Surgery Today, 2014, 44, 1336-1342. | 1.5 | 12 |
| 265 | The combinatorial complexity of cancer precision medicine. Oncoscience, 2014, 1, 504-509. | 2.2 | 48 |
| 266 | Potential clinical implications of <i>BRAF</i> mutations in histiocytic proliferations. Oncotarget, 2014, 5, 4060-4070. | 1.8 | 78 |
| 267 | Targeted ultra-deep sequencing reveals recurrent and mutually exclusive mutations of cancer genes in blastic plasmacytoid dendritic cell neoplasm. Oncotarget, 2014, 5, 6404-6413. | 1.8 | 82 |
| 268 | Co-expression of MET and CD47 is a novel prognosticator for survival of luminal-type breast cancer patients. Oncotarget, 2014, 5, 8147-8160. | 1.8 | 83 |
| 269 | Mutational profiles in triple-negative breast cancer defined by ultradeep multigene sequencing show high rates of PI3K pathway alterations and clinically relevant entity subgroup specific differences. Oncotarget, 2014, 5, 9952-9965. | 1.8 | 58 |
| 270 | Molecular Diagnostic Profiling of Lung Cancer Specimens with a Semiconductor-Based Massive Parallel Sequencing Approach. Journal of Molecular Diagnostics, 2013, 15, 765-775. | 2.8 | 107 |

| # | Article | IF | CITATIONS |
|-----|--|------|-----------|
| 271 | Radiosensitization by histone deacetylase inhibition in an osteosarcoma mouse model. Strahlentherapie Und Onkologie, 2013, 189, 957-966. | 2.0 | 14 |
| 272 | Identification of a population of blood circulating tumor cells from breast cancer patients that initiates metastasis in a xenograft assay. Nature Biotechnology, 2013, 31, 539-544. | 17.5 | 920 |
| 273 | Follicleâ€stimulating hormone receptor expression in soft tissue sarcomas. Histopathology, 2013, 63, 29-35. | 2.9 | 16 |
| 274 | Correlation of radio- and histomorphological pattern of pulmonary adenocarcinoma. European Respiratory Journal, 2013, 41, 943-951. | 6.7 | 105 |
| 275 | DNA Index as a Strong Prognostic Factor in Patients With Adenocarcinoma of the Pancreatic Head. Pancreas, 2013, 42, 807-812. | 1.1 | 14 |
| 276 | Fibroblast Growth Factor Receptor 1 as a Putative Therapy Target in Colorectal Cancer. Digestion, 2013, 88, 172-181. | 2.3 | 25 |
| 277 | Prognostic impact of tumour-infiltrating immune cells on biliary tract cancer. British Journal of Cancer, 2013, 109, 2665-2674. | 6.4 | 209 |
| 278 | High SIRT1 expression is a negative prognosticator in pancreatic ductal adenocarcinoma. BMC Cancer, 2013, 13, 450. | 2.6 | 63 |
| 279 | Survival of Patients with Oral Cavity Cancer in Germany. PLoS ONE, 2013, 8, e53415. | 2.5 | 69 |
| 280 | Absence Of BRAF and KRAS Hotspot Mutations In Primary Mediastinal and Other Diffuse Large B-Cell Lymphoma. Blood, 2013, 122, 4325-4325. | 1.4 | 0 |
| 281 | Prolyl Hydroxylase Domain 2 Protein Is a Strong Prognostic Marker in Human Gastric Cancer. Pathobiology, 2012, 79, 11-17. | 3.8 | 13 |
| 282 | Interobserver variability in the application of the novel IASLC/ATS/ERS classification for pulmonary adenocarcinomas. European Respiratory Journal, 2012, 40, 1221-1227. | 6.7 | 97 |
| 283 | BRAFV600E mutant protein is expressed in cells of variable maturation in Langerhans cell histiocytosis. Blood, 2012, 120, e28-e34. | 1.4 | 199 |
| 284 | Quantitative Analysis of Diagnostic Guidelines for HER2-Status Assessment. Journal of Molecular Diagnostics, 2012, 14, 199-205. | 2.8 | 18 |
| 285 | The Novel Histologic International Association for the Study of Lung Cancer/American Thoracic Society/European Respiratory Society Classification System of Lung Adenocarcinoma Is a Stage-Independent Predictor of Survival. Journal of Clinical Oncology, 2012, 30, 1438-1446. | 1.6 | 606 |
| 286 | Training increases concordance in classifying pulmonary adenocarcinomas according to the novel IASLC/ATS/ERS classification. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2012, 461, 185-193. | 2.8 | 55 |
| 287 | Reversion-inducing cysteine-rich protein with Kazal motif (RECK) expression: an independent prognostic marker of survival in colorectal cancer. Human Pathology, 2012, 43, 1314-1321. | 2.0 | 15 |
| 288 | High extracellular matrix metalloproteinase inducer/CD147 expression is strongly and independently associated with poor prognosis in colorectal cancer. Human Pathology, 2012, 43, 1471-1481. | 2.0 | 30 |

| # | Article | IF | CITATIONS |
|-----|--|-----|-----------|
| 289 | Postoperative Complications Deteriorate Long-Term Outcome in Pancreatic Cancer Patients. Annals of Surgical Oncology, 2012, 19, 856-863. | 1.5 | 78 |
| 290 | Phenotyping of pulmonary carcinoids and a Ki-67-based grading approach. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2012, 460, 299-308. | 2.8 | 60 |
| 291 | Visceral leishmaniasis in a patient with AIDS: early pathological diagnosis using conventional histology, PCR and electron microscopy is the key for adequate treatment. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2012, 460, 357-360. | 2.8 | 5 |
| 292 | High nuclear polyâ€(ADPâ€ribose)â€polymerase expression is prognostic of improved survival in pancreatic cancer. Histopathology, 2012, 61, 409-416. | 2.9 | 31 |
| 293 | Downâ€regulation of the microRNA processing enzyme Dicer is a prognostic factor in human colorectal cancer. Histopathology, 2012, 61, 552-561. | 2.9 | 44 |
| 294 | Who Is at Risk for Diagnostic Discrepancies? Comparison of Pre- and Postmortal Diagnoses in 1800 Patients of 3 Medical Decades in East and West Berlin. PLoS ONE, 2012, 7, e37460. | 2.5 | 53 |
| 295 | The presence of circulating tumor cells (CTCs) correlates with lymph node metastasis in nonresectable squamous cell carcinoma of the head and neck region (SCCHN). Annals of Oncology, 2011, 22, 1878-1885. | 1.2 | 112 |
| 296 | Decreased RECK and Increased EMMPRIN Expression in Urothelial Carcinoma of the Bladder Are Associated with Tumor Aggressiveness. Pathobiology, 2011, 78, 123-131. | 3.8 | 13 |
| 297 | Expression of Amphiregulin and EGFRvIII Affect Outcome of Patients with Squamous Cell Carcinoma of the Head and Neck Receiving Cetuximab–Docetaxel Treatment. Clinical Cancer Research, 2011, 17, 5197-5204. | 7.0 | 85 |
| 298 | mTOR expression and activity patterns in gastroenteropancreatic neuroendocrine tumours. Endocrine-Related Cancer, 2011, 18, 181-192. | 3.1 | 90 |
| 299 | KRAS Genotyping of Paraffin-Embedded Colorectal Cancer Tissue in Routine Diagnostics. Journal of Molecular Diagnostics, 2010, 12, 35-42. | 2.8 | 94 |
| 300 | Chapter 6 Cell and Molecular Biology of the Novel Protein Tyrosineâ€Phosphataseâ€Interacting Protein 51. International Review of Cell and Molecular Biology, 2009, 275, 183-246. | 3.2 | 25 |
| 301 | Expression profile of PTPIP51 in mouse brain. Journal of Comparative Neurology, 2009, 517, 892-905. | 1.6 | 18 |
| 302 | The novel protein PTPIP51 is expressed in human keratinocyte carcinomas and their surrounding stroma. Journal of Cellular and Molecular Medicine, 2008, 12, 2083-2095. | 3.6 | 16 |
| 303 | Epidermal Growth Factor-, Transforming Growth Factor-β-, Retinoic Acid- and 1,25-Dihydroxyvitamin D ₃ -Regulated Expression of the Novel Protein PTPIP51 in Keratinocytes. Cells Tissues Organs, 2006, 184, 76-87. | 2.3 | 21 |
| 304 | The novel protein PTPIP51 exhibits tissue- and cell-specific expression. Histochemistry and Cell Biology, 2005, 123, 19-28. | 1.7 | 26 |