

Vanesa GarcÃ-a-BarberÃ;n

List of Publications by Year in descending order

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Version: 2024-02-01

31
papers

1,388
citations

623734

14
h-index

580821

25
g-index

32
all docs

32
docs citations

32
times ranked

3361
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 1 | Cancer Risks Associated With <i>BRCA1</i> and <i>BRCA2</i> Pathogenic Variants. <i>Journal of Clinical Oncology</i> , 2022, 40, 1529-1541. | 1.6 | 90 |
| 2 | Abstract P2-01-18: Orthogonal assessment of <i>PIK3CA</i> and <i>ESR1</i> mutation detection in longitudinal cfDNA samples from endocrine-resistant HR+/HER2- advanced breast cancer patients using dPCR and NGS-based SafeSEQ technology. <i>Cancer Research</i> , 2022, 82, P2-01-18-P2-01-18. | 0.9 | 0 |
| 3 | Abstract P3-05-06: Genome-wide DNA methylation analysis identifies novel biomarkers associated with risk of relapse beyond oncoTYPE DX recurrence-score risk assessment within HR+/HER2- early-stage breast cancer patients. <i>Cancer Research</i> , 2022, 82, P3-05-06-P3-05-06. | 0.9 | 0 |
| 4 | Detection of IDH1 Mutations in Plasma Using BEAMing Technology in Patients with Gliomas. <i>Cancers</i> , 2022, 14, 2891. | 3.7 | 8 |
| 5 | <i>BRIP1</i> , a Gene Potentially Implicated in Familial Colorectal Cancer Type X. <i>Cancer Prevention Research</i> , 2021, 14, 185-194. | 1.5 | 7 |
| 6 | A case-only study to identify genetic modifiers of breast cancer risk for <i>BRCA1/BRCA2</i> mutation carriers. <i>Nature Communications</i> , 2021, 12, 1078. | 12.8 | 19 |
| 7 | Mapping of Genomic Vulnerabilities in the Post-Translational Ubiquitination, SUMOylation and Neddylation Machinery in Breast Cancer. <i>Cancers</i> , 2021, 13, 833. | 3.7 | 11 |
| 8 | Cancer-associated fibroblast-derived gene signatures determine prognosis in colon cancer patients. <i>Molecular Cancer</i> , 2021, 20, 73. | 19.2 | 44 |
| 9 | Genomic Mapping of Splicing-Related Genes Identify Amplifications in <i>LSM1</i> , <i>CLNS1A</i> , and <i>ILF2</i> in Luminal Breast Cancer. <i>Cancers</i> , 2021, 13, 4118. | 3.7 | 7 |
| 10 | Association of Genomic Domains in <i>BRCA1</i> and <i>BRCA2</i> with Prostate Cancer Risk and Aggressiveness. <i>Cancer Research</i> , 2020, 80, 624-638. | 0.9 | 39 |
| 11 | Plasma PD-L1 levels according to histologic grade and IDH status in patients with gliomas.. <i>Journal of Clinical Oncology</i> , 2020, 38, 69-69. | 1.6 | 2 |
| 12 | Monitoring of <i>PIK3CA</i> and <i>ESR1</i> mutations in circulating tumor DNA as predictive and prognostic biomarkers in patients with endocrine-resistant ER+/HER2- advanced breast cancer.. <i>Journal of Clinical Oncology</i> , 2020, 38, e13045-e13045. | 1.6 | 0 |
| 13 | Comparison of circulating tumor cells and cell-free DNA in the molecular characterization of patients with head and neck cancer.. <i>Journal of Clinical Oncology</i> , 2020, 38, e18521-e18521. | 1.6 | 0 |
| 14 | Contribution of New Adenomatous Polyposis Predisposition Genes in an Unexplained Attenuated Spanish Cohort by Multigene Panel Testing. <i>Scientific Reports</i> , 2019, 9, 9814. | 3.3 | 9 |
| 15 | Comparison of the Clinical Sensitivity of the Idylla Platform and the OncoBEAM RAS CRC Assay for <i>KRAS</i> Mutation Detection in Liquid Biopsy Samples. <i>Scientific Reports</i> , 2019, 9, 8976. | 3.3 | 34 |
| 16 | Circulating biomarkers for early detection and clinical management of colorectal cancer. <i>Molecular Aspects of Medicine</i> , 2019, 69, 107-122. | 6.4 | 214 |
| 17 | Alternative splicing and ACMG-AMP-2015-based classification of <i>PALB2</i> genetic variants: an ENIGMA report. <i>Journal of Medical Genetics</i> , 2019, 56, 453-460. | 3.2 | 30 |
| 18 | A Snapshot of The Tumor Microenvironment in Colorectal Cancer: The Liquid Biopsy. <i>International Journal of Molecular Sciences</i> , 2019, 20, 6016. | 4.1 | 27 |

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|----|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 19 | Height and Body Mass Index as Modifiers of Breast Cancer Risk in <i>BRCA1</i> Mutation Carriers: A Mendelian Randomization Study. <i>Journal of the National Cancer Institute</i> , 2019, 111, 350-364. | 6.3 | 30 |
| 20 | Mutational profile of dysplastic lesions evolving to laryngeal cancer.. <i>Journal of Clinical Oncology</i> , 2019, 37, e17551-e17551. | 1.6 | 1 |
| 21 | RAS analysis of circulating tumor cells from advanced colorectal cancer using BEAMing technology.. <i>Journal of Clinical Oncology</i> , 2019, 37, e15151-e15151. | 1.6 | 0 |
| 22 | Novel genetic mutations detected by multigene panel are associated with hereditary colorectal cancer predisposition. <i>PLoS ONE</i> , 2018, 13, e0203885. | 2.5 | 24 |
| 23 | Differential distribution and enrichment of non-coding RNAs in exosomes from normal and Cancer-associated fibroblasts in colorectal cancer. <i>Molecular Cancer</i> , 2018, 17, 114. | 19.2 | 61 |
| 24 | Evaluation of the sensitivity of RAS mutation detection of the Idylla platform in comparison to the OncoBEAM RAS CRC assay.. <i>Journal of Clinical Oncology</i> , 2018, 36, 592-592. | 1.6 | 2 |
| 25 | Tumor burden monitoring using cell-free tumor DNA could be limited by tumor heterogeneity in advanced breast cancer and should be evaluated together with radiographic imaging. <i>BMC Cancer</i> , 2017, 17, 210. | 2.6 | 59 |
| 26 | Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691. | 21.4 | 356 |
| 27 | Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017, 49, 1767-1778. | 21.4 | 289 |
| 28 | Role of GALNT12 in the genetic predisposition to attenuated adenomatous polyposis syndrome. <i>PLoS ONE</i> , 2017, 12, e0187312. | 2.5 | 10 |
| 29 | A novel TP53 germline inframe deletion identified in a Spanish series of Li-fraumeni syndrome suspected families. <i>Familial Cancer</i> , 2017, 16, 567-575. | 1.9 | 5 |
| 30 | Noninvasive EGFR testing in plasma circulating free DNA (cfDNA) by a new diagnostic method to detect point mutations, deletions and insertions associated to non small cell lung cancer: <i>CLART CMA EGFR LB</i> .. <i>Journal of Clinical Oncology</i> , 2017, 35, e20008-e20008. | 1.6 | 0 |
| 31 | Fine-Scale Mapping at 9p22.2 Identifies Candidate Causal Variants That Modify Ovarian Cancer Risk in <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>PLoS ONE</i> , 2016, 11, e0158801. | 2.5 | 10 |