## Francis Barany

List of Publications by Year in descending order

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Version: 2024-02-01

201674 206112 3,121 49 27 48 citations h-index g-index papers 50 50 50 3912 docs citations times ranked citing authors all docs

| #  | Article  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Solid-phase XRN1 reactions for RNA cleavage: application in single-molecule sequencing. Nucleic Acids Research, 2021, 49, e41-e41.   | 14.5 | 6         |
| 2  | PROTACs: Promising Approaches for Epigenetic Strategies to Overcome Drug Resistance. Current Cancer Drug Targets, 2021, 21, 306-325.   | 1.6  | 4         |
| 3  | MGMT Epigenetics: The Influence of Gene Body Methylation and Other Insights Derived from Integrated Methylomic, Transcriptomic, and Chromatin Analyses in Various Cancer Types. Current Cancer Drug Targets, 2021, 21, 360-374.                              | 1.6  | 5         |
| 4  | A Unified Transcriptional, Pharmacogenomic, and Gene Dependency Approach to Decipher the Biology, Diagnostic Markers, and Therapeutic Targets Associated with Prostate Cancer Metastasis. Cancers, 2021, 13, 5158.   | 3.7  | 3         |
| 5  | Novel, Self-Assembling Dimeric Inhibitors of Human $\hat{l}^2$ Tryptase. Journal of Medicinal Chemistry, 2020, 63, 3004-3027.  | 6.4  | 29        |
| 6  | Singleâ€molecule detection of cancer mutations using a novel PCRâ€LDRâ€qPCR assay. Human Mutation, 2020, 41, 1051-1068.  | 2.5  | 8         |
| 7  | Application of Multiplex Bisulfite PCR–Ligase Detection Reaction–Real-Time Quantitative PCR Assay in Interrogating Bioinformatically Identified, Blood-Based Methylation Markers for Colorectal Cancer. Journal of Molecular Diagnostics, 2020, 22, 885-900. | 2.8  | 5         |
| 8  | Prediction of blood-based biomarkers and subsequent design of bisulfite PCR-LDR-qPCR assay for breast cancer detection. BMC Cancer, 2020, 20, 85.  | 2.6  | 12        |
| 9  | Can CpG methylation serve as surrogate markers for immune infiltration in cancer?. Advances in Cancer Research, 2019, 143, 351-384.  | 5.0  | 19        |
| 10 | Pathways- and epigenetic-based assessment of relative immune infiltration in various types of solid tumors. Advances in Cancer Research, 2019, 142, 107-143.   | 5.0  | 10        |
| 11 | A Multiplex PCR/LDR Assay for Viral Agents of Diarrhea with the Capacity to Genotype Rotavirus.<br>Scientific Reports, 2018, 8, 13215.   | 3.3  | 3         |
| 12 | Target-Directed Self-Assembly of Homodimeric Drugs Against $\hat{l}^2$ -Tryptase. ACS Medicinal Chemistry Letters, 2018, 9, 827-831.   | 2.8  | 5         |
| 13 | Reversible Linkage of Two Distinct Small Molecule Inhibitors of Myc Generates a Dimeric Inhibitor with Improved Potency That Is Active in Myc Over-Expressing Cancer Cell Lines. PLoS ONE, 2015, 10, e0121793.   | 2.5  | 14        |
| 14 | A Multiplex PCR/LDR Assay for the Simultaneous Identification of Category A Infectious Pathogens: Agents of Viral Hemorrhagic Fever and Variola Virus. PLoS ONE, 2015, 10, e0138484.   | 2.5  | 15        |
| 15 | Abstract B43: Reversible linkage of two distinct small molecule inhibitors of MYC generates a more potent and selective dimeric inhibitor that is active in cancer cell lines over-expressing MYC., 2015,,.  |      | О         |
| 16 | A multiplex PCR/LDR assay for simultaneous detection and identification of the NIAID category B bacterial food and water-borne pathogens. Diagnostic Microbiology and Infectious Disease, 2014, 79, 135-140.   | 1.8  | 12        |
| 17 | EndoV/DNA ligase mutation scanning assay using microchip capillary electrophoresis and dual-color laser-induced fluorescence detection. Analytical Methods, 2012, 4, 58-64.  | 2.7  | 34        |
| 18 | Modular microfluidic system fabricated in thermoplastics for the strain-specific detection of bacterial pathogens. Lab on A Chip, 2012, 12, 3348.  | 6.0  | 31        |

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|----|--|--------------|-----------|
| 19 | High-Throughput Selection, Enumeration, Electrokinetic Manipulation, and Molecular Profiling of Low-Abundance Circulating Tumor Cells Using a Microfluidic System. Analytical Chemistry, 2011, 83, 2301-2309.                    | 6.5          | 168       |
| 20 | Molecular Profiling of Colon Tumors: The Search for Clinically Relevant Biomarkers of Progression, Prognosis, Therapeutics, and Predisposition. Annals of Surgical Oncology, 2011, 18, 3694-3700.                                | 1.5          | 51        |
| 21 | Gene Dysregulations Driven by Somatic Copy Number Aberrations-Biological and Clinical Implications in Colon Tumors. Journal of Molecular Diagnostics, 2010, 12, 552-561.   | 2.8          | 23        |
| 22 | PCR/LDR/Universal Array Platforms for the Diagnosis of Infectious Disease. Methods in Molecular Biology, 2010, 632, 141-157.   | 0.9          | 6         |
| 23 | Emerging Paradigms in Cancer Genetics: Some Important Findings from High-Density Single Nucleotide Polymorphism Array Studies: Fig. 1 Cancer Research, 2009, 69, 723-727.  | 0.9          | 46        |
| 24 | Genomeâ€wide autozygosity mapping in human populations. Genetic Epidemiology, 2009, 33, 172-180.   | 1.3          | 58        |
| 25 | Association of survival and disease progression with chromosomal instability: A genomic exploration of colorectal cancer. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 7131-7136. | 7.1          | 326       |
| 26 | Ligase detection reaction for the analysis of point mutations using freeâ€solution conjugate electrophoresis in a polymer microfluidic device. Electrophoresis, 2008, 29, 4751-4760.   | 2.4          | 24        |
| 27 | CpG Island Methylator Phenotype Associates with Low-Degree Chromosomal Abnormalities in Colorectal Cancer. Clinical Cancer Research, 2008, 14, 6005-6013.  | 7.0          | 101       |
| 28 | The Signatures of Autozygosity among Patients with Colorectal Cancer. Cancer Research, 2008, 68, 2610-2621.  | 0.9          | 47        |
| 29 | MDM2 Gene Amplification Is Correlated to Tumor Progression but not to the Presence of SNP309 or TP53 Mutational Status in Primary Colorectal Cancers. Molecular Cancer Research, 2008, 6, 205-211.                               | 3.4          | 47        |
| 30 | Multiplexed Identification of Blood-Borne Bacterial Pathogens by Use of a Novel 16S rRNA Gene PCR-Ligase Detection Reaction-Capillary Electrophoresis Assay. Journal of Clinical Microbiology, 2007, 45, 1927-1935.              | 3.9          | 44        |
| 31 | Serial processing of biological reactions using flow-through microfluidic devices: coupled PCR/LDR for the detection of low-abundant DNA point mutations. Analyst, The, 2007, 132, 913.  | <b>3.</b> 5  | 46        |
| 32 | Harnessing asymmetrical substrate recognition by thermostable EndoV to achieve balanced linear amplification in multiplexed SNP typing. Biochemistry and Cell Biology, 2006, 84, 232-242.  | 2.0          | 4         |
| 33 | Polymerase chain reaction/ligase detection reaction/hybridization assays using flow-through microfluidic devices for the detection of low-abundant DNA point mutations. Biosensors and Bioelectronics, 2006, 21, 1915-1923.      | 10.1         | 125       |
| 34 | Relationship of Gene Expression and Chromosomal Abnormalities in Colorectal Cancer. Cancer Research, 2006, 66, 2129-2137.  | 0.9          | 231       |
| 35 | Fabrication of DNA microarrays onto poly(methyl methacrylate) with ultraviolet patterning and microfluidics for the detection of low-abundant point mutations. Analytical Biochemistry, 2005, 340, 123-135.                      | 2.4          | 81        |
| 36 | Multiplexed profiling of candidate genes for CpG island methylation status using a flexible PCR/LDR/Universal Array assay. Genome Research, 2005, 16, 282-289.   | 5 <b>.</b> 5 | 36        |

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|----|--|------|-----------|
| 37 | Ligase Detection Reaction/Hybridization Assays Using Three-Dimensional Microfluidic Networks for the Detection of Low-Abundant DNA Point Mutations. Analytical Chemistry, 2005, 77, 3243-3255.                       | 6.5  | 77        |
| 38 | Fabrication of DNA microarrays onto polymer substrates using UV modification protocols with integration into microfluidic platforms for the sensing of low-abundant DNA point mutations. Methods, 2005, 37, 103-113. | 3.8  | 42        |
| 39 | High sensitivity EndoV mutation scanning through real-time ligase proofreading. Nucleic Acids Research, 2004, 32, e148-e148.   | 14.5 | 23        |
| 40 | Harmonized microarray/mutation scanning analysis of TP53 mutations in undissected colorectal tumors. Human Mutation, 2004, 24, 63-75.  | 2.5  | 20        |
| 41 | Microarrays Assembled in Microfluidic Chips Fabricated from Poly(methyl methacrylate) for the Detection of Low-Abundant DNA Mutations. Analytical Chemistry, 2003, 75, 1130-1140.                                    | 6.5  | 145       |
| 42 | Single nucleotide polymorphism seeking long term association with complex disease. Nucleic Acids Research, 2002, 30, 3295-3311.  | 14.5 | 157       |
| 43 | An endonuclease/ligase based mutation scanning method especially suited for analysis of neoplastic tissue. Oncogene, 2002, 21, 1909-1921.  | 5.9  | 40        |
| 44 | Universal DNA array detection of small insertions and deletions in BRCA1 and BRCA2. Nature Biotechnology, 2000, 18, 561-564.   | 17.5 | 200       |
| 45 | Biochemical properties of a high fidelity DNA ligase from Thermus species AK16D. Nucleic Acids Research, 1999, 27, 788-794.  | 14.5 | 89        |
| 46 | Multiplex PCR/LDR for detection of K-ras mutations in primary colon tumors. Oncogene, 1999, 18, 27-38.   | 5.9  | 166       |
| 47 | Ligase detection reaction for identification of low abundance mutations. Clinical Biochemistry, 1999, 32, 287-290.   | 1.9  | 50        |
| 48 | Universal DNA microarray method for multiplex detection of low abundance point mutations 1 1Edited by K. Yamamoto. Journal of Molecular Biology, 1999, 292, 251-262.   | 4.2  | 329       |
| 49 | Cloning, overexpression and nucleotide sequence of a thermostable DNA ligase-encoding gene. Gene, 1991, 109, 1-11.   | 2.2  | 104       |