# Georgina L Hold

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/1232025/publications.pdf

Version: 2024-02-01

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|---|---|---|---|
| 145 papers | 15,555 citations | 28274 55 h-index | 17592 121 g-index |
| 149 all docs | 149 docs citations | 149 times ranked | 21769 citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | The gut microbiota, bacterial metabolites and colorectal cancer. Nature Reviews Microbiology, 2014, 12, 661-672. | 28.6 | 2,007 |
| 2 | Fusobacterium nucleatum Potentiates Intestinal Tumorigenesis and Modulates the Tumor-Immune Microenvironment. Cell Host and Microbe, 2013, 14, 207-215. | 11.0 | 1,913 |
| 3 | The gut microbiota and host health: a new clinical frontier. Gut, 2016, 65, 330-339. | 12.1 | 1,719 |
| 4 | The microbiology of butyrate formation in the human colon. FEMS Microbiology Letters, 2002, 217, 133-139. | 1.8 | 1,105 |
| 5 | IBDâ€"what role do Proteobacteria play?. Nature Reviews Gastroenterology and Hepatology, 2012, 9, 219-230. | 17.8 | 587 |
| 6 | Growth requirements and fermentation products of Fusobacterium prausnitzii, and a proposal to reclassify it as Faecalibacterium prausnitzii gen. nov., comb. nov.. International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 2141-2146. | 1.7 | 479 |
| 7 | Assessment of microbial diversity in human colonic samples by 16S rDNA sequence analysis. FEMS Microbiology Ecology, 2002, 39, 33-39. | 2.7 | 324 |
| 8 | Inflammation and Cancer II. Role of chronic inflammation and cytokine gene polymorphisms in the pathogenesis of gastrointestinal malignancy. American Journal of Physiology - Renal Physiology, 2004, 286, G515-G520. | 3.4 | 302 |
| 9 | Role of the gut microbiota in inflammatory bowel disease pathogenesis: What have we learnt in the past 10 years?. World Journal of Gastroenterology, 2014, 20, 1192. | 3.3 | 293 |
| 10 | Roseburia intestinalis sp. nov., a novel saccharolytic, butyrate-producing bacterium from human faeces.. International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 1615-1620. | 1.7 | 285 |
| 11 | Oligonucleotide Probes That Detect Quantitatively Significant Groups of Butyrate-Producing Bacteria in Human Feces. Applied and Environmental Microbiology, 2003, 69, 4320-4324. | 3.1 | 284 |
| 12 | A Functional Polymorphism of Toll-Like Receptor 4 Gene Increases Risk of Gastric Carcinoma and Its Precursors. Gastroenterology, 2007, 132, 905-912. | 1.3 | 247 |
| 13 | Microbiota of De-Novo Pediatric IBD: Increased Faecalibacterium Prausnitzii and Reduced Bacterial Diversity in Crohn's But Not in Ulcerative Colitis. American Journal of Gastroenterology, 2012, 107, 1913-1922. | 0.4 | 245 |
| 14 | The Impact of Different DNA Extraction Kits and Laboratories upon the Assessment of Human Gut Microbiota Composition by 16S rRNA Gene Sequencing. PLoS ONE, 2014, 9, e88982. | 2.5 | 236 |
| 15 | Polymorphisms in Toll-like receptor genes and risk of cancer. Oncogene, 2008, 27, 244-252. | 5.9 | 218 |
| 16 | Vacuolating Cytotoxin and Variants in Atg16L1 That Disrupt Autophagy Promote Helicobacter pylori Infection in Humans. Gastroenterology, 2012, 142, 1160-1171. | 1.3 | 190 |
| 17 | The Inflammatory Microenvironment in Colorectal Neoplasia. PLoS ONE, 2011, 6, e15366. | 2.5 | 151 |
| 18 | Anaerostipes caccae gen. nov., sp. nov., a New Saccharolytic, Acetate-utilising, Butyrate-producing Bacterium from Human Faeces. Systematic and Applied Microbiology, 2002, 25, 46-51. | 2.8 | 150 |

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|---|---|---|---|
| 19 | Review article: the gut microbiome in inflammatory bowel diseaseâ€"avenues for microbial management. Alimentary Pharmacology and Therapeutics, 2018, 47, 26-42. | 3.7 | 147 |
| 20 | Efficacy of different faecal microbiota transplantation protocols for Clostridium difficile infection: A systematic review and metaâ€analysis. United European Gastroenterology Journal, 2018, 6, 1232-1244. | 3.8 | 137 |
| 21 | First-Pass Meconium Samples from Healthy Term Vaginally-Delivered Neonates: An Analysis of the Microbiota. PLoS ONE, 2015, 10, e0133320. | 2.5 | 134 |
| 22 | The gut virome: the â€˜missing linkâ€™ between gut bacteria and host immunity?. Therapeutic Advances in Gastroenterology, 2019, 12, 175628481983662. | 3.2 | 127 |
| 23 | Characterisation of bacterial communities associated with toxic and non-toxic dinoflagellates: Alexandrium spp. and Scrippsiella trochoidea. FEMS Microbiology Ecology, 2001, 37, 161-173. | 2.7 | 126 |
| 24 | Detection of Campylobacter concisus and Other Campylobacter Species in Colonic Biopsies from Adults with Ulcerative Colitis. PLoS ONE, 2011, 6, e21490. | 2.5 | 124 |
| 25 | Screening of colorectal cancer: present and future. Expert Review of Anticancer Therapy, 2017, 17, 1131-1146. | 2.4 | 123 |
| 26 | Growth requirements and fermentation products of Fusobacterium prausnitzii, and a proposal to reclassify it as Faecalibacterium prausnitzii gen. nov., comb. nov. International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 2141-2146. | 1.7 | 122 |
| 27 | Role of the polymorphic ILâ€1B , ILâ€1RN and TNFâ€A genes in distal gastric cancer in Mexico. International Journal of Cancer, 2005, 114, 237-241. | 5.1 | 117 |
| 28 | Genetic aspects of inflammation and cancer. Biochemical Journal, 2008, 410, 225-235. | 3.7 | 116 |
| 29 | Use of Restriction Fragment Length Polymorphism To Distinguish between Salmon Species. Journal of Agricultural and Food Chemistry, 2000, 48, 2184-2188. | 5.2 | 115 |
| 30 | Systematic review: gastric microbiota in health and disease. Alimentary Pharmacology and Therapeutics, 2020, 51, 582-602. | 3.7 | 113 |
| 31 | Sporadic colorectal cancer Ã¢Â€Â“ role of the commensal microbiota. FEMS Microbiology Letters, 2005, 244, 1-7. | 1.8 | 104 |
| 32 | The role of infection in the aetiology of inflammatory bowel disease. Journal of Gastroenterology, 2010, 45, 266-276. | 5.1 | 104 |
| 33 | Roseburia intestinalis sp. nov., a novel saccharolytic, butyrate-producing bacterium from human faeces. International Journal of Systematic and Evolutionary Microbiology, 2002, 52, 1615-1620. | 1.7 | 102 |
| 34 | Multi-omics differentially classify disease state and treatment outcome in pediatric Crohnâ€™s disease. Microbiome, 2018, 6, 13. | 11.1 | 94 |
| 35 | Increase in NF-ÎºB Binding Affinity of the Variant C Allele of the Toll-Like Receptor 9 â^'1237T/C Polymorphism Is Associated with Helicobacter pylori -Induced Gastric Disease. Infection and Immunity, 2010, 78, 1345-1352. | 2.2 | 93 |
| 36 | Anticancer effects of bioactive berry compounds. Phytochemistry Reviews, 2014, 13, 295-322. | 6.5 | 91 |

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|---|---|---|---|
| 37 | Two-stage Genome-wide Methylation Profiling in Childhood-onset CrohnÊ¼s Disease Implicates Epigenetic Alterations at the VMP1/MIR21 and HLA Loci. Inflammatory Bowel Diseases, 2014, 20, 1784-1793. | 1.9 | 84 |
| 38 | The Effect of Vitamin D on Intestinal Inflammation and Faecal Microbiota in Patients with Ulcerative Colitis. Journal of Crohn's and Colitis, 2018, 12, 963-972. | 1.3 | 78 |
| 39 | Comparison of paralytic shellfish toxin (PST) production by the dinoflagellates Alexandrium lusitanicum NEPCC 253 and Alexandrium tamarense NEPCC 407 in the presence and absence of bacteria. FEMS Microbiology Ecology, 2001, 36, 223-234. | 2.7 | 77 |
| 40 | A Comprehensive Evaluation of Colonic Mucosal Isolates of Sutterella wadsworthensis from Inflammatory Bowel Disease. PLoS ONE, 2011, 6, e27076. | 2.5 | 76 |
| 41 | The Role of Cytokine Gene Polymorphisms in Colorectal Cancer and Their Interaction with Aspirin Use in the Northeast of Scotland. Cancer Epidemiology Biomarkers and Prevention, 2005, 14, 1613-1618. | 2.5 | 75 |
| 42 | Enterohepatic Helicobacter in Ulcerative Colitis: Potential Pathogenic Entities?. PLoS ONE, 2011, 6, e17184. | 2.5 | 75 |
| 43 | Could Helicobacter organisms cause inflammatory bowel disease?. FEMS Immunology and Medical Microbiology, 2011, 61, 1-14. | 2.7 | 73 |
| 44 | Genetic variation in C20orf54, PLCE1 and MUC1 and the risk of upper gastrointestinal cancers in Caucasian populations. European Journal of Cancer Prevention, 2012, 21, 541-544. | 1.3 | 72 |
| 45 | Oral iron exacerbates colitis and influences the intestinal microbiome. PLoS ONE, 2018, 13, e0202460. | 2.5 | 71 |
| 46 | Identification of Hake Species (MerlucciusGenus) Using Sequencing and PCRâ^'RFLP Analysis of Mitochondrial DNA Control Region Sequences. Journal of Agricultural and Food Chemistry, 2001, 49, 5108-5114. | 5.2 | 70 |
| 47 | Genetic Variation in the Prostate Stem Cell Antigen Gene and Upper Gastrointestinal Cancer in White Individuals. Gastroenterology, 2011, 140, 435-441. | 1.3 | 70 |
| 48 | Oceanicaulis alexandrii gen. nov., sp. nov., a novel stalked bacterium isolated from a culture of the dinoflagellate Alexandrium tamarense (Lebour) Balech. International Journal of Systematic and Evolutionary Microbiology, 2003, 53, 1901-1906. | 1.7 | 69 |
| 49 | The fungal microbiota of de-novo paediatric inflammatory bowel disease. Microbes and Infection, 2015, 17, 304-310. | 1.9 | 67 |
| 50 | A standardised model for stool banking for faecal microbiota transplantation: a consensus report from a multidisciplinary UEG working group. United European Gastroenterology Journal, 2021, 9, 229-247. | 3.8 | 66 |
| 51 | The Microaerophilic Microbiota of De-Novo Paediatric Inflammatory Bowel Disease: The BISCUIT Study. PLoS ONE, 2013, 8, e58825. | 2.5 | 63 |
| 52 | Gastrointestinal Microbiota and Colon Cancer. Digestive Diseases, 2016, 34, 244-250. | 1.9 | 61 |
| 53 | Autism Spectrum Disorder and the Gut Microbiota in Children: A Systematic Review. Annals of Nutrition and Metabolism, 2020, 76, 16-29. | 1.9 | 61 |
| 54 | Development of a DNA-Based Method Aimed at Identifying the Fish Species Present in Food Products. Journal of Agricultural and Food Chemistry, 2001, 49, 1175-1179. | 5.2 | 60 |

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|---|---|---|---|
| 55 | Identification of Flatfish (Pleuronectiforme) Species Using DNA-Based Techniques. Journal of Agricultural and Food Chemistry, 2001, 49, 4562-4569. | 5.2 | 60 |
| 56 | Identification of Cephalopod Species (Ommastrephidae and Loliginidae) in Seafood Products by Forensically Informative Nucleotide Sequencing (FINS). Journal of Food Science, 2002, 67, 1672-1676. | 3.1 | 58 |
| 57 | Mucosal Microbiome in Patients with Recurrent Aphthous Stomatitis. Journal of Dental Research, 2015, 94, 87S-94S. | 5.2 | 57 |
| 58 | Volatile organic compounds emitted from faeces as a biomarker for colorectal cancer. Alimentary Pharmacology and Therapeutics, 2019, 49, 1005-1012. | 3.7 | 57 |
| 59 | Identification of gadoid fish species using DNA-based techniques. European Food Research and Technology, 2003, 217, 259-264. | 3.3 | 48 |
| 60 | CD14-159C/T and TLR9-1237T/C polymorphisms are not associated with gastric cancer risk in Caucasian populations. European Journal of Cancer Prevention, 2009, 18, 117-119. | 1.3 | 46 |
| 61 | Western lifestyle: a â€˜masterâ€™ manipulator of the intestinal microbiota?. Gut, 2014, 63, 5-6. | 12.1 | 46 |
| 62 | Development of a method for the quantification of haddock (Melanogrammus aeglefinus) in commercial products using real-time PCR. European Food Research and Technology, 2005, 220, 633-637. | 3.3 | 45 |
| 63 | Extending colonic mucosal microbiome analysisâ€”assessment of colonic lavage as a proxy for endoscopic colonic biopsies. Microbiome, 2016, 4, 61. | 11.1 | 43 |
| 64 | Inflammation associated ethanolamine facilitates infection by Crohn's disease-linked adherent-invasive Escherichia coli. EBioMedicine, 2019, 43, 325-332. | 6.1 | 42 |
| 65 | Propionic Acid Promotes the Virulent Phenotype of Crohnâ€™s Disease-Associated Adherent-Invasive Escherichia coli. Cell Reports, 2020, 30, 2297-2305.e5. | 6.4 | 42 |
| 66 | The gut microbiota, dietary extremes and exercise. Gut, 2014, 63, 1838-1839. | 12.1 | 41 |
| 67 | The Impact of NOD2 Variants on Fecal Microbiota in Crohnâ€™s Disease and Controls Without Gastrointestinal Disease. Inflammatory Bowel Diseases, 2018, 24, 583-592. | 1.9 | 40 |
| 68 | Systematic review: ileoanal pouch microbiota in health and disease. Alimentary Pharmacology and Therapeutics, 2018, 47, 466-477. | 3.7 | 38 |
| 69 | Validation of a PCR-RFLP based method for the identification of salmon species in food products. European Food Research and Technology, 2001, 212, 385-389. | 3.3 | 36 |
| 70 | Differentiation of raw or processed eel by PCR-based techniques: restriction fragment length polymorphism analysis (RFLP) and single strand conformation polymorphism analysis (SSCP). European Food Research and Technology, 2002, 214, 171-177. | 3.3 | 36 |
| 71 | Role of host genetics in fibrosis. Fibrogenesis and Tissue Repair, 2009, 2, 6. | 3.4 | 35 |
| 72 | A network metaâ€anal‌ysis of randomized controlled trials exploring the role of fecal microbiota transplantation in recurrent Clostridium difficile infection. United European Gastroenterology Journal, 2019, 7, 1051-1063. | 3.8 | 35 |

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|---|---|---|---|
| 73 | Systematic review with metaâ€analysis: dietary intake in adults with inflammatory bowel disease. Alimentary Pharmacology and Therapeutics, 2021, 54, 742-754. | 3.7 | 30 |
| 74 | The Role of Microbiota in Gastrointestinal Cancer and Cancer Treatment: Chance or Curse?. Cellular and Molecular Gastroenterology and Hepatology, 2022, 13, 857-874. | 4.5 | 30 |
| 75 | Microbiota organizationâ€"a key to understanding CRC development. Nature Reviews Gastroenterology and Hepatology, 2015, 12, 128-129. | 17.8 | 28 |
| 76 | Changing molecular epidemiology of rotavirus infection after introduction of monovalent rotavirus vaccination in Scotland. Vaccine, 2017, 35, 156-163. | 3.8 | 28 |
| 77 | COXâ€2 expression in sporadic colorectal adenomatous polyps is linked to adenoma characteristics. Histopathology, 2008, 52, 806-815. | 2.9 | 26 |
| 78 | Cytokine gene polymorphisms, cytokine levels and the risk of colorectal neoplasia in a screened population of Northeast Scotland. European Journal of Cancer Prevention, 2015, 24, 296-304. | 1.3 | 26 |
| 79 | Comparative genomics of Campylobacter concisus: Analysis of clinical strains reveals genome diversity and pathogenic potential. Emerging Microbes and Infections, 2018, 7, 1-17. | 6.5 | 25 |
| 80 | Expression of Genes for Drug Transporters in the Human Female Genital Tract and Modulatory Effect of Antiretroviral Drugs. PLoS ONE, 2015, 10, e0131405. | 2.5 | 25 |
| 81 | Impact of the Gastrointestinal Microbiome in Health and Disease: Co-evolution with the Host Immune System. Current Topics in Microbiology and Immunology, 2019, 421, 303-318. | 1.1 | 24 |
| 82 | The microbiology of butyrate formation in the human colon. FEMS Microbiology Letters, 2002, 217, 133-139. | 1.8 | 24 |
| 83 | The influence of early research experience in medical school on the decision to intercalate and future career in clinical academia: a questionnaire study. BMC Medical Education, 2017, 17, 245. | 2.4 | 23 |
| 84 | The Impact of NOD2 Genetic Variants on the Gut Mycobiota in Crohnâ€™s Disease Patients in Remission and in Individuals Without Gastrointestinal Inflammation. Journal of Crohn's and Colitis, 2021, 15, 800-812. | 1.3 | 22 |
| 85 | Review article: the future of microbiomeâ€based therapeutics. Alimentary Pharmacology and Therapeutics, 2022, 56, 192-208. | 3.7 | 21 |
| 86 | Isolation, growth on prebiotics and probiotic potential of novel bifidobacteria from pigs. Anaerobe, 2004, 10, 33-39. | 2.1 | 19 |
| 87 | Biochemical Characterization of Sinorhizobium meliloti Mutants Reveals Gene Products Involved in the Biosynthesis of the Unusual Lipid A Very Long-chain Fatty Acid. Journal of Biological Chemistry, 2011, 286, 17455-17466. | 3.4 | 19 |
| 88 | The TLR4 D299G and T399I SNPs Are Constitutively Active to Up-Regulate Expression of Trif-Dependent Genes. PLoS ONE, 2014, 9, e111460. | 2.5 | 19 |
| 89 | Adaptive response of neonatal sepsis-derived Group B Streptococcus to bilirubin. Scientific Reports, 2018, 8, 6470. | 3.3 | 18 |
| 90 | Expression of neutrophil gelatinase-associated lipocalin in colorectal neoplastic progression: a marker of malignant potential?. British Journal of Cancer, 2013, 108, 2537-2541. | 6.4 | 17 |

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|---|---|---|---|
| 91 | Development of real-time PCR assays for the detection of Atlantic cod (Gadus morhua), Atlantic salmon (Salmo salar) and European plaice (Pleuronectes platessa) in complex food samples. European Food Research and Technology, 2012, 234, 127-136. | 3.3 | 16 |
| 92 | Drug transporter gene expression in human colorectal tissue and cell lines: modulation with antiretrovirals for microbicide optimization. Journal of Antimicrobial Chemotherapy, 2016, 71, 372-386. | 3.0 | 16 |
| 93 | Possible association between a genetic polymorphism at 8q24 and risk of upper gastrointestinal cancer. European Journal of Cancer Prevention, 2011, 20, 54-57. | 1.3 | 15 |
| 94 | Innate Immune Sensors and Gastrointestinal Bacterial Infections. Clinical and Developmental Immunology, 2011, 2011, 1-11. | 3.3 | 14 |
| 95 | Gut microbial biofilm composition and organisation holds the key to CRC. Nature Reviews Gastroenterology and Hepatology, 2019, 16, 329-330. | 17.8 | 14 |
| 96 | Redefining intestinal immunity with single-cell transcriptomics. Mucosal Immunology, 2022, 15, 531-541. | 6.0 | 12 |
| 97 | The Other Helicobacters. Helicobacter, 2011, 16, 70-75. | 3.5 | 11 |
| 98 | The Molecular Basis of Lipid A and Toll-Like Receptor 4 Interactions. , 2011, , 371-387. | | 9 |
| 99 | Lack of association between the rs2294008 polymorphism in the prostate stem cell antigen gene and colorectal neoplasia: a case-control and immunohistochemical study. BMC Research Notes, 2012, 5, 371. | 1.4 | 9 |
| 100 | Comparative genomics and genome biology of Campylobacter showae. Emerging Microbes and Infections, 2019, 8, 827-840. | 6.5 | 8 |
| 101 | Long-Term Iron Deficiency and Dietary Iron Excess Exacerbate Acute Dextran Sodium Sulphate-Induced Colitis and Are Associated with Significant Dysbiosis. International Journal of Molecular Sciences, 2021, 22, 3646. | 4.1 | 8 |
| 102 | Faecal microbiota transplantation as a treatment for inflammatory bowel disease: a national survey of adult and paediatric gastroenterologists in the UK. Frontline Gastroenterology, 2018, 9, 250-255. | 1.8 | 7 |
| 103 | Characterisation of bacterial communities associated with toxic and non-toxic dinoflagellates: Alexandrium spp. and Scrippsiella trochoidea. FEMS Microbiology Ecology, 2001, 37, 161-173. | 2.7 | 7 |
| 104 | Refining a Protocol for Faecal Microbiota Engraftment in Animal Models After Successful Antibiotic-Induced Gut Decontamination. Frontiers in Medicine, 2022, 9, 770017. | 2.6 | 7 |
| 105 | A functional toll-like receptor 4 polymorphism increases risk of H. pylori-induced premalignant changes. Gastroenterology, 2003, 124, A19-A20. | 1.3 | 6 |
| 106 | Transporters for Antiretroviral Drugs in Colorectal CD4+ T Cells and Circulating Î±4Î²7 Integrin CD4+ T Cells: Implications for HIV Microbicides. Molecular Pharmaceutics, 2016, 13, 3334-3340. | 4.6 | 6 |
| 107 | Molecular Analysis of the Microbiome in Colorectal Cancer. Methods in Molecular Biology, 2018, 1765, 139-153. | 0.9 | 6 |
| 108 | Australia IBD Microbiome (AIM) Study: protocol for a multicentre longitudinal prospective cohort study. BMJ Open, 2021, 11, e042493. | 1.9 | 6 |

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|---|---|---|---|
| 109 | Inflammatory bowel disease and the gut microbiota. Proceedings of the Nutrition Society, 2021, , 1-11. | 1.0 | 6 |
| 110 | Other Helicobacters and the gastric microbiome. Helicobacter, 2018, 23, e12521. | 3.5 | 5 |
| 111 | Colonic mucosal bacterial diversity of de novo extensive paediatric ulcerative colitis by next-generation sequencing. Gut, 2011, 60, A146-A147. | 12.1 | 4 |
| 112 | MicroRNAs in gastrointestinal malignancy. European Journal of Cancer Prevention, 2014, 23, 540-549. | 1.3 | 4 |
| 113 | Novel Campylobacter concisus lipooligosaccharide is a determinant of inflammatory potential and virulence. Journal of Lipid Research, 2018, 59, 1893-1905. | 4.2 | 4 |
| 114 | Gut Mucosal Microbiome Signatures of Colorectal Cancer Differ According to BMI Status. Frontiers in Medicine, 2021, 8, 800566. | 2.6 | 4 |
| 115 | W1208 Variable Detection of Entero-Hepatic Helicobacter Species in Colonic Mucosal Pinch Biopsies By Different Molecular Techniques. Gastroenterology, 2008, 134, A-655. | 1.3 | 3 |
| 116 | The role of microaerophilic colonic mucosal bacteria in de novo paediatric inflammatory bowel disease. Gut, 2011, 60, A147-A147. | 12.1 | 3 |
| 117 | Assessment of microbial diversity in human colonic samples by 16S rDNA sequence analysis. FEMS Microbiology Ecology, 2002, 39, 33-39. | 2.7 | 3 |
| 118 | Microbiome Understanding in Maternity Study (MUMS), an Australian prospective longitudinal cohort study of maternal and infant microbiota: study protocol. BMJ Open, 2020, 10, e040189. | 1.9 | 3 |
| 119 | Bilirubin Has Anti-Bacterial Properties Against Gram-Positive Bacteria: A Potential Benefit of Physiological Jaundice?. Gastroenterology, 2011, 140, S-941. | 1.3 | 2 |
| 120 | Biopsy Sampling in Upper Gastrointestinal Endoscopy: A Survey from 10 Tertiary Referral Centres Across Europe. Digestive Diseases, 2021, 39, 179-189. | 1.9 | 2 |
| 121 | Mo1584 Genetic Variation in C20orf54, PLCE1 and MUC1 and Risk of Upper Gastrointestinal Cancers in Caucasian Populations. Gastroenterology, 2012, 142, S-634. | 1.3 | 1 |
| 122 | Sa1972 Assessment of Bacterial Diversity in Colorectal Adenomatous Polyps. Gastroenterology, 2013, 144, S-348. | 1.3 | 1 |
| 123 | Expression of Drug Transporters in Cervicovaginal Cell Lines and Modulatory Effect of Candidate Anti-retroviral Microbicides. AIDS Research and Human Retroviruses, 2014, 30, A201-A201. | 1.1 | 1 |
| 124 | Characterisation of Drug Transporter Gene Expression in Colorectal Tissue and Cell Lines: Induction with Anti-retrovirals for Microbicide Optimization. AIDS Research and Human Retroviruses, 2014, 30, A201-A201. | 1.1 | 1 |
| 125 | Gastric cancer relatives have a high prevalence of IL-18 and TGF-B1 proinflammatory genotypes. Gastroenterology, 2003, 124, A6. | 1.3 | 0 |
| 126 | Role of interleukin-1 beta and other potential genetic markers as indicators of gastric cancer risk. , 2003, , 215-223. | | 0 |

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|---|---|---|---|
| 127 | PP-012â€…Cytokine gene polymorphisms, cytokine levels and risk of colorectal neoplasia in the screened population of northeast Scotland. Gut, 2010, 59, A44.3-A45. | 12.1 | 0 |
| 128 | OC-048â€…Impact of the TLR4 Asp299gly polymorphism on induction of the inflammatory response followingHelicobacter pyloriinfection. Gut, 2010, 59, A20.1-A20. | 12.1 | 0 |
| 129 | OC-053â€…CanHelicobacter pylorilipopolysaccharide lipid a composition affect its ability to induce an inflammatory response through Toll-like receptor 4: Abstract OC-053. Gut, 2010, 59, A22.1-A22. | 12.1 | 0 |
| 130 | S1640 Can Helicobacter pylori Lipopolysaccharide Lipid a Composition Affect Its Ability to Induce an Inflammatory Response Through Toll Like Receptor 4. Gastroenterology, 2010, 138, S-244. | 1.3 | 0 |
| 131 | T2009 Assessment of Novel Genetic Polymorphisms and Risk of Upper Gastrointestinal Carcinoma. Gastroenterology, 2010, 138, S-612. | 1.3 | 0 |
| 132 | W1738 Role of TLR4 in Carcinogenesis and Tumor Progression of Colorectal Cancer. Gastroenterology, 2010, 138, S-730. | 1.3 | 0 |
| 133 | S1641 Impact of the TLR4 Asp299Gly Polymorphism on Induction of the Inflammatory Response Following H. pylori Infection. Gastroenterology, 2010, 138, S-244. | 1.3 | 0 |
| 134 | T2012 Cytokine Gene Polymorphisms, Cytokine Levels and Risk of Colorectal Neoplasia in the Screened Population of Northeast Scotland. Gastroenterology, 2010, 138, S-613. | 1.3 | 0 |
| 135 | Detection of Campylobacter Concisus in Colonic Biopsies From Adult Patients With Ulcerative Colitis. Gastroenterology, 2011, 140, S-268. | 1.3 | 0 |
| 136 | Bacterial Diversity of the Colonic Microbiota in De-Novo Extensive Paediatric Ulcerative Colitis by Next-Generation Sequencing. Gastroenterology, 2011, 140, S-196. | 1.3 | 0 |
| 137 | The Role of the Microaerophilic Colonic Microbiota in De-Novo Paediatric Inflammatory Bowel Disease. Gastroenterology, 2011, 140, S-512. | 1.3 | 0 |
| 138 | Detection of campylobacter concisus in colonic biopsies from adult patients with ulcerative colitis. Gut, 2011, 60, A211-A211. | 12.1 | 0 |
| 139 | 946 Genome-Wide Analysis of DNA Methylation in Low-Grade Colorectal Adenomas and Normal Colonic Mucosa. Gastroenterology, 2014, 146, S-165. | 1.3 | 0 |
| 140 | Influence of Host Gene Polymorphisms on Development of Gastroduodenal Diseases. , 2016, , 339-362. | | 0 |
| 141 | The Effect of Lactulose on the Faecal Microbiota of Patients with Minimal Hepatic Encephalopathy. Gastroenterology, 2017, 152, S1049. | 1.3 | 0 |
| 142 | PWE-039â€…FMT as a treatment for IBD: a national survey of gastroenterologists in the UK. , 2018, , . | | 0 |
| 143 | Changes in Gut Microbiota Due to Gastrointestinal Surgery. , 2021, , 139-139. | | 0 |
| 144 | Next-generation sequencing as a clinical laboratory tool for describing different microbiotas: an urgent need for future paediatric practice. Archives of Disease in Childhood, 2021, 106, 1035-1035. | 1.9 | 0 |

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|---|---|---|---|
| 145 | Comparison of paralytic shellfish toxin (PST) production by the dinoflagellates Alexandrium lusitanicum NEPCC 253 and Alexandrium tamarense NEPCC 407 in the presence and absence of bacteria. FEMS Microbiology Ecology, 2001, 36, 223-234. | 2.7 | 0 |