CRISPR/Cas-Based Genome Editing for Human Gut Cor

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Citation Report

#	Article	IF	CITATIONS
1	Engineering the human gut commensal Bacteroides thetaiotaomicron with synthetic biology. Current Opinion in Chemical Biology, 2022, 70, 102178.	2.8	5
2	Methods of DNA introduction for the engineering of commensal microbes. Engineering Microbiology, 2022, 2, 100048.	2.2	6
3	Novel technologies to characterize and engineer the microbiome in inflammatory bowel disease. Gut Microbes, 2022, 14, .	4.3	4
4	Intestinal Engineered Probiotics as Living Therapeutics: Chassis Selection, Colonization Enhancement, Gene Circuit Design, and Biocontainment. ACS Synthetic Biology, 2022, 11, 3134-3153.	1.9	9
6	Play the plug: How bacteria modify recognition by host receptors?. Frontiers in Microbiology, 0, 13, .	1.5	0
7	Microbiome and Human Health: Current Understanding, Engineering, and Enabling Technologies. Chemical Reviews, 2023, 123, 31-72.	23.0	54
8	Construction and characterization of a genome-scale ordered mutant collection of Bacteroides thetaiotaomicron. BMC Biology, 2022, 20, .	1.7	8
9	Revealing the hidden heights of microbial metabolites on reproductive physiology. , 2023, , 217-248.		0
10	Highly efficient CRISPRâ€mediated base editing for the gut <i>Bacteroides</i> spp. with pnCasBSâ€CBE. Biotechnology Journal, 2023, 18, .	1.8	2
23	Novel Techniques and Models for Studying the Role of the Gut Microbiota in Drug Metabolism. European Journal of Drug Metabolism and Pharmacokinetics, 2024, 49, 131-147.	0.6	0