

Identifier	Author(s)	Title	Journal or Publication	Volume	Issue	Publisher	Year	Pages	DOI
Benjamin 2005	Benjamin, Ivor J., and Michael D. Schneider.	Learning from failure: congestive heart failure in the postgenomic age	<i>Journal of clinical investigation</i>	115	3		2005	495-499	https://doi.org/10.1172/jci24477
Bui 2009	Bui, Minh, and Zhongchi Liu.	Simple allele-discriminating PCR for cost-effective and rapid genotyping and mapping	<i>Plant Methods</i>	5	1		2009	1	https://doi.org/10.1186/1746-4811-5-1
Bui 2009b	Bui, Minh	LEUNIG, LEUNIG HOMOLOG, and SEUSS are transcriptional co-repressors that regulate flower development, mucilage secretion, and pathogen resistance				University of Maryland, College Park	2009		https://www.proquest.com/openview/b3480855618427130165ce76e01386e2/1?pq-origsite=gscholar&cbl=18750
Dahl 2005	Dahl, Fredrik, et al.	Multiplex amplification enabled by selective circularization of large sets of	<i>Nucleic acids research</i>	33	8		2005	e71-e71	https://doi.org/10.1093/nar/gni070

		genomic DNA fragments							
Dahl 2005b	Dahl, Fredrik	Selector Technology: For Multiplex DNA Analysis	<i>Diss. Acta Universitatis Upsaliensis</i>				2005		
Hall 2007	Hall, C. Dennis, et al.	Organophosphorus Chemistry: Volume 35				Royal Society of Chemistry	2007		-
Howell 2006	Howell, W. Mathias, et al.	Detection Techniques for Single Nucleotide Polymorphisms	<i>Nucleic Acid Testing for Human Disease</i>			CRC Press	2006	139-169	https://doi.org/10.1201/9781420016888-6
Jin 2011	Jin, Tao, et al.	Molecular basis for resistance to tribenuron in shepherd's purse (<i>Capsella bursa-pastoris</i> (L.) Medik.).	<i>Pesticide Biochemistry and Physiology</i>	100	2		2011	160-164	https://doi.org/10.1016/j.pestbp.2011.03.005
Kaur 2018	Kaur, Rajdeep	Allele specific primer designing for SNPs of ACHE gene related to oxidative stress	<i>JETIR</i>	5			2018		-

Kim 2005	Kim, Namshin, Seokmin Shin, and Sanghyuk Lee	ECgene: genome-based EST clustering and gene modeling for alternative splicing	<i>Genome research</i>	15	4		2005	566-576	https://doi.org/10.1101/gr.3030405
Li 2006	Li, Li, et al.	SNP genotyping by multiplex amplification and microarrays assay for forensic application	<i>Forensic science international</i>	162	1-3		2006	74-79	https://doi.org/10.1016/j.forsciint.2006.06.010
Liu 2012	Liu, Jing, et al.	An improved allele-specific PCR primer design method for SNP marker analysis and its application	<i>Plant methods</i>	8	1		2012	34	https://doi.org/10.1186/1746-4811-8-34
Liu 2024	Liu, Ying, et al.	Multiplex asymmetric PCR by combining the amplification refractory mutation system with the homo-tag-assisted nondimer system	<i>Analytical Chemistry</i>	96	22		2024	9200-9208	https://doi.org/10.1021/acs.analchem.4c01322
Loakes 2007	Loakes, David	Nucleotides and nucleic acids; oligo-and polynucleotides	<i>Organophosphorus Chemistry</i>	36			2007	232-312	https://doi.org/10.1039/b603852m

Strömqvist Meuzelaar 2007	Meuzelaar, Linda Strömqvist, et al.	MegaPlex PCR: a strategy for multiplex amplification	<i>Nature methods</i>	4	10		2007	835- 837	https://doi.org/10.1038/nmeth1091
Morin 2007	Morin, Phillip A., and Melissa McCarthy	Highly accurate SNP genotyping from historical and low-quality samples	<i>Molecular Ecology Notes</i>	7	6		2007	937- 946	https://doi.org/10.1111/j.1471-8286.2007.01804.x
Palka- Santini 2009	Palka-Santini, Maria, et al.	Large scale multiplex PCR improves pathogen detection by DNA microarrays	<i>BMC microbiology</i>	9	1		2009	1	https://doi.org/10.1186/1471-2180-9-1
Ronaghi 2005	Ronaghi, Mostafa, and Taimour Langae	Single nucleotide polymorphisms: discovery, detection and analysis	<i>Personalized Medicine</i>	2	2		2005	111- 125	https://doi.org/10.1517/17410541.2.2.111
Stenberg 2006	Stenberg, Johan	Software Tools for Design of Reagents for Multiplex Genetic Analyses	<i>Diss. Acta Universitatis Upsaliensis</i>				2006		-
Van Baren 2006	Van Baren, Marijke J., and Michael R. Brent	Iterative gene prediction and pseudogene removal improves genome annotation	<i>Genome research</i>	16	5		2006	678- 685	https://doi.org/10.1101/gr.4766206

Vaniusheva 2005	Vaniusheva, O.V., et al.	Development of a Technique for Identification of Alleles for Single-Nucleotide Polymorphous Markers based on an Allele-specific PCR using and Automatic DNA Analyzer	<i>Biotechnology in Russia</i>	4			2005	28-39	-
Wang 2006	Wang, Weipeng, Kun-yi Ni, and Guohua Zhou	Multiplex single nucleotide polymorphism genotyping by adapter ligation-mediated allele-specific amplification	<i>Analytical Biochemistry</i>	355	2		2006	240-248	https://doi.org/10.1016/j.ab.2006.04.022
Wang 2006b	Wang, Weipeng, Kunyi Ni, and Guohua Zhou	Association of IL1B polymorphisms with gastric cancer in a Chinese population	<i>Clinical biochemistry</i>	40	3-4		2006	218-225	https://doi.org/10.1016/j.clinbiochem.2006.10.018
Wu 2005	Wu, Thomas D., and Colin K. Watanabe	GMAP: a genomic mapping and alignment program for mRNA and EST sequences	<i>Bioinformatics</i>	21	9		2005	1859-1875	https://doi.org/10.1093/bioinformatics/bti310

Zakharova 2008	Zakharova, Natalia V., et al.	Simultaneous qualitative and quantitative identification of multiple targets by polymerase chain reaction on microarray	<i>arXiv:0811.1 263</i>				2008		https://doi.org/10.48550/arxiv. 0811.1263
Zhou 2016	Zhou, Guohua, et al.	Improvement of Pyrosequencing to Allow Multiplex SNP Typing in a Pyrogram	<i>Advances and Clinical Practice in Pyrosequenci ng</i>			Springer	2016	129- 143	https://doi.org/10.1007/978-1- 4939-3308-2_11
Zhou 2005	Zhou, Guo- Hua, et al.	Multiplex SNP typing by bioluminometric assay coupled with terminator incorporation (BATI)	<i>Nucleic Acids Research</i>	33	15		2005	e133- e133	https://doi.org/10.1093/nar/gn i132