

Transposon vectors containing non-antibiotic resistance  
and stable chromosomal insertion of foreign genes in gr

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Mini-Tn5 transposon derivatives for insertion mutagenesis, promoter probing, and chromosomal insertion of cloned DNA in gram-negative eubacteria. <i>Journal of Bacteriology</i> , 1990, 172, 6568-6572.	1.0	1,465
2	Mutagenesis of <i>Brucella abortus</i> : comparative efficiency of three transposon delivery systems. <i>Microbial Pathogenesis</i> , 1991, 11, 443-446.	1.3	30
3	Antibiotic resistance in the bacteria of a remote upland river catchment. <i>Letters in Applied Microbiology</i> , 1991, 13, 145-149.	1.0	14
4	[19] Genetic techniques in <i>Rhizobium meliloti</i> . <i>Methods in Enzymology</i> , 1991, 204, 398-418.	0.4	122
5	An upstream XylR- and IHF-induced nucleoprotein complex regulates the sigma 54-dependent Pu promoter of TOL plasmid.. <i>EMBO Journal</i> , 1991, 10, 1159-1167.	3.5	150
6	Identification and characterization of a gene responsible for inhibiting propagation of methylated DNA sequences in <i>mcrA mcrB1 Escherichia coli</i> strains. <i>Journal of Bacteriology</i> , 1991, 173, 4707-4716.	1.0	30
7	Molecular approaches for understanding biological control mechanisms in bacteria: Studies of the interaction of <i>Enterobacter cloacae</i> with <i>Pythium ultimum</i> . <i>Canadian Journal of Plant Pathology</i> , 1992, 14, 106-114.	0.8	19
8	Chromosome transfer in <i>Rhodobacter sphaeroides</i> : Hfr formation and genetic evidence for two unique circular chromosomes. <i>Journal of Bacteriology</i> , 1992, 174, 1135-1145.	1.0	68
9	Evidence for proliferation of <i>Enterobacter cloacae</i> on carbohydrates in cucumber and pea spermosphere. <i>Canadian Journal of Microbiology</i> , 1992, 38, 1128-1134.	0.8	30
10	Expression of <i>Bordetella pertussis</i> filamentous hemagglutinin in <i>Escherichia coli</i> using a two cistron system. <i>Microbial Pathogenesis</i> , 1992, 12, 383-389.	1.3	2
11	An improved suicide vector for construction of chromosomal insertion mutations in bacteria. <i>Gene</i> , 1992, 118, 145-146.	1.0	284
12	Mercury biotransformations and their potential for remediation of mercury contamination. <i>Biodegradation</i> , 1992, 3, 147-159.	1.5	49
13	A general system to integrate lacZ fusions into the chromosomes of gram-negative eubacteria: regulation of the Pm promoter of the TOL plasmid studied with all controlling elements in monocopy. <i>Molecular Genetics and Genomics</i> , 1992, 233, 293-301.	2.4	285
14	Cloning of a DNA fragment involved in pigment production in <i>Streptomyces avermitilis</i> . <i>FEMS Microbiology Letters</i> , 1992, 91, 9-13.	0.7	7
15	Genetic engineering strategies for environmental applications. <i>Current Opinion in Biotechnology</i> , 1992, 3, 227-231.	3.3	23
16	How do non-differentiating bacteria adapt to starvation?. <i>Antonie Van Leeuwenhoek</i> , 1993, 63, 333-341.	0.7	166
17	Analysis of the DNA damage-mediated induction of <i>Pseudomonas putida</i> and <i>Pseudomonas aeruginosa</i> <i>lexA</i> genes. <i>FEMS Microbiology Letters</i> , 1993, 110, 65-70.	0.7	14
18	A DNA module encoding <i>bph</i> genes for the degradation of polychlorinated biphenyls (PCBs). <i>FEMS Microbiology Letters</i> , 1993, 113, 149-154.	0.7	32

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19	Microbial retention of mercury from waste streams in a laboratory column containing merAgene bacteria. <i>FEMS Microbiology Reviews</i> , 1993, 11, 145-152.	3.9	55
20	Engineering of alkyl- and haloaromatic-responsive gene expression with mini-transposons containing regulated promoters of biodegradative pathways of <i>Pseudomonas</i> . <i>Gene</i> , 1993, 130, 41-46.	1.0	113
21	A T7 RNA polymerase-based system for the construction of <i>Pseudomonas</i> strains with phenotypes dependent on TOL-meta pathway effectors. <i>Gene</i> , 1993, 134, 103-106.	1.0	36
22	Analysis of <i>Pseudomonas</i> gene products using lacIq/P <sub>trp</sub> -lac plasmids and transposons that confer conditional phenotypes. <i>Gene</i> , 1993, 123, 17-24.	1.0	429
23	Three different 2,3-dihydroxybiphenyl-1,2-dioxygenase genes in the gram-positive polychlorobiphenyl-degrading bacterium <i>Rhodococcus globerulus</i> P6. <i>Journal of Bacteriology</i> , 1993, 175, 4631-4640.	1.0	163
24	Transposon mutagenesis in <i>Acinetobacter calcoaceticus</i> RAG-1. <i>Journal of Bacteriology</i> , 1993, 175, 1838-1840.	1.0	20
25	Isolation and characterization of <i>Bordetella bronchiseptica</i> mutants deficient in siderophore activity. <i>Journal of Bacteriology</i> , 1993, 175, 1144-1152.	1.0	43
26	Transposon mutagenesis in <i>Actinobacillus pleuropneumoniae</i> with a Tn10 derivative. <i>Journal of Bacteriology</i> , 1993, 175, 5717-5722.	1.0	62
27	Early and late responses of TOL promoters to pathway inducers: identification of postexponential promoters in <i>Pseudomonas putida</i> with lacZ-tet bicistronic reporters. <i>Journal of Bacteriology</i> , 1993, 175, 6902-6907.	1.0	92
28	BIOCHEMICAL AND GENETIC ANALYSIS OF SIDEROPHORES PRODUCED BY PLANT-ASSOCIATED PSEUDOMONAS AND ERWINIA SPECIES. , 1993, , 27-73.		5
30	Detection of Introduced Bacteria in the Rhizosphere Using Marker Genes and DNA Probes. , 0, , 29-47.		7
31	Genetic evidence that the XylS regulator of the <i>Pseudomonas</i> TOL meta operon controls the P <sub>m</sub> promoter through weak DNA-protein interactions. <i>Journal of Bacteriology</i> , 1994, 176, 3171-3176.	1.0	44
32	Chromosomal gene capture mediated by the <i>Pseudomonas putida</i> TOL catabolic plasmid. <i>Journal of Bacteriology</i> , 1994, 176, 4635-4641.	1.0	46
33	Genetic Strategies to Engineer Expression Systems Responsive to Relevant Environmental Signals. , 0, , 91-101.		0
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35	Molecular cloning of bacterial DNA in vivo using a transposable R6K ori and a P1vir phage. <i>Journal of Bacteriology</i> , 1994, 176, 1188-1191.	1.0	3
36	Transformation of <i>Vibrio vulnificus</i> by electroporation. <i>Current Microbiology</i> , 1994, 28, 289-291.	1.0	8
37	The organization of the P <sub>m</sub> promoter of the TOL plasmid reflects the structure of its cognate activator protein XylS. <i>Molecular Genetics and Genomics</i> , 1994, 244, 596-605.	2.4	27

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38	Designing microbial systems for gene expression in the field. <i>Trends in Biotechnology</i> , 1994, 12, 365-371.	4.9	52
39	Controlled expression of click beetle luciferase using a bacterial operator-repressor system. <i>FEMS Microbiology Letters</i> , 1994, 121, 11-18.	0.7	5
40	The Behavior of Bacteria Designed for Biodegradation. <i>Nature Biotechnology</i> , 1994, 12, 1349-1356.	9.4	76
41	Universal barrier to lateral spread of specific genes among microorganisms. <i>Molecular Microbiology</i> , 1994, 13, 855-861.	1.2	75
42	Analysis of the multimer resolution system encoded by the parCBA operon of broad-host-range plasmid RP4. <i>Molecular Microbiology</i> , 1994, 12, 131-141.	1.2	91
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44	Catabolism of aromatics in <i>Pseudomonas putida</i> U. Formal evidence that phenylacetic acid and 4-hydroxyphenylacetic acid are catabolized by two unrelated pathways. <i>FEBS Journal</i> , 1994, 221, 375-381.	0.2	34
45	Role of colonization in biocontrol: studies with <i>Enterobacter cloacae</i> . <i>Plant Science</i> , 1994, 101, 83-89.	1.7	22
46	Transformation of <i>Acinetobacter calcoaceticus</i> RAG-1 by electroporation. <i>Canadian Journal of Microbiology</i> , 1994, 40, 233-236.	0.8	13
47	Use of the rep technique for allele replacement to construct new <i>Escherichia coli</i> hosts for maintenance of R6K $\lambda$ origin plasmids at different copy numbers. <i>Gene</i> , 1994, 138, 1-7.	1.0	184
48	<i>Escherichia coli</i> genome targeting I. Cre-Zox-mediated in vitro generation of ori $\lambda$ <sup>+</sup> plasmids and their in vivo chromosomal integration and retrieval. <i>Gene</i> , 1994, 150, 51-56.	1.0	59
49	[31] Analysis and construction of stable phenotypes in gram-negative bacteria with Tn5- and Tn10-derived minitransposons. <i>Methods in Enzymology</i> , 1994, 235, 386-405.	0.4	852
50	Designing Microorganisms for the Treatment of Toxic Wastes. <i>Annual Review of Microbiology</i> , 1994, 48, 525-557.	2.9	174
51	Transposon Tn5 mutagenesis of <i>Actinobacillus actinomycetemcomitans</i> via conjugation. <i>Oral Microbiology and Immunology</i> , 1994, 9, 290-296.	2.8	15
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53	[33] Identification of bacterial cell-surface virulence determinants with TnpH $\alpha$ . <i>Methods in Enzymology</i> , 1994, 235, 426-448.	0.4	18
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55	Identification of a locus required for the regulation of bvg-repressed genes in <i>Bordetella pertussis</i> . <i>Journal of Bacteriology</i> , 1995, 177, 2727-2736.	1.0	66

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56	Site-specific deletions of chromosomally located DNA segments with the multimer resolution system of broad-host-range plasmid RP4. <i>Journal of Bacteriology</i> , 1995, 177, 52-58.	1.0	122
57	A carbon starvation survival gene of <i>Pseudomonas putida</i> is regulated by sigma 54. <i>Journal of Bacteriology</i> , 1995, 177, 1850-1859.	1.0	51
58	Site directed chromosomal marking of a fluorescent pseudomonad isolated from the phytosphere of sugar beet; stability and potential for marker gene transfer.. <i>Molecular Ecology</i> , 1995, 4, 755-764.	2.0	127
59	A lipase of <i>Aeromonas hydrophila</i> showing nonhemolytic phospholipase C activity. <i>Current Microbiology</i> , 1995, 31, 28-33.	1.0	20
60	Regulation of citrate-dependent iron transport of <i>Escherichia coli</i> : FecR is required for transcription activation by Fecl. <i>Molecular Microbiology</i> , 1995, 15, 119-132.	1.2	108
61	Induction of phospholipase- and flagellar synthesis in <i>Serratia liquefaciens</i> is controlled by expression of the flagellar master operon flhD. <i>Molecular Microbiology</i> , 1995, 15, 445-454.	1.2	96
62	Expansion and deletion of CTG repeats from human disease genes are determined by the direction of replication in <i>E. coli</i> . <i>Nature Genetics</i> , 1995, 10, 213-218.	9.4	356
63	Use of ans-triazine nitrogen source to select for and isolate a recombinant chlorobenzoate-degrading <i>Pseudomonas</i> . <i>FEMS Microbiology Letters</i> , 1995, 133, 47-52.	0.7	7
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65	Use of conjugative and thermosensitive cloning vectors for transposon delivery to <i>Mycobacterium smegmatis</i> . <i>FEMS Microbiology Letters</i> , 1995, 127, 35-39.	0.7	14
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69	Isolation of a carbon starvation regulatory mutant in a marine <i>Vibrio</i> strain. <i>Journal of Bacteriology</i> , 1995, 177, 6978-6982.	1.0	21
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73	Specific binding of the replication protein of plasmid pPS10 to direct and inverted repeats is mediated by an HTH motif. <i>Nucleic Acids Research</i> , 1995, 23, 5048-5054.	6.5	23

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75	Mini-Tn10 transposon derivatives for insertion mutagenesis and gene delivery into the chromosome of Gram-negative bacteria. <i>Gene</i> , 1995, 160, 59-62.	1.0	104
76	An improved TnMax mini-transposon system suitable for sequencing shuttle mutagenesis and gene fusions. <i>Gene</i> , 1995, 167, 53-57.	1.0	68
77	Molecular techniques for the study of rhizobial ecology in the field. <i>Soil Biology and Biochemistry</i> , 1995, 27, 501-514.	4.2	62
78	A versatile most probable number system to quantify lacZY-marked pseudomonads present at low cell numbers in the rhizosphere. <i>Letters in Applied Microbiology</i> , 1995, 20, 220-224.	1.0	2
79	Transcription of repA, the Gene of the Initiation Protein of the Pseudomonas Plasmid pPS10, is Autoregulated by Interactions of the RepA Protein at a Symmetrical Operator. <i>Journal of Molecular Biology</i> , 1995, 247, 211-223.	2.0	29
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81	Rapid Detection of Mutagens by Induction of Luciferase-Bearing Prophage in Escherichia coli. <i>Environmental Science &amp; Technology</i> , 1996, 30, 2478-2483.	4.6	6
82	Recombinant Toluene-4-monooxygenase: Catalytic and Massbauer Studies of the Purified Diiron and Rieske Components of a Four-Protein Complex. <i>Biochemistry</i> , 1996, 35, 9106-9119.	1.2	180
83	Amino acids as reduced carbon sources for Enterobacter cloacae during colonization of the spermospheres of crop plants. <i>Soil Biology and Biochemistry</i> , 1996, 28, 1015-1020.	4.2	20
84	Bacterial plasmid conjugation on semi-solid surfaces monitored with the green fluorescent protein (GFP) from Aequorea victoria as a marker. <i>Gene</i> , 1996, 173, 59-65.	1.0	115
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87	Relationship Between Escherichia coli Growth and Deletions of CTG-CAG Triplet Repeats in Plasmids. <i>Journal of Molecular Biology</i> , 1996, 264, 82-96.	2.0	65
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91	Evaluation of the biological containment system based on the Escherichia coli gef gene in Pseudomonas aeruginosa W51D. <i>Applied Microbiology and Biotechnology</i> , 1996, 46, 549-553.	1.7	7

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93	Designing bacteria for the degradation of nitro- and chloroaromatic pollutants. <i>Die Naturwissenschaften</i> , 1996, 83, 201-213.	0.6	12
94	Development of techniques for the genetic manipulation of the gliding bacteria <i>Lysobacter enzymogenes</i> and <i>Lysobacter brunescens</i> . <i>Canadian Journal of Microbiology</i> , 1996, 42, 896-902.	0.8	13
95	Ferric rhizoferrin uptake into <i>Morganella morganii</i> : characterization of genes involved in the uptake of a polyhydroxycarboxylate siderophore. <i>Journal of Bacteriology</i> , 1996, 178, 496-504.	1.0	37
96	Characterization of an OprL null mutant of <i>Pseudomonas putida</i> . <i>Journal of Bacteriology</i> , 1996, 178, 5836-5840.	1.0	29
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100	Characterization and expression of the plasmid-borne <i>bedD</i> gene from <i>Pseudomonas putida</i> ML2, which codes for a NAD <sup>+</sup> -dependent cis-benzene dihydrodiol dehydrogenase. <i>Journal of Bacteriology</i> , 1996, 178, 5592-5601.	1.0	24
101	Development of techniques for the genetic manipulation of the gliding bacterium <i>Cytophaga johnsonae</i> . <i>Journal of Bacteriology</i> , 1996, 178, 583-590.	1.0	113
102	Tn10 insertional mutagenesis in <i>Pasteurella multocida</i> . <i>Veterinary Microbiology</i> , 1996, 50, 143-148.	0.8	11
103	Efficiency of MucAB and <i>Escherichia coli</i> UmuDC proteins in quinolone and UV mutagenesis in <i>Salmonella typhimurium</i> : effect of MucA and UmuD processing. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 1996, 349, 201-208.	0.4	4
104	Mini-TnhlyAs: a new tool for the construction of secreted fusion proteins. <i>Molecular Genetics and Genomics</i> , 1996, 252, 266-274.	2.4	6
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108	Chemotactic motility is required for invasion of the host by the fish pathogen <i>Vibrio anguillarum</i> . <i>Molecular Microbiology</i> , 1996, 19, 625-637.	1.2	160
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111	Construction of a modified mini-Tn5lacZYnon-antibiotic marker cassette: ecological evaluation of alacZYmarked <i>Pseudomonas</i> strain in the sugarbeet rhizosphere. <i>FEMS Microbiology Letters</i> , 1996, 135, 251-257.	0.7	15
112	Linker insertion analysis of the FimH adhesin of type 1 fimbriae in an <i>Escherichia coli</i> fimH-null background. <i>FEMS Microbiology Letters</i> , 1996, 137, 257-263.	0.7	56
113	Characterisation of carbon dioxide-inducible genes of the marine bacterium, <i>pseudomonas</i> sp. S91. <i>FEMS Microbiology Letters</i> , 1996, 140, 37-42.	0.7	19
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115	Construction of GFP vectors for use in Gram-negative bacteria other than <i>Escherichia coli</i> . <i>FEMS Microbiology Letters</i> , 1996, 145, 87-94.	0.7	124
116	Tn5-induced <i>Xenorhabdus bovienii</i> lecithinase mutants demonstrate reduced virulence for <i>Galleria mellonella</i> larvae. <i>Journal of Applied Bacteriology</i> , 1996, 80, 411-417.	1.1	7
117	An <i>Escherichia coli</i> hemolysin transport system-based vector for the export of polypeptides: Export of shiga-like toxin IIeB subunit by <i>Salmonella typhimurium</i> aroA. <i>Nature Biotechnology</i> , 1996, 14, 765-769.	9.4	75
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120	Modulation of gene expression through chromosomal positioning in <i>Escherichia coli</i> . <i>Microbiology (United Kingdom)</i> , 1997, 143, 2071-2078.	0.7	118
121	Effector Specificity Mutants of the Transcriptional Activator NahR of Naphthalene Degrading <i>Pseudomonas</i> Define Protein Sites Involved in Binding of Aromatic Inducers. <i>Journal of Biological Chemistry</i> , 1997, 272, 3986-3992.	1.6	87
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123	Maintenance of broad-host-range incompatibility group P and group Q plasmids and transposition of Tn5 in <i>Bartonella henselae</i> following conjugal plasmid transfer from <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 1997, 179, 538-540.	1.0	262
124	Structural and functional analysis of the phosphoenolpyruvate carboxylase gene from the purple nonsulfur bacterium <i>Rhodospseudomonas palustris</i> No. 7. <i>Journal of Bacteriology</i> , 1997, 179, 4942-4945.	1.0	20
125	Construction and characterization of genetically-marked bivalent anti- <i>Shigella dysenteriae</i> 1 and anti- <i>Shigella flexneri</i> Y live vaccine candidates. <i>Microbial Pathogenesis</i> , 1997, 22, 363-376.	1.3	10
126	Identification and molecular characterization of a 27 kDa <i>Shigella flexneri</i> invasion plasmid antigen, IpaJ. <i>Microbial Pathogenesis</i> , 1997, 23, 357-369.	1.3	12
127	Measurement of the competitiveness index of <i>Rhizobium tropici</i> strain CIAT899 derivatives marked with the <i>gusA</i> gene. <i>Soil Biology and Biochemistry</i> , 1997, 29, 1099-1110.	4.2	33



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128	Chromosomal insertion of the entire <i>Escherichia coli</i> lactose operon, into two strains of <i>Pseudomonas</i> , using a modified mini-Tn5 delivery system. <i>Gene</i> , 1997, 186, 167-173.	1.0	51
129	Green fluorescent protein-based reporter systems for genetic analysis of bacteria including monocopy applications. <i>Gene</i> , 1997, 196, 69-74.	1.0	130
130	Insertional mutagenesis by a modified in vitro Ty1 transposition system. <i>Gene</i> , 1997, 198, 27-35.	1.0	27
131	Genetic characterization of insertion sequence ISJP4 on plasmid pJP4 from <i>Ralstonia eutropha</i> JMP134. <i>Gene</i> , 1997, 202, 103-114.	1.0	33
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1544	The Role of Intercellular Signaling in the Regulation of Bacterial Adaptive Proliferation. <i>International Journal of Molecular Sciences</i> , 2023, 24, 7266.	1.8	1
1546	Evolution and the Role of SXT-Related Integrative Conjugative Elements in Multidrug-Resistant <i>Vibrio cholerae</i> . , 2023, , 1-17.		0

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1552	Evolution and the Role of SXT-Related Integrative Conjugative Elements in Multidrug-Resistant <i>Vibrio cholerae</i> . , 2023, , 465-481.		0
1554	Identification and Characterization of Some Genes, Enzymes, and Metabolic Intermediates Belonging to the Bile Acid Aerobic Catabolic Pathway from <i>Pseudomonas</i> . <i>Methods in Molecular Biology</i> , 2023, , 51-83.	0.4	0