

Mervyn J Bibb

List of Publications by Year in descending order

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7421
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#	ARTICLE	IF	CITATIONS
1	<i>Streptomyces venezuelae</i> NRRL B-65442: genome sequence of a model strain used to study morphological differentiation in filamentous actinobacteria. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2021, , .	1.4	14
2	Genome editing reveals that pSCL4 is required for chromosome linearity in <i>Streptomyces clavuligerus</i> . <i>Microbial Genomics</i> , 2021, 7, .	1.0	2
3	<i>In Situ</i> Activation and Heterologous Production of a Cryptic Lantibiotic from an African Plant Ant-Derived <i>Saccharopolyspora</i> Species. <i>Applied and Environmental Microbiology</i> , 2020, 86, .	1.4	22
4	New Molecular Tools for Regulation and Improvement of A40926 Glycopeptide Antibiotic Production in <i>Nonomuraea gerenzanensis</i> ATCC 39727. <i>Frontiers in Microbiology</i> , 2020, 11, 8.	1.5	19
5	Heterologous Expression of a Cryptic Gene Cluster from <i>Streptomyces leeuwenhoekii</i> C34 Yields a Novel Lasso Peptide, LEEPPTIN. <i>Applied and Environmental Microbiology</i> , 2019, 85, .	1.4	20
6	The "gifted" actinomycete <i>Streptomyces leeuwenhoekii</i> . <i>Antonie Van Leeuwenhoek</i> , 2018, 111, 1433-1448. 0.7	0.7	24
7	Structures of DPAGT1 Explain Glycosylation Disease Mechanisms and Advance TB Antibiotic Design. <i>Cell</i> , 2018, 175, 1045-1058.e16.	13.5	67
8	Analysis of the Tunicamycin Biosynthetic Gene Cluster of <i>Streptomyces chartreusis</i> Reveals New Insights into Tunicamycin Production and Immunity. <i>Antimicrobial Agents and Chemotherapy</i> , 2018, 62, .	1.4	19
9	A novel mechanism of immunity controls the onset of cinnamycin biosynthesis in <i>Streptomyces cinnamoneus</i> DSM 40646. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2017, 44, 563-572.	1.4	21
10	Watasemycin biosynthesis in <i>Streptomyces venezuelae</i> : thiazoline C-methylation by a type B radical-SAM methylase homologue. <i>Chemical Science</i> , 2017, 8, 2823-2831.	3.7	42
11	Next Generation Sequencing of Actinobacteria for the Discovery of Novel Natural Products. <i>Marine Drugs</i> , 2016, 14, 78.	2.2	118
12	Discovery of Unusual Biaryl Polyketides by Activation of a Silent <i>Streptomyces venezuelae</i> Biosynthetic Gene Cluster. <i>ChemBioChem</i> , 2016, 17, 2189-2198.	1.3	50
13	Two Master Switch Regulators Trigger A40926 Biosynthesis in <i>Nonomuraea</i> sp. Strain ATCC 39727. <i>Journal of Bacteriology</i> , 2015, 197, 2536-2544.	1.0	36
14	A <i>relA</i> -dependent regulatory cascade for autoinduction of microbisporicin production in <i>Micromonospora corallina</i> . <i>Molecular Microbiology</i> , 2015, 97, 502-514.	1.2	28
15	A <i>Streptomyces coelicolor</i> host for the heterologous expression of Type III polyketide synthase genes. <i>Microbial Cell Factories</i> , 2015, 14, 145.	1.9	34
16	The <i>Streptomyces leeuwenhoekii</i> genome: de novo sequencing and assembly in single contigs of the chromosome, circular plasmid pSLE1 and linear plasmid pSLE2. <i>BMC Genomics</i> , 2015, 16, 485.	1.2	61
17	Identification and Heterologous Expression of the Chaxamycin Biosynthesis Gene Cluster from <i>Streptomyces leeuwenhoekii</i> . <i>Applied and Environmental Microbiology</i> , 2015, 81, 5820-5831.	1.4	38
18	New Insights into Chloramphenicol Biosynthesis in <i>Streptomyces venezuelae</i> ATCC 10712. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 7441-7450.	1.4	74

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19	Heterologous expression of natural product biosynthetic gene clusters in <i>Streptomyces coelicolor</i> : from genome mining to manipulation of biosynthetic pathways. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2014, 41, 425-431.	1.4	122
20	Relationship between Glycopeptide Production and Resistance in the Actinomycete <i>Nonomuraea</i> sp. ATCC 39727. <i>Antimicrobial Agents and Chemotherapy</i> , 2014, 58, 5191-5201.	1.4	24
21	Use of the Meganuclease I-SceI of <i>Saccharomyces cerevisiae</i> to select for gene deletions in actinomycetes. <i>Scientific Reports</i> , 2014, 4, 7100.	1.6	57
22	Ribosomally synthesized and post-translationally modified peptide natural products: overview and recommendations for a universal nomenclature. <i>Natural Product Reports</i> , 2013, 30, 108-160.	5.2	1,692
23	High resolution crystal structure of Sco5413, a widespread actinomycete MarR family transcriptional regulator of unknown function. <i>Proteins: Structure, Function and Bioinformatics</i> , 2013, 81, 176-182.	1.5	7
24	Understanding and manipulating antibiotic production in actinomycetes. <i>Biochemical Society Transactions</i> , 2013, 41, 1355-1364.	1.6	59
25	Cloning and Analysis of the Planosporicin Lantibiotic Biosynthetic Gene Cluster of <i>Planomonospora alba</i> . <i>Journal of Bacteriology</i> , 2013, 195, 2309-2321.	1.0	42
26	Investigation of DNA sequence recognition by a streptomycete MarR family transcriptional regulator through surface plasmon resonance and X-ray crystallography. <i>Nucleic Acids Research</i> , 2013, 41, 7009-7022.	6.5	39
27	The antibiotic planosporicin coordinates its own production in the actinomycete <i>Planomonospora alba</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E2500-9.	3.3	78
28	Synthetic RNA Silencing of Actinorhodin Biosynthesis in <i>Streptomyces coelicolor</i> A3(2). <i>PLoS ONE</i> , 2013, 8, e67509.	1.1	18
29	Phage P1-Derived Artificial Chromosomes Facilitate Heterologous Expression of the FK506 Gene Cluster. <i>PLoS ONE</i> , 2013, 8, e69319.	1.1	80
30	Structure and biosynthesis of the unusual polyketide alkaloid coelimycin P1, a metabolic product of the cpk gene cluster of <i>Streptomyces coelicolor</i> M145. <i>Chemical Science</i> , 2012, 3, 2716.	3.7	152
31	<i>Streptomyces coelicolor</i> as an Expression Host for Heterologous Gene Clusters. <i>Methods in Enzymology</i> , 2012, 517, 279-300.	0.4	43
32	Biosynthesis of the tunicamycin antibiotics proceeds via unique exo-glycal intermediates. <i>Nature Chemistry</i> , 2012, 4, 539-546.	6.6	79
33	Posttranslational $\hat{\text{I}}^2$ -methylation and macrolactamidation in the biosynthesis of the bottromycin complex of ribosomal peptide antibiotics. <i>Chemical Science</i> , 2012, 3, 3522.	3.7	67
34	Genome Sequence of the Abyssomicin- and Proximicin-Producing Marine Actinomycete <i>Verrucosipora maris</i> AB-18-032. <i>Journal of Bacteriology</i> , 2011, 193, 3391-3392.	1.0	24
35	Engineering <i>Streptomyces coelicolor</i> for heterologous expression of secondary metabolite gene clusters. <i>Microbial Biotechnology</i> , 2011, 4, 207-215.	2.0	439
36	Genome-wide analysis of the role of GlnR in <i>Streptomyces venezuelae</i> provides new insights into global nitrogen regulation in actinomycetes. <i>BMC Genomics</i> , 2011, 12, 175.	1.2	127

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37	Abyssomicin Biosynthesis: Formation of an Unusual Polyketide, Antibiotic Feeding Studies and Genetic Analysis. <i>ChemBioChem</i> , 2011, 12, 1401-1410.	1.3	66
38	Biosynthesis and Regulation of Grisemycin, a New Member of the Linaridin Family of Ribosomally Synthesized Peptides Produced by <i>Streptomyces griseus</i> IFO 13350. <i>Journal of Bacteriology</i> , 2011, 193, 2510-2516.	1.0	63
39	Feed-Forward Regulation of Microbisporicin Biosynthesis in <i>Microbispora corallina</i> . <i>Journal of Bacteriology</i> , 2011, 193, 3064-3071.	1.0	39
40	A system for the targeted amplification of bacterial gene clusters multiplies antibiotic yield in <i>Streptomyces coelicolor</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 16020-16025.	3.3	66
41	ZouA, a Putative Relaxase, Is Essential for DNA Amplification in <i>Streptomyces kanamyceticus</i> . <i>Journal of Bacteriology</i> , 2011, 193, 1815-1822.	1.0	14
42	Draft Genome Sequence of <i>Streptomyces</i> Strain S4, a Symbiont of the Leaf-Cutting Ant <i>Acromyrmex octospinosus</i> . <i>Journal of Bacteriology</i> , 2011, 193, 4270-4271.	1.0	27
43	Methods for the genetic manipulation of <i>Nonomuraea</i> sp. ATCC 39727. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2010, 37, 1097-1103.	1.4	26
44	Heterologous expression of the biosynthetic gene clusters of coumermycin A ₁ , clorobiocin and caprazamycins in genetically modified <i>Streptomyces coelicolor</i> strains. <i>Biopolymers</i> , 2010, 93, 823-832.	1.2	39
45	A mixed community of actinomycetes produce multiple antibiotics for the fungus farming ant <i>Acromyrmex octospinosus</i> . <i>BMC Biology</i> , 2010, 8, 109.	1.7	211
46	Analysis of the phosphoproteome of the multicellular bacterium <i>Streptomyces coelicolor</i> A3(2) by protein/peptide fractionation, phosphopeptide enrichment and high accuracy mass spectrometry. <i>Proteomics</i> , 2010, 10, 2486-2497.	1.3	68
47	Novel Mechanism of Glycopeptide Resistance in the A40926 Producer <i>Nonomuraea</i> sp. ATCC 39727. <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 2465-2472.	1.4	43
48	Deletion of a regulatory gene within the <i>cpk</i> gene cluster reveals novel antibacterial activity in <i>Streptomyces coelicolor</i> A3(2). <i>Microbiology (United Kingdom)</i> , 2010, 156, 2343-2353.	0.7	143
49	Microbisporicin gene cluster reveals unusual features of lantibiotic biosynthesis in actinomycetes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 13461-13466.	3.3	141
50	Genome mining and genetic analysis of cypemycin biosynthesis reveal an unusual class of posttranslationally modified peptides. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 16297-16302.	3.3	123
51	Discovery of Unique Lanthionine Synthetases Reveals New Mechanistic and Evolutionary Insights. <i>PLoS Biology</i> , 2010, 8, e1000339.	2.6	186
52	Dissecting tunicamycin biosynthesis by genome mining: cloning and heterologous expression of a minimal gene cluster. <i>Chemical Science</i> , 2010, 1, 581.	3.7	58
53	The role of <i>absC</i> , a novel regulatory gene for secondary metabolism, in zinc-dependent antibiotic production in <i>Streptomyces coelicolor</i> A3(2). <i>Molecular Microbiology</i> , 2009, 74, 1427-1444.	1.2	63
54	Chapter 4 Analyzing the Regulation of Antibiotic Production in Streptomycetes. <i>Methods in Enzymology</i> , 2009, 458, 93-116.	0.4	36

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55	A complex role for the γ -butyrolactone SCB1 in regulating antibiotic production in <i>Streptomyces coelicolor</i> A3(2). <i>Molecular Microbiology</i> , 2008, 41, 1015-1028.	1.2	211
56	Manipulating and understanding antibiotic production in <i>Streptomyces coelicolor</i> A3(2) with decoy oligonucleotides. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 1020-1025.	3.3	25
57	Elongation Factor Tu3 (EF-Tu3) from the Kirromycin Producer <i>Streptomyces ramocissimus</i> Is Resistant to Three Classes of EF-Tu-Specific Inhibitors. <i>Journal of Bacteriology</i> , 2007, 189, 3581-3590.	1.0	15
58	A New Piece of an Old Jigsaw: Glucose Kinase Is Activated Posttranslationally in a Glucose Transport-Dependent Manner in <i>Streptomyces coelicolor</i> A3(2). <i>Journal of Molecular Microbiology and Biotechnology</i> , 2007, 12, 67-74.	1.0	57
59	In vivo DNase I sensitivity of the <i>Streptomyces coelicolor</i> chromosome correlates with gene expression: implications for bacterial chromosome structure. <i>Nucleic Acids Research</i> , 2006, 34, 5395-5401.	6.5	9
60	Amplification of the entire kanamycin biosynthetic gene cluster during empirical strain improvement of <i>Streptomyces kanamyceticus</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 9661-9666.	3.3	95
61	EshA Accentuates ppGpp Accumulation and Is Conditionally Required for Antibiotic Production in <i>Streptomyces coelicolor</i> A3(2). <i>Journal of Bacteriology</i> , 2006, 188, 4952-4961.	1.0	42
62	A bacterial hormone (the SCB1) directly controls the expression of a pathway-specific regulatory gene in the cryptic type I polyketide biosynthetic gene cluster of <i>Streptomyces coelicolor</i> . <i>Molecular Microbiology</i> , 2005, 56, 465-479.	1.2	146
63	Regulation of secondary metabolism in streptomycetes. <i>Current Opinion in Microbiology</i> , 2005, 8, 208-215.	2.3	672
64	A rare leucine codon in <i>adpA</i> is implicated in the morphological defect of <i>bldA</i> mutants of <i>Streptomyces coelicolor</i> . <i>Molecular Microbiology</i> , 2003, 50, 475-486.	1.2	114
65	Engineering of Primary Carbon Metabolism for Improved Antibiotic Production in <i>Streptomyces lividans</i> . <i>Applied and Environmental Microbiology</i> , 2002, 68, 4731-4739.	1.4	79
66	Induction of ppGpp synthesis in <i>Streptomyces coelicolor</i> A3(2) grown under conditions of nutritional sufficiency elicits <i>actII-ORF4</i> transcription and actinorhodin biosynthesis. <i>Molecular Microbiology</i> , 2001, 39, 136-144.	1.2	76
67	Analysis of the prodiginine biosynthesis gene cluster of <i>Streptomyces coelicolor</i> A3(2): new mechanisms for chain initiation and termination in modular multienzymes. <i>Chemistry and Biology</i> , 2001, 8, 817-829.	6.2	164
68	Functional Analysis of <i>relA</i> and <i>rshA</i> , Two <i>relA/spoT</i> Homologues of <i>Streptomyces coelicolor</i> A3(2). <i>Journal of Bacteriology</i> , 2001, 183, 3488-3498.	1.0	71
69	A single amino acid substitution in region 1.2 of the principal sigma factor of <i>Streptomyces coelicolor</i> A3(2) results in pleiotropic loss of antibiotic production. <i>Molecular Microbiology</i> , 2000, 37, 995-1004.	1.2	45
70	Glucose kinase of <i>Streptomyces coelicolor</i> A3(2): large-scale purification and biochemical analysis. <i>Antonie Van Leeuwenhoek</i> , 2000, 78, 253-261.	0.7	45
71	Purification and Structural Determination of SCB1, a γ -Butyrolactone That Elicits Antibiotic Production in <i>Streptomyces coelicolor</i> A3(2). <i>Journal of Biological Chemistry</i> , 2000, 275, 11010-11016.	1.6	154
72	Analysis of <i>aptsH</i> homologue from <i>Streptomyces coelicolor</i> A3(2). <i>FEMS Microbiology Letters</i> , 1999, 177, 279-288.	0.7	24

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73	The Linear Plasmid SCP1 of <i>Streptomyces coelicolor</i> A3(2) Possesses a Centrally Located Replication Origin and Shows Significant Homology to the Transposon Tn4811. <i>Plasmid</i> , 1999, 42, 174-185.	0.4	34
74	Actinorhodin and undecylprodigiosin production in wild-type and mutant strains of <i>Streptomyces coelicolor</i> A3(2) grown in continuous culture. <i>FEMS Microbiology Letters</i> , 1998, 168, 221-226.	0.7	60
75	A novel family of proteins that regulates antibiotic production in streptomycetes appears to contain an OmpR-like DNA-binding fold. <i>Molecular Microbiology</i> , 1997, 25, 1181-1184.	1.2	287
76	Substrate induction and glucose repression of maltose utilization by <i>Streptomyces coelicolor</i> A3(2) is controlled by malR, a member of the lacI-galR family of regulatory genes. <i>Molecular Microbiology</i> , 1997, 23, 537-549.	1.2	95
77	A novel plasmid vector that uses the glucose kinase gene (glkA) for the positive selection of stable gene disruptants in <i>Streptomyces</i> . <i>Gene</i> , 1996, 182, 229-230.	1.0	21
78	Cloning, characterization and disruption of a (p)ppGpp synthetase gene (relA) of <i>Streptomyces coelicolor</i> A3(2). <i>Molecular Microbiology</i> , 1996, 19, 357-368.	1.2	83
79	afsR is a pleiotropic but conditionally required regulatory gene for antibiotic production in <i>Streptomyces coelicolor</i> A3(2). <i>Molecular Microbiology</i> , 1996, 21, 385-396.	1.2	202
80	Construction of thiostrepton-inducible, high-copy-number expression vectors for use in <i>Streptomyces</i> spp.. <i>Gene</i> , 1995, 166, 133-137.	1.0	133
81	Glucose repression in <i>Streptomyces coelicolor</i> A3(2): a likely regulatory role for glucose kinase. <i>Molecular Genetics and Genomics</i> , 1994, 244, 135-143.	2.4	106
82	The mRNA for the 23S rRNA methylase encoded by the ermE gene of <i>Saccharopolyspora erythraea</i> is translated in the absence of a conventional ribosome-binding site. <i>Molecular Microbiology</i> , 1994, 14, 533-545.	1.2	178
83	The Stringent Response, ppGpp and Antibiotic Production in <i>Streptomyces coelicolor</i> A3(2).. <i>Nihon Hosenkin Gakkai Shi = Actinomycetologica</i> , 1994, 8, 1-16.	0.3	32
84	Stationary-phase production of the antibiotic actinorhodin in <i>Streptomyces coelicolor</i> A3(2) is transcriptionally regulated. <i>Molecular Microbiology</i> , 1993, 7, 837-845.	1.2	194
85	Derivatives of pUC18 that have BglII sites flanking a modified multiple cloning site and that retain the ability to identify recombinant clones by visual screening of <i>Escherichia coli</i> colonies. <i>Gene</i> , 1993, 124, 133-134.	1.0	234
86	Codon usage in the G+C-rich <i>Streptomyces</i> genome. <i>Gene</i> , 1992, 113, 55-65.	1.0	417
87	A simple and reliable turbidimetric and kinetic assay for alpha-amylase that is readily applied to culture supernatants and cell extracts. <i>Journal of Industrial Microbiology</i> , 1990, 5, 295-301.	0.9	10
88	Tandem promoters, tsrp1 and tsrp2, direct transcription of the thiostrepton resistance gene (tsr) of <i>Streptomyces azureus</i> : Transcriptional initiation from tsrp2 occurs after deletion of the 35 region. <i>Molecular Genetics and Genomics</i> , 1990, 221, 339-346.	2.4	15
89	<i>Streptomyces</i> promoter-probe plasmids that utilise the xylE gene of <i>Pseudomonas putida</i> . <i>Nucleic Acids Research</i> , 1990, 18, 1077-1077.	6.5	77
90	A cassette containing the barg gene of <i>Streptomyces hygrosopicus</i> : a selectable marker for plant transformation. <i>Nucleic Acids Research</i> , 1990, 18, 1062-1062.	6.5	78

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91	Transcriptional analysis of the repressor gene of the temperate <i>Streptomyces</i> phage ϕ C31. <i>Gene</i> , 1989, 85, 275-282.	1.0	16
92	Organisation of the ribosomal RNA genes in <i>Streptomyces coelicolor</i> A3(2). <i>Molecular Genetics and Genomics</i> , 1988, 211, 191-196.	2.4	48
93	The repressor gene (c) of the <i>Streptomyces</i> temperate phage ϕ C31: Nucleotide sequence, analysis and functional cloning. <i>Molecular Genetics and Genomics</i> , 1988, 213, 269-277.	2.4	38
94	At least three different RNA polymerase holoenzymes direct transcription of the agarase gene (dagA) of <i>Streptomyces coelicolor</i> A3(2). <i>Cell</i> , 1988, 52, 599-607.	13.5	153
95	Cloning, characterisation and regulation of an α -amylase gene from <i>Streptomyces venezuelae</i> . <i>Gene</i> , 1988, 74, 321-334.	1.0	73
96	The nucleotide sequence of a 16S rRNA gene from <i>Streptomyces coelicolor</i> A3(2). <i>Nucleic Acids Research</i> , 1987, 15, 7176-7176.	6.5	52
97	[9] Plasmid and phage vectors for gene cloning and analysis in <i>Streptomyces</i> . <i>Methods in Enzymology</i> , 1987, 153, 116-166.	0.4	74
98	The agarase gene (dagA) of <i>Streptomyces coelicolor</i> A3(2): nucleotide sequence and transcriptional analysis. <i>Molecular Genetics and Genomics</i> , 1987, 209, 101-109.	2.4	157
99	Cloning and analysis of the promoter region of the erythromycin-resistance gene (ermE) of <i>Streptomyces erythraeus</i> . <i>Gene</i> , 1986, 41, E357-E368.	1.0	261
100	Construction and characterisation of a series of multi-copy promoter-probe plasmid vectors for <i>Streptomyces</i> using the aminoglycoside phosphotransferase gene from Tn5 as indicator. <i>Molecular Genetics and Genomics</i> , 1986, 203, 468-478.	2.4	405
101	Nucleotide sequences encoding and promoting expression of three antibiotic resistance genes indigenous to <i>Streptomyces</i> . <i>Molecular Genetics and Genomics</i> , 1985, 199, 26-36.	2.4	164
102	The nucleotide sequence of the tyrosinase gene from <i>Streptomyces antibioticus</i> and characterization of the gene product. <i>Gene</i> , 1985, 37, 101-110.	1.0	177
103	Cloning and analysis of the promoter region of the erythromycin resistance gene (ermE) of <i>Streptomyces erythraeus</i> . <i>Gene</i> , 1985, 38, 215-226.	1.0	312
104	Dissecting the <i>Streptomyces</i> genome. <i>Biochemical Society Transactions</i> , 1984, 12, 584-586.	1.6	4
105	Cloning <i>Streptomyces</i> genes for antibiotic production. <i>Trends in Biotechnology</i> , 1983, 1, 42-48.	4.9	43
106	Gene expression in <i>Streptomyces</i> : Construction and application of promoter-probe plasmid vectors in <i>Streptomyces lividans</i> . <i>Molecular Genetics and Genomics</i> , 1982, 187, 265-277.	2.4	322
107	Excision of chromosomal DNA sequences from <i>Streptomyces coelicolor</i> forms a novel family of plasmids detectable in <i>Streptomyces lividans</i> . <i>Molecular Genetics and Genomics</i> , 1981, 184, 230-240.	2.4	135
108	A DNA cloning system for interspecies gene transfer in antibiotic-producing <i>Streptomyces</i> . <i>Nature</i> , 1980, 284, 526-531.	13.7	171

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109	Transformation of plasmid DNA into <i>Streptomyces</i> at high frequency. <i>Nature</i> , 1978, 274, 398-400.	13.7	306
110	Physical and genetical characterisation of a second sex factor, SCP2, for <i>Streptomyces coelicolor</i> A3(2). <i>Molecular Genetics and Genomics</i> , 1977, 154, 155-166.	2.4	222