

# CITATION REPORT

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**ICEberg: a web-based resource for integrative and conjugative elements found in Bacteria**

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#	Paper	IF	Citations
192	On the Power of Penetrating into Space—The Telescopes of William Herschel. <b>1976</b> , 7, 75-108		32
191	Complete genome sequence of <i>Klebsiella pneumoniae</i> subsp. <i>pneumoniae</i> HS11286, a multidrug-resistant strain isolated from human sputum. <b>2012</b> , 194, 1841-2		112
190	Characterization of a <i>Streptococcus suis</i> tet(O/W/32/O)-carrying element transferable to major streptococcal pathogens. <b>2012</b> , 56, 4697-702		50
189	The <i>Bacillus subtilis</i> conjugative transposon ICEBs1 mobilizes plasmids lacking dedicated mobilization functions. <b>2012</b> , 194, 3165-72		63
188	Data mining for microbiologists. <b>2012</b> , 39, 27-79		7
187	The phn Island: A New Genomic Island Encoding Catabolism of Polynuclear Aromatic Hydrocarbons. <i>Frontiers in Microbiology</i> , <b>2012</b> , 3, 125	5.7	36
186	CRISPR-Cas systems target a diverse collection of invasive mobile genetic elements in human microbiomes. <b>2013</b> , 14, R40		45
185	Type II Toxin-Antitoxin Loci: Phylogeny. <b>2013</b> , 239-247		
184	Mobile Genetic Elements (MGEs) Carrying Catabolic Genes. <b>2013</b> , 167-214		9
183	Generation and analysis of an ICE R391 deletion library identifies genes involved in the element encoded UV-inducible cell-sensitising function. <i>FEMS Microbiology Letters</i> , <b>2013</b> , 342, 45-53	2.9	12
182	A conserved helicase processivity factor is needed for conjugation and replication of an integrative and conjugative element. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003198	6	32
181	Interspecies mobilization of an ermT-carrying plasmid of <i>Streptococcus dysgalactiae</i> subsp. <i>equisimilis</i> by a coresident ICE of the ICESa2603 family. <i>Journal of Antimicrobial Chemotherapy</i> , <b>2013</b> , 68, 23-6	5.1	49
180	SecReT4: a web-based bacterial type IV secretion system resource. <i>Nucleic Acids Research</i> , <b>2013</b> , 41, D660-61	5.1	61
179	Functional validation of putative toxin-antitoxin genes from the Gram-positive pathogen <i>Streptococcus pneumoniae</i> : phd-doc is the fourth bona-fide operon. <i>Frontiers in Microbiology</i> , <b>2014</b> , 5, 677	5.7	28
178	The use of genome wide association methods to investigate pathogenicity, population structure and serovar in <i>Haemophilus parasuis</i> . <b>2014</b> , 15, 1179		29
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176	The genome of the intracellular bacterium of the coastal bivalve, <i>Solemya velum</i> : a blueprint for thriving in and out of symbiosis. <b>2014</b> , 15, 924		22

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172	Characterization of the core and accessory genomes of <i>Pseudomonas aeruginosa</i> using bioinformatic tools Spine and AGEnt. <b>2014</b> , 15, 737		105
171	An overview of the domestication and impact of the <i>Salmonella mobilome</i> . <b>2014</b> , 40, 63-75		9
170	Complete genome sequence of <i>Ornithobacterium rhinotracheale</i> strain ORT-UMN 88. <b>2014</b> , 9, 16		8
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