## Abraham L Sonenshein

# List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/6756219/abraham-l-sonenshein-publications-by-year.pdf

Version: 2024-04-10

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

198
papers

9,099
citations

54
h-index
g-index

200
ext. papers

6.1
avg, IF
L-index

| #   | Paper  | IF                   | Citations |
|-----|--|----------------------|-----------|
| 198 | In vivo commensal control of Clostridioides difficile virulence. <i>Cell Host and Microbe</i> , <b>2021</b> , 29, 1693-170   | 0&. <del>g</del> .74 | 14        |
| 197 | Genome-wide identification of Listeria monocytogenes CodY-binding sites. <i>Molecular Microbiology</i> , <b>2020</b> , 113, 841-858  | 4.1                  | 4         |
| 196 | Role of GlnR in Controlling Expression of Nitrogen Metabolism Genes in. <i>Journal of Bacteriology</i> , <b>2020</b> , 202,  | 3.5                  | 1         |
| 195 | Impact of CodY protein on metabolism, sporulation and virulence in Clostridioides difficile ribotype 027. <i>PLoS ONE</i> , <b>2019</b> , 14, e0206896   | 3.7                  | 13        |
| 194 | Role of the global regulator Rex in control of NAD -regeneration in Clostridioides (Clostridium) difficile. <i>Molecular Microbiology</i> , <b>2019</b> , 111, 1671-1688   | 4.1                  | 15        |
| 193 | DdlR, an essential transcriptional regulator of peptidoglycan biosynthesis in Clostridioides difficile. <i>Molecular Microbiology</i> , <b>2019</b> , 112, 1453-1470   | 4.1                  | 1         |
| 192 | Oral Immunization with Nontoxigenic Clostridium difficile Strains Expressing Chimeric Fragments of TcdA and TcdB Elicits Protective Immunity against C. difficile Infection in Both Mice and Hamsters. <i>Infection and Immunity</i> , <b>2018</b> , 86, | 3.7                  | 6         |
| 191 | A Mutation in the Bacillus subtilis rsbU Gene That Limits RNA Synthesis during Sporulation. <i>Journal of Bacteriology</i> , <b>2017</b> , 199,  | 3.5                  | 3         |
| 190 | Structure of the Branched-chain Amino Acid and GTP-sensing Global Regulator, CodY, from. <i>Journal of Biological Chemistry</i> , <b>2017</b> , 292, 2714-2728   | 5.4                  | 21        |
| 189 | Interplay of CodY and ScoC in the Regulation of Major Extracellular Protease Genes of Bacillus subtilis. <i>Journal of Bacteriology</i> , <b>2016</b> , 198, 907-20  | 3.5                  | 26        |
| 188 | CodY regulates expression of the Bacillus subtilis extracellular proteases Vpr and Mpr. <i>Journal of Bacteriology</i> , <b>2015</b> , 197, 1423-32  | 3.5                  | 14        |
| 187 | Effects of surotomycin on Clostridium difficile viability and toxin production in vitro. <i>Antimicrobial Agents and Chemotherapy</i> , <b>2015</b> , 59, 4199-205   | 5.9                  | 22        |
| 186 | Three cellulosomal xylanase genes in Clostridium thermocellum are regulated by both vegetative SigA ((A)) and alternative SigI6 ((16)) factors. <i>FEBS Letters</i> , <b>2015</b> , 589, 3133-40   | 3.8                  | 14        |
| 185 | Integration of metabolism and virulence in Clostridium difficile. <i>Research in Microbiology</i> , <b>2015</b> , 166, 37  | 5 <sub>z</sub> β3    | 74        |
| 184 | Interactive regulation by the Bacillus subtilis global regulators CodY and ScoC. <i>Molecular Microbiology</i> , <b>2015</b> , 97, 698-716   | 4.1                  | 15        |
| 183 | Regulating the Intersection of Metabolism and Pathogenesis in Gram-positive Bacteria. <i>Microbiology Spectrum</i> , <b>2015</b> , 3,  | 8.9                  | 68        |
| 182 | Regulating the Intersection of Metabolism and Pathogenesis in Gram-positive Bacteria <b>2015</b> , 129-165   |                      | 2         |

### (2014-2015)

| 181 | Intermediate Levels of Bacillus subtilis CodY Activity Are Required for Derepression of the Branched-Chain Amino Acid Permease, BraB. <i>PLoS Genetics</i> , <b>2015</b> , 11, e1005600   | 6   | 13  |
|-----|---|-----|-----|
| 180 | The metabolic regulator CodY links Listeria monocytogenes metabolism to virulence by directly activating the virulence regulatory gene prfA. <i>Molecular Microbiology</i> , <b>2015</b> , 95, 624-44   | 4.1 | 56  |
| 179 | CodY-mediated regulation of the Staphylococcus aureus Agr system integrates nutritional and population density signals. <i>Journal of Bacteriology</i> , <b>2014</b> , 196, 1184-96   | 3.5 | 52  |
| 178 | Multiple regulatory mechanisms control the expression of the Geobacillus stearothermophilus gene for extracellular xylanase. <i>Journal of Biological Chemistry</i> , <b>2014</b> , 289, 25957-75   | 5.4 | 22  |
| 177 | GTP dysregulation in Bacillus subtilis cells lacking (p)ppGpp results in phenotypic amino acid auxotrophy and failure to adapt to nutrient downshift and regulate biosynthesis genes. <i>Journal of Bacteriology</i> , <b>2014</b> , 196, 189-201 | 3.5 | 67  |
| 176 | Biosynthesis of Riboflavin, Biotin, Folic Acid, and Cobalamin <b>2014</b> , 319-334   |     | 15  |
| 175 | Systematics and Ecology of Bacillus <b>2014</b> , 1-16  |     | 70  |
| 174 | Lactococcus and Lactobacillus <b>2014</b> , 65-82   |     | 10  |
| 173 | Spore Germination and Outgrowth <b>2014</b> , 537-548   |     | 73  |
| 172 | Purine and Pyrimidine Salvage Pathways <b>2014</b> , 359-378  |     | 27  |
| 171 | Integrational Vectors for Genetic Manipulation in Bacillus subtilis <b>2014</b> , 615-624   |     | 119 |
| 170 | Synthesis of Serine, Glycine, Cysteine, and Methionine <b>2014</b> , 245-254  |     | 24  |
| 169 | The Bacillus subtilis Genome, Genes, and Functions <b>2014</b> , 7-11   |     |     |
| 168 | The Dynamic Architecture of the Bacillus Cell <b>2014</b> , 13-20   |     | 3   |
| 167 | Structure and Synthesis of Cell Wall, Spore Cortex, Teichoic Acids, S-Layers, and Capsules <b>2014</b> , 21-41  |     | 55  |
| 166 | Commercial Production of Extracellular Enzymes <b>2014</b> , 917-937  |     | 37  |
| 165 | Proteins of the Spore Core and Coat <b>2014</b> , 527-535   |     | 42  |
| 164 | Biosynthesis of Aromatic Amino Acids <b>2014</b> , 269-280  |     | 12  |

| 163        | Fermentation of Bacillus <b>2014</b> , 869-895  | 8                              |
|------------|---|--------------------------------|
| 162        | RNA Polymerase and Sigma Factors <b>2014</b> , 287-312  | 72                             |
| 161        | DNA Repair Systems <b>2014</b> , 529-537  | 34                             |
| 160        | Endospore-Forming Bacteria: an Overview <b>2014</b> , 131-150   | 9                              |
| 159        | Biosynthesis of the Branched-Chain Amino Acids <b>2014</b> , 307-317  | 14                             |
| 158        | Peptide Antibiotics <b>2014</b> , 897-916   | 40                             |
| 157        | Biosynthesis of Glutamine and Glutamate and the Assimilation of Ammonia <b>2014</b> , 281-298   | 29                             |
| 156        | Transformation and Recombination <b>2014</b> , 453-471  | 26                             |
| 155        | Bacillus anthracis <b>2014</b> , 113-124  | 35                             |
| 154        | Carbohydrate Catabolism: Pathways, Enzymes, Genetic Regulation, and Evolution <b>2014</b> , 157-170   | 25                             |
| 153        | The Genetic Map of Bacillus megaterium <b>2014</b> , 475-481  | 4                              |
| 152        |   |                                |
|            | Carbon Source-Mediated Catabolite Repression <b>2014</b> , 213-219  | 37                             |
| 151        | Carbon Source-Mediated Catabolite Repression <b>2014</b> , 213-219  General View of the Bacillus subtilis Chromosome <b>2014</b> , 552-569  | 37                             |
| 151<br>150 |   | <ul><li>37</li><li>6</li></ul> |
|            | General View of the Bacillus subtilis Chromosome <b>2014</b> , 552-569  |                                |
| 150        | General View of the Bacillus subtilis Chromosome <b>2014</b> , 552-569  Overall Transport Capabilities of Bacillus subtilis <b>2014</b> , 111-128   | 6                              |
| 150<br>149 | General View of the Bacillus subtilis Chromosome 2014, 552-569  Overall Transport Capabilities of Bacillus subtilis 2014, 111-128  Cell Division during Growth and Sporulation 2014, 97-109 | 6<br>7                         |

### (2011-2014)

| 145 | Hierarchical expression of genes controlled by the Bacillus subtilis global regulatory protein CodY. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2014</b> , 111, 8227-32 | 11.5 | 64  |
|-----|--|------|-----|
| 144 | Carvacrol and trans-cinnamaldehyde reduce Clostridium difficile toxin production and cytotoxicity in vitro. <i>International Journal of Molecular Sciences</i> , <b>2014</b> , 15, 4415-30                               | 6.3  | 38  |
| 143 | Use of a mariner-based transposon mutagenesis system to isolate Clostridium perfringens mutants deficient in gliding motility. <i>Journal of Bacteriology</i> , <b>2013</b> , 195, 629-36                                | 3.5  | 30  |
| 142 | Fidaxomicin inhibits toxin production in Clostridium difficile. <i>Journal of Antimicrobial Chemotherapy</i> , <b>2013</b> , 68, 515-22  | 5.1  | 54  |
| 141 | Genome-wide identification of Bacillus subtilis CodY-binding sites at single-nucleotide resolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2013</b> , 110, 7026-31  | 11.5 | 70  |
| 140 | Dual role of CcpC protein in regulation of aconitase gene expression in Listeria monocytogenes and Bacillus subtilis. <i>Microbiology (United Kingdom)</i> , <b>2013</b> , 159, 68-76                                    | 2.9  | 12  |
| 139 | Two roles for aconitase in the regulation of tricarboxylic acid branch gene expression in Bacillus subtilis. <i>Journal of Bacteriology</i> , <b>2013</b> , 195, 1525-37   | 3.5  | 20  |
| 138 | Proline-dependent regulation of Clostridium difficile Stickland metabolism. <i>Journal of Bacteriology</i> , <b>2013</b> , 195, 844-54   | 3.5  | 88  |
| 137 | Comment on the Howy JacobsSEditorial "Yes we can, but do we?". EMBO Reports, 2013, 14, 1027  | 6.5  |     |
| 136 | CodY deletion enhances in vivo virulence of community-associated methicillin-resistant Staphylococcus aureus clone USA300. <i>Infection and Immunity</i> , <b>2012</b> , 80, 2382-9                                      | 3.7  | 47  |
| 135 | Fidaxomicin inhibits spore production in Clostridium difficile. <i>Clinical Infectious Diseases</i> , <b>2012</b> , 55 Suppl 2, S162-9   | 11.6 | 88  |
| 134 | Proline utilization by Bacillus subtilis: uptake and catabolism. <i>Journal of Bacteriology</i> , <b>2012</b> , 194, 745-58  | 3.5  | 49  |
| 133 | Roadblock repression of transcription by Bacillus subtilis CodY. <i>Journal of Molecular Biology</i> , <b>2011</b> , 411, 729-43   | 6.5  | 53  |
| 132 | Dissecting complex metabolic integration provides direct genetic evidence for CodY activation by guanine nucleotides. <i>Journal of Bacteriology</i> , <b>2011</b> , 193, 5637-48  | 3.5  | 26  |
| 131 | Contributions of multiple binding sites and effector-independent binding to CodY-mediated regulation in Bacillus subtilis. <i>Journal of Bacteriology</i> , <b>2011</b> , 193, 473-84                                    | 3.5  | 33  |
| 130 | CodY-mediated regulation of guanosine uptake in Bacillus subtilis. <i>Journal of Bacteriology</i> , <b>2011</b> , 193, 6276-87   | 3.5  | 13  |
| 129 | The dlt operon confers resistance to cationic antimicrobial peptides in Clostridium difficile. <i>Microbiology (United Kingdom)</i> , <b>2011</b> , 157, 1457-1465   | 2.9  | 108 |
| 128 | Identification of a genetic locus responsible for antimicrobial peptide resistance in Clostridium difficile. <i>Infection and Immunity</i> , <b>2011</b> , 79, 167-76  | 3.7  | 87  |

| 127 | Integration of metabolism and virulence by Clostridium difficile CodY. <i>Journal of Bacteriology</i> , <b>2010</b> , 192, 5350-62   | 3.5                     | 134 |
|-----|--|-------------------------|-----|
| 126 | Inhibiting the initiation of Clostridium difficile spore germination using analogs of chenodeoxycholic acid, a bile acid. <i>Journal of Bacteriology</i> , <b>2010</b> , 192, 4983-90        | 3.5                     | 216 |
| 125 | Direct targets of CodY in Staphylococcus aureus. <i>Journal of Bacteriology</i> , <b>2010</b> , 192, 2861-77   | 3.5                     | 148 |
| 124 | Regulation of CodY activity through modulation of intracellular branched-chain amino acid pools.<br>Journal of Bacteriology, <b>2010</b> , 192, 6357-68                                      | 3.5                     | 51  |
| 123 | CcpC-dependent regulation of citrate synthase gene expression in Listeria monocytogenes. <i>Journal of Bacteriology</i> , <b>2009</b> , 191, 862-72  | 3.5                     | 6   |
| 122 | The global regulator CodY regulates toxin gene expression in Bacillus anthracis and is required for full virulence. <i>Infection and Immunity</i> , <b>2009</b> , 77, 4437-45                | 3.7                     | 70  |
| 121 | Chenodeoxycholate is an inhibitor of Clostridium difficile spore germination. <i>Journal of Bacteriology</i> , <b>2009</b> , 191, 1115-7   | 3.5                     | 141 |
| 120 | Genetic and biochemical analysis of the interaction of Bacillus subtilis CodY with branched-chain amino acids. <i>Journal of Bacteriology</i> , <b>2009</b> , 191, 6865-76                   | 3.5                     | 36  |
| 119 | Structural rearrangement accompanying ligand binding in the GAF domain of CodY from Bacillus subtilis. <i>Journal of Molecular Biology</i> , <b>2009</b> , 390, 1007-18                      | 6.5                     | 32  |
| 118 | Control of nitrogen metabolism by Bacillus subtilis glutamine synthetase. <i>Molecular Microbiology</i> , <b>2008</b> , 68, 242-245  | 4.1                     | 5   |
| 117 | Genetic and biochemical analysis of CodY-binding sites in Bacillus subtilis. <i>Journal of Bacteriology</i> , <b>2008</b> , 190, 1224-36   | 3.5                     | 76  |
| 116 | Bile salts and glycine as cogerminants for Clostridium difficile spores. <i>Journal of Bacteriology</i> , <b>2008</b> , 190, 2505-12   | 3.5                     | 445 |
| 115 | Interaction of Bacillus subtilis CodY with GTP. Journal of Bacteriology, 2008, 190, 798-806  | 3.5                     | 86  |
| 114 | Staphylococcus aureus CodY negatively regulates virulence gene expression. <i>Journal of Bacteriology</i> , <b>2008</b> , 190, 2257-65   | 3.5                     | 137 |
| 113 | Control of key metabolic intersections in Bacillus subtilis. <i>Nature Reviews Microbiology</i> , <b>2007</b> , 5, 917-27  | 22.2                    | 291 |
| 112 | Characterization of relA and codY mutants of Listeria monocytogenes: identification of the CodY regulon and its role in virulence. <i>Molecular Microbiology</i> , <b>2007</b> , 63, 1453-67 | 4.1                     | 128 |
| 111 | Repression of Clostridium difficile toxin gene expression by CodY. <i>Molecular Microbiology</i> , <b>2007</b> , 66, 206   | 6 <b>-</b> 41. <b>9</b> | 175 |
| 110 | Molecular mechanism of the regulation of Bacillus subtilis gltAB expression by GltC. <i>Journal of Molecular Biology</i> , <b>2007</b> , 365, 1298-313                                       | 6.5                     | 59  |

### (2002-2006)

| 109 | CcpC-dependent regulation of citB and lmo0847 in Listeria monocytogenes. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 179-90  | 3.5           | 21  |
|-----|--|---------------|-----|
| 108 | Bacillus subtilis aconitase is required for efficient late-sporulation gene expression. <i>Journal of Bacteriology</i> , <b>2006</b> , 188, 6396-405   | 3.5           | 45  |
| 107 | The structure of CodY, a GTP- and isoleucine-responsive regulator of stationary phase and virulence in gram-positive bacteria. <i>Journal of Biological Chemistry</i> , <b>2006</b> , 281, 11366-73            | 5.4           | 78  |
| 106 | Positive regulation of Bacillus subtilis ackA by CodY and CcpA: establishing a potential hierarchy in carbon flow. <i>Molecular Microbiology</i> , <b>2006</b> , 62, 811-22                                    | 4.1           | 82  |
| 105 | CodY, a global regulator of stationary phase and virulence in Gram-positive bacteria. <i>Current Opinion in Microbiology</i> , <b>2005</b> , 8, 203-7  | 7.9           | 267 |
| 104 | Bacillus subtilis ilvB operon: an intersection of global regulons. <i>Molecular Microbiology</i> , <b>2005</b> , 56, 1549-5  | 5 <b>2</b> .1 | 67  |
| 103 | A region of Bacillus subtilis CodY protein required for interaction with DNA. <i>Journal of Bacteriology</i> , <b>2005</b> , 187, 4127-39  | 3.5           | 40  |
| 102 | CcpA-dependent regulation of Bacillus subtilis glutamate dehydrogenase gene expression. <i>Journal of Bacteriology</i> , <b>2004</b> , 186, 3392-8   | 3.5           | 45  |
| 101 | Modulation of activity of Bacillus subtilis regulatory proteins GltC and TnrA by glutamate dehydrogenase. <i>Journal of Bacteriology</i> , <b>2004</b> , 186, 3399-407   | 3.5           | 36  |
| 100 | Activation of the Bacillus subtilis global regulator CodY by direct interaction with branched-chain amino acids. <i>Molecular Microbiology</i> , <b>2004</b> , 53, 599-611                                     | 4.1           | 175 |
| 99  | Complex regulation of the Bacillus subtilis aconitase gene. <i>Journal of Bacteriology</i> , <b>2003</b> , 185, 1672-80  | 3.5           | 47  |
| 98  | Efficient sporulation in Clostridium difficile requires disruption of the sigmaK gene. <i>Molecular Microbiology</i> , <b>2003</b> , 48, 811-21  | 4.1           | 104 |
| 97  | Mechanism of repression by Bacillus subtilis CcpC, a LysR family regulator. <i>Journal of Molecular Biology</i> , <b>2003</b> , 334, 609-24  | 6.5           | 31  |
| 96  | Specificity of the interaction of RocR with the rocG-rocA intergenic region in Bacillus subtilis. <i>Microbiology (United Kingdom)</i> , <b>2003</b> , 149, 739-750  | 2.9           | 14  |
| 95  | Additional targets of the Bacillus subtilis global regulator CodY identified by chromatin immunoprecipitation and genome-wide transcript analysis. <i>Journal of Bacteriology</i> , <b>2003</b> , 185, 1911-22 | 3.5           | 233 |
| 94  | Developmental biology: regulation by selective gene localization. <i>Current Biology</i> , <b>2002</b> , 12, R90-2   | 6.3           | 2   |
| 93  | Regulation of the bacillus subtilis ccpC gene by ccpA and ccpC. <i>Molecular Microbiology</i> , <b>2002</b> , 43, 399-41   | 04.1          | 28  |
| 92  | Direct and indirect roles of CcpA in regulation of Bacillus subtilis Krebs cycle genes. <i>Molecular Microbiology</i> , <b>2002</b> , 45, 179-90   | 4.1           | 62  |

| 91 | Environmental response and autoregulation of Clostridium difficile TxeR, a sigma factor for toxin gene expression. <i>Journal of Bacteriology</i> , <b>2002</b> , 184, 5971-8                                  | 3.5   | 104 |
|----|--|-------|-----|
| 90 | Molecular biology. Turning gene regulation on its head. <i>Science</i> , <b>2001</b> , 293, 2018-9   | 33.3  | 5   |
| 89 | Bacillus subtilis CodY represses early-stationary-phase genes by sensing GTP levels. <i>Genes and Development</i> , <b>2001</b> , 15, 1093-103   | 12.6  | 268 |
| 88 | Role of TnrA in nitrogen source-dependent repression of Bacillus subtilis glutamate synthase gene expression. <i>Journal of Bacteriology</i> , <b>2000</b> , 182, 5939-47                                      | 3.5   | 73  |
| 87 | CcpC, a novel regulator of the LysR family required for glucose repression of the citB gene in Bacillus subtilis. <i>Journal of Molecular Biology</i> , <b>2000</b> , 295, 865-78                              | 6.5   | 92  |
| 86 | Metabolic imbalance and sporulation in an isocitrate dehydrogenase mutant of Bacillus subtilis.<br>Journal of Bacteriology, <b>1999</b> , 181, 3382-91   | 3.5   | 33  |
| 85 | Regulated transcription of Clostridium difficile toxin genes. <i>Molecular Microbiology</i> , <b>1998</b> , 27, 107-20   | 4.1   | 217 |
| 84 | Role and regulation of Bacillus subtilis glutamate dehydrogenase genes. <i>Journal of Bacteriology</i> , <b>1998</b> , 180, 6298-305   | 3.5   | 124 |
| 83 | Interaction of CodY, a novel Bacillus subtilis DNA-binding protein, with the dpp promoter region. <i>Molecular Microbiology</i> , <b>1996</b> , 20, 843-52   | 4.1   | 65  |
| 82 | A gene required for nutritional repression of the Bacillus subtilis dipeptide permease operon. <i>Molecular Microbiology</i> , <b>1995</b> , 15, 689-702   | 4.1   | 145 |
| 81 | Mechanism of initiation of transcription by Bacillus subtilis RNA polymerase at several promoters.<br>Journal of Molecular Biology, <b>1992</b> , 223, 399-414   | 6.5   | 69  |
| 80 | Regulation of Bacillus subtilis glutamine synthetase gene expression by the product of the glnR gene. <i>Journal of Molecular Biology</i> , <b>1989</b> , 210, 51-63   | 6.5   | 107 |
| 79 | Glutamine synthetase gene of Bacillus subtilis. <i>Gene</i> , <b>1984</b> , 32, 427-38   | 3.8   | 53  |
| 78 | Initiation of transcription in vitro inhibited by lipiarmycin. <i>Journal of Molecular Biology</i> , <b>1979</b> , 127, 55-72  | 2 6.5 | 36  |
| 77 | Isolation and characterization of rifampin-resistant and streptolydigin-resistant mutants of Bacillus subtilis with altered sporulation properties. <i>Journal of Bacteriology</i> , <b>1974</b> , 120, 253-65 | 3.5   | 118 |
| 76 | RNA polymerase mutants blocked in sporulation. <i>Nature</i> , <b>1970</b> , 227, 906-9  | 50.4  | 80  |
| 75 | Change in the template specificity of RNA polymerase during sporulation of Bacillus subtilis. <i>Nature</i> , <b>1969</b> , 224, 35-7  | 50.4  | 185 |
| 74 | The course of phage phi-e infection in sporulating cells of Bacillus subtilis strain 3610. <i>Virology</i> , <b>1969</b> , 39, 265-75  | 3.6   | 100 |

| 73 | Respiratory Cytochromes, Other Heme Proteins, and Heme Biosynthesis163-179                          | 11 |
|----|---|----|
| 72 | Nitrogen Source Utilization and Its Regulation181-191   | 21 |
| 71 | The Pho Regulon193-201  | 38 |
| 70 | Biosynthesis of Amino Acids of the Glutamate and Aspartate Families, Alanine, and Polyamines203-231 | 34 |
| 69 | Aromatic Amino Acid Metabolism in Bacillus subtilis233-244  | 13 |
| 68 | Purine, Pyrimidine, and Pyridine Nucleotide Metabolism255-269                                       | 13 |
| 67 | Vitamin Biosynthesis271-286   | 24 |
| 66 | Ribosomes, Protein Synthesis Factors, and tRNA Synthetases313-322                                   | 8  |
| 65 | Regulation by Termination-Antitermination: a Genomic Approach323-336                                | 8  |
| 64 | Protein Transport Pathways in Bacillus subtilis: a Genome-Based Road Map337-355                     | 9  |
| 63 | Regulation and Function of Heat-Inducible Genes in Bacillus subtilis357-368                         | 32 |
| 62 | General Stress Response369-384  | 60 |
| 61 | Adaptation to Changing Osmolanty385-391   | 42 |
| 60 | Anaerobiosis393-404   | 17 |
| 59 | Metal Ion Uptake and Oxidative Stress405-414  | 20 |
| 58 | Nonribosomal Assembly of Peptide Antibiotics on Modular Protein Templates415-435                    | 6  |
| 57 | Chemotaxis and Motility437-452  | 27 |
| 56 | Two-Component Systems, Phosphorelays, and Regulation of Their Activities by Phosphatases473-481     | 55 |

| 55 | Sporulation Genes and Intercompartmental Regulation483-517                          | 108 |
|----|---|-----|
| 54 | A Gene Odyssey: Exploring the Genomes of Endospore-Forming Bacteria519-525          | 43  |
| 53 | Biosynthesis and Function of Membrane Lipids43-55                                   | 34  |
| 52 | Cell Wall-Anchored Surface Proteins and Lipoproteins of Gram-Positive Bacteria57-70 | 1   |
| 51 | Chromosome Replication and Segregation71-86   | 7   |
| 50 | Termination of Chromosome Replication87-95  | 8   |
| 49 | Transport Mechanisms133-156   | 14  |
| 48 | Glycolysis171-180   | 14  |
| 47 | The Krebs Citric Acid Cycle181-197  | 26  |
| 46 | Respiratory Chains199-212   | 10  |
| 45 | Utilization of Amino Acids and Other Nitrogen-Containing Compounds221-228           | 15  |
| 44 | Regulation of Phosphorus Metabolism229-235  | 4   |
| 43 | Biosynthesis of the Aspartate Family of Amino Acids237-267                          | 18  |
| 42 | Staphylococcus17-33   | 2   |
| 41 | Biosynthesis of Arginine, Proline, and Related Compounds299-306                     | 14  |
| 40 | De Novo Purine Nucleotide Synthesis335-341  | 7   |
| 39 | De Novo Pyrimidine Nucleotide Synthesis343-358                                      | 4   |
| 38 | Cell Wall Structure, Synthesis, and Turnover379-410                                 | 28  |

| 37 | Biosynthesis and Function of Membrane Lipids411-421  | 28           |
|----|--|--------------|
| 36 | The Genetic Map of Bacillus subtilis423-461  | 9            |
| 35 | Clostridium35-52   | 8            |
| 34 | Physical Map of the Bacillus subtilis 168 Chromosome463-471  | 5            |
| 33 | An Ordered Collection of Bacillus subtilis DNA Segments in Yeast Artificial Chromosomes473-474               | 2            |
| 32 | The Chromosome Map of Streptomyces coelicolor A3(2)497-504   | 4            |
| 31 | Initiation and Termination of Chromosome Replication505-528  | 14           |
| 30 | Restriction/Modification and Methylation Systems in Bacillus subtilis, Related Species, and Their Phages539- | ·55 <b>g</b> |
| 29 | Genetic Exchange and Homologous Recombination553-584   | 23           |
| 28 | Transposons and Their Applications585-596  | 5            |
| 27 | Plasmids625-644  | 37           |
| 26 | RNA Polymerase and Transcription Factors651-667  | 25           |
| 25 | Ribosomal Structure and Genetics669-682  | 3            |
| 24 | tRNA, tRNA Processing, and Aminoacyl-tRNA Synthetases683-698   | 7            |
| 23 | Translation and Its Regulation699-711  | 19           |
| 22 | Protein Secretion713-726   | 23           |
| 21 | Two-Component Regulatory Systems727-745  | 21           |
| 20 | spoO Genes, the Phosphorelay, and the Initiation of Sporulation747-755                                       | 30           |

| 19               | AbrB, a Transition State Regulator757-764   | 36               |
|------------------|---|------------------|
| 18               | Motility and Chemotaxis765-784  | 27               |
| 17               | Regulatory Proteins That Control Late-Growth Development785-800   | 40               |
| 16               | Spore Structural Proteins801-809  | 19               |
| 15               | SPO1 and Related Bacteriophages811-829  | 3                |
| 14               | Temperate Bacteriophages831-842   | 8                |
| 13               | Replication and Transcription of Bacteriophage ?29 DNA843-857   | 9                |
| 12               | Morphogenesis of Bacteriophage ?29859-867   | 14               |
| 11               | Streptomyces83-99   | 22               |
|                  |   |                  |
| 10               | Proteases939-952  | 18               |
| 10               | Proteases939-952 Insecticidal Toxins953-963   | 18<br>7          |
|                  |   |                  |
| 9                | Insecticidal Toxins953-963  | 7                |
| 9                | Insecticidal Toxins953-963  Pasteuria, Metchnikoff, 1888101-111   | 7                |
| 9 8 7            | Insecticidal Toxins953-963  Pasteuria, Metchnikoff, 1888101-111  Introduction to Metabolic Pathways125-132  | 7<br>6<br>5      |
| 9<br>8<br>7      | Insecticidal Toxins953-963  Pasteuria, Metchnikoff, 1888101-111  Introduction to Metabolic Pathways125-132  In vivo commensal control of Clostridioides difficile virulence  The branched chain aminotransferase IlvE promotes growth, stress resistance and pathogenesis                           | 7<br>6<br>5<br>4 |
| 9<br>8<br>7<br>6 | Insecticidal Toxins953-963  Pasteuria, Metchnikoff, 1888101-111  Introduction to Metabolic Pathways125-132  In vivo commensal control of Clostridioides difficile virulence  The branched chain aminotransferase IlvE promotes growth, stress resistance and pathogenesis of Listeria monocytogenes | 7<br>6<br>5<br>4 |

#### LIST OF PUBLICATIONS

1 Temperate Phage Vectors645-650