

Recently Published Papers in the Field of Molecular Evolution

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GENERAL ASPECTS

Journal of Theoretical Biology

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Phylogenetic Trees Constructed from Hydrophobicity Values of Protein Sequences. Leunissen, J.A.M. and DeJong, W.W. (Nijmegen, NL) - p. 189 [771]

The Classification of Amino Acid Conservation. Taylor, W.R. (London, UK) - p. 205 [772]

Molecular Biology and Evolution

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Temporal Scaling of Molecular Evolution in Primates and Other Mammals. Gingerich, P.D. (Ann Arbor, USA) - p. 205 [773]

Mammalian Phylogeny: Comparison of Morphological and Molecular Results. Shoshani, J. (Detroit, USA) - p. 222 [774]

Nature

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Phylogenetic Distribution of *Antennapedia*-Like Homoeo Boxes. Holland, P.W.H. and Hogan, B.L.M. (London, UK) - p. 251 [775]

Origins of Life

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Trace Elements in Chemical Evolution. I. Kobayashi, K. and Ponnampereuma, C. (College Park, USA) - p. 41 [776]

Trace Elements in Chemical Evolution. II. Synthesis of Amino Acids Under Simulated Primitive Earth Conditions in the Presence of Trace Elements. Kobayashi, K. and Ponnampereuma, C. (College Park, USA) - p. 57 [777]

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RNA Catalysis and the Origin of Life. Pace, N.R. and Marsh, T.L. (Bloomington, USA) - p. 97 [778]

Chirally Selective, Intramolecular Interactions Observed in an Aminoacyl Adenylate Anhydride. Lacey, J.C. et al. (Birmingham, USA) - p. 151 [779]

Conservation of the Secondary Structure of Protein During Evolution and the Role of the Genetic Code. Soto, M.A. et al. (Santiago, Chile) - p. 157 [780]

Proceedings of the National Academy of Sciences of the USA

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Proteins of *Escherichia coli* Come in Sizes that are Multiples of 14 kDa: Domain Concepts and Evolutionary Implications. Savageau, M.A. (Ann Arbor, USA) - p. 1198 [781]

Speculations on the Early Course of Evolution. Darnell, J.E. and Doolittle, W.F. (New York, USA) - p. 1271 [782]

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Origin of Eukaryotic Introns: a Hypothesis, Based on Codon Distribution Statistics in Genes, and Its Implication. Senapathy, P. (Bethesda, USA) - p. 2133 [783]

Multigene Families and Vestigial Sequences. Loomis, W.F. and Gilpin, M.E. (La Jolla, USA) - p. 2143 [784]

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A Model for the RNA-Catalyzed Replication of RNA. Cech, T.R. (Boulder, USA) - p. 4360 [785]

Systematics and Applied Microbiology

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Comparative Studies on Structure and Function of Archaeobacterial Elongation Factors Indicate the

Phylogenetic Diversity of the Urkingdom. Gehrman, R. et al. (Kiel, FRG) - p. 115 [786]

Mapping Evolution with 3 Dimensional Ribosome Structure. Lake, J.A. et al. (Los Angeles, USA) - p. 131 [787]

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Biochemistry

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Molecular Structure and Polymorphic Map of the Human Phenylalanine Hydroxylase Gene. DiLella, A. et al. (Houston, USA) - p. 743 [788]

25 No. 7 1986

Characterization of a cDNA Coding for Human Factor XII (Hageman Factor). Que, B.G. and Davie, E.W. (Seattle, USA) - p. 1525 [789]

25 No. 8 1986

Chicken U2 and U1 RNA Genes are Found in Very Different Genomic Environments but Have Similar Promoter Structures. Korf, G.M. and Stumph, W.E. (San Diego, USA) - p. 2041 [790]

Amino Acid Sequence of Human Histidine-Rich Glycoprotein Derived from the Nucleotide Sequence of Its cDNA. Koide, T. et al. (Seattle, USA) - p. 2220 [791]

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cDNA and Protein Structure for the α Subunit of Human Liver Alcohol Dehydrogenase. Von Bahr-Lindström, H. et al. (Stockholm, Sweden) - p. 2465 [792]

Glutathione Reductase from *Escherichia coli*: Cloning and Sequence Analysis of the Gene and Relationship to Other Flavoprotein Disulfide Oxidoreductases. Greer, S. and Perham, R.N. (Cambridge, UK) - p. 2736 [793]

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cDNA Sequences for Human von Willebrand Factor Reveal Five Types of Repeated Domains and Five Possible Protein Sequence Polymorphisms. Shelton-Inloes, B.B. et al. (St. Louis, USA) - p. 3164 [794]

Biosynthetic *alr* Alanine Racemase from *Salmonella typhimurium*: DNA and Protein Sequence Determination. Galakatos, N.G. et al. (Cambridge, USA) - p. 3255 [795]

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Structure, Evolution, and Tissue-Specific Synthesis of Human Apolipoprotein AIV. Karathanasis, S.K. et al. (Boston, USA) - p. 3962 [796]

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Detailed Analysis of the Mouse H-2K^b Promoter: Enhancer-Like Sequences and Their Role in the Regulation of Class I Gene Expression. Kimura, A. et al. (Paris, France) - p. 261 [797]

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Two Ca⁺⁺ ATPase Genes: Homologies and Mechanistic Implications of Deduced Amino Acid Sequences. Brandl, C.J. et al. (Toronto, Canada) - p. 597 [798]

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A Tandemly Repeated Sequence Determines the Binding Domain for an Erythrocyte Receptor Binding Protein of *E. faeciparum*. Kochan, J. et al. (New York, USA) - p. 689 [799]

An Opsin Gene Expressed in Only One Photoreceptor Cell Type of the *Drosophila* Eye. Cowman, A.F. et al. (Berkeley, USA) - p. 705 [800]

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Molecular Characterization of the Human Excision Repair Gene ERCC-1: cDNA Cloning and Amino Acid Homology with the Yeast DNA Repair Gene RAD10. Van Duin, M. et al. (Rotterdam, NL) - p. 913 [802]

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cdc25⁺ Functions as an Inducer in the Mitotic Control of Fission Yeast. Russell, P. and Nurse, P. (London, UK) - p. 145 [803]

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Characterization of the Intron in the Phage T4 Thymidylate Synthase Gene and Evidence for Its Self-Excision from the Primary Transcript. Chu, F.K. et al. (Albany, USA) - p. 157 [804]

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Nucleotide Sequence of Mason-Pfizer Monkey Virus: An Immunosuppressive D-Type Retrovirus. Sonigo, P. et al. (Paris, France) - p. 375 [805]

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Identification and Characterization of Conserved and Variable Regions in the Envelope Gene of HTLV-III/LAV, the Retrovirus of AIDS. Starcich, B.R. et al. (Bethesda, USA) - p. 637 [806]

Isolation of the Bovine and Human Genes for Mullerian Inhibiting Substance and Expression of the Human Gene in Animal Cells. Cate, R.L. et al. (Cambridge, USA) - p. 685 [807]

Isolation and Characterization of a New Cellular Oncogene Encoding a Protein with Multiple Potential Transmembrane Domains. Young, D. et al. (Cold Spring Harbor, USA) - p. 711 [808]

Diversity, Rearrangement, and Expression of Murine T Cell γ Genes. Garman, R.D. et al. (Cambridge, USA)-p.733[809]

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A Single Gene from Yeast for Both Nuclear and Cytoplasmic Polyadenylate-Binding Proteins: Domain Structure and Expression. Sachs, A.B. et al. (Stanford, USA)-p.827[810]

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The Triosephosphate Isomerase Gene from Maize: Introns Antedate the Plant-Animal Divergence. Marchionni, M. and Gilbert, W. (Cambridge, USA) - p. 133 [811]

Nucleotide Sequence of the Triosephosphate Isomerase Gene from *Aspergillus nidulans*: Implications for a Differential Loss of Introns. McKnight, G.L. et al. (Seattle, USA) - p. 143 [812]

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The Structure and Expression of Maize Genes Encoding the Major Heat Shock Protein, hsp 70. Rochester, D.E. et al. (St. Louis, USA) - p. 451 [813]

Isolation and Characterization of Genes Encoding Two Chitinase Enzymes from *Serratia marcescens*. Jones, J.D.G. et al. (Oakland, USA) - p. 467 [814]

Gene Conversion-Like Mechanisms May Generate Polymorphism in Human Class I Genes. Seemann, G.H.A. et al. (Amsterdam, NL) - p. 547 [815]

The Chromosomal Gene Structure and Two mRNAs for Human Granulocyte Colony-Stimulating Factor. Nagata, S. et al. (Tokyo, Japan) - p. 575 [816]

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The *Rhizobium leguminosarum* Nodulation Gene *nodF* Encodes a Polypeptide Similar to Acyl-Carrier Protein and Is Regulated by *nodD* Plus a Factor in Pea Root Exudate. Shearman, C.A. et al. (Norwich, UK) - p. 647 [817]

Regulated Expression of Repetitive Sequences Including the Identifier Sequence During Myotube Formation in Culture. Herget, T. et al. (Köln, FRG) - p. 659 [818]

Structural Organization and Sequence of the Homeotic Gene *Antennapedia* of *Drosophila melanogaster*. Schneuwly, S. et al. (Basel, CH) - p. 733 [819]

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Molecular Analysis of the En/Spm Transposable Element System of *Zea mays*. Pereira, A. et al. (Köln, FRG)-p.835[820]

The Chicken Oestrogen Receptor Sequence: Homology with *v-erbA* and the Human Oestrogen and Glucocorticoid Receptors. Krust, A. et al. (Strasbourg, France)-p.891[821]

Sequence, Topography, and Protein Coding Potential of Mouse *int-2*: a Putative Oncogene Activated by Mouse Mammary Tumour Virus. Moore, R. et al. (London, UK)-p.919[822]

Stage-Specific Expression of a Homeo Box-Containing Gene in the Non-Segmented Sea Urchin Embryo. Dolecki, G.J. et al. (Honolulu, USA) - p. 925 [823]

Chromosome Walking Shows a Highly Homologous Repetitive Sequence Present in All the Centromere Regions of Fission Yeast. Nakaseko, Y. et al. (Kyoto, Japan) - p. 1011 [824]

Two Tandemly Linked Identical Genes Code for the Glycosomal Glyceraldehyde-Phosphate Dehydrogenase in *Trypanosoma brucei*. Michels, P.A.M. et al. (Brussels, Belgium) - p. 1049 [825]

Archaeobacteria: Transcription and Processing of Ribosomal RNA Sequences in *Halobacterium cutirubrum*. Chant, J. and Dennis, P. (Vancouver, Canada) - p. 1091 [826]

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The 'Light' and 'Medium' Subunits of the Photosynthetic Reaction Centre from *Rhodospseudomonas viridis*: Isolation of the Genes, Nucleotide and Amino Acid Sequence. Michel, H. et al. (Martinsried, FRG) - p. 1149 [827]

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Nucleotide Sequence of the Coding Region of the Mouse *N-myc* Gene. Taya, Y. et al. (Tokyo, Japan) - p. 1215 [829]

The Structure of the Mouse Glutathione Peroxidase Gene: The Selenocysteine in the Active Site is Encoded by the 'Termination' Codon, TGA. Chambers, I. et al. (Glasgow, UK) - p. 1221 [830]

Isolation of a Mouse cDNA Coding for a Developmentally Regulated, Testis-Specific Transcript Containing Homeo Box Homology. Wolgemuth, D.J. et al. (New York, USA) - p. 1229 [831]

T38 pre-mRNA is Transcribed from a Non-TATA Promoter and is Alternatively Spliced in Human T Cells. Tunnaciffe, A. et al. (Cambridge, UK) - p. 1245 [832]

Characterization of the Gene for the Microbody (Glycosomal) Triosephosphate Isomerase of *Trypanosoma brucei*. Swinkels, B.W. et al. (Amsterdam, NL)-p.1291[833]

Dramatic Events in Ciliate Evolution: Alteration of UAA and UAG Termination Codons to Glutamine Codons Due to Anticodon Mutations in Two *Tetrahymena* tRNAs. Hanyu, N. et al. (Tokyo, Japan) - p. 1307 [834]

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Primary Structure and Differential Expression of Glutamine Synthetase Genes in Nodules, Roots and Leaves of *Phaseolus vulgaris*. Gebhardt, C. et al. (Harpden, UK)-p.1429[836]

Evidence from Molecular Cloning that SPARC, a Product of Mouse Embryo Parietal Endoderm, is Related to an Endothelial Cell 'Culture Shock' Glycoprotein of M_r 43 000. Mason, I.J. et al. (London, UK) - p. 1465 [837]

Complete Sequence and Transcript Regulation of a Cell Adhesion Protein from Aggregating *Dictyostelium* Cells. Noegel, A. et al. (Martinsried, FRG) - p. 1473 [838]

Structure and Expression of the Chicken δ Nerve Growth Factor Gene. Ebendal, T. et al. (Uppsala, Sweden) - p.1483 [839]

The Primary Structure of Bovine Chromogranin A: a Representative of a Class of Acidic Secretory Proteins Common to a Variety of Peptidergic Cells. Benedum, U.M. et al. (Heidelberg, FRG) - p. 1495 [840]

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Genes Coding for the Elongation Factor EF-1 in *Artemia*. Lenstra, J.A. et al. (Leiden, NL) - p. 475 [844]

Nucleotide Sequence of the Promoter and Amino-Terminal Encoding Region of the *Escherichia coli* *pepN* Gene. Bally, M. et al. (Marseille, France) - p. 565 [845]

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Cloning and Nucleotide Sequence of a cDNA Encoding the Precursor of the Barley Toxin α -Hordeothionin. Ponz, F. et al. (Madrid, Spain) - p. 131 [846]

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Structure and Expression of Mouse Aldolase Genes. Brain-Specific Aldolase C Amino Acid Sequence is Closely Related to Aldolase A. Paolella, G. et al. (Napoli, Italy) - p.229 [847]

Nucleotide Sequence of the Bacteriophage T5 DNA Fragment Which Contains the Gene for tRNA^{ASP}. Shlyapnikov, M.G. et al. (Pushchino, USSR) - p. 285 [848]

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The Nucleotide Sequence of the *HEM1* Gene and Evidence for a Precursor Form of the Mitochondrial 5-Aminolevulinic Synthase in *Saccharomyces cerevisiae*. Urban-Grimal, D. et al. (Paris, France) - p. 511 [850]

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Nucleotide Sequence of the *ARGRII* Regulatory Gene and Amino Acid Sequence Homologies Between *ARGRII*, *PPRI* and *GAL4* Regulatory Proteins. Messenguy, F. et al. (Bruxelles, Belgium) - p. 77 [851]

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Cloning and Structural Characterization of the *Salmonella typhimurium* *pyrC* Gene Encoding Dihydroorotase. Neuhard, J. et al. (Kobenhavn, Denmark) - p. 335 [853]

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Nucleotide Sequence Characterization of a *Drosophila* Retrotransposon, *412*. Yuki, S. et al. (Fukuoka, Japan) - p. 403 [854]

Rainbow Trout Protamines. Amino Acid Sequences of Six Distinct Proteins from a Single Testis. McKay, D.J. et al. (Calgary, Canada) - p. 361 [855]

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Structure of a *Cephalosporium acremonium* mtDNA Replicator. Penalva, M.A. et al. (Madrid, Spain) - p. 92 [857]

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The Human Osteosarcoma Cell Line U-2 OS Expresses a 3.8 Kilobase mRNA which Codes for the Sequence of the PDGF-B Chain. Weich, H.A. et al. (Braunschweig, FRG)-p.344[858]

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Protease Gene Structure and *env* Gene Variability of the AIDS Virus. Yasunaga, T. et al. (Saitama, Japan)-p.145[859]

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A Tobacco Chloroplast DNA Sequence Possibly Coding for a Polypeptide Similar to *E. coli* RNA Polymerase β -Subunit. Ohme, M. et al. (Magoya, Japan) - p. 87 [860]

The Nucleotide Sequences of Chloroplast 4.5S rRNAs from Four Species of Plants, Celery (*Aquium graveolus*), Barley (*Hordeum vulgare*), Chinese Chive (*Allium tuberosum*) and Dayflower (*Commelina communis*). Zhen-Qi, C. et al. (Beijing, China) - p. 193 [861]

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Tick-Borne Encephalitis Virus Genome. The Nucleotide Sequence Coding for Virion Structural Proteins. Pletnev, A.G. et al. (Novosibirsk, USSR) - p. 317 [862]

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Characteristics of Two Types of Legumin Genes in the Field Bean (*Vicia faba* L. var. minor) Genome as Revealed by cDNA Analysis. Wobus, U. et al. (Gatersleben, GDR) - p. 74 [863]

Predicted Sequence of the Host-Protective Immunogen of Infectious Bursal Disease Virus. Hudson, P.J. et al. (Parkville, Australia) - p. 143 [864]

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Pig Kidney Na^+, K^+ -ATPase. Primary Structure and Spatial Organization. Ovchinnikov, Y.A. et al. (Moscow, USSR) - p. 237 [865]

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Molecular Cloning of cDNA of S100 α Subunit mRNA. Kuwano, R. et al. (Niigata, Japan) - p. 97 [866]

Characteristic Anticodon Sequences of Major tRNA Species from an Extreme Thermophile, *Thermus thermophilus* HB8. Hara-Yokoyama, M. et al. (Tokyo, Japan) - p. 149 [867]

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Isolation of Partial cDNAs for Rat Liver and Muscle Glycogen Phosphorylase Isozymes. Osawa, S. et al. (Worcester, USA) - p. 282 [868]

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Isolation and Partial Characterization of the Calcitonin Gene in a Lower Vertebrate. Predicted Structure of Avian Calcitonin Gene-Related Peptide. Minivielle, S. et al. (Paris, France) - p. 7 [869]

Mutations in the Guinea Pig Preproglucagon Gene are Restricted to a Specific Portion of the Prohormone Sequence. Seino, S. et al. (Chicago, USA) - p. 25 [870]

Molecular Cloning of the Avian β -Nerve Growth Factor Gene: Transcription in Brain. Wion, D. et al. (Paris, France) - p. 82 [871]

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Cloning of Rat Brain Protein Kinase C Complementary DNA. Ono, Y. et al. (Osaka, Japan) - p. 111 [872]

Isolation of a Human Ceruloplasmin cDNA Clone that Includes the N-Terminal Leader Sequence. Mercer, J.F.B. and Grimes, A. (Parkville, Australia) - p. 185 [873]

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Cloning, Sequencing and Transcription of an Inactivated Copy of *Bacillus amyloliquefaciens* Extracellular Ribonuclease (Barnase). Paddon, C.J. and Hartley, R.W. (Bethesda, USA) - p. 231 [874]

Nucleotide Sequence and Genome Organization of Bacteriophage S13 DNA. Lau, P.C.K. and Spencer, J.H. (Kingston, Canada) - p. 273 [875]

The Complete Sequence of *Bacillus* Phage ϕ 29 Gene 16: A Protein Required for the Genome Encapsidation Reaction. Garvey, K.J. et al. (Tucson, USA) - p. 311 [876]

The Sequence of Foot-and-Mouth Disease Virus RNA to the 5' Side of the Poly(C) Tract. Newton, S.E. et al. (Woking, UK) - p. 331 [877]

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The GC Clusters of the Mitochondrial Genome of Yeast and Their Evolutionary Origin. De Zamaroczy, M. and Bernardi, G. (Paris, France) - p. 1 [878]

Complete Nucleotide Sequence of an *EcoRI* 1.35-kb Repeated Element (ER-1) of Mouse: Homology with the Intervening Region Between Two Intracisternal A-Particle Genes. Yang, A.D. et al. (Baltimore, USA) - p. 33 [879]

Primary Structure of the Maltase Gene of the *MAL6* Locus of *Saccharomyces cerevisiae*. Hong, S.H. and Marmur, J. (Bronx, USA) - p. 75 [880]

Segmental Homology Between the Promoter Region of the Human Renin Gene and the Mouse *ren1* and *ren2* Promoter Regions. Soubrier, F. et al. (Paris, France) - p. 85 [881]

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Androgen Induction of Alcohol Dehydrogenase in Mouse Kidney. Studies with a cDNA Probe Confirmed by Nucleotide Sequence Analysis. Ceci, J.D. et al. (Columbia, USA) - p. 217 [882]

Nucleotide Sequence and Derived Amino Acid Sequence of a cDNA Encoding Human Muscle Carbonic Anhydrase. Lloyd, J. et al. (London, UK) - p. 233 [883]

Primary Structure of Human Salivary α -Amylase Gene. Nishide, T. et al. (Osaka, Japan) - p. 299 [884]

Cloning, Characterization and Nucleotide Sequences of Two cDNAs Encoding Human Pancreatic Trypsinogens. Emi, M. et al. (Osaka, Japan) - p. 305 [885]

Isolation and Characterization of Human Blood-Coagulation Factor X cDNA. Kaul, R.K. et al. (Chicago, USA)-p.311[886]

Cloning and Characterization of the *Schizosaccharomyces pombe* DNA Ligase Gene *CDC17*. Johnston, L.H. et al. (London, UK) - p. 321 [887]

Nucleotide Sequence of the R26 Chloramphenicol-Resistance Determinant and Identification of Its Gene Product. Dorman, C.J. et al. (Dundee, UK) - p. 349 [888]

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Structural and Functional Analysis of the Goat ϵ -Globin Genes. Menon, A.G. and Lingrel, J.B. (Cincinnati, USA) - p. 141 [889]

Cloning and Sequencing of mRNAs Coding for the Adult α -Globin Chains of the Salamander *Pleurodeles waltlii*. Flavin, M. et al. (Toulouse, France) - p. 159 [890]

The Nucleotide Sequence of the *malT* Gene Encoding the Positive Regulator of the *Escherichia coli* Maltose Regulon. Cole, S.T. and Raibaud, O. (Paris, France) - p. 201 [891]

Genomic Organization and Nucleotide Sequence of Two Corn Histone H4 Genes. Philipps, G. et al. (Strasbourg, France) - p. 225 [892]

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Cloning, Nucleotide Sequencing, and Expression of Tetanus Toxin Fragment C in *Escherichia coli*. Fairweather, N.F. et al. (Beckenham, UK) - p. 21 [893]

Evolution of the Regulatory Isozymes of 3-Deoxy-D-Arabinose-Heptulosonate 7-Phosphate Synthase Present in the *Escherichia coli* Genealogy. Ahmad, S. et al. (Binghamton, USA) - p. 146 [894]

Nucleotide Sequence Corresponding to Five Chemotaxis Genes in *Escherichia coli*. Mutoh, N. and Simon, M.I. (Pasadena, USA) - p. 161 [895]

Nucleotide Sequence of the Transcription Unit Containing the *aroL* and *aroM* Genes from *Escherichia coli* K-12. DeFeyter, R.C. et al. (Parkville, Australia)-p.233[896]

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Genes for *Bacillus megaterium* Small, Acid-Soluble Spore Proteins: Cloning and Nucleotide Sequence of Three Additional Genes from this Multigene Family. Fliss, E.R. et al. (Farmington, USA) - p. 467 [897]

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Isolation and Sequence Analysis of the Gene (*cpdB*) Encoding Periplasmic 2',3'-Cyclic Phosphodiesterase. Lin, J. et al. (Brisbane, Australia) - p. 1002 [898]

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Nucleotide Sequence of the Enterotoxin B Gene from *Staphylococcus aureus*. Jones, C.L. and Khan, S.A. (Pittsburgh, USA) - p. 29 [899]

Structural Features of Multiple *nifH*-Like Sequences and Very Biased Codon Usage in Nitrogenase Genes of *Clostridium pasteurianum*. Chen, K.C.-K. et al. (Blacksburg, USA) - p. 162 [900]

Characterization of the Developmentally Regulated *Bacillus subtilis* Glucose Dehydrogenase Gene. Lample, K.A. et al. (Bethesda, USA) - p. 238 [901]

Nucleotide Sequence of the *Escherichia coli* *motB* Gene and Site-Limited Incorporation of Its Product into the Cytoplasmic Membrane. Stader, J. et al. (Chicago, USA) - p. 244 [902]

Chromosomal Location and Nucleotide Sequence of the *Escherichia coli* *dapA* Gene. Richaud, F. et al. (Orsay, France) - p. 297 [903]

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Nucleotide Sequences of the R1-19 Plasmid Transfer Genes *traM*, *finP*, *traJ*, and *traY* and the *traY2* Promoter. Finlay, B.B. et al. (Edmonton, Canada) - p. 368 [904]

Lysogenic Conversion of Staphylococcal Lipase is Caused by Insertion of the Bacteriophage L54a Genome into the Lipase Structural Gene. Lee, C.Y. and Iandolo, J.J. (Manhattan, USA) - p. 385 [905]

Cloning and Nucleotide Sequencing of Genes for Three Small, Acid-Soluble Proteins from *Bacillus subtilis* Spores. Connors, M.J. et al. (Farmington, USA) - p. 417 [906]

Cloning and Characterization of *livH*, the Structural Gene Encoding a Component of the Leucine Transport System in *Escherichia coli*. Nazos, P.M. et al. (Ann Arbor, USA) - p. 565 [907]

Structural Genes Encoding the Thermophilic α -Amylases of *Bacillus stearothermophilus* and *Bacillus licheniformis*. Gray, G.L. et al. (San Francisco, USA) - p. 635 [908]

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Nucleotide Sequence of the Surface Exclusion Genes *traS* and *traT* from the IncF₁ *lac* Plasmid pED208. Finlay, B.B. and Paranchych, W. (Edmonton, Canada) - p. 713 [909]

Structure of a β -Galactosidase Gene of *Bacillus stearothermophilus*. Hirata, H. et al. (Osaka, Japan) - p. 722 [910]

Structural Similarity Between the Lepidoptera- and Diptera-Specific Insecticidal Endotoxin Genes of *Bacillus thuringiensis* subsp. "kurstak" and "israelensis". Thorne, L. et al. (San Diego, USA) - p. 801 [911]

Nucleotide Sequence, Transcript Mapping, and Regulation of the *RAD2* Gene of *Saccharomyces cerevisiae*. Madura, K. and Prakash, S. (Rochester, USA) - p. 914 [912]

Sequence of the *flaA* (*cheC*) Locus of *Escherichia coli* and Discovery of a New Gene. Kuo, S.C. and Koshland, D.E. (Berkeley, USA) - p. 1007 [913]

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