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| <p>A Comparative Transcription Map of the Murine Bare Patches (<i>Bpa</i>) and Striated (<i>Str</i>) Critical Regions and Human Xq28</p> | <p>Michael L. Levin, Aurobindo Chatterjee, Antonella Pragliola, Kim C. Worley, Manfred Wehnert, Olga Zhuchenko, Randall F. Smith, Cheng Chi Lee, and Gail E. Herman</p> | <p>465</p> |
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| <p>Novel Gene Transcripts Preferentially Expressed in Human Muscles Revealed by Quantitative Hybridization of a High Density cDNA Array</p> | <p>Geneviève Piétu, Olivier Alibert, Valérie Guichard, Bernard Lamy, Florence Bois, Elisabeth Leroy, Régine Mariage-Samson, Rémi Houlgatte, Pascal Soularue, and Charles Auffray</p> | <p>492</p> |
| <p>Construction of a YAC Contig Encompassing the Usher Syndrome Type IC and Familial Hyperinsulinism Loci on Chromosome 11p14–15.1</p> | <p>Radha Ayyagari, Ann Nestorowicz, Yan Li, Settara Chandrasekharappa, Craig Chinault, Peter van Tuinen, Richard J.H. Smith, J. Fielding Hejtmancik, and M. Alan Permutt</p> | <p>504</p> |
| <p>The Region Surrounding the <i>PKD1</i> Gene: A 700-kb PI Contig from a YAC-deficient Interval</p> | <p>William R. Dackowski, Timothy D. Connors, Ann E. Bowe, Vincent Stanton Jr., David Housman, Norman A. Doggett, Gregory M. Landes, and Katherine W. Klinger</p> | <p>515</p> |
| <p>Generation of a Transcriptional Map for a 700-kb Region Surrounding the Polycystic Kidney Disease Type 1 (<i>PKD1</i>) and Tuberous Sclerosis Type 2 (<i>TSC2</i>) Disease Genes on Human Chromosome 16p13.3</p> | <p>Timothy C. Burn, Timothy D. Connors, Terence J. Van Raay, William R. Dackowski, John M. Millholland, Katherine W. Klinger, and Gregory M. Landes</p> | <p>525</p> |
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COVER Expression profile of novel genes in human muscles. (*Top two panels*) Hybridization signals of 1339 human skeletal muscle cDNA clone inserts spotted on a high-density filter. (*Bottom two panels*) Distribution of the signal intensities after hybridization with the muscle cDNA probe. (For details, see Piétu et al., p. 492.)