Texas Genetics Society

Twenty-seventh Annual Meeting

Schedule of Events and Presentations

Thursday evening, March 23, 2000

5:00 - 8:00 Registration: Gallery

7:30 - 8:30 Keynote Address: Dr. William R. Brinkley, Dean, Graduate School of

Biomedical Sciences, Baylor College of Medicine, Houston. Graduate

Education in the Post-Genome Era

8:30 - 10:30 Reception: Houston I Room

Friday, March 24, 2000

8:00 - 8:15 Welcome and Announcements

8:15 - 9:15 Contributed Paper Session I: Stephen P. Daiger, Chair

8:15 [1] Recombination (gene conversion) mediated expansions of TRS occur independently of TRS orientation. J.P. Jakupciak and R.D. Wells. Institute of Biosciences and Technology; Center for Genome Research; Texas A&M University Health Science Center System; Texas Medical Center, Houston, Texas. [G]

8:30 [2] **Does p53 status predict genome-wide instability throughout development?** A. Maleki, ¹ L.C. Strong, ² B.W. Brown, ³ and M.J. Siciliano. ¹ Departments of ¹ Molecular Genetics, ² Experimental Pediatrics, and ³ Biomathematics, The University of Texas M.D. Anderson Cancer Center, Houston, Texas. [G]

8:45 [3] Mutations in a novel photoreceptor-pineal gene on 17p cause Leber congenital amaurosis (LCA4). Melanie M. Sohocki, Sara J. Bowne, Lori S. Sullivan, Seth Blackshaw, Constance L. Cepko, Annette M. Payne, Shomi S. Bhattacharya, Shagufta Khaliq, S. Qasim Mehdi, David G. Birch, Wilbur R. Harrison, Frederick F.B. Elder, John R. Heckenlively, and Stephen P. Daiger. Human Genetics Center, School of Public Health, and Department of Ophthalmology and Visual Science, the University of Texas-Houston Science Center, Houston, Texas; Department of Genetics and Howard Hughes Medical Institute, Harvard Medical School, Boston, Massachusetts; Department of Molecular Genetics, Institute of Ophthalmology, University College, London, United Kingdom; Dr. A.Q. Khan Research Laboratories, Biomedical and Genetic Engineering Division, Islamabad, Pakistan; Retina Foundation of the Southwest, Dallas, Texas; Department of Pathology and Laboratory Medicine,

University of Texas-Houston Health Science Center, Houston, Texas; ⁸Jules Stein Eye Institute, University of California, Los Angeles, California. [P]

9:00 [4] Homomeric channels of P932L mutations in muscle chloride channel gene (hClC-1) is associated with a severe myotomic myopathy. Shinichiro Nagamitsu, 1,2 Tohru Matsuura, 1,2 Mehrdad Khajavi, 1,2 Richard Armstrong, 1 Clifton Gooch, 1 and Tetsuo Ashizawa. 1,2 Department of Neurology, Baylor College of Medicine and 2Veterans Affairs Medical Center, Houston, Texas. 9:15 - 10:00 Invited Speaker: Dr. Richard Gibbs, Baylor College of Medicine, Houston. The Status of the Human Genome Sequencing Effort

10:00 - 10:30 Coffee Break

10:30 - 12:00 Contributed Paper Session II: James E. Womack, Chair

10:30 [5] **DNA base excision repair: soma vs. germline.** Gabriel W. Intano, ¹ C. Alex McMahan, ² John R. McCarrey, ³ Ronald B. Walter, ⁴ and Christi A. Walter. ¹ Departments of Cellular and Structural Biology and ² Pathology, The University of Texas Health Science Center at San Antonio, San Antonio, Texas; ³ Department of Genetics, Southwest Foundation for Biomedical Research, San Antonio, Texas; ⁴ Department of Chemistry, Southwest Texas State University, San Marcos, Texas. [G]

- 10:45 [6] Construction of a BAC contig across the NRC-1 region of chromosomal band 3p12. J.M. McDonald, S.T. Lott, M. Lovell, S. Naylor, and A.M. Killary. Division of Pathology and Laboratory Medicine, The University of Texas M.D. Anderson Cancer Center, Houston, Texas; Department of Cellular and Structural Biology, University of Texas Health Science Center at San Antonio, San Antonio, Texas. [G]
- 11:00 [7] **Development of a bacterial artificial chromosome contig at the putative tumor suppressor locus 5q13.3.** P.D. Castro, J.C. Liang, and L. Nagarajan. Department of Molecular Genetics, and the Division of Laboratory Medicine, The University of Texas M.D. Anderson Cancer Center, Houston, Texas. [G]
- 11:15 [8] Chromosome microdissection as a tool for the generation of high resolution marker maps at cattle trypanotolerant QTL regions on BTA 5q21-q24 and 7q14-q22. T. Goldammer, R.M. Brunner, S. Kang'a, P. Nilsson, J.E. Womack, O. Hanotte, and M. Schwerin. Department of Molecular Biology, Research Institute for the Biology of Farm Animals, Dummerstorf, Germany; Department of Molecular Biology, International Livestock Research Institute, Nairobi, Kenya; Department of Veterinary Pathobiology, Texas A&M University, College Station, Texas. [P]

- 11:30 [9] **Genetic control of diversity within species.** <u>Huang-Mo Sung</u> and Ronald E. Yasbin. *Department of Molecular and Cell Biology, University of Texas at Dallas, Richardson, Texas.* [G]
- 11:45 [10] Chronic exposure to high levels of ionizing radiation: what are the genetic consequences? <u>Jeffrey K. Wickliffe</u>, ¹ Brenda E. Rodgers, ¹ Ronald K. Chesser, ² and Robert J. Baker. ¹ Department of Biological Sciences, Texas Tech University, Lubbock, Texas; ² Department of Genetics, University of Georgia, Athens, Georgia. [G]
- 12:00 1:30 Lunch
- 1:30 2:15 Invited Speaker: Dr. Ronald B. Walter, Southwest Texas State University. **The Genetic Determinants of Tumorigenesis as Developed in** *Xiphophorus* **Fish**
- 2:15 3:15 Contributed Paper Session III: Charleen M. Moore, Chair
- 2:15 [11] Genetic variability in island populations: a comparison among congeneric species of migratory and endemic songbirds. Patrick W. Zwartjes. Department of Genetics, Southwest Foundation for Biomedical Research, San Antonio, Texas. [P]
- 2:30 [12] Evidence for genetic subdivisions in the desert shrew: *Notiosorex crawfordi*. Mark B. O'Neill and Robert J. Baker *Department of Biological Sciences, Texas Tech University, Lubbock, Texas.* [G]
- 2:45 [13] **Polynesian origins: new insights from the Y-chromosome.** Bing Su, ¹ Li Jin, ^{1,2} Peter Underhill, ³ Jeremy Martinson, ⁴ Nilmani Saha, ⁵ Stephen T. McGarvey, ⁶ Mark D. Shriver, ⁷ Jiayou Chu, ⁸ Peter Oefner, ³ Ranajit Chakraborty, ¹ and Ranjan Deka. ⁹ Human Genetics Center, University of Texas Health Science Center, Houston, Texas; ² Institute of Genetics, Fudan University, Shanghai, China; ³ Department of Genetics, Stanford University, Stanford, California; ⁴ Department of Anthropology, Oxford University, Oxford, England; ⁵ Department of Molecular Medicine, Australian National University, Canberra, Australia; ⁶ Department of Medicine and International Health Institute, Brown University School of Medicine, Providence, Rhode Island; ⁷ Department of Anthropology, Pennsylvania State University, University Park, Pennsylvania; ⁸ Institute of Medical Biology, Chinese Academy of Medical Sciences, Kunming, China; ⁹ Department of Environmental Health, University of Cincinnati, Cincinnati, Ohio. [P]
- 3:00 [14] Mitochondrial DNA sequence variation in baboons (*Papio hamadryas*): phylogeny and the evolution of social organization. <u>Timothy K. Newman</u>. Department of Genetics, Southwest Foundation for Biomedical Research, San Antonio, Texas. [P]

- 3:15 3:45 Coffee Break
- 3:45 4:30 Poster Session: San Antonio Room
- 4:30 5:15 Contributed Paper Session IV: Rodney S. Nairn, Chair
- 4:30 [15] **Multiple-trait-locus model for quantitative trait analysis.** <u>Jinying Zhao</u> and Momiao Xiong. *Human Genetics Center, The University of Texas at Houston, Houston, Texas.* [P]
- 4:45 [16] Linkage disequilibrium mapping of quantitative trait loci: multiple trait analysis. Joshua Akey, Li Jin, and Momiao Xiong. *Human Genetics Center, University of Texas, Houston, Texas.* [G]
- 5:00 [17] Characterization and mapping of a Xiphophorus fish RAB gene. K. Kelnar, D. Rains, S. Kazianis, and R. Walter. Department of Chemistry and Biochemistry, Southwest Texas State University, San Marcos, Texas; Research Division, The University of Texas M.D. Anderson Cancer Center, Science Park, Smithville, Texas. [G]
- 6:30 7:30 Reception (cash bar)
- 7:30 8:30 Banquet
- 8:30 9:30 Barbara H. Bowman Distinguished Texas Geneticist Award: Dr. Robert E. Ferrell

Saturday morning, March 25, 2000

- 8:15 9:00 Invited Speaker: Dr. David L. Rainwater, Southwest Foundation for Biomedical Research. **Genetics of Cardiovascular Disease**
- 9:00 10:00 Contributed Paper Session V: Candace M. Kammerer, Chair
- 9:00 [18] **Family history of stroke predicts stroke risk: the ARIC study.** A.C. Morrison, ¹ M. Fornage, ¹ D. Liao, ² and E. Boerwinkle. ¹ Human Genetics Center and Institute of Molecular Medicine, University of Texas Houston, Houston, Texas; ² Department of Health Evaluation Sciences, Penn State Hershey Medical Center, Hershey, Pennsylvania. [G]
- 9:15 [19] Nucleotide sequence of cyclophilin 40 and mapping of two cyclophilin-like loci in *Xiphophorus* fish. C. Daniels, D. Rains, L. Della Coletta, S. Kazianis, B. McIntire, R. Nairn, D. Morizot, and R. Walter. Department of Chemistry and Biochemistry, Southwest Texas State University,

San Marcos, Texas; ²Research Division, The University of Texas M.D. Anderson Cancer Center, Science Park, Smithville, Texas. [G]

- 9:30 [20] **Sequence analysis resources at the Baylor Human Genome Sequencing Center.** Michael P. McLeod, ^{1,2} Kim C. Worley, ¹ John Bouck, ¹ K. James Durbin, ¹ George M. Weinstock, ^{1,2} Richard A. Gibbs. ¹ Department of Molecular and Human Genetics, Human Genome Sequencing Center, Baylor College of Medicine, Houston, Texas; ² Department of Microbiology and Molecular Genetics, University of Texas, Houston, Texas. [G]
- 9:45 [21] **Partial nucleotide sequence determination and mapping of a** *Xiphophorus* **fish** *NF1* **gene.** <u>A. Wheeler</u>, ¹ D. Rains, ¹ C. Daniels, ¹ S. Kazianis, ² and R. Walter. ¹ Department of Chemistry and Biochemistry, Southwest Texas State University, San Marcos, Texas; ²Research Division, The University of Texas M.D. Anderson Cancer Center, Science Park, Smithville, Texas. [G]
- 10:00 10:30 Coffee Break
- 10:30 12:00 Contributed Paper Session VI: Chair to be named
- 10:30 [22] **E12** is a novel gene isolated from a differential screen of the early mouse embryo. Qiyong Hu and Richard Behringer. Department of Molecular Genetics, The University of Texas M.D. Anderson Cancer Center, Houston, Texas. [G]
- 10:45 [23] **Characterization of the PMP22 gene promoters.** M. Hai, S.I. Bidichandani, and P.I. Patel. Departments of Molecular and Human Genetics, Neurology, and Division of Neuroscience, Baylor College of Medicine, Houston, Texas. [G]
- 11:00 [24] **Analysis of the human cartilage oligomeric matrix protein** (**COMP**) **promoter region.** Kerry B. Gunning, ¹ Michelle Deere, ¹ Catherine Rhoades Hall, ¹ Xiaowen Liang, ¹ Amy L. Ridall, ² and Jacqueline T. Hecht. ¹ Department of Pediatrics, University of Texas Medical School at Houston, Houston, Texas; ² Departments of Basic Sciences and Prosthodontics, Dental Branch, University of Texas Health Science Center, Houston, Texas. [P]
- 11:15 12:00 Invited Speaker: Dr. Jacqueline T. Hecht, University of Texas Medical School at Houston, Houston. **Pseudoachondroplasia: Too Much of a Good Thing**
- 12:00 1:00 Business meeting; awards for best papers.
- 1:00 Adjourn