

Other publications from data generated from NASR:

Yang H, Chen X and Wong W-H. "Completely Phased Genome Sequencing Through Chromosome Sorting." PNAS 108 (2011): 12-17. [Acknowledgement of Yan PS for sequencing, advice and discussion.] Full text: <http://www.pnas.org/content/108/1/12.full>

Hsu P-Y, et al. "Estrogen-Mediated Epigenetic Repression of Large Chromosomal Regions Through DNA Looping." Genome Res 20 (2010): 733-744. Full text: <http://genome.cshlp.org/content/20/6/733.full>

Taslim C, et al. "Comparative Study on ChIP-seq Data: Normalization and Binding Pattern Characterization." Bioinformatics 25 (2009): 2334. Full text: <http://bioinformatics.oxfordjournals.org/content/25/18/2334.full>

Camerlengo T, et al. "Enabling Data Analysis on High-Throughput Data in Large Data Depository Using Web-Based Analysis Platform – A Case Study on Integrating QUEST with GenePattern in Epigenetics Research." IEEE International Conference on Bioinformatics and Biomedicine November 2009.

Reichert B, et al. "Concerted Action of Aldehyde Dehydrogenases Influences Depot-Specific Fat Formation" Molecular Endocrinology. 2011 May;25(5):799-809. Epub 2011 Mar 24. (NASR provided all the Quantitative PCR [qPCR] data).

Ben Rodriguez and Pearly Yan gave a 2-hour tutorial on 'Next Generation Sequencing: Biological and Technological Information Crucial to Effective Data Analysis' at the **Great Lakes Bioinformatics Conference 2011**, May 2-4, 2011, at Ohio University.