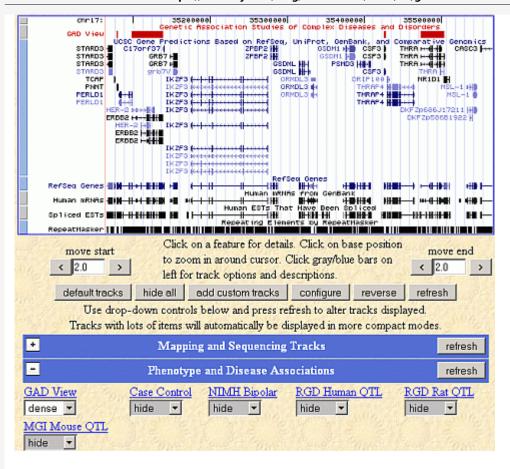
ALLERGY, SOFTWARE

GENOME BROWSER NOW WITH GADVIEW TRACK

3.06.2008

Just reveived an email from the creators of the Genetic Association Database (GAD)

... Your published genetic association study has been included in the recent update of the NIH based, Genetic Association Database (GAD), the database of <u>human genetic</u> association studies. This update is in cooperation with the CDC HuGENet. This increases the gene-based content of GAD from 28,000 to approximately 40,000 records, thus making GAD one of the largest gene based collections of medically relevant disease-gene based information publicly available. Along with this update we have added MeSH disease and geographic annotation, as well as links to the Comparative ToxicoGenomics Database. In addition, in collaboration with UCSC we have developed a public track for the Genetic Association Database in the <u>UCSC Genome Browser</u> system, called GAD View, in their new section on Phenotype and Disease Associations. This GAD track allows viewing of disease based genetic association information in the context of cross species genome sequence, human polymorphism, and other annotation found in the UCSC genomic browser. Could you please check your record in the GAD, http://geneticassociationdb.nih.gov, correct any errors, and add any missing information that may be available. Accurate information of this type is of critical importance in the understanding and analysis of the genetics of common human disease...



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