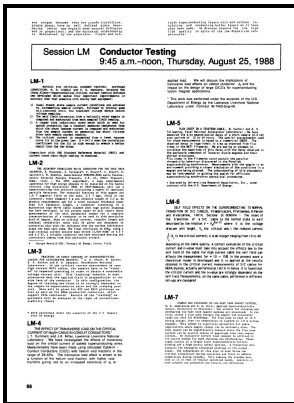


Voltage noise properties of bi-epitaxial and proximity effect Josephson junctions fabricated from yttrium barium copper oxide

University of Birmingham - Bioinformatics and Genomics @ CRG



Description: -

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Notes: Thesis (Ph.D) - University of Birmingham, School of Physics and Space Research, Faculty of Science.
This edition was published in 1996



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Tags: #Bioinformatics #and #Genomics #@ #CRG

Bioinformatics and Genomics @ CRG

To maximize reliability and coverage, we retrieved gene trees from PhylomeDB, Ensembl, TreeFam and Orthogroups.

Bioinformatics and Genomics @ CRG

We use existing tools, but we also design and implement novel computational tools and databases, and we are also interested in basic research in algorithms. This first published genome of a basal hemimetabolous insect, provides an out-group for comparison with other sequenced insects, and paves the way to elucidate molecular mechanisms in Negative selection prevents the fixation of deleterious alleles. .

Bioinformatics and Genomics @ CRG

In an letter to Nature published today we show that for mt-tRNAs this is not always the case. Two of the groups in our programme Gabaldón and Guigó have participated in the analysis of the genomic sequence of the pea aphid *Acyrtosiphon pisum*, published this week in PLoS Biology. In addition we built maximum likelihood trees for orthologous gene families in OrthoMCL, COG, and EggNOG.

Bioinformatics and Genomics @ CRG

The Groups at CRG's Bioinformatics and Genomics Program have strong collaborations with the CRG's Bioinformatics core and with other experimental groups at the CRG. It contains phylomes from model Human, Yeast, *Drosophila*, *Arabidopsis*, but also alternative models such as the fungal pathogen *Candida albicans*, or the insects *T.*

Bioinformatics and Genomics @ CRG

A review article detailing and contrasting different theories on the evolution of gene duplications has been in Nature Reviews Genetics.

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