

International Conference on Advances in Interdisciplinary Statistics and Combinatorics,
October 12–14, 2007, University of North Carolina at Greensboro, Greensboro, North
Carolina, USA

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**SIMULTANEOUS CONFIDENCE INTERVALS FOR ODDS
RATIOS IN CANDIDATE GENE STUDIES.**

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Candidate gene studies routinely involve construction of multiple confidence intervals for odds ratios for the various haplotypes. Confidence intervals are often preferred to hypothesis tests so that the practical significance of any observed significances can also be determined. Unfortunately, these one-at-a-time intervals are often utilized to make overall conclusions, such as that one particular haplotype has the largest odds of disease. Overall conclusions such as these require an adjustment for multiplicity. We provide several methods which adjust for multiplicity and compare their performance.

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