A Comparison Study of Sib-Trait Linkage Analysis Methods

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Abstract

Many procedures have been created to test linkage between a quantitative trait and a marker locus in sibling pairs. Past research has proven some methods to be stronger than other methods based on certain parameters. This paper analyszes four procedures and tests their type I error and power. The four methods we analyzed are the original Haseman-Elston methods using the squared differences (1972), Wrights's squared sums method (1997), the revisited Haseman-Elston method using the cross product (2000) and the Unified Haseman-Elston method using weighted slope estimates.