



**BAYLOR
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July 27, 2004

Dr. John H. Werren
Professor, Dept of Biology
University of Rochester
Rochester, NY 14627-0211

Dear Jack,

I am writing this letter to express my support for the sequencing of the *Nasonia* genome. The sequencing of this genome is compelling because it complements the honeybee genome sequence already produced at Baylor College of Medicine, provides insight into the parasitic wasps, important for biological control of agricultural pests, and finally, has the potential to be an excellent model for the study of polygenetic traits due to the ease of genetic manipulation of a haploid-diploid organism.

I wish you the best of luck in the approval of this white paper.

Sincerely,

A handwritten signature in black ink, appearing to read "R. Gibbs".

Richard A. Gibbs, PhD
Director, Human Genome Sequencing Center
Wofford Cain Professor of Molecular and Human Genetics