

Statistical method will analyze important, poorly studied areas of human genome

Each year, more and more pieces of the human genome puzzle fall into place, but large holes still remain. Researchers at the University of Wisconsin-Madison hope to fill in many more pieces with a new \$1.1 million grant from the National Human Genome Research Institute (NHGRI). The grant will support a School of Medicine and Public Health team of researchers who have created new computational tools to analyze important yet poorly studied areas of the human genome.

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