

New approach for efficient analysis of emerging genetic data

With the ability to sequence human genes comes an onslaught of raw material about the genetic characteristics that distinguish us, and wading through these reserves of data poses a major challenge for life scientists. Researchers at Roswell Park Cancer Institute (RPCI) and the Center for Human Genome Variation at Duke University Medical Center (DUMC) have developed an approach for analyzing data that can help researchers studying genetic factors in disease to quickly cull out relevant genetic patterns and identify variants that lead to particular disorders.

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