

大型矩阵求逆的实现与应用

宋端磊, 鹿 涛, 乔园园*

南开大学化学学院, 天津 300071

E-mail: yuanyuanqiao@nankai.edu.cn

矩阵求逆在科学计算中具有重要应用价值。由于计算机软硬件的限制, 传统的计算方法在 PC 上很难实现大型矩阵求逆运算。本文提出了一种求解方法, 一方面用内存映射文件存储数据, 另一方面用 OpenMP 并行库加快求解速度。运算表明求解速度显著提升, 求解矩阵规模明显增大。作为关键的计算步骤, 成功地运用在乳腺癌相关基因的研究中。

Table 1. Maximum size of matrix using different methods

Method	Gauss-Jordan	MatLab PINV()	Our Method
Maximum Matrix	2,000 x 2,000	8,000 x 8,000	15,000 x 15,000

Table 2. Performance of Our method

Matrix size	7,000 x 7,000	9,000 x 9,000	13,000 x 13,000	15,000 x 15,000
CPU-time /minute	34	70	203	310

关键词: 矩阵求逆; 内存映射; OpenMP

参考文献:

- [1] 徐士良, C 语言常用算法程序集. 北京: 清华大学出版社出版, 2004
- [2] The OpenMP specification for parallel programming. <http://www.openmp.org/>. March 27, 2008
- [3] 王祥雒, 李毅, Linux 中基于 mmap() 的共享存储实现研究, 计算机应用, 2006, 26(z2):307-309,320

Inverse Matrix by Memory Map and OpenMP

Dunlei Song, Tao Lu, Yuanyuan Qiao*

College of Chemistry, Nankai University, Tianjin 300071

Matrix inversion is important for scientific computaion. Traditional methods are limited by the specifications of hardware and operating system. Here, a combined method using mameory map with OpenMP that improves the space and speed of the inversion computation, and increases calculated size of martrix significantly. As a key step in calculation, it was successfully applied to explore genes correlatons for breast cancer research.