专论与综述

大规模酵母双杂交技术研究蛋白质相互作用的应用

吴志豪, 王 建, 贺福初

(军事医学科学院放射与辐射医学研究所, 北京 100850)

收稿日期 2006-1-22 修回日期 2006-3-28 网络版发布日期 2006-11-13 接受日期

摘要

简介了酵母双杂交技术原理,总结了酵母双杂交技术大规模筛选蛋白质相互作用的基础、应用及存在的问题。因为大规模酵母双杂交技术结果有大量假阳性及假阴性问题,因此,有条件情况下有必要同时开展其他方法的大规模蛋白相互作用研究,以构建规模更大可信度更高的蛋白质相互作用网络图。

关键词 大规模 酵母双杂交 蛋白质相互作用

分类号 0933

The Application of Large-scale Yeast Two-hybrid to Study Protein — Protein Interaction

WU Zhi-Hao, WANG Jian, HE Fu-Chu

(Beijing Institute of Radiation Medicine, Beijing 100850, China)

Abstract

<P>The principle of yeast two-hybrid were summarized, the basis for large scale yeast two-hybrid screening of protein-protein interaction was given and the main application of yeast two-hybrid in large scale screening of protein-protein interaction was presented. The problems of large scale yeast two-hybrid screening of protein-protein interaction were also discussed. Because there were a lot of false positives and false negatives in the result of protein-protein interaction obtained by large-scale yeast two-hybrid screening, the employment of other methods to study protein-protein interaction in large scale in parallel was proposed if possible.

Key words <u>large scale</u> <u>yeast</u> <u>two-hybrid</u> <u>protein-protein interaction</u>

DOI: 10.1360/yc-006-1627

扩展功能

本文信息

- ▶ Supporting info
- ▶ <u>PDF</u>(0KB)
- **▶[HTML全文]**(0KB)
- ▶参考文献

服务与反馈

- ▶把本文推荐给朋友
- ▶加入我的书架
- ▶加入引用管理器
- ▶复制索引
- ▶ Email Alert
- ▶文章反馈
- ▶浏览反馈信息

相关信息

▶ <u>本刊中 包含"大规模"的</u> 相关文章

▶本文作者相关文章

- 吴志豪
- 王 建
- 贺福初