研究报告

利用放射杂交克隆板进行猪*FMR*1等5个基因间的连锁分析

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培更

放射杂交技术在人类图谱(包括ESTs、STSs和微卫星)的构建中已证明是一种非常有效的方法。根据人类基因组X染色体上FMR1、IDS、FATE、BGN、F8A等5个基因的信息资源,用已构建的猪/仓鼠96个放射杂交细胞系分析猪染色体该5个基因间的连锁关系。结果显示:当LOD值为4时,FMR1、IDS、FATE、BGN、F8A都处于同一个连锁群内;当LOD值为5时,FMR1、IDS处于同一个连锁群,FATE和BGN在同一个连锁群内,而F8A单独处于一个连锁群中。

关键词 <u>猪;放射杂交;杂交体克隆板;基因连锁分析</u> 分类号 **0953**

Linkage Analysis of Five Genes in Pigs Using Radiation Hybrid Clone Panel

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Abstract

<DIV align=center>RH (radiation hybrid) has proved to be an effective method in constructing human genome maps (including ESTs, STSs and microsatellites). In this study, based on the information of five human genes (<I>FMR</I>1, <I>IDS</I>, <I>FATE</I>, <I>BGN</I>, <I>F8A</I>) on the X chromosome, the linkage relationship of these five genes in pigs were analyzed by a panel of 96 radiation hybrid cell lines. The results showed that <I>FMR1, IDS, FATE, BGN, F8A</I> were in the same linkage group, when LOD was set at 4. When LOD was set at 5, <I>FMR</I>1 and <I>IDS</I> were in the one group, <I> FATE</I> and <I>BGN</I> in the other group, and <I>F8A</I> was in a group by itself.

Key words pig radiation hybrid hybrid clone panel gene linkage analysis

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