

用染色体原位抑制杂交法研究人和猕猴染色体同源性

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摘要 本文用生物素标记的人类1号、2号、4号染色体NDA探针进行染色体原位抑制(chromosomal in situ suppression, 简称CISS)杂交以研究人和猕猴染色体的同源性。结果表明:人1号染色体与猕猴1号染色体同源。其中与猕猴1pter→1q33的同源程度高,与猕猴1q33→1qter的同源程度相对较低;人2号染色体与猕猴13号染色体长臂、9号染色体长臂和部分短臂同源;人4号染色体与猕猴2号染色体同源。结合染色体带型比较分析,本文对人和猕猴染色体的演化关系进行了探讨,该研究进一步证明了染色体重排可能是灵长类染色体进化的主要机制。

关键词 [生物素标记探针,原位杂交,染色体,染色体进化](#)

分类号

Stuies of The Homology of Chromosomes Between Human Being and Rhesus Monkey with Chromosomal In situ Suppression Hybridization

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Abstract

Chromosomal in situ suppression (CISS) hybridization of biotin labeled DNA libraries for human chromosomes 1,2 and 4 was used to investigate chromosome homology between human being and rhesus monkey. The results demonstrate that chromosome 1 in human being and rhesus monkey is homologous, of which the 1pter→1q33 of rhesus monkey is highly homologous with a cognate region of chromosome 1 in human, while the 1q33→1qter is relatively low; Chromosome 2 in human shows homology with the long arm of chromosome 13 as well as the long arm and the partial short arm of chromosome 9 in the rhesus, and chromosome 4 in human is identical to chromosome 2 in rhesus monkey. Combined with a comparative analysis of banding patterns of chromosomes, derivation relationship of chromosomes between man and the rhesus are discussed. Our present data provide a definitive proof that chromosomal rearrangements may be a major mechanism of chromosomal evolution in primates.

Key words [Biotinylated probe](#) [In situ hybridization](#) [Chromosome](#) [Chromosomal evolution](#)

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