长期人工选择对群体遗传结构的影响——一项计算机模拟研究

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摘要 运用数字计算机模拟研究了长期人工选择的反应。考虑了不同的基因作用模型(加性、显性 与超显性)、群体含量(小群体Nf=Nm=10, Ne=20; 大群体Nf=Nm=100, Ne=200)、遗传力水 平以及位点间连锁强度等因素共54个组合,270个群体各49代的选择反应。结果表明,对于 三种基因作用模型,只有强烈的连锁(相邻位点之间交换率为r=0.01)在小群体中对选择反 应有显著影响,它可以阻碍有利基因的固定,加速它们的丢失,从而降低选择反应。中等强 度的连锁(r=0.1)或大群体中的紧密连锁都没有显著的作用。大群体中的选择反应要明显 地高于小群体中的,特别是遗传力低的性状。本文还讨论了不同选择系杂交以获致进一步进 展的问题,讨论了长期选择理论研究结果与实际模拟的偏差。

关键词 长期选择,选择反应,计算机模拟

分类号

Effects of Longterm Selection on the Genetic Structure of Populations —A Simu lation Study I.Simulation Models and Selection Responses

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Abstract

Monte Carlo method has been utilized to inverstigate responses to longterm selec tion. The simulation involves 3 gene effect models (additive, dominant, and over-dom inant), two population sizes, three linkage intensities, and three heritability lev els, thus making 54 combinations. For each combination, 5 replicate populations are s elected over 49 consecutive non-overlapping generations. The results demonstrate that under all the gene effect models, linkage is important only in the case of s mall populations and intense linkage, when it reduces responses significantly due to hampering the fixation of favourable alleles and accelerating their loss. Lin kage has no apparent effect if it is not very tight or if the population is larg e. Much greater responses are achieved in the large populations than in the small ones, especially for traits with low heritability. Among those discussed are prob lems of crossing selected lines to obtain extra responses, and discrepancy of som e theoretic results on selection limits.

Key words longterm selection Responses Computer simulation

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