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畜牧—研究报告

鸡IL-2基因的克隆、序列分析及蛋白结构预测

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摘要:

根据GenBank中已发表的鸡白细胞介素2(IL-2)mRNA基因序列,利用Premier 5.0软件设计一对特异性引物,采用RT-PCR技术,以ConA刺激的鸡外周血淋巴细胞为材料,从总RNA中扩增出鸡IL-2基因。经琼脂糖凝胶电泳显示扩增片段约为432 bp,分离纯化片段,克隆入pMD-18T载体,转化DH5a感受态细胞,获得阳性重组质粒,经酶切鉴定,测序结果显示,我们得到了长432bp的基因,含有一个 432bp的开放阅读框,编码143个氨基酸。生物软件分析结果表明该序列与GenBank中发表的鸡的IL-2的核苷酸序列同源性为98.8%,与黄牛、马、鸡、绵羊、牛、犬、人、山羊、鼠、水牛的核苷酸同源性分别为31.2%、28.2%、27.3%、30.6%、26.2%、31.7%、30.1%、30.8%、26.6%、30.6%。与GenBank中发表的鸡IL-2编码的氨基酸序列同源性为95.8%,与上述动物的氨基酸序列同源性分别为7.6%、7.6%、9.7%、7.6%、6.9%、10.4%、11.8%、7.6%、9.0%、7.6%、10.4%。这表明鸡的IL-2基因被成功克隆,它存在着种属特异性。这为进一步研究该基因的生物学作用特别是利用IL-2增强DNA疫苗的免疫效果奠定了基础。

关键词: 蛋白结构预测

Cloning and sequence analysis of gallus IL-2 gene and prediction of protein structure

Abstract:

According to GenBank in the IL-2 cDNA sequences, using Premier 5.0 software design a pairs of specific primer, the local Dorking inject 100 μ g / ml ConA, total RNA was extracted from feeding 24 hours lymphocyte from spleen, by RT-PCR cloning colony IL-2 cDNA fragment. The DNA fragments were analyzed use DNAStar software between different species identity for comparison. Sequence analysis showed that IL-2 gene has an open reading frames for 432 nucleotide acids, encoding 143 amino acids and most of signal peptide, compared with published on Genbank, the nucleotide identity was 98.8%、31.2%、28.2%、27.3%、30.6%、26.2%、31.7%、30.1%、30.8%、26.6%、30.6%、compared with published on Genbank, the Amino acids was 95.8%, 7.6%、7.6%、9.7%、7.6%、6.9%、10.4% 11.8%, 7.6%, 9.0%, 7.6%, 10.4% with cattle, horse, cat, sheep, bovine, canis, people, capra, mice, cow of IL-2 sequence comparison The gallus IL-2 gene was successfully cloned, there is a species-specific. And construct a based on further study the biological effects of IL-2gene in particular the use of IL-2 enhance the immune effect of DNA vaccine.

Keywords: Prediction of protein structure

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