

研究简报

中国寄生虫学与寄生虫病杂志

CHINESE JOURNAL OF PARASITOLOGY AND PARASITIC DISEASES

ISSN 1000-7423 CN 31-1248/R

主管: 中华人民共和国国家卫生和

主办:中华预防医学会

中国疾病预防控制中心寄生虫病 预防控制所

★ 返回首页 期刊介绍 | 编 多

期刊介绍 | 编 委 会 | 稿约 | 欢迎订阅 | 广告合作 | 获奖情况 | 检索库收录情况 | 联系我们 | Englis

最新目录 | 下期目录 | 过刊浏览 | 高级检索

中国寄生虫学与寄生虫病杂志 » 2013, Vol. 31 » Issue (6):447-449 DOI:

<< Previous Articles | Next Articles >>

刚地弓形虫ROP11基因的克隆及生物信息学分析

张晓磊,张进顺,贾晓晖,徐云鹏,张颖,王春苗,王燕,卢致民,赵建玲,贾天军

河北北方学院病原生物与免疫学研究所,张家口 075000

Cloning and Bioinformatics Analysis of Rhoptry Protein 11 of Toxoplasma gondii

ZHANG Xiao-lei, ZHANG Jin-shun, JIA Xiao-hui, XU Yun-peng, ZHANG Ying, WANG Chun-miao, WANG Yan, LU Zhi-min, ZHAO Jian-ling, JIA Tian-jun

Institute of Pathogen Microbes and Immunology, Hebei North University, Zhangjiakou 075000, China

摘要 参考文献

相关文章

Download: PDF (999KB) HTML 1KB Export: BibTeX or EndNote (RIS) Supporting Info

摘要 提取刚地弓形虫(Toxoplasma~gondii)RH株速殖子总RNA,根据棒状体蛋白11(ROP11)全长编码序列(登录号为DQ077905)的开放阅读框设计引物并进行逆转录PCR(RT-PCR)扩增,PCR产物经EcoR I 和Not I 酶切后与原核表达载体pGEX-6P-2连接,重组质粒转化大肠埃希菌(E.~coli)XL-Blue,阳性菌落经PCR和双酶切鉴定,并测序。对所得序列进行生物信息学分析。结果显示,RT-PCR扩增产物约为1500 bp。菌落PCR及双酶切结果正确。测序结果显示,获得的ROP11基因片段为1548 bp(登录号为KC456639),与GenBank上已有的弓形虫ROP11序列相比,序列一致性为99%。生物信息学分析发现,ROP11编码蛋白质的预期相对分子质量为 $M_{\rm F}$ 57020,包括有12个保守结构区域,其前26个氨基酸残基构成信号肽,丝氨酸/苏氨酸蛋白激酶催化区域位于170~511氨基酸,且有2个潜在的N-糖基化位点。

关键词: 刚地弓形虫 棒状体蛋白11 原核表达 生物信息学分析

Abstract: Total RNA was extracted from tachyzoites of RH strain of *Toxoplasma gondii*. The open reading frame of ROP11 gene was amplified by using a pair of specific primers designed according to the coding sequence of ROP11 gene (Accession No. DQ077905). The RT-PCR product was digested by restriction enzyme *Eco*R I and *Not* I, and then ligated into a pGEX-6P-2 vector. The recombinant plasmid was transferred into *E. coli* XL-Blue. The positive clones was selected by colony PCR, and confirmed by the double restriction enzyme digestion and sequencing. The RT-PCR product was 1 548 bp. The recombinant plasmid was confirmed by colony PCR and double restriction enzyme digestion. Sequencing results showed that the obtained ROP11 gene was 1 548 bp. (Accession No. KC456639). There was a high sequence consistency (99%) between the obtained ROP11 gene sequence and the Toxoplasma ROP11 gene from GenBank. Bioinformatics analysis showed that the ROP11 protein (Mr 57 020) consisted of the signal peptide (amino acids 1-26), 12 conservative domains, a serine/threonine protein kinase catalytic domain (amino acids 170-511), and two potential N-glycosylation sites.

Keywords: Toxoplasma gondii Rhoptry protein 11 Prokaryotic expression Bioinformatics analysis

Service

- ▶ 把本文推荐给朋友
- ▶ 加入我的书架
- ▶ 加入引用管理器
- ▶ Email Alert
- ▶ RSS

作者相关文章

- ▶ 张晓磊
- ▶ 张进顺
- ▶ 贾晓晖
- ▶ 徐云鹏
- 张颖
- ▶ 王春苗
- ▶ 王燕
- ▶卢致民
- ▶ 赵建玲
- ▶ 贾天军

引用本文:

张晓磊, 张进顺, 贾晓晖, 徐云鹏, 张颖, 王春苗, 王燕, 卢致民, 赵建玲, 贾天军.刚地弓形虫ROP11基因的克隆及生物信息学分析[J] 中国寄生虫学与寄生虫病杂志, 2013, V31(6): 447-449

ZHANG Xiao-Lei, ZHANG Jin-Shun, JIA Xiao-Hui, XU Yun-Peng, ZHANG Ying, WANG Chun-Miao, WANG Yan, LU Zhi-Min, ZHAO Jian-Ling, JIA Tian-Jun. Cloning and Bioinformatics Analysis of Rhoptry Protein 11 of *Toxoplasma gondii*[J], 2013, V31(6): 447-449

Copyright 2010 by 中国寄生虫学与寄生虫病杂志