

英文

首页 | 期刊介绍 | 投稿指南 | 排行榜 | 光荣榜 | 编委会 | 期刊订阅 | 留言板 | 联系我们 | 自荐编委/审稿人 | 广告合作

徐景野,闫鹏,杨元斌,章丹阳,胡荣华.宁波地区食品中致病菌监测与流行株分析[J].中国食品卫生杂志,2015,27(5):562-568.

宁波地区食品中致病菌监测与流行株分析

Inspection of pathogenic bacteria in food and analysis of epidemic strains in Ningbo

投稿时间 : 2015-06-19

DOI :

中文关键词: 食品 食源性致病菌 鉴定 分型 耐药性 流行株 宁波**Key Words:** Food foodborne pathogen identification genotyping resistance epidemic strains Ningbo**基金项目:**宁波市重大(重点)项目(2013C51014) ;宁波市创新团队项目(2012B82018)

二维码 (扫一下试试看 !)

作者	单位	E-mail
徐景野	宁波市疾病预防控制中心浙江宁波 315010	xujy@nbcdc.org.cn
闫鹏	宁波市疾病预防控制中心浙江宁波 315010	
杨元斌	宁波市疾病预防控制中心浙江宁波 315010	
章丹阳	宁波市疾病预防控制中心浙江宁波 315010	
胡荣华	宁波市卫生监督所浙江宁波 315010	

摘要点击次数: 448**全文下载次数: 486****中文摘要:**

了解宁波地区食品中致病菌检出情况和菌株的耐药性,发现其流行优势菌。方法 致病菌检测采用直接分离与增菌分离相结合的方法;细菌鉴定采用生化筛选和API等方法;细菌分型采用诊断血清和PFGE基因分型;药敏试验采用K-B法,耐药基因检测采用PCR法。结果 6 812份食品样品中检出目标菌7类12种,共2 331份,检出率为34.22%。致病性弧菌检出数最高,其次为沙门菌和致病性气单胞菌。副溶血性弧菌与其他致病菌差异有统计学意义($P<0.01$),分离出10个血清群和29个PFGE型,其中O6、O5血清群和PFGE 1型是副溶血性弧菌的主要优势流行型。检出的致病菌对大多数抗生素敏感,其中3株气单胞菌为带aacc耐药基因的多重耐药菌。结论 宁波地区食品中致病菌种类较多,易引起食源性疾病;各致病菌均有流行优势株,副溶血性弧菌是最主要的流行优势株;血清分型和PFGE型能发现优势菌,但均有一定的局限性。

Abstract:

To study the contamination of pathogenic bacteria in food and their resistance to antibiotics, and to find the dominant epidemic strains in Ningbo. Methods Direct separation and culture enrichment were both used for pathogen isolation. The identification of bacteria was performed by Vitek 2 compact and API method. Sub-typing of bacterial was performed by serology and PFGE. Antibiotics resistances were tested by Kirby-Bauer method. Antibiotic resistance genes were detected by PCR. Results 7 categories and 12 kinds of bacteria (2 331 strains) were detected from 6 812 food samples. Most of them were pathogenic Vibrio, the second prevalent was Salmonella, and the third was Aeromonas. Vibrio parahaemolyticus was the major epidemic foodborne pathogenic strain in Ningbo. The detection rate had significant difference ($P<0.01$) compared to other pathogens. 10 serum groups and 29 PFGE types were identified, in which O6 and O5 serum groups and type 1 of PFGE were the dominant epidemic. Pathogenic bacteria detected were sensitive to most antibiotics. There were three Aeromonas strains which were resistant to multi-drug with aacc genes. Conclusion Foodborne pathogenic bacteria in Ningbo were various. All categories of pathogenic bacteria had their dominant epidemic strains, while Vibrio parahaemolyticus was the major contributor of foodborne diseases. Serotype and PFGE typing of bacteria could be applied to find their dominant epidemic strains, but either of them had certain limitations. The broad-spectrum antibiotic treatment was effective for foodborne bacterial illness.

[查看全文](#) [查看/发表评论](#) [下载PDF阅读器](#)

您是第27862810位访问者 今日一共访问41次

版权所有 : 《中国食品卫生杂志》编辑部 京ICP备12013786号-3

地址 : 北京市朝阳区广渠路37号院2号楼501室 邮编:100022

E-mail:spws462@163.com 电话/传真 : 010-52165456/5441 (编辑室) 010-52165556 (主编室)

未经授权禁止复制或建立镜像

技术支持:北京勤云科技有限公司

