

[1]尚伟龙,胡启文,胡珍,等.重庆地区耐甲氧西林金黄色葡萄球菌主要流行克隆更替的机制研究[J].第三军医大学学报,2014,36(08):735-739.

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重庆地区耐甲氧西林金黄色葡萄球菌主要流行克隆

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Title: Mechanisms in replacement of major methicillin-resistant *Staphylococcus aureus* epidemic clones in Chongqing

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关键词: [耐甲氧西林金黄色葡萄球菌](#); [克隆替代](#); [生长曲线](#); [利福平耐药](#); [复方新诺明耐药](#)

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摘要: 目的 探讨重庆地区耐甲氧西林金黄色葡萄球菌菌株ST239-MRSA-III-t030替代ST239-MRSA-III-t037克隆的机制。方法 选择重庆地区分离的ST239-MRSA-III-t030和ST239-MRSA-III-t037临床株,检测各菌株的生长曲线,观察菌落形态,分析菌株的药物敏感谱;再通过体外、体内竞争实验和交叉抑制实验深入探讨ST239-MRSA-III-t030替代ST239-MRSA-III-t037的机制。结果 ST239-MRSA-III-t030菌株较ST239-MRSA-III-t037有更短的生长迟缓期,后者在固体培养基上生长的菌落相对较小,且都对复方新诺明耐药,对利福平敏感,这与ST239-MRSA-III-t030菌株不同(大多数对复方新诺明敏感,对利福平都耐药)。ST239-MRSA-III-t030菌株在体外和体内都表现出比ST239-MRSA-III-t037菌株更强的生存竞争优势,但两克隆的菌株间并不存在明显的交叉抑制现象。结论 ST239-MRSA-III-t030克隆的生存竞争优势和独特的耐药表型可能是其成功替代ST239-MRSA-III-t037而成为第一流行克隆的原因。

Abstract: Objective To explore the mechanisms underlying the replacement of methicillin-resistant *Staphylococcus aureus* (MRSA) strain ST239-MRSA-III-t037 by ST239-MRSA-III-t030 in Chongqing, China. Methods Eight ST239-MRSA-III-

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t030 and ST239-MRSA-III-t037 isolates collected from Chongqing were divided into four t030/t037 pairs. The independent growth rates of all isolates were then determined. The competitive growth rates of the bacterial pairs were detected *in vitro* and *in vivo*, and cross inhibition assays were also performed.

Results The ST239-MRSA-III-t030 isolates were rifampicin-resistant, whereas the ST239-MRSA-III-t037 strains were all rifampicin-sensitive and trimethoprim/sulfamethoxazole-resistant. The ST239-MRSA-III-t030 strains had shorter lag phases than the ST239-MRSA-III-t037 isolates. The ST239-MRSA-III-t030 strains could out-compete their rivals in a competition assay *in vitro* as well as in a mouse model. No significant cross-inhibition effect was observed between ST239-MRSA-III-t030 and ST239-MRSA-III-t037. The growth of ST239-MRSA-III-t037 strains on the BHI nonselective plates had smaller colonies as compared to ST239-MRSA-III-t030 strains. **Conclusion** ST239-MRSA-III-t030 replacing ST239-MRSA-III-t037 as the first predominant clone may result from its higher growth rate and the specific drug resistance profiles.

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