

实验方法

酪氨酸激酶抑制剂伊马替尼耐药K562细胞系的建立及其耐药特征

冯帆, 张琼, 刘洪英, 李松, 王莉莉

军事医学科学院毒物药物研究所药物合成研究室, 北京 100850

收稿日期 2011-12-26 修回日期 2012-5-26 网络版发布日期 2012-8-21 接受日期

摘要 目的 建立酪氨酸激酶抑制剂伊马替尼(STI-571)耐药K562细胞系并研究其耐药特征。方法采用含递增浓度伊马替尼的培养基培养K562细胞,诱导耐伊马替尼K562细胞系(K562R)。通过细胞生长曲线、细胞形态和细胞凋亡分析确定耐药细胞是否形成;采用MTT法观察耐药细胞的耐药谱;通过半定量PCR和Western印迹法检测相关基因及蛋白的表达,并通过基因测序分析K562R细胞B细胞受体-c-Abelson (BCR-ABL)激酶区基因序列。结果 成功地将伊马替尼高敏感的K562细胞诱导成K562R,伊马替尼抑制K562和K562R细胞存活的IC₅₀值分别为0.01±0.00和(2.35±0.01) μmol·L⁻¹,耐药倍数为235.0倍。K562R具有一定的交叉耐药性,对高三尖杉酯碱、长春新碱和对柔红霉素的耐药倍数分别为13.2,63.2和11.8倍。K562R可对抗伊马替尼诱导的细胞凋亡,伊马替尼1.0 μmol·L⁻¹培养24 h K562和K562R细胞凋亡率分别为72.1%和18.2%。耐药形成机制研究表明,与K562细胞相比,K562R细胞BCR-ABL基因、多药耐药基因(MDR)和p-糖蛋白基因(p-GP)的表达均明显增加。此外,K562R细胞BCR-ABL基因的第696位发生A→C点突变,该位点突变导致激酶区第231位氨基酸由赖氨酸取代原有的天冬酰胺。结论 成功建立了耐伊马替尼细胞系K562R。K562R细胞具有交叉耐药性和对抗伊马替尼诱导的细胞凋亡等特征。K562R细胞BCR-ABL基因序列发生点突变,MDR和p-GP等基因表达亦明显升高。

关键词 [伊马替尼](#) [酪氨酸激酶抑制剂](#) [K562细胞](#) [抗药性, 肿瘤](#)

分类号 [R965.2](#)

扩展功能

本文信息

- ▶ [Supporting info](#)
- ▶ [PDF\(1966KB\)](#)
- ▶ [\[HTML全文\]\(0KB\)](#)

参考文献

服务与反馈

- ▶ [把本文推荐给朋友](#)
- ▶ [加入我的书架](#)
- ▶ [加入引用管理器](#)
- ▶ [复制索引](#)
- ▶ [Email Alert](#)
- ▶ [文章反馈](#)
- ▶ [浏览反馈信息](#)

相关信息

- ▶ [本刊中包含“伊马替尼”的相关文章](#)

本文作者相关文章

- [冯帆](#)
- [张琼](#)
- [刘洪英](#)
- [李松](#)
- [王莉莉](#)

Establishment of human K562 cell line resistant to tyrosine kinase inhibitor imatinib and its resistance characters

FENG Fan, ZHANG Qiong, LIU Hong-ying, LI Song, WANG Li-li

Department of Drug Design and Synthesis, Institute of Pharmacology and Toxicology, Academy of Military Medical Sciences, Beijing 100850, China

Abstract

OBJECTIVE To screen imatinib(STI-571) resistant human K562 cell line and to study the imatinib resistance mechanisms. **METHODS** Human K562 cells were induced by culture in the medium with gradually increasing imatinib concentrations to establish the imatinib-resistant K562 (K562R) cells. Biological and pharmacological characters of the cells were examined by MTT method, and cell phenotype and apoptosis were determined. The expression of B cell receptor-c-Abelson (BCR-ABL), multi-drug resistance (MDR) and p-glycoprotein(p-GP) genes was examined by semi-QPCR and Western blotting. **RESULTS** Compared to K562 cells ($IC_{50}=0.01 \mu\text{mol} \cdot \text{L}^{-1}$), K562R cells exhibited 235.0 fold resistance to imatinib ($IC_{50}=2.35 \mu\text{mol} \cdot \text{L}^{-1}$). Besides, K562R also became resistant to homoharringtonine (13.2 folds), vincristine (63.2 folds) and daunorubicin (11.8 folds). These resistance abilities consisted with the expression of BCR-ABL, MDR and p-GP genes. PCR results showed that the expression of BCR-ABL, MDR and p-GP genes increased in K562R cells compared with K562 cells. The DNA sequencing analysis showed that a A→C point-mutant occurred in the BCR-ABL gene sequence of K562R cells, and asparagine was submitted by lysine in the 231st site of kinase-domain. **CONCLUSION** K562R cells are successfully constructed. The expression of BCR-ABL, MDR and p-GP genes is up-regulated in K562R cells. There is a point-mutant in BCR-ABL gene of K562R cells.

Key words [imatinib](#) [tyrosine kinase inhibitor](#) [K562 cells](#) [drug resistance](#) [neoplasm](#)

DOI: 10.3867/j.issn.1000-3002.2012.04.017

通讯作者 王莉莉, E-mail: wangll63@126.com wangll63@126.com