密切相关。但是要证实 MTB139 基因与结核病耐多药性的相关性仍需扩大样本量及菌株来源。同时本课题组今后将深入探索 MTB139 基因的功能，以鉴定其是否参与结核分支杆菌的耐多药性，并阐明机制。如果这一推测得到印证，将有助于进一步揭示耐多药结核菌的基因表达特征及结核耐多药产生的基因调节机制，为广大结核患者带来福音。

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更 正

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