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External links Category:Ceramic materials Category:Engineering polymer materials Category:Inorganic polymers Category:Lithium compounds Category:Monomers Category:Transition metal oxides Category:E-number additives[Identification of the isolates of Mycobacterium tuberculosis complex by RFLP]. To study the prevalence of RFLP types of Mycobacterium tuberculosis complex (MTBC) in an area with a high rate of human TB in the southeast region of China. A total of 1 732 clinical isolates of MTBC from patients were collected from the 16 sentinel hospitals in 8 counties of 6 districts of Dongguan and 1 114 clinically isolated strains were used as the reference strains for this study. RFLP was performed using restriction enzymes, Hinf I and Bgll I, and the fragments were analyzed using genetic analysis software. Of the 1 732 isolates, 1250 (73.2%) were identified as M. tuberculosis. RFLP was performed on these isolates with the Bgll I and Hinf I restriction enzymes. With the Hinf I restriction enzyme, a total of 1 015 isolates (72.2%) were successfully genotyped into nine types and 111 strains (7.0%) into seven types; with the Bgll I restriction enzyme, a total of 1 042 isolates (66.7%) were successfully genotyped into 16 types and 274 strains (18.0%) into 15 types. Type 3 was the most prevalent type, followed by type 2, type 9, and type 8, which were all non-tuberculous Mycobacterium except for one isolate. Type 6, 1, 12 and 7 were identified in the Beijing family. Among patients with single TB, 95.2% of the cases were of type 3 and 9.6% of the cases were of type 2. Thirty-three patients harbored multiple TB strains. Twelve of them had mixed type 4, and the remaining twenty-one cases had multiple type 2 strains. In all, 90 patients were positive for tuberculous lymphadenitis. Among them, 29 (32.2%) were of type 1 and 64 (70.3%) were of type 2. The results of our study suggest that the strains causing multiple types of TB are isolated from different patients. The reference strains in our study are isolated from the Beijing family. These results indicate that the c6a93da74d

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