

台灣牛隻牛結核病流行現況

生物研究組

黃春申 助理研究員

摘要

牛結核病(bovine tuberculosis, bTB)是 *Mycobacterium bovis* (*M. bovis*)感染所造成的重要人畜共通傳染病，牛為主要宿主，但可感染多數哺乳動物。感染後依感染途徑會於頭胸部淋巴結或腹腔臟器產生特徵性結節病灶。皮內結核菌素試驗(Intradermal tuberculin test, ITT)是國際間貿易用的標準牛結核病檢驗方法，也是我國標準檢測方法。統計台灣 2019 年至 2021 年牛結核病陽性場案例，2020 年下半年起屠宰場發現多起病例，陽性場大增，再將 *M. bovis* 分離株以結核菌群常用之兩種基因分型方法 Spoligotyping 及 MIRU/VNTR (ETR-A、ETR-B 及 Qub11b 三個位置)進行分子流行病學分析，2021 年共得 12 個基因型別，為歷年之最。

Epidemiology of bovine tuberculosis in Taiwan cattle

Chen-Shen Huang

Abstract

Bovine tuberculosis (bTB) is an important zoonotic disease caused by *Mycobacterium bovis*. Although, cattle are considered to be the main hosts, most mammals can be infected by *M. bovis*. Depending on the route of infection, characteristic nodular lesions can be observed in lymph nodes of the head and chest, or in abdominal organs. The intradermal tuberculin test (ITT) is the standard method for testing bovine tuberculosis, in particular, to ensure the safety of the international trade market. It is also the standard method for bovine tuberculosis detection in Taiwan. We analyzed bTB positive cases in Taiwan from 2019 to 2021, and discovered, that since the second half of 2020, many cases have been found in slaughterhouses, and that the number of farms with positive cases increased. The *M. bovis* isolates were genotyped by two genotyping methods commonly used within the *Mycobacterium tuberculosis* complex for molecular epidemiological analysis: spoligotyping and the use of MIRU/VNTR typing (three loci, ETR-A, ETR-B, and Qub11b). A total of 12 genotypes were obtained in 2021, the most in history.