

Supplementary Material:

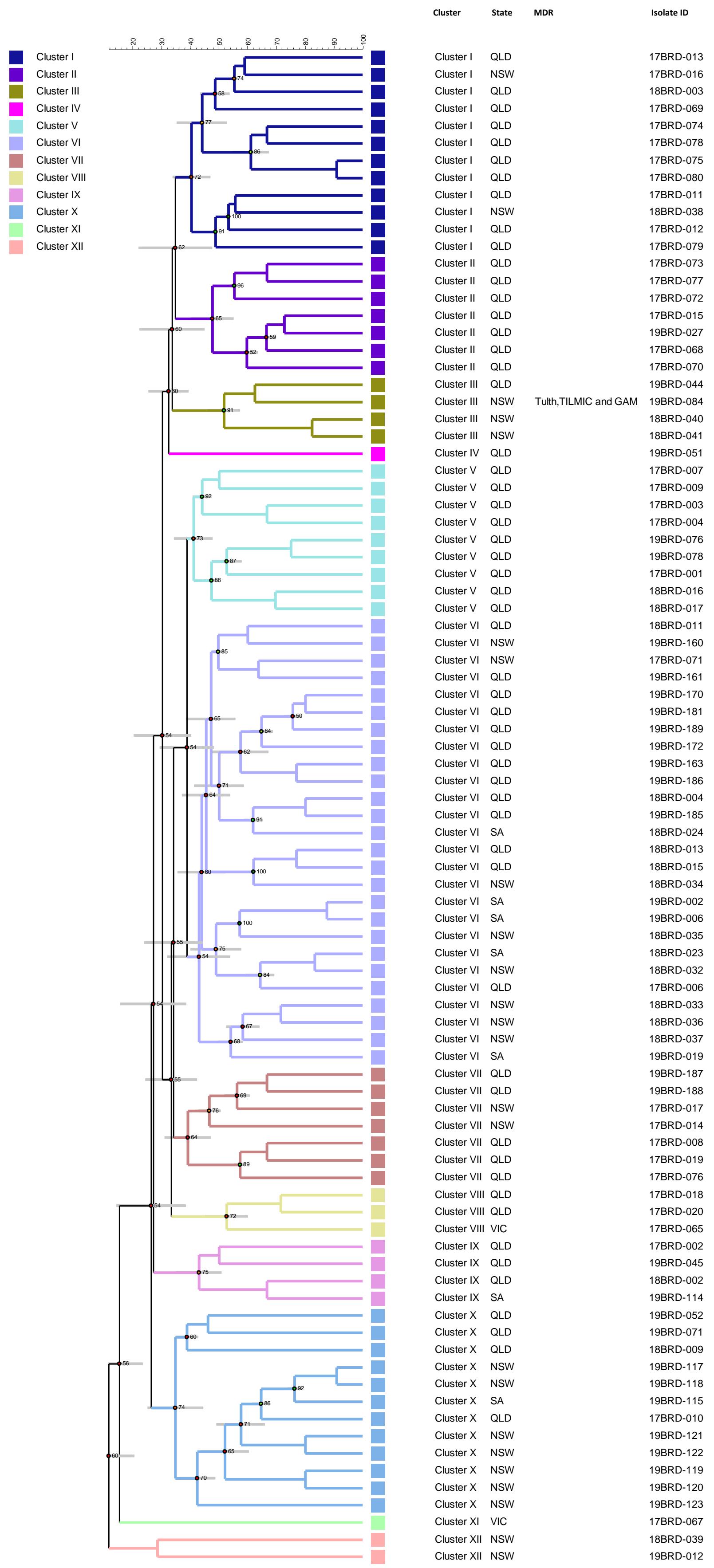


Figure S1. Random amplified polymorphic profiles and antimicrobial susceptibility profiles of 88 *M. haemolytica* strains. The dendrogram was generated with BioNumerics (Applied Maths) based on the Dice coefficient. The cluster number, Australian state of origin (NSW, New South Wales; QLD, Queensland; VIC, Victoria; SA, South Australia), multidrug-resistance (MDR) status (Tulth, tulthromycin, TLMIC, tilmicosin and GAM, gamithromycin resistant) and isolate identification number, are from right to left side, respectively.

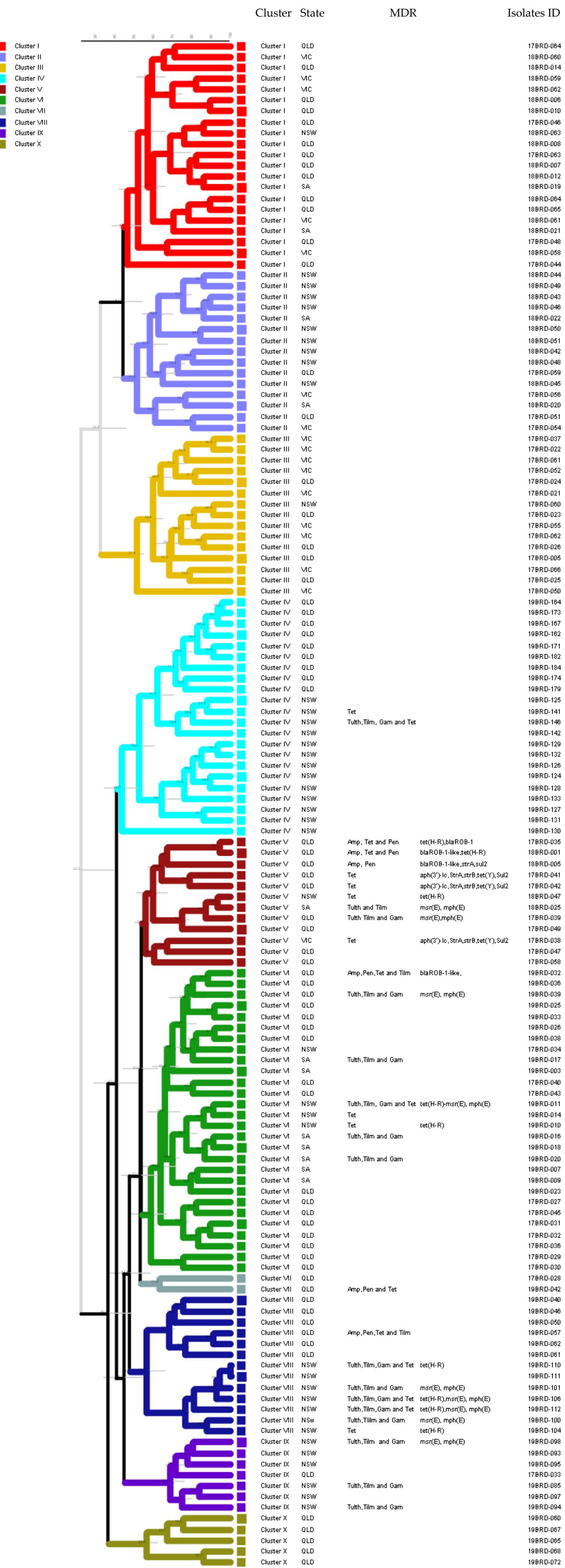


Figure S2. Random amplified polymorphic profiles and antimicrobial susceptibility profiles and antimicrobial susceptibility profiles of 140 *P. multocida* strains. The dendrogram was generated with BioNumerics program (Applied Maths) based on the Dice coefficient. The cluster number, Australian state of origin (NSW, New South Wales; QLD, Queensland; VIC, Victoria; SA, South Australia), multidrug-resistance (MDR) status (Tulth, tulthromycin, TLMIC, tilmicosin, GAM, gamithromycin, AMP, ampicilin and Pen, penicillin resistant) and isolate identification number, are from right to left side, respectively.