Table S2. Process of assigning genetic groups/populations to potential lineages within Queensland.

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| Number of potential lineages | Explanation | Graphical representation | Populations included |
| 2 | Major division within Qld | 12523411234561234567813212345678910123456789101112345678910111213 | 1 = MAR, TSFN, TSFS, TL, BRB, BER2 = HB, BER, TEW, LW, GC |
| 3 | Major division within top portion of tree |  | 1 = MAR, TSFN, TSFS2 = TSFS, TL, BRB, BER3 = HB, BER, TEW, LW, GC |
| 5 | Major divisions within bottom portion of tree |  | 1 = MAR, TSFN, TSFS, TL, BRB, BER2 = BER3 = HB4 = TEW, LW5 = GC |
| 6 | Major divisions within both top and bottom portion of tree |  | 1 = MAR, TSFN, TSFS2 = TSFS, TL, BRB, BER3 = BER4 = HB5 = TEW, LW6 = GC |
| 8 | Minor divisions within both top and bottom portion of tree |  | 1 = MAR2 = TSFN, TSFS3 = TSFS4 = TL, BRB, BER5 = BER6 = HB7 = TEW, LW8 = GC |
| 10 | All geographic localities separated within bottom portion of tree |  | 1 = MAR2 = TSFN, TSFS3 = TSFS4 = TL, BRB, BER5 = BER6 = HB7 = TEW8 = LW9 = GC110 = GC2 |
| 11 | All geographic localities separated within top portion of tree |  | 1 = MAR2 = TSFN3 = TSFS4 = TSFS5 = TL6 = BRB7 = BER8 = BER9 = HB10 = TEW, LW11 = GC |
| 13 | All geographic localities separated within entire tree |  | 1 = MAR2 = TSFN3 = TSFS4 = TSFS5 = TL6 = BRB7 = BER8 = BER9 = HB10 = TEW11 = LW12 = GC113= GC2 |