

Table 5 Pairwise Nei's genetic distance (below the diagonal) and F_{ST} (above the diagonal) between RM (r) and YM (y) subpopulations

| | CTr | CTy | BHr | BHy | ZJr | ZJy |
|-----|-------|-------|-------|--------|--------|--------|
| CTr | - | 0.004 | 0.033 | 0.018 | 0.034 | 0.009 |
| CTy | 0.216 | - | 0.072 | 0.052 | 0.076 | 0.011 |
| BHr | 0.108 | 0.293 | - | -0.010 | -0.003 | 0.003 |
| BHy | 0.167 | 0.331 | 0.124 | - | -0.009 | -0.012 |
| ZJr | 0.135 | 0.326 | 0.076 | 0.122 | - | 0.006 |
| ZJy | 0.161 | 0.291 | 0.136 | 0.181 | 0.131 | - |