

| | <i>M. tb</i> (n=41) | | <i>M. afr-M. mic-M. bov</i> and subspecies (n=56) | | | | <i>M. can</i> (n=2) and STB (n=2) | | | | | |
|----------------------------------|---------------------|----------------------|---|-----------------------|----|----------------------|-----------------------------------|--------------------|--|----|--|----------------------------|
| | PE | PGRS17 | PE | PGRS18 | PE | PGRS17 | PE | PGRS18 | | | | |
| No. of gene copies | | 41 | | 41 | | 56 | | 56 | | 3 | | 4 |
| No. haplotypes | | 3 | | 13 | | 7 | | 5 | | 1 | | 3 |
| No. of polymorphic sites | | 2 | | 18 | | 8 | | 9 | | 21 | | 14 |
| Number of observed transitions | | 2 | | 9 | | 5 | | 5 | | 10 | | 7 |
| Number of observed transversions | | 0 | | 9 | | 3 | | 3 | | 10 | | 6 |
| Pi(s) | | 0,00429 | | 0,02656 | | 0,003 | | 0,008 | | 0 | | 0.02035 |
| Pi(a) | | 0,00171 | | 0,00612 | | 0,003 | | 0,003 | | 0 | | 0.00528 |
| Mean Ka (Range) | | 0.0013 (0 to 0.0026) | | 0,006 (0 to 0,012) | | 0,00321 (0 to 0,008) | | 0,003 (0 to 0,006) | | 0 | | 0.0053 (0.004 to 0.008) |
| Mean Ks (Range) | | 0.0065 (0.0065) | | 0,027 (0,006 to 0,06) | | 0,00306 (0 to 0,005) | | 0,008 (0 to 0,015) | | 0 | | 0.02063 (0.0206 to 0.0207) |
| Mean Ka/Ks (Range) | | 0.2 (0 to 0.4) | | 0,273 (0 to 1,569) | | 1,019 (0 to 1,588) | | 0,293 (0 to 0,386) | | NA | | 0.258 (0.193 to 0.388) |