a- Royal Y-chromosome Lineage Mummies

| Sample No. | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | YGATA | DYS | DYS | DYS | Predicted |
|---------------|-----|------|-----|-------|-----|-----|--------|-----|-----|-----|-----|-----|-------|-----|-----|-----|-----------|
| | 456 | 3891 | 390 | 38911 | 458 | 19 | 385a,b | 393 | 391 | 439 | 635 | 392 | H4 | 437 | 438 | 448 | haplotype |
| Yuya | 15 | 14 | - | 29 | - | - | - | - | - | - | 20 | - | 11 | - | - | 24 | G2a* |
| Amenhotep III | 15 | 13 | 24 | 30 | 16 | - | -,14 | 13ª | 11 | - | - | - | 11ª | 14 | - | - | R1b** |
| KV55 | 15 | 13 | 24 | 30 | 16 | 14 | 11,14 | 13ª | 11 | 10 | 23 | 13 | 11ª | 14 | 12 | 19 | R1b* |
| (Akhenaten) | | | | | | | | | | | | | | | | | |
| Tutankhamun | 15 | 13 | 24 | - | 16 | - | 11,- | 13ª | 11 | | 23 | - | 11ª | 14 | - | - | R1b** |

b- Reference Mummies

| Sample No. | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | YGATA | DYS | DYS | DYS | Predicted |
|---------------------------|-----|------|-----|-------|-----|-----|--------|-----|-----|-----|-----|-----|-------|-----|-----|-----|------------------------|
| | 456 | 3891 | 390 | 38911 | 458 | 19 | 385a,b | 393 | 391 | 439 | 635 | 392 | H4 | 437 | 438 | 448 | haplotype ¹ |
| TT320- | 15 | 12 | 19 | - | 18 | 13 | - | 9ª | 8 | | 20 | - | 9ª | 15 | - | - | L* |
| CCG61065 | | | | | | | | | | | | | | | | | |
| Ramesses III ^b | 13 | 13 | 21 | 33 | - | 19 | 20 | 8 | 8 | | - | 17 | 13 | 14 | 10 | 20 | E1b1a* |
| Unknown ManE b | 13 | 13 | 21 | 33 | - | 19 | 20 | 8 | 8 | - | - | 17 | 13 | 14 | 10 | 20 | E1b1a* |

c- DNA Controls

| Sample No. | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | DYS | YGATA | DYS | DYS | DYS | Predicted |
|------------|-----|------|-----|-------|-----|-----|--------|-----|-----|-----|-----|-----|-------|-----|-----|-----|------------|
| | 456 | 3891 | 390 | 38911 | 458 | 19 | 385a,b | 393 | 391 | 439 | 635 | 392 | H4 | 437 | 438 | 448 | haplotype1 |
| DNA 007 | 15 | 13 | 24 | 29 | 17 | 15 | 11,14 | 13 | 11 | 12 | 24 | 13 | 13 | 15 | 12 | 19 | R1b*** |
| Staff 1 | 16 | 13 | 24 | 29 | 15 | 14 | 11,11 | 12 | 11 | 12 | 23 | 13 | 12 | 15 | 12 | 19 | R1b*** |
| Staff 2 | 14 | 11 | 24 | 27 | 19 | 13 | 13,17 | 12 | 10 | 12 | 25 | 11 | 11 | 14 | 11 | 19 | Q*** |
| Staff 3 | 16 | 13 | 24 | 29 | 15 | 14 | 11,11 | 12 | 11 | 12 | 23 | 13 | 12 | 15 | 12 | 19 | R1b*** |

Note: All female mummies were also tested with Y-chromosomal markers and did not yield any positive results.

selection. All probabilities exceeded 95%.

a Data already published in Hawass et al. (2010)

^{*}The haplogroup (hg) prediction was carried out through the website: www.hprg.com/hapest5/, by matching the concluded ancient Y-STR profiles versus those of the Mediterranean populations (meditp) as well as of the equal priors (eqp) selection (Athey 2006, 34-39). The predicted hg was suggested based on the probability that is more than 85%. The actual probabilities for the detected 4 ancient haplotypes were for the R1b hg (meditp: 99.9%; eqp: 99.6%), the G2a hg (meditp: 94.9%; eqp: 89.2%), the L hg (meditp: 92.9%; eqp: 98.4%) and E1b1a hg (meditp: 86.7%; eqp: 98.3%).

*** Since the weights of the various alleles and loci in haplogroup prediction are apparently unequal, the Amenhotep III-KV55-Tutankhamun male lineage haplotype was predicted based on the complete KV55 data rather than the partial data of Tutankhamun and Amenhotep III which gave different probabilities. Amenhotep III partial profile gave (meditp: R1a: 55.5%, R1b: 31.5% and E1b1b: 12.5%; eqp: R1a: 84.1%, R1b: 9.3% and E1b1b: 5.4%), while that of Tutankhamun gave (meditp: R1b: 71.7%, R1a: 28% and E1b1b: 0%; eqp: R1a: 65.9%, R1b: 32.9% and E1b1b: 0%).

***The contemporary staff samples were matched only to the Mediterranean populations selection and the 007 profile to the Northwest European