

Table S5. Bottleneck analysis in lion populations using the standardized differences test and the stepwise mutation model (SMM).

	<u>Standardized differences test</u>
	<u>SMM</u>
GIR	*
UGA	ns
KEN	ns
SER-I	***
SER-II	**
SER-III	***
NGC	*
KRU	*
BOT-I	*
BOT-II	ns
NAM	ns

ns - not significant; \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ .