



Fig. S5 Stutter analysis was performed on a subset of 50 samples that were selected according to the total number of reads (selection criteria: category I (blue rhombuses): $\leq 63,500$ reads; category II (green circles): $\geq 199,000$ reads). Mean stutter ratios in percent are shown for all STR markers (error bars denote stutter ranges). The x-axis of all figures relate to marker specific alleles, e.g. “D22S1045 alleles”. Allele calls without symbol indicate alleles without detectable stutter products.





