

Figure S1. Probability of obtaining the correct genotype at 3 or more replicates for a homozygous locus. using Ethanol and NAP as a preservation method. Calculations are based on the amplification success (AS) and the genotyping success (GS) calculated for each preservation method (**Table 1**). Horizontal line marks probability of 0.95.

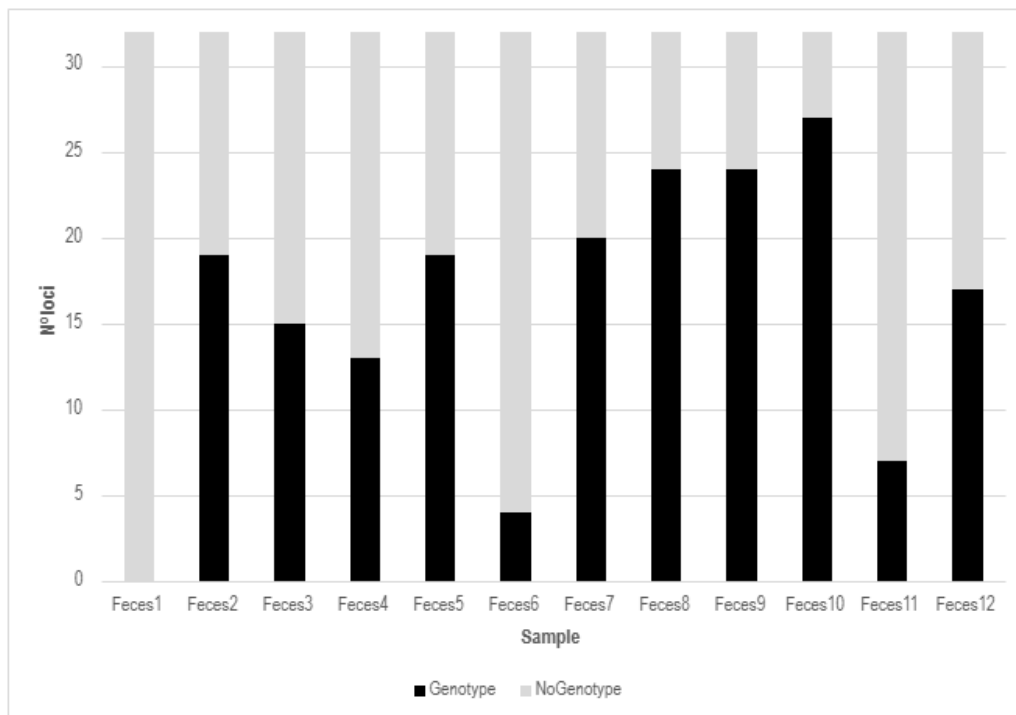


Figure S2. Completeness of the reference genotype obtained for the microsatellite panel for each fecal sample. Number of loci with a genotype out of the 32 microsatellite loci in the panel for each sample is indicated with the black bar (**Table S4**). Loci without a genotype, marked in grey, were those with no data or considered as 'ambiguous' (see Methods).