

Suppl. Figure 1.

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Amplification of hypervariable MIRU-VNTR loci (1982, 3232, 3820, 4102) from samples with Beijing and non-Beijing lineage spoligotypes
Lanes 1 to 5, Gabon samples with Beijing lineage spoligotype SIT1; Lane 7, Gabon sample with Beijing lineage spoligotype SIT260; lanes 6, 8, 10, 11 and 12, Gabon strains with spoligotypes not belonging to the Beijing lineage; Lane 9 was left intentionally empty; lanes A and B, strains from Paris Bichat-Claude Bernard Hospital with spoligotypes belonging to the Beijing lineage; lanes C and D, strains from Paris Bichat-Claude Bernard Hospital with spoligotypes not of the Beijing lineage

