# Distribution and Risk of Mycolactone-Producing Mycobacteria Transmission within Buruli Ulcer Endemic Communities in Côte d'Ivoire

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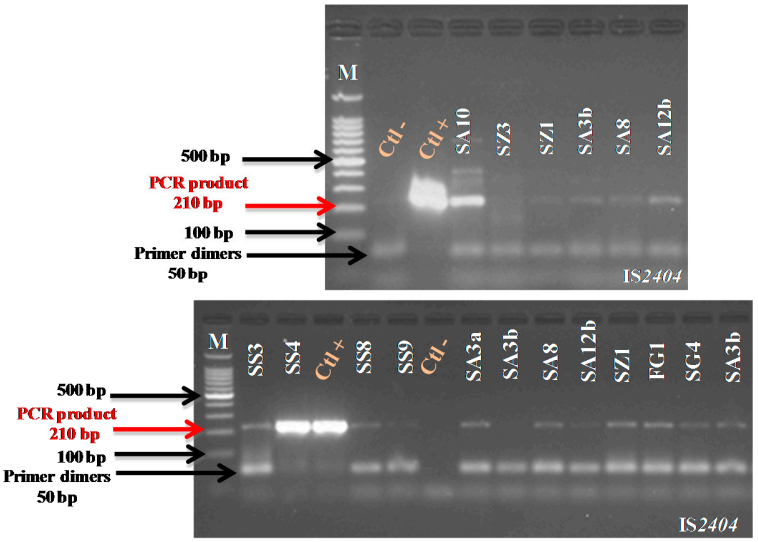


**Figure S1.** Amplification of IS*2404* from some clinical samples. M: 100 bp DNA ladder; Ctl+: positive control (*M. ulcerans* strain); Ctl-: negative control (sterile water); SA10-SA12b, SS3, SS4, SS8, SS9, SA3b-SG4: clinical samples tested.

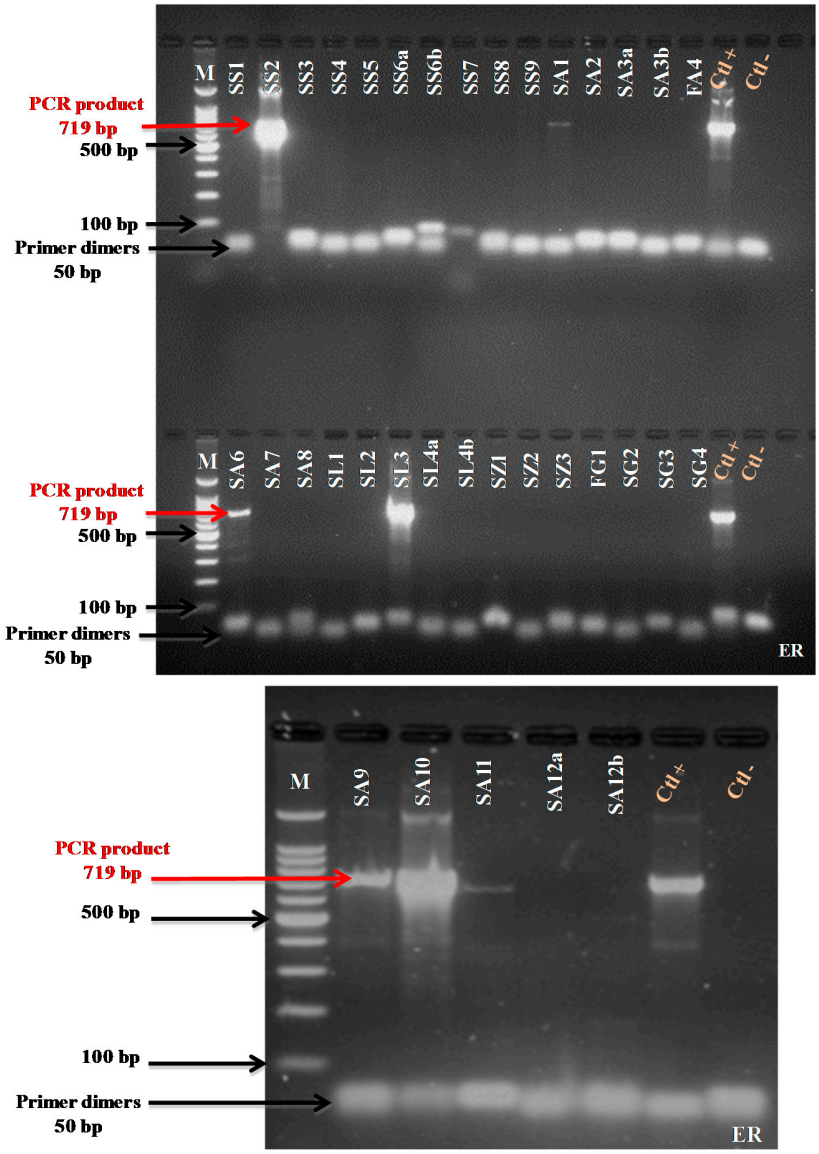


**Figure S2.** Amplification of enoyl reductase (ER) gene from clinical samples. M: 100 bp DNA ladder; Ctl+: positive control (*M. ulcerans* strain); Ctl-: negative control (sterile water); SS1-FA4, SA6-SG4, SA9-SA12b: clinical samples tested.

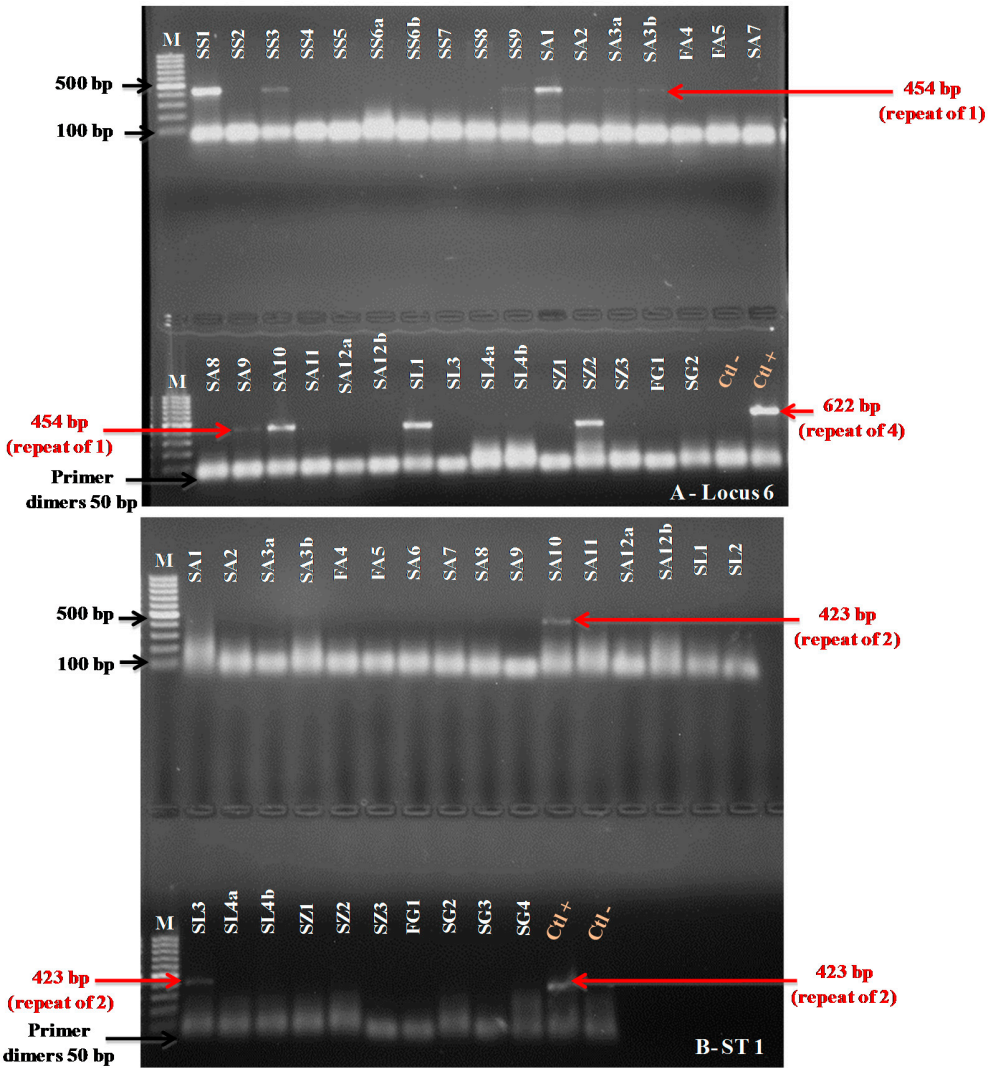


**Figure S3.** Amplification of VNTR loci from clinical samples. A: Locus 6 amplification; B: ST1 amplification. M: 100 bp DNA ladder; Ctl+: positive control (*M. Marinum* DL strain); Ctl-: negative control (sterile water); SS1-SG4: clinical samples tested.

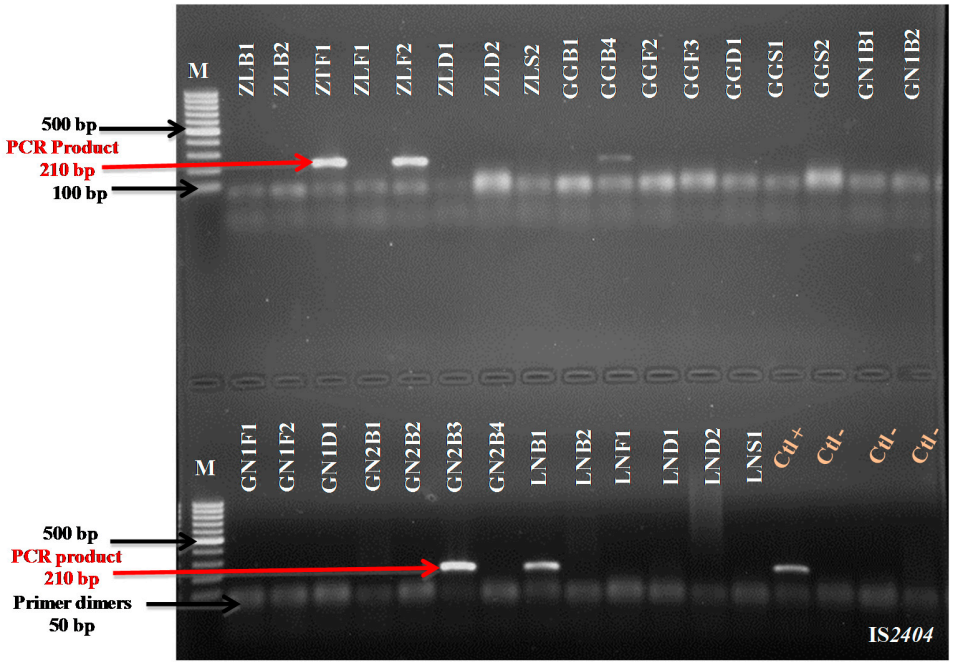


**Figure S4.** Amplification of IS*2404* from positive 16S rRNA environmental samples. M: 100 bp DNA ladder; Ctl+: positive control (*M. Marinum* strain); Ctl-: negative controls (sterile water); ZLB1-LNS1: environmental samples tested.

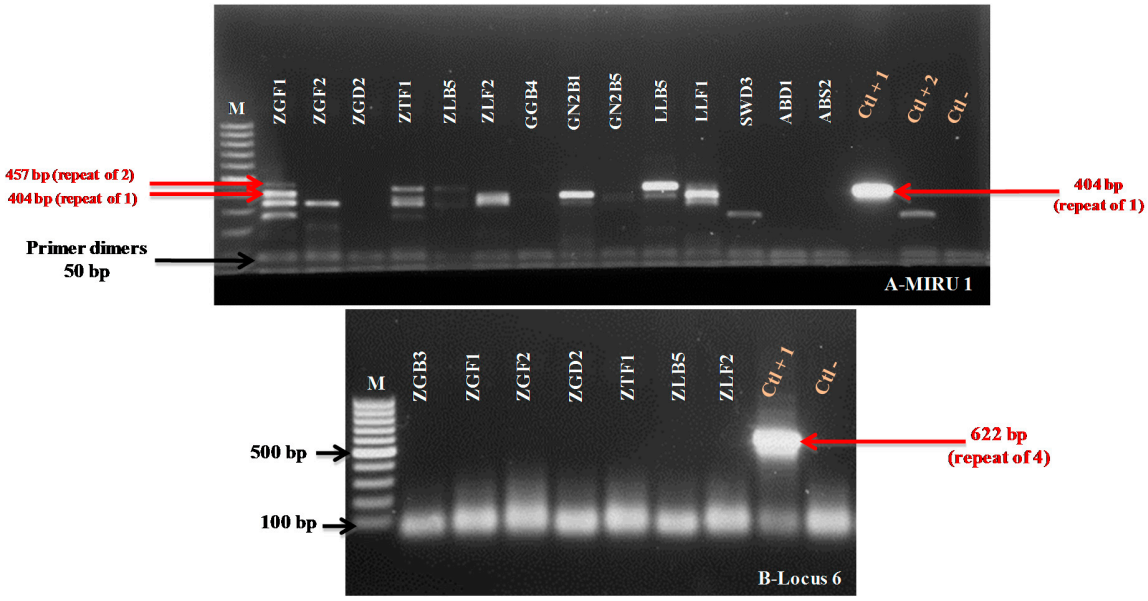


**Figure S5.** Amplification of VNTR loci from environmental samples. A: MIRU 1 amplification; B: Locus 6 amplification. M: 100 bp DNA ladder; Ctl+ 1: positive control (*M. Marinum* DL strain); Ctl+ 2: positive control (*M. ulcerans* strain); Ctl-: negative control (sterile water); ZGF1-ABS2 (A), ZGB31-ZLF2 (B): environmental samples tested.

**Table S1.** Primers used for identification of MPMs.

| Primer Name | Forward and Reverse Primer Sequences | PCR Product Size | Reference |
|---|---|---|---|
| 16S rRNA | PA:5'-AGAGTTTGATCCTGGCTCAG-3'<br>MSHA: 5′-AAAAAGCGACAAACCTACGAG-3' | 620 bp | [1] |
| IS*2404* (nested 1) | pGp1:5′-AGGGCAGCGCGGTGATACGG-3′<br>pGp2: 5′-CAGTGGATTGGTGCCGATCGAG-3′ | 400 bp | [2] |
| IS*2404* (nested 2) | pGp3: 5′-GGCGCAGATCAACTTCGCGGT-3′<br>pGp4: 5′-CTGCGTGGTGCTTTACGCGC-3 | 210 bp | [2] |
| ER | F:5′-GAGATCGGTCCCGACGTCTAC-3′<br>R:5′-GGCTTGACTCATGTCACGTAAG-3′ | 719 bp | [3] |
| Locus 6 | F-5′-GACCGTCATGTCGTTCGATCCTAGT-3′<br>R-5′-GACATCGAAGAGGTGTGCCGTCT-3′ | variable | [3] |
| Locus 19 | F-5′-CCGACGGATGAATCTGTAGGT-3′<br>R-5′-TGGCGACGATCGAGTCTC-3′ | variable | [3] |
| MIRU 1 | F-5′-GCTGGTTCATGCGTGGAAG-3′<br>R-5′-GCCCTCGGGAATGTGGTT-3′ | variable | [3] |
| ST1 | F-5′-CTGAGGGGATTTCACGACCAG-3′<br>R-5'-CGCCACCCGCGGACACAGTCG-3' | variable | [3] |

**Table S2.** Size of PCR product of VNTR loci and associated repeat number.

| VNTR Loci | Repeat Lenght (bp) | PCR Product Size in bp (Associated Repeat Number) | | | | | | | | |
|---|---|---|---|---|---|---|---|---|---|---|
| MIRU 1 | 53 | 404 (1) | 457 (2) | 510 (3) | 563 (4) | 616 (5) | 669 (6) | 722 (7) | 775 (8) | 828 (9) |
| Locus 6 | 56 | 454 (1) | 510 (2) | 566 (3) | 622 (4) | 678 (5) | 734 (6) | 790 (7) | 846 (8) | 902 (9) |
| ST1 | 54 | 369 (1) | 423 (2) | 477 (3) | 531 (4) | 585 (5) | 639 (6) | 693(7) | 747 (8) | 801 (9) |
| Locus 19 | 56 | 288 (1) | 344 (2) | 400 (3) | 456 (4) | 512 (5) | 568 (6) | 624 (7) | 680 (8) | 736 (9) |

## References


1. Hughes, M.S.; Skuce, R.A.; Beck, L.A.; Neill, S.D. Identification of mycobacteria from animals by restriction enzyme analysis and direct DNA cycle sequencing of polymerase chain reaction-amplified 16S rRNA gene sequences. *J. Clin. Microbiol.* **1993**, *31*, 3216–3222.
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3. Williamson, H.R.; Benbow, M.E.; Nguyen, K.D.; Beachboard, D.C.; Kimbirauskas, R.K.; McIntosh, M.D.; Quaye, C.; Ampadu, E.O.; Boakye, D.; Merritt, R.W.; et al. Distribution of *Mycobacterium ulcerans* in Buruli ulcer endemic and non-endemic aquatic sites in Ghana. *PLoS Negl. Trop. Dis.* **2008**, *2*, e205.