

| Locus | Lineage (Family) | LTRr location | Chromatin type | Condition | % Div | Analysis | WEST AFRICA | | EAST AFRICA | |
|----------|---------------------|---------------------------------|-------------------|-----------|-------|-----------|-------------|-------------|-------------|-------------|
| | | | | | | | M | S | Asembo | Jego |
| Locus 1 | CsRn1 (G52_AG) | 2R: 4043596- 4047772 (8A) | NPE | Proviral | 0.0 | Screening | 0.0 (0/5) | 0.0 (0/5) | 0.0 (0/5) | 0.0 (0/5) |
| Locus 2 | CsRn1 (G50_AG) | 2R: 18580773- 18584954 (11A) | NPE | Proviral | 0.0 | Screening | 0.0 (0/5) | 0.0 (0/5) | 0.0 (0/5) | 0.0 (0/5) |
| Locus 3 | CsRn1 (G50_AG) | 2R: 27194388- 27194533 (12E) | NPE | Solo-LTR | 0.0 | Screening | 0.0 (0/5) | 0.0 (0/5) | 0.0 (0/5) | 0.0 (0/5) |
| Locus 4 | CsRn1 (G51_AG) | 2R: 30309166- 30315068 (13D) | NPE | Proviral | 0.6 | Screening | 0.0 (0/5) | 0.0 (0/5) | 0.0 (0/5) | 0.0 (0/5) |
| Locus 5 | CsRn1 (G50_AG) | 3L: 4901607- 4901752 (38C) | IH | Proviral | 5.7 | Screening | 1.0 (5/5) | 1.0 (5/5) | 1.0 (5/5) | 1.0 (5/5) |
| Locus 6 | Mag | 2L: 19195392- 19195689 (22F) | NPE | Solo-LTR | 7.3 | Screening | 1.0 (5/5) | 1.0 (5/5) | 1.0 (5/5) | 1.0 (5/5) |
| GQ468823 | (G57_AG) | | | | | Extensive | 1.0 (50/50) | 1.0 (48/48) | 1.0 (46/46) | 1.0 (19/19) |
| Locus 7 | Mdg3 | UNK | n. d. | Solo-LTR | 7.9 | Screening | 1.0 (5/5) | 1.0 (5/5) | 1.0 (5/5) | 1.0 (5/5) |
| GQ468822 | (8967) | | | | | Extensive | 1.0 (50/50) | 1.0 (48/48) | 1.0 (46/46) | 1.0 (19/19) |
| Locus 8 | Mdg3 | 3R: 43554406- 43554585 (35C) | NPE | Solo-LTR | 8.2 | Screening | 1.0 (5/5) | 1.0 (5/5) | 1.0 (5/5) | 1.0 (5/5) |
| GQ468821 | (G31_AG) | | | | | Extensive | 1.0 (50/50) | 1.0 (48/48) | 1.0 (46/46) | 1.0 (19/19) |