

Detection of numerous HIV-1/MO recombinants in France: implications for biological and clinical practice

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Supplementary Data

Alternative RT-PCRs

Alternative RT-PCRs were performed as follows: viral RNA was extracted from 200 µl of plasma using EZ1 DSP Virus Kit, eluted in 60 µL elution buffer and 10 µl of extract were used in a 50 µL RT-PCR using SuperScript™ III One-Step RT-PCR System with Platinum® Taq DNA polymerase (Invitrogen), 20 pmol of each primer (supplementary Table 2) and 1.5 mM MgSO₄. Two microliters of amplification product were used in a 50 µl nested PCR using HotStar Taq Master Mix (Qiagen), 20 pmol of each primer (supplementary Table 2) and 1.5 mM MgCl₂. (RT-)PCRs were run on a PerkinElmer Gene Amp 9700 thermal cycler and cycling conditions are described in Supplementary Table 2.

Phylogenetic analyses

Phylogenetic analyses were performed using the neighbor-joining method. Genetic distances were calculated with the Kimura two-parameter method and 1000-replicate parametric bootstrap analysis was performed to assess the reliability of branching order. Sequences of available co-infections HIV-1/M+O and recombinants HIV-1/MO previously described were included. All the accession numbers of the previously and newly described sequences are listed in Supplementary Table 3.

Supplementary Table 1. Patient's characteristics

| Patients | Gender | Age | CD4/mm ³ | Year of sampling | Origin |
|----------|--------|-----|---------------------|------------------|----------|
| RBF216 | F | 35 | 25 | 2009 | Cameroon |
| BCF212 | F | 61 | 231 | 2006 | Cameroon |
| RBF221 | F | 62 | N/A | 2010 | Cameroon |
| BCF204 | F | 55 | N/A | 2013 | Cameroon |
| BCF174 | F | 30 | 363 | 2005 | Cameroon |
| RBF222 | M | 45 | 306 | 2010 | Cameroon |
| RBF237 | F | 37 | 399 | 2013 | Cameroon |
| RBF240 | M | 35 | 390 | 2016 | Cameroon |
| RBF243 | F | 44 | 130 | 2017 | Cameroon |
| RBF235 | F | 46 | 84 | 2013 | Chad |

N/A: Not Available

Supplementary Table 2. Alternative primers and PCR conditions used to characterize HIV-1/MO recombinants

| Sample | Genomic region | Step | Primer | Primer sequence (5'-3') | Fragment size (bp) | Cycling conditions | | |
|----------------|----------------|--------|-------------------------|-------------------------------------|--------------------|----------------------|-----------------|--------|
| | | | | | | AT ³ (°C) | ET ⁴ | Cycles |
| RBF235 | RT-PT | RT PCR | PROTO (S ¹) | TTCAAYTGTTGGRAAAGAGGGAC | 1392 | 55 | 1' 30" | 35 |
| | | | MJ4 (AS ²) | CTGTTAGTGCTTTGGTTCCTCT | | | | |
| | | NESTED | PROT4 (S) | CAGCCCCACCRATGGAGG | 1163 | 50 | 1' 30" | 35 |
| | | | NE1 (AS) | CCTACTAACTTCTGTATGTCATTGACAGTCCAGCT | | | | |
| RBF240 | RT-PT | RT PCR | 40U (S) | GGYTYTGGAAATGTGG | 1761 | 55 | 2' | 35 |
| | | | 90L (AS) | GAATCCAGGTRGCTGCC | | | | |
| | | NESTED | 43U (S) | ATGTGGAMAGGAAGWCAYCA | 1334 | 50 | 1'30" | 35 |
| | | | 85L (AS) | CCTGGATARATYTGACTTGC | | | | |
| RBF221 (REC A) | ACC | RT PCR | 127U (S) | GGGTYTATTACAGRGACAGCAGAG | 1561 | 50 | 2" | 35 |
| | | | 167L (AS) | GGGTCTGTGGGWACACAGGC | | | | |
| | | NESTED | 127U (S) | GGGTYTATTACAGRGACAGCAGAG | 1240 | 50 | 1'30" | 35 |
| | | | MVPU (AS) | TACTATRGTCACACAACATATKGCT | | | | |
| RBF221 (REC B) | ACC | RT PCR | 127U (S) | GGGTYTATTACAGRGACAGCAGAG | 1561 | 50 | 2' | 35 |
| | | | 167L (AS) | GGGTCTGTGGGWACACAGGC | | | | |
| | | NESTED | 128U (S) | AGAGAYCCWATTGGAAAGGACC | 1514 | 55 | 2' | 35 |
| | | | 307L (AS) | TTGTGMTGCCCAAATATTATG | | | | |

¹ sense primer² antisense primer³ annealing temperature⁴ extension time

Supplementary Table 3. GenBank accession numbers of previously described and new HIV-1/MO recombinants and HIV-1/M+O dual infections

| Sample | GenBank accession number | Reference | |
|---------------|---------------------------------|--------------------------------|-----------------|
| 97CA.MP645MO | AJ239083 | Peeters <i>et al.</i> 1999 | |
| DSC1320 | AY489738 | Yamaguchi <i>et al.</i> 2004 | |
| RBF140_GPO | KY359367 | Plantier <i>et al.</i> 2004 | |
| RBF140_GPM | KY359368 | | |
| RBF140_polM | KY359375 | | |
| RBF140_polO | KY359376 | | |
| RBF208 | GQ351296 | Vessière <i>et al.</i> 2010 | |
| REC003 | KM438031 | Ngoupo <i>et al.</i> 2016 | |
| REC024 | KM438032 | | |
| YBF301_polM | KX398152 | De Oliveira <i>et al.</i> 2017 | |
| YBF301_polO | KX398153 | | |
| YBF320_polM | KX398154 | | |
| YBF320_polO | KX398155 | | |
| YBF211_polO | KX398156 | | |
| YBF211_polM | KX398157 | | |
| YBF212_polO | KX398158 | | |
| YBF212_polM | KX398159 | | |
| YBF205_polO | KX398160 | | |
| YBF205_polM | KX398161 | | |
| YBF298_polM | KX398162 | | |
| YBF301_GPO | KX398163 | | |
| YBF301_GPM | KX398164 | | |
| YBF320_GPO | KX398165 | | |
| YBF320_GPM | KX398166 | | |
| YBF211_GPO | KX398167 | | |
| YBF211_GPM | KX398168 | | |
| YBF212_GPM | KX398169 | | |
| YBF212_GPO | KX398170 | | |
| YBF205_GPO | KX398171 | | |
| YBF205_GPM | KX398172 | | |
| YBF298_GPM | KX398173 | | |
| YBF282_polM | KX398174 | | |
| YBF282_GPO | KX398175 | | |
| YBF280_polO | KX398176 | | |
| YBF280_polM | KX398177 | | |
| YBF280_GPO | KX398178 | | |
| YBF280_GPM | KX398179 | | |
| YBF274_polO | KX398180 | | |
| YBF274_GPO | KX398181 | | |
| YBF274_GPM | KX398182 | | |
| YBF221_polO | KX398183 | | |
| YBF221_polM | KX398184 | | |
| YBF221_GPO | KX398185 | | |
| YBF221_GPM | KX398186 | | |
| RBF235_GPO | KY359362 | | New submissions |
| RBF235_GPM | KY359363 | | |
| BCF174_GPO | KY359364 | | |
| RBF221_GPO | KY359365 | | |
| RBF221_GPM | KY359366 | | |
| RBF216_GPO | KY359369 | | |
| RBF216_GPM | KY359370 | | |
| RBF235_polOM | KY359371 | | |
| RBF235_polO | KY359372 | | |
| BCF174_polM | KY359373 | | |
| RBF221_polM | KY359374 | | |
| RBF216_polM | KY359377 | | |
| RBF216_polO | KY359378 | | |
| BCF212_GPO | KY359379 | | |
| BCF212 | KY359380 | | |
| RBF237 | KY359381 | | |
| RBF222 | KY359382 | | |
| BCF174 | KY359383 | | |
| RBF240 | KY359384 | | |
| BCF204 | KY359385 | | |
| RBF243 | KY995542 | | |