

Supplementary Sequence information:

Sequence alignment of the regions where the two arms of the construction differ (Oxa7 vs. oxa 11).
The indices are relative to the center of the construction. The numbers in bold are the indices corresponding to fig 5A and 5B

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Oxa7 arm      GTTCGCAATCTGGGTATGTTTGTTCCTGAGCGCGGTTGGTGCTGTTGGCGTTCGGT 2280
Oxa11 arm     GTTCGCAATCTGGGTATGTTTGTTCCTGAGCGCGGTTGGTGCTGTTGGCGTTCGGT 2280
*****

Oxa7 arm      CAGTGCCGGACTGTCCAGTGGGCTTTTCTGTTCGTTTCATCCATTACCACCTTAACCGCC 2340
Oxa11 arm     CAGTGCCGGACTGTCCAGTGGGCTTTTCTGTTCGTTTCATCCATTACCACCTTAACCGCC 2340
*****

Oxa7 arm      TTTGGCGTTGCAGCAAGCGTTTCAGACGTGCTGTTGGTTGCACTGCT--GAAAACATTTG 2398
Oxa11 arm     TTTGGCGTTGCAGCAAGCGTTTCAGACGTGCTGTTGGTTGCACTGCTATGAAAACATTTG 2400
*****

Oxa7 arm      CCGCATATGTAATTACTGCGTGTCTTTCAAGTACGGCATTAGCTAGTTCAATTACAGAAA 2458
Oxa11 arm     CCGCATATGTAATTATCGCGTGTCTTTGAGTACGGCATTAGCTGGTTCAATTACAGAAA 2460
*****

Oxa7 arm      ATACGTTTTGGAACAAAGAGTTCTCTGCCGAAGCCGTC AATGGTGTTCGTTGCTTTGTA 2518
Oxa11 arm     ATACGTTCTTGGGAACAAAGAGTTCTCTGCCGAAGCCGTC AATGGTGTTCGTTGCTTTGTA 2520
*****

Oxa7 arm      AAAGTAGCAGTAAATCCTGCGCTACCAATAA CTTAGCTCGTGCATCAAAGGAATATCTTC 2578
Oxa11 arm     AAAGTAGCAGTAAATCCTGCGCTACCAATGACTTAGCTCGTGCATCAAAGGAATATCTTC 2580
*****

Oxa7 arm      CAGCATCAACATTTAAGATCCCCAACGCAATTATCGGCCTAGAAACTGGTGT CATAAAGA 2638
Oxa11 arm     CAGCATCAACATTTAAGATCCCCAACGCAATTATCGGCCTAGAAACTGGTGT CATAAAGA 2640
*****

Oxa7 arm      ATGAGCATCAGATTTTCAAATGGGACGGAAAGCCAAGAGCCATGAAACAATGGGAAAGAG 2698
Oxa11 arm     ATGAGCATCAGGTTTTCAAATGGGACGGAAAGCCAAGAGCCATGAAGCAATGGGAAAGAG 2700
*****

Oxa7 arm      ACTTGAGCTTAAGAGGGGCAATACAAGTTTCAGCGGTTCCCGTATTTCAACAAATCGCCA 2758
Oxa11 arm     ACTTGACCTTAAGAGGGGCAATACAAGTTTCAGCTGTTCCCGTATTTCAACAAATCGCCA 2760
*****
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2790 2805 2811 2813
Oxa7 arm GAGAAGTTGGCGAAGTAAGAATGCAGAAATATCTTAAAAAATTTTCATATGGTAACCAGA 2818
Oxa11 arm GAGAAGTTGGCGAAGTAAGAATGCAGAAATACCTTAAAAAATTTTCCTATGGCAGCCAGA 2820

2853 2855
Oxa7 arm ATATCAGTGGTGGCATTGACAAATCTGGTTGGAGGGTCAGCTTAGAATTTCCGCAGTTA 2878
Oxa11 arm ATATCAGTGGTGGCATTGACAAATCTGGTTGGAAGACCAGCTTAGAATTTCCGCAGTTA 2880

2906 2928 2937
Oxa7 arm ATCAAGTGGAGTTTCTAGAGTCTCTATTTTTAAATAAAATGTGCAGCATCAAAGAAAATC 2938
Oxa11 arm ATCAAGTGGAGTTTCTAGAGTCTCTATTTTTAAATAAAATGTGCAGCATCTAAAGAAAACC 2940

2973 2976 2988
Oxa7 arm AGCTAATAGTAAAAGAGGCTTTGGTAACGGAGGCTGCGCCTGAATATCTTGTGCATTCAA 2998
Oxa11 arm AGCTAATAGTAAAAGAGGCTTTGGTAACGGAGGCGGCACCTGAATATCTAGTGCATTCAA 3000

3057
Oxa7 arm AAAGTGGTTTTCTGGTGTGGGAAGTGGTCAAATCCTGGTGTGCATGGTGGGTTGGTT 3058
Oxa11 arm AAAGTGGTTTTCTGGTGTGGGAAGTGGTCAAATCCTGGTGTGCATGGTGGGTTGGTT 3060

3071
Oxa7 arm GGGTTGAGAAGGGAGCAGAGGTTTACTTTTTTCGCCTTTAACATGGATATAGACAACGAAA 3118
Oxa11 arm GGGTTGAGAAGGGAGCAGAGGTTTACTTTTTTCGCCTTTAACATGGATATAGACAACGAAA 3120

3119 3161
Oxa7 arm ATAAGTTGCCGCTAAGAAAATCCATTCACCAAAAATCATGGCAAGTGAGGGCATCATTG 3178
Oxa11 arm GTAAGTTGCCGCTAAGAAAATCCATTCACCAAAAATCATGGAAAGTGAGGGCATCATTG 3180

3187
Oxa7 arm GTGGCTAAGAGCTGCACTATCCCCTTTCTCGTTGTGTCCGCATCCTCAAGCGCGACAGCT 3238
Oxa11 arm GTGGCTAA-AGCTGCACTATCCCCTTTCTCGTTGTGTCCGCATCCTCAAGCGCGACAGCT 3239
