

### Supplementary Figure 1B

3'NCR of all nimbus (*BgI*) elements arranged in the order they occur in the BAC clone (truncated element cassettes A, B and C and full-length elements I, II and III).

Element	bp	3' NCR conserved region	16 bp 3' RE	3' NCR conserved region	tandem repeat (TCMn or C Mm)	Start of next Sequence (with or - TSD)	bp
A.1	612	TGAACATGCACTATTTACATTAGATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATATTACTGTATAGAATCTGGCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCAAAATCA	ACTATTTAATAATGTGACCCCTGGAAAGGTACTGGC	862
A.2	911	TGAACATGCACTATTTACATTAGATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATATGACTGTATAGAATCTGCCCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCAAAATCA	GACTGCGAGAACCGATGAGGAAGCTGAGACTCAGCAAGG	1155
A.3	1491	TGAACATGCACTATTTACATTAGATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATATGACTGTATAGAATCTGCCCCTTGTGTTT--AGAGAGAGAGGGATCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCAAAATCA	ACCAGGGATGAAATGACCGGGGATGAAAGCGACCAAGG	1739
II	34060	TGAACATGCACTATTTACATTAGATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATATTACTGTATAGAATCTGGCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCAAAATCA	ACATGACATGACATTTACATTAGATTTTACCAAA	34033
B.1	34270	TGAACATGCACTATTTACATTAGATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATATGACTGTATAGAATCTGGCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCAAAATCA	ACCAGGGATGTCAGGGGAAACATTTTAGAGCCACTAA	34512
B.2	34607	TGAACATGCACTATTTACATTAGATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATGACTGTATAGAATCTGGCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCAAAATCA	GAACACGTCCTCATCTTCTGTTGTTCTGTT	34841
III	43821	TGAACATGCACTATTTACATTAGATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATATTACTGTATAGAATCTGGCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCAAAATCA	CAACCGACTGGGAGCTGACAGAGATACCTTGCTGC	44068
C.1	45402	TGAACATGCACTATTTACATTAAATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATATTACTGTATAGAATCTGGCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCAAAATCA	CAACCGACTGGGAGCTGACAGAGATACCTTGCTGC	45649
C.2	46983	TGAACATGCACTATTTACATTAAATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATATTACTGTATAGAATCTGGCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCAAAATCA	CAACCGACTGGGAGCTGACAGAGATACCTTGCTGC	47230
C.3	48565	TGAACATGCACTATTTACATTAGATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATGACTGTATAGAATCTGGCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCAAAATCA	CAAGCTTGGGGGGGATGAGGACTGAGCATCTTATTG	48812
C.4	49385	TGAACATGCACTATTTACATTAGATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATATTACTGTATAGAATCTGGCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCAAAATCA	ATGAGAGAACCCATGCCAAATTTTAACCCACCTATTG	49640
C.5	50609	TGAACATGCACTATTTACATTAGATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATGACTGTATAGAATCTGGCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCA	TCACAAAGCTATTGGGCTCTTATTGGGCTTTAAAAA	50848
I	59274	TGAACATGCACTATTTACATTAGATTTTACCAAAATA	TTAAATTTACTACCTTACTATTTA	ACTGTGAATAGACCTTAATTITAATTATATTACTGTATAGAATCTGGCCTTGTGTTT--AGAGAGAGAGTAGCTCCCTAGGGACTCGAGCAGACATGCC	CTAAATCACAAATCA	ACCATTGACCCACAAAGGGGATGAGGACTCACAACT	59518

5'NCR of all full-length nimbus (*BgI*) elements arranged in the order they occur in the BAC clone (Elements I, II and III)

Element	bp	End of previous sequence with 5' TSD	5' NCR conserved region	16 bp 5' RE	5' NCR conserved region	bp
II	28401	GTTGGTATCCGTAACGAACCGTCTCGTCCCGAGACATGACGTTAACCTCGTCCCGAGACATGACATTAAACTGOGCTCCTCTTCTTCTT-AGCACTTTGGAGCTCAAAGTGCCCTACTCTTCTTCTATCATTTGTCGCCCCTCTCTTCTTCTAAGTCTGCCCCATTTGATCATCACACTGTCTCTTAACTTTAAATTCTCACTACGTGCTTAGACCACTCCGTTGCGTGACTGCGACCGGTAACGGGOGA	28659			
III	39342	GTTATCAGGCTCACTTGACAGAGA <sup>CCAG</sup> CAAAGGTTAGGCANTATCAGAGTCGTCCCGAGACATGACGTTAACCTGOGCTCCTCTTCTTCTTCTAAGTCTGCCCCATTTGATCATCACACTGTCTCTTAACTTTAAATTCTCACTACGTGCTTAGACCACTCCGTTGCGTGACTGCGACGGTAACGGGOGA	39601			
I	53586	TCAATAATTATGATGGACCTAGGCGACCGCTGGCAAATCACCATTGACCCCTCGTCCCGAGACATGACGTTAACCTGOGCTCCTCTTCTTCTAAGTCTGCCCCATTTGATCATCACACTGTCTCTTAACTTTAAATTCTCACTACGTGCTTAGACCACTCCGTTGCGTGACTGCGACGGTAACGGGOGA	53845			