

pIND/Hygro Multiple Cloning Site

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12  AGATCTCGGC  CGCATATTAA  GTGCATTGTT  CTCGATACCG  CTAAGTGCAT
                                     E/GRE
62  TGTTCCTCGTT  AGCTCGATGG  ACAAGTGCAT  TGTTCCTCTTG  CTGAAAGCTC
                                     E/GRE
112 GATGGACAAG  TGCATTGTTC  TCTTGCTGAA  AGCTCGATGG  ACAAGTGCAT
                                     E/GRE
162 TGTTCCTCTTG  CTGAAAGCTC  AGTACCCGGG  AGTACCCTCG  ACCGCCGGAG
                                     ↓ 5' end minimal heat shock promoter
212 TATAAATAGA  GGCGCTTCGT  CTACGGAGCG  ACAATTCAAT  TCAAACAAGC
                                     ↗ Start of transcription
262 AAAGTGAACA  CGTCGCTAAG  CGAAAGCTAA  GCAAATAAAC  AAGCGCAGCT
312 GAACAAGCTA  AACAATCTGC  AGTAAAGTGC  AAGTTAAAGT  GAATCAATTA
362 AAAGTAACCA  GCAACCAAGT  AAATCAACTG  CAACTACTGA  AATCTGCCAA
412 GAAGTAATTA  TTGAATACAA  GAAGAGAACT  CTGAATACTT  TCAACAAGTT
462 ACCGAGAAAG  AAGAACTCAC  ACACAGCTAG  CGTTTAAACT  TAAGCTTGTT
                                     |       |       |       |       |
                                     Nhe I   Pme I  Afl II  Hind III  Asp718 I
512 ACCGAGCTCG  GATCCACTAG  TCCAGTGTGG  TGGAATTCTG  CAGATATCCA
       |       |       |       |       |       |
       Kpn I   Ecl136 II  Sac I   BamHI  Spe I       BstX I*       EcoR V
562 GCACAGTGGC  GGCCGCTCGA  GTCTAGAGGG  CCCGTTTAAA  CCCGCTGATC
                                     |       |       |       |       |
                                     BstX I*  Not I   Xho I   Xba I   Dra II  Apa I   Pme I
612 AGCCTCGACT  GTGCCTTCTA  GTTGCCAGCC  ATCTGTTGTT  TGCCCCCTCCC
                                     BGH Reverse priming site

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*Note that there are two *BstX I* sites in the polylinker.