

pIND(SP1)/Hygro Multiple Cloning Site

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12 AGATCTCGGC CGCATATTAAGTGCATTGTT CTCGATACCG CTAAGTGCAT
      E/GRE
62 TGTTCCTCGTT AGCTCGATGG ACAAGTGCAT TGTTCTCTTG CTGAAAGCTC
      E/GRE
112 GATGGACAAG TGCATTGTTC TCTTGCTGAA AGCTCGATGG ACAAGTGCAT
      E/GRE
162 TGTTCTCTTG CTGAAAGCTC AGTACCCGGG TCGGAGTACT GCCCCGCCCC
      SP1
212 TAGCGATTAG CCCC GGCCCC GCATAGCTCC GCCCCGGGAG TACCCTCGAC
      SP1
      5' end minimal heat shock promoter
262 CGCCGGAGTA TAAATAGAGG CGCTTCGTCT ACGGAGCGAC AATTCAATTC
      Start of transcription
312 AAACAAGCAA AGTGAACACG TCGCTAAGCG AAAGCTAAGC AAATAAACAA
362 GCGCAGCTGA ACAAGCTAAA CAATCTGCAG TAAAGTGCAA GTTAAAGTGA
412 ATCAATTAAA AGTAACCAGC AACCAAGTAA ATCAACTGCA ACTACTGAAA
462 TCTGCCAAGA AGTAATTATT GAATACAAGA AGAGA ACTCT GAATACTTTC
512 AAC AAGTTAC CGAGAAAGAA GAACTCACAC ACAGCTAGCG TTTAAACTTA
      Nhe I Pme I Afl II
      Hind III Asp718 I Kpn I Ecl136 II Sac I BamH I Spe I BstX I*
562 AGCTTGGTAC CGAGCTCGGA TCCACTAGTC CAGTGTGGTG GAATTCTGCA
      EcoR V BstX I* Not I Xho I Xba I Dra II Apa I Pme I
612 GATATCCAGC ACAGTGGCGG CCGCTCGAGT CTAGAGGGCC CGTTTAAACC
      BGH Reverse priming site
662 CGCTGATCAG CCTCGACTGT GCCTTCTAGT TGCCAGCCAT CTGTTGTTTG
712 CCCCTCCC

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