BDP	BDP_sequence	Sequence_length_bp	Chr	Upstream_ORF	Upstream_TATA	Upstream_ORF_Start	Upstream_TSS	NDR_edge_minus	NDR_center	NDR_edge_plus	Downstream_TSS	Downstream_ORF_Start	Downstream_ORF	Downstream_TATA	BDP_length_bp	NDR_width_bp
BP724	tcttgtgtggagcttgaaagtttatctttcgtatacaaatagagagaattacggtcaagcatggaattggaatggaaa gaggttaactgatttttttctttctatctcacctgtttatatttgcgattgccatgctccggtttttccgctctttttta aaggcccctttatcccctctaacgccttcgcaatcccctttcactcggctgcaaacgacggcgtttgtaagcagc taagctttctagcctagcaatgaaatagaaggacagtcagacgatttgacgtcaaaaaatgagtgccgctctctttc tttctttaggccaactagacagccgtatgatgaaaat	362	ChrXVI	YPL004C	Yes	551657	551694	551752	551857	551983	552008	552020	YPL003W	No	362	211
BP306	tgtttacaaacacgccctatatacttatgaggggtagccgagacagtgccgctgtatgctgaatttgtactcgt agatgttagcaagcgtagcaaatgtgatctcctgctacagcatttatttttctccttctaaaatatacactcaagcaagaa gcccttaatacacagttaaccagactaatcagtcgactttttccgcgaatgtagaaaagaaacttttctatcgccgctgc aaacttaatctatccctccaccaaagcaaaatgcaaaacaagccatactgggaaaatctgaaaaaaaatgggt aggatacaaaagaaaagaaaaaaaaaggttaccctgcagtttgtatagtgggtaacatttggccctttctcttg atttgatatattaccccgatttacccctcatctttggcagtgcccgctttatttcccccgccaatcggctattaacggctt tacgtcattcgtgggcgggtcaagcgagcgccttcctgtttgtttgctcacgcaaaaccgaaggctcaaccaaa actaaggccatcatatatatatatgtcggctgtgctgtgtatctcccggatatatgtgtgctgcaatttgaatgtatg gagaaaattttcttccctttcatttacgggcggaaatacttcatatataaaaaaaaagaatatcaatctagtctttaacgctata cgcatacagcatttcaagacacatatgaaacacaaaacctatatttta	754	ChrVII	YGR241C	No	976581	976748	976769	976966	977164	977276	977336	YGR243W	Yes	754	395
BP670	tagggggctttctatatttttacacgaatatacacctactaagtttattgaggtcatgatgagctggaagaaaacaacaaa aacaaaatagtgtatctaatttctgtcatttcatacttaaaagcactctagtgacaatcaccttgaaagttgttgtatacccgt gaaacaaatttcatcggccgggtggctatgcgtaaagccgcgccttgatagaatatgagtgagtgagctttatactat atacctgatgctacaaaagcatatatatgctagcaatcaaaaaccaaaaaaagatatcattagcgttatgaaaatcgg gaatt	318	ChrXV	YOR219C	No	755009	755065	755139	755229	755319	755304	755328	YOR220W	No	318	180
BP565	tgaggttgccctgtggttatagtcttgtggtttttgaatagaagcaatagaatatataaagccaggacctgcaatcaacgtt tgtatgttttatttgtgtaaaaagtgaagtggcgaatagccaacgctatttaaaactttatgtctagctaattatcatgaat gttgtcctgtccacatgatgactcccctgagttgccgttgctgctgctgaagggagttaattaattccagggccaccg cgcgaagccccgcgcagcaatgttatacctgagggaagaaggcccttttgagaagctgccaatttgcaacttcta tcgagtgagtgatgttacaatacgctgttcatttgaggaagtag	356	ChrXIII	YMR261C	No	793369	793452	793532	793586	793640	793716	793726	YMR262W	No	356	108
BP511	ctttcttcttaattaacagttcttttgtagttgtctcctctctattgcagatcctcttctaaacctcggtaatctaatcatatgcta ggaatatacgactctatgagtgtgcttttaatatacctgtgcgatccctttcacgccgtcttgttttcctccctctccttatttt ccttcagttcaaagatccctaattctttccccttctaagtgtgaggccggcgatatgtctgctttaccgcgctaatgtgtg gaaaagtgtgaaaaagtgaaaatgtgatatttctagcgagaaaaacgaaaacaacaagtcgtgtcaagcagaggtagt gttttgggaaagacgtcatatgagaaagcgccgcaaagctgttgaagaagtggggaaataaagatgtaag	387	ChrXIII	YML128C	No	16676	16778	16834	16924	17014	16984	17064	YML127W	No	387	180
BP535	agcgcttgttgtatatatgtcgaaaaatgatgtgtgacggaaagattaatattaaatagcttttctaagtgggcaacc aaaagtgttttgatttgggcattttacaaaattcacgtttacaacctctctctgtctctcctatgctttcctgatcaaat taagaattttcatcaagaatagagagcgaagccaagccaagagagaaaaagaaaggagagaaagaaagaaaa gggatggaagaggtgatggtaaaagagcgaaaactgagtttgtatgaaagaaattgcgtgaagaagtaaaagtaaa atttccattcaaatgaatgtatttaatataatttctatcatttaaaaggtgggaaaaattacgtatatgaaaaacc gtgtgtgtgcagtgctatctatttctgctgtgtgtatgtatatttcgtccacagtcgaaaaaactcaaaataagtatt accccgatgcgaggctgtggccatttagccgccgtctgtcgaccatccaccaatggtgatgatacatcgggtgatca gccctttttggtgtattactatccctatctcatcctcaaaaaacacgtatatatcatgacgagatcgacgatacgtcg ctgtgttaacgccaactgtgtgtgtagcgtttgtgtcaatttagccaaatgtgtttcttgaaaaaagcaacgggaa aaaaaaggtgtgatagcaatatatgtacagctgccctctctgagcaaaagctagccgccgtcgaagtaaatgat acgctgatgtgtcgttgcagcaatgatatgtgcaatttttcttttgtttccttacatttatccccgcatatctcgccgtaagtatatt gaata														
BP76	tactcagctagaaacgcacactctgcgttattctcctccgcattgatatatgatagcgttccatattattgcatttttatc agggggatctctcaagaaaggaacttttttaggaaaaaaaaaaataatcatcttttatacatgtgtaatatctcaagca atcataaacaagagttaaagaagcaaagttctgctgtagtgtgtgtctccaatactctgcacgacaaagttttagttat cccctctacgtgtatgttctcctccgtgtgtcgattgttcaaagatgggagctctttcgtgtttgcactaagacgcatac tgttaatccgcaaaggccactaaaacttacctgcattttgtctactaaagatacattccaactatttacattcatttttattct aaaagagcattctgaacctcggcaatttatttcatactcctttctcagaatagggaaa	471	ChrIII	YCL052C	No	35393	35426	35444	35496	35549	35584	35865	YCL051W	No	471	105
BP565：有 BbsI、未见 BsaI																
BP535：有 BbsI、未见 BsaI																
BP670：未见常用 BsaI/BbsI 酶显问题																
BP724：有 HindIII、但 HindIII 不影响 BsaI/BbsI Golden Gate																
BP306：未见常用 BsaI/BbsI 酶显问题																
BP511：有 HindIII、但不影响 Golden Gate																