

Table A2: Oligonucleotides

Primer Name	Sequence	Sites
Degenerate Primers for Cloning Cbf3 genes		
CgCBF3A/1.2	TAYCCIAAYAAYGARGARATG	
CgCBF3A/2	ATTAAAGTATGATACAAAATC	
CgCBF3B/1	TTCTGGCAAAGCTATGAA	
CgCBF3B/2	CAAATATGCTGCATGTTT	
CgCBF3C/1.2	ATIGTIAAYTTRGARGARATG	
CgCBF3C/2.3	ACRTRRTTIATYTTIATCCA	
CgMIF2/1	AAAGATGAGTATAGTATGGAGAA	
CgMIF2/2.3	TAYTGIATYTTYTCRTTICGCCA	
CgCSE4/1.2	GAYGTIGAYGARGAYAGRTGG	
CgCSE4/2.2	AACTGTCCTCTTATTCTCCTA	
Primers for Gene Deletions		
CgA1/1	CGGGGTACCGGGCCAGTCTTCGGG CTT	<i>KpnI</i>
CgA1/2	TTAACTCGAGAATAGGCCATTTGAT CACACA	<i>XhoI</i>
CgA2/1	GGGCTCGAGGAATGGTATACGCCA GATCCA	<i>XhoI</i>
CgA2/2	GTGAGTTCTAGATAGGGCACT	<i>XbaI</i>
CgB1/1	CGGGGTACCAGTATTATAAATTATC AATAG	<i>KpnI</i>
CgB1/2	TTAACTCGAGTCTCTTTCGACAATG CCCGCA	<i>XhoI</i>
CgB2/1	GGGCTCGAGGTAAATCTACCTCCG ATATTT	<i>XhoI</i>
CgB2/2	AAATCTAGATCTTCAATAGTGGATG ATGTT	<i>XbaI</i>
CgC1/1	CGGGGTACCTTCTATATACTTTTCG CATATTC	<i>KpnI</i>
CgC1/2	TTAACTCGAGGTAAACCAACTTCCT GACCTC	<i>XhoI</i>
CgC2/1	GGGCTCGAGTGGATTAAGATTAAT AATGTG	<i>XhoI</i>
CgC2/2	AAATCTAGATAAATCAGAACCAAC AAATGA	<i>XbaI</i>

Primer for amplifying open reading frames		
GST-MIF2.1	CGGGATCCATGGATTACATGAACCT AGGG	<i>Bam</i> HI
GST-MIF2.2	CAATGAATTCCTCAATAATGTTTCAG AATGAAAGT	<i>Eco</i> RI
GST-3D1	CGGGATCCATGGACAAGAAGGGCA AGTACGT	<i>Bam</i> HI
GST-3D2	CAATGAATTCCTTAGCGGTCCTCCG CCCA	<i>Eco</i> RI
GST-3C-1	CGGATCCATGGTTGAGGAGCGGTT CAAT	<i>Bam</i> HI
GST-3C-2	CAATGAATTCCTTAAATCACAAC TATTACTATGAAC	<i>Eco</i> RI
Cg3B.1	CGCGAATTCAAGTATGAAAATAAT GAAGTT	<i>Eco</i> RI
CgB5.2	CGCGGATCCTTAGTCAATGACTCTA TCAAT	<i>Bam</i> HI
GST-3B-1.2	TCCCCCGGGATGTTCCAGAGGCCAT TGCC	<i>Sma</i> I
GST-3B-2.1	CCGCTCGAGCTATTAGTCAATGACT CTATCAAT	<i>Xho</i> I
GST-p110-1	TCCCCCGGGATGATCTCCAAGGAC AATGAG	<i>Sma</i> I
GST-P110-2	CCGCTCGAGTTAGGTGTGCTGTATC TGATT	<i>Xho</i> I
CgD2	CCCTCGAGGATCCTTAGCGGTCCTC CGCC	<i>Xho</i> I/ <i>Bam</i> HI
CgD3	GGCTGCAGAATTCAACATTGCACG CTA	<i>Pst</i> I/ <i>Eco</i> RI
Primers for amplifying promoters		
Cg1-P.1	TTA TAC CGA GCT CAC AAA CAA ATC ACT CTA CAC G	<i>Sac</i> I
Cg1.P.2	CCA ATT AAG CTT CCT AGA ACG GTT GCT TC	<i>Hind</i> III
CgA-4	GGG TCC CCC GGG TGG TCT TCT TGC GGT GTA	<i>Sma</i> I
CgA-5.2	CGG GAT CCT TGG AGA TCA TGA TTG	<i>Bam</i> HI
3B-1	TCC CCC GGG CGA TAT AGG CGC CAG	<i>Sma</i> I
3B-2	GGA ATT CCA TAT GTA TCT TCC AAT ATA TAT TGA T	<i>Nde</i> I
3B-3	CGC GGA TCC GCA GAA TAA CTT TCA GCA TCT	<i>Bam</i> HI

3B-4	TTC CAA TGC ATT GGC TGC AGC TTG TAA GAG TTG ATT CTT CT	<i>Pst</i> I
3C-1	TCC CCC GGG ATG GTG GTG GCA CTA CGT A	<i>Sma</i> I
3C-2	GGA ATT CAT TAA TAC GCG AAA ACT TCT TGG CTT	<i>Ase</i> I
3C-3	CGC GGA TCC CTG AAA TAT ATA ATT ACC ATG	<i>Bam</i> H1
3C-4	TTC CAA TGC ATT GGC TGC AGT TGG ACT CCT GGA CGA CTC AC	<i>Pst</i> I
Primers for HA-tagging and ChIP analysis		
CgA6.1.2	GAA GAT CTA CCA GAG AAT TCA TTA T	<i>Bgl</i> II
CgA6.2	CCT TGG CAT ATG GGT GTG CTG TAT CTG ATT CAA	<i>Nde</i> I
CgB6.1.2	GCT CTA GAG TTG CAA CGT TTA	<i>Xba</i> I
CgB6.2	CCT TGG CAT ATG GTC AAT GAC TCT ATC AAT A	<i>Nde</i> I
CgC6.1	CGG GAT CCA TGA GGA TAG CAA TGA ATC	<i>Bam</i> H1
CgC6.2	CCT TGG CAT ATG AAT CAC AAC TAT ATT ACT ATG	<i>Nde</i> I
CgC6.3	CGG GGT ACC CTG AAA TAT ATA ATT ACC ATG	<i>Kpn</i> I
CgC6.5	GGT TAC GTA ACC AAC TGC AGG TAC CCA AAA TAT AAA TCA GA	<i>Pst</i> I
HA-F	GGA ATT CCA TAT GGT TGT TTA CCC ATA CGA T	<i>Nde</i> I
CgHIS3R	GGG GTA CCA TAC CAG ATC ATT TTC AAT AT	<i>Kpn</i> I
CgCEN1	TAA CGC AGA GCA CAT CTT TC	
CgCEN2	CGC GTA ACT AAT GAT GCA AT	
CgACT1	CGG GGT ACC ATG GAT TCT GGT ATG TTC GA	
CgACT4	TAT TCA CAC ACG CAG GGA CAG	
CgLEU1	CAA GAC AAT TGT AGT TCC ACC	
CgLEU2	ACC AAC AGC ACC CAG AAG C	