

BF2_2_GAGCACC	7	6	6	1	2	1	1	2	1	2	0	5	4	4	3	4	3	1	2	1	3	2	1	2	1	3	3	2	2	1	1	1	2	2	2	2	2	3	3	2	2	0	0	0	1	1	1	3	0	1	1	1	2	
BF2_2_GAGCCCC	6	7	6	2	3	2	1	2	1	2	0	4	5	4	4	5	4	1	2	1	3	1	2	3	2	3	2	2	2	2	2	2	2	2	2	2	1	2	2	2	2	1	1	1	1	1	1	2	0	1	1	1	1	
BF2_2_GAGCTCC	6	6	7	1	2	1	1	2	1	3	0	4	4	5	3	4	3	1	2	1	3	1	1	2	1	3	2	2	3	1	1	1	3	3	3	1	2	2	2	3	0	0	0	2	2	2	2	2	1	2	2	2	1	
BF2_3_AGTGCAC	1	2	1	7	6	6	1	2	1	4	2	2	3	2	4	3	3	1	2	1	1	2	4	3	3	1	2	2	2	3	2	2	0	0	0	2	3	1	1	1	3	2	2	1	1	1	1	1	1	0	0	0	1	
BF2_3_AGTGCCC	2	3	2	6	7	6	2	3	2	4	1	3	4	3	3	4	3	2	3	2	1	1	3	4	3	1	2	2	2	2	2	2	1	1	1	2	2	1	1	1	2	2	2	2	2	2	2	1	0	1	1	1	1	
BF2_3_AGTGCTC	1	2	1	6	6	7	1	2	1	4	1	2	3	2	3	3	4	1	2	1	1	1	3	3	4	1	2	2	2	2	2	3	0	0	0	3	2	1	1	1	2	2	3	1	1	1	2	0	0	0	0	2		
BF2_4_CTTTGCA	1	1	1	1	2	1	7	6	6	0	2	2	2	2	0	1	0	4	3	3	3	0	1	2	1	1	1	2	1	1	1	1	1	1	3	2	2	1	1	1	4	3	3	2	0	4	3	3	0					
BF2_4_CTTTGCC	2	2	2	2	3	2	6	7	6	1	2	3	3	3	1	2	1	3	4	3	4	0	2	3	2	2	1	2	1	1	1	1	1	1	2	2	1	1	1	3	3	3	3	0	3	3	3	0						
BF2_4_CTTTGCT	1	1	1	1	2	1	6	6	7	0	2	2	2	2	0	1	0	3	3	4	3	0	1	2	1	1	1	2	1	1	1	1	1	1	2	2	3	1	2	1	2	1	1	1	3	3	4	2	1	3	3	4	1	
BF2_5_AGGGTGC	2	2	3	4	4	4	0	1	0	7	0	3	3	4	2	2	2	1	2	1	3	1	2	2	2	2	2	2	3	2	3	2	1	1	1	2	1	1	1	2	1	2	1	2	2	2	2	1	1	1	1	1		
BF2_6_TCTAGAG	0	0	0	2	1	1	2	2	2	0	7	0	0	0	1	0	0	2	2	2	1	3	2	1	1	2	2	3	2	3	2	2	1	2	1	1	2	1	2	3	4	3	3	2	2	0	1	0	1	4	1	2	1	1
BF2_B_ATGCACC	5	4	4	2	3	2	2	3	2	3	0	7	6	6	1	2	1	3	4	3	4	1	1	2	1	1	3	2	2	1	1	1	2	2	2	2	1	2	1	1	1	1	1	1	4	0	1	1	1	2				
BF2_B_ATGCCCC	4	5	4	3	4	3	2	3	2	3	0	6	7	6	2	3	2	3	4	3	4	0	2	3	2	1	2	2	2	2	2	2	2	2	2	2	1	0	1	1	1	2	2	2	1	1	1	3	0	1	1	1	1	
BF2_B_ATGCTCC	4	4	5	2	3	2	2	3	2	4	0	6	6	7	1	2	1	3	4	3	4	0	1	2	1	1	2	2	3	1	1	1	3	3	3	1	0	1	1	2	1	1	1	2	2	2	3	1	2	2	2	1		
BF2_C_GACGCAC	3	4	3	4	3	3	0	1	0	2	1	1	2	1	7	6	6	1	2	1	1	4	4	3	3	4	1	1	1	2	1	1	1	1	1	1	2	4	1	1	1	2	1	1	1	1	1	1	0	0	0	0		
BF2_C_GACGCCC	4	5	4	3	4	3	1	2	1	2	0	2	3	2	6	7	6	2	3	2	1	3	3	4	3	4	1	1	1	1	1	1	2	2	2	2	3	1	1	1	1	1	2	2	2	1	0	1	1	1	0			
BF2_C_GACGCTC	3	4	3	3	3	4	0	1	0	2	0	1	2	1	6	6	7	1	2	1	1	3	3	3	4	4	1	1	1	1	1	2	1	1	1	3	3	1	1	1	1	1	2	1	1	1	2	0	0	0	0	1		
BF2_D_ATCAGCA	1	1	1	1	2	1	4	3	3	1	2	3	3	3	1	2	1	7	6	6	2	1	0	1	0	3	1	2	1	0	0	0	4	3	3	1	0	0	1	0	1	1	1	3	2	2	2	1	2	1	1	0		
BF2_D_ATCAGCC	2	2	2	2	3	2	3	4	3	2	2	4	4	4	2	3	2	6	7	6	3	1	1	2	1	4	1	2	1	0	0	0	3	3	3	2	0	0	1	0	1	1	1	2	2	2	3	1	1	1	1	0		
BF2_D_ATCAGCT	1	1	1	1	2	1	3	3	4	1	2	3	3	3	1	2	1	6	6	7	2	1	0	1	0	3	1	2	1	0	0	0	3	3	4	1	1	0	1	0	1	1	1	2	2	3	2	2	1	1	2	1		
BF2_E_CTGCGGC	3	3	3	1	1	1	3	4	3	3	1	4	4	4	1	1	1	2	3	2	7	0	2	2	2	3	2	3	2	1	2	1	1	1	1	1	1	0	2	3	2	0	1	0	1	1	1	4	0	1	1	1	1	
BF2_F_TACGAAG	2	1	1	2	1	1	0	0	0	1	3	1	0	0	4	3	3	1	1	1	0	7	2	1	1	2	3	2	2	3	2	2	0	1	0	3	4	2	1	1	2	1	1	2	1	2	0	1	0	2				
BF2_W_CCAGCAC	1	2	1	4	3	3	1	2	1	2	2	1	2	1	4	3	3	0	1	0	2	2	7	6	6	1	0	0	0	2	1	1	1	1	1	1	2	1	1	1	4	3	3	1	1	1	2	3	3	3	3	1		
BF2_W_CCAGCCC	2	3	2	3	4	3	2	3	2	2	1	2	3	2	3	4	3	1	2	1	2	1	6	7	6	1	0	0	0	1	1	1	2	2	2	1	1	1	1	3	3	3	2	2	2	2	2	4	4	4	1			
BF2_W_CCAGCTC	1	2	1	3	3	4	1	2	1	2	1	1	2	1	3	3	4	0	1	0	2	1	6	6	7	1	0	0	0	1	1	2	1	1	1	2	1	1	1	3	3	4	1	1	1	3	2	3	3	3	2			
BF2_Y_GACAGGC	3	3	3	1	1	1	1	2	1	2	2	1	1	1	4	4	4	3	4	3	3	2	1	1	1	7	2	3	2	0	1	0	2	2	2	2	2	2	2	2	3	2	0	1	0	1	1	1	2	1	0	0	0	
BR2_2_AATCAGG	3	2	2	2	2	2	1	1	1	2	2	3	2	2	1	1	1	1	1	1	2	3	0	0	0	2	7	6	6	1	2	1	0	1	0	1	3	5	4	4	2	3	2	0	1	0	1	0	0	1	0	2		
BR2_2_AATCGGG	2	2	2	2	2	2	2	2	2	2	3	2	2	2	1	1	1	2	2	2	3	2	0	0	0	3	6	7	6	1	2	1	0	1	0	0	2	4	5	4	2	3	2	0	1	0	0	0	0	1	0	1		
BR2_2_AATCTGG	2	2	3	2	2	2	1	1	1	3	2	2	2	3	1	1	1	1	1	1	2	2	0	0	0	2	6	6	7	1	2	1	1	2	1	0	2	4	4	5	2	3	2	1	2	1	0	1	1	2	1	1		
BR2_3_TGGTCAG	1	2	1	3	2	2	1	1	1	2	3	1	2	1	2	1	1	0	0	0	1	3	2	1	1	0	1	1	1	7	6	6	0	1	0	3	1	1	1	1	4	3	3	2	3	2	1	2	1	2	1	2		
BR2_3_TGGTCGG	1	2	1	2	2	2	1	1	1	3	2	1	2	1	1	1	1	0	0	0	2	2	1	1	1	1	2	2	2	6	7	6	0	1	0	3	0	2	2	2	3	4	3	2	3	2	1	1	1	2	1	2		
BR2_3_TGGTCTG	1	2	1	2	2	3	1	1	1	2	2	1	2	1	1	1	2	0	0	0	1	2	1	1	2	0	1	1	1	6	6	7	0	1	0	4	0	1	1	1	3	3	4	2	3	2	2	1	1	2	1	3		
BR2_4_GTAATCA	2	2	3	0	1	0	3	2	2	1	1	2	2	3	1	2	1	4	3	3	1	0	1	2	1	2	0	0	1	0	0	0	7	6	6	0	1	1	1	2	1	1	1	3	2	2	2	3	4	3	3	1		
BR2_4_GTAATCG	2	2	3	0	1	0	2	2	2	1	2	2	2	3	1	2	1	3	3	3	1	1	1	2	1	2	1	1	2	6	7	6	0	1	2	2	3	2	2	2	2	2	3	2	2	3	3	4	3	1				
BR2_4_GTAATCT	2	2	3	0	1	0	2	2	3	1	1	2	2	3	1	2	1	3	3	4	1	0	1	2	1	2	0	0	1	0	0	0	6	6	7	0	2	1	1	2	1	1	1	2	2	3	2	4	3	3	4	2		
BR2_5_TGCTATC	2	1	1	2	2	3	1	2	1	2	1	2	1	1	2	2	3	1	2	1	1	3	1	1	2	2	1	0	0	3	3	4	0	0	0	7	1	1	0	0	1	1	2	3	3	3	3	1	1	1	1	4		
BR2_6_GATGAAT	3	2	2	3	2	2	1	1	2	1	2	1	0	0	4	3	3	0	0	1	0	4	2	1	1	2	2	3	2	2	1	0	0	1	1	7	3	2	2	1	0	0	0	0	0	1	1	2	0	0	1	2		
BR2_B_GCTCAGG	3	2	2	1	1	1	1	1	1	1	3	2	1	1	1	1	1	0	0	0	2	2	1	1	1	2	5	4	4	1	2	1	1	2	1	1																		