

Additional file 18. Alignment of Acala Ultima (AU) and TM-1 (Ref) alleles of Gh\_A07G2049. Primer locations for *CFBd0004* are highlighted yellow.

AU	AAT	CTT	CATT	TCT	GCT	TC	CT	TTG	G	G	T	AT	GG	TT	CG	ATT	CTT	TTG	GA	AT	GAG	AT	GAT	TT	ACC	AG	TA	TT	GCC	CT	TC	CAT	CT	GGAA	90		
Ref	AAT	CTT	CATT	TCT	GCT	TC	CT	TTG	G	G	T	AT	GG	TT	CG	ATT	CTT	TTG	GA	AT	GAG	AT	GAT	TT	ACC	AG	TA	TT	GCC	CT	TC	CAT	CT	GGAA	90		
AU	ATC	GAAT	GCT	TCAC	TTTC	CGA	TCCA	ACAC	CT	G	A	CAG	T	ATG	AGA	AT	TG	AGA	AT	TG	AGA	AT	TG	GGT	TTA	AT	ACCG	G	T	CT	GT	CT	ACT	G	TTT	CA	180
Ref	ATC	GAAT	GCT	TCAC	TTTC	CGA	TCCA	ACAC	CT	G	A	CAG	T	ATG	AGA	AT	TG	AGA	AT	TG	AGA	AT	TG	GGT	TTA	AT	ACCG	G	T	CT	GT	CT	ACT	G	TTT	CA	180
AU	TCAC	CTT	CAT	CTTC	CAT	CT	TTT	GCT	CT	T	C	T	CT	GGT	G	T	TT	CTT	C	CT	T	GGT	TT	AT	AC	CC	A	G	T	TT	CA	AT	TTT	AG	CT	270	
Ref	TCAC	CTT	CAT	CTTC	CAT	CT	TTT	GCT	CT	T	C	T	CT	GGT	G	T	TT	CTT	C	CT	T	GGT	TT	AT	AC	CC	A	G	T	TT	CA	AT	TTT	AG	CT	270	
AU	CCTT	GT	TAAT	C	T	CT	TTG	TGGA	T	C	T	TTT	GGAGT	A	G	T	C	T	A	T	C	A	T	A	G	T	C	T	G	T	T	T	C	G	360		
Ref	CCTT	GT	TAAT	C	T	CT	TTG	TGGA	T	C	T	TTT	GGAGT	A	G	T	C	T	A	T	C	A	T	A	G	T	C	T	G	T	T	T	C	G	360		
AU	CCTT	CATT	CT	TG	G	CT	TTT	CC	TT	A	T	AT	AT	CGG	T	C	G	T	CT	G	T	CT	G	AT	AT	AT	T	C	T	T	CC	A	450				
Ref	CCTT	CATT	CT	TG	G	CT	TTT	CC	TT	A	T	AT	AT	CGG	T	C	G	T	CT	G	T	CT	G	AT	AT	AT	T	C	T	T	CC	A	450				
AU	ACA	ATTT	TG	CT	GAGG	TTTC	AG	AA	G	C	T	A	T	C	T	G	T	TT	GG	AA	CT	TT	G	CC	A	TT	G	AT	TT	T	GC	ACT	CA	540			
Ref	ACA	ATTT	TG	CT	GAGG	TTTC	AG	AA	G	C	T	A	T	C	T	G	T	TT	GG	AA	CT	TT	G	CC	A	TT	G	AT	TT	T	GC	ACT	CA	540			
AU	ATAGA	AT	CT	T	C	T	CT	CC	T	GG	A	A	T	T	G	T	CC	AA	T	G	T	A	T	T	G	A	T	T	G	T	A	T	T	G	630		
Ref	ATAGA	AT	CT	T	C	T	CT	CC	T	GG	A	A	T	T	G	T	CC	AA	T	G	T	A	T	T	G	A	T	T	G	T	A	T	T	G	630		
AU	TTT	AG	ACT	CC	AT	GG	AT	GG	AC	CT	T	TT	TT	TT	AG	CC	C	T	TT	AA	CT	AC	TT	TT	CA	TT	TT	CT	AT	TT	CT	720					
Ref	TTT	AG	ACT	CC	AT	GG	AT	GG	AC	CT	T	TT	TT	TT	AG	CC	C	T	TT	AA	CT	AC	TT	TT	CA	TT	TT	CT	AT	TT	CT	720					
AU	AAT	GT	GAT	AT	C	T	TTT	TT	CT	TT	A	T	A	T	G	T	CT	T	TT	C	AT	T	C	T	TT	TT	CA	TT	TT	CA	AC	810					
Ref	AAT	GT	GAT	AT	C	T	TTT	TT	CT	TT	A	T	A	T	G	T	CT	T	TT	C	AT	T	C	T	TT	TT	CA	TT	TT	CA	AC	810					
AU	GGT	T	CT	TT	TC	TT	TG	AT	TG	TC	TT	GC	AG	CT	TT	GG	AA	GG	T	GG	CT	TT	GG	AA	AT	CA	AG	AT	CA	CT	TT	AG	CA	900			
Ref	GGT	T	CT	TT	TC	TT	TG	AT	TG	TC	TT	GC	AG	CT	TT	GG	AA	GG	T	GG	CT	TT	GG	AA	AT	CA	AG	AT	CA	CT	TT	AG	CA	900			
AU	AGG	CA	AG	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	990				
Ref	AGG	CA	AG	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	AA	990				
AU	CTT	TT	GT	TT	TT	TT	TT	GT	TA	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1080				
Ref	CTT	TT	GT	TT	TT	TT	TT	GT	TA	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1080				
AU	ATT	TA	AG	TT	AG	CA	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1170				
Ref	ATT	TA	AG	TT	AG	CA	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1170				
AU	TTA	AA	GG	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1260				
Ref	TTA	AA	GG	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1260				
AU	TT	CC	CT	G	CT	CC	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	1350				
Ref	TT	CC	CT	G	CT	CC	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	AT	1350				
AU	TG	C	AT	T	CT	AC	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1440				
Ref	TG	C	AT	T	CT	AC	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1440				
AU	CG	TG	TC	GA	GT	AA	TT	CC	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1530				
Ref	CG	TG	TC	GA	GT	AA	TT	CC	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1530				
AU	CT	CC	TT	GATT	TT	GA	TTT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1620				
Ref	CT	CC	TT	GATT	TT	GA	TTT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1620				
AU	TTC	AG	CC	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1710				
Ref	TTC	AG	CC	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	TT	1710				
AU	ATC	AA	TT	CC</td																																	

	2,260	2,280	2,300	2,320	2,340					
AU	TTGGCCACTC	AGTTAGTGGT	TATCGAAGAC	AATAATAGAA	AGGAGAACATGA	ATAAGAAAATA	CATGCAATTAA	TTAAGGTTTT	GAGCTAGTAT	2340
Ref	TTGGCCACTC	AGTTAGTGGT	TATCGAAGAC	AATAATAGAA	AGGAGAACATGA	ATAAGAAAATA	CATGCAATTAA	TTAAGGTTTT	GAGCTAGTAT	2340
	2,360	2,380	2,400	2,420						
AU	TGCTTACCTC	TTCTTTGAA	ACTTCATTC	GTATGATTTC	ATTTCTTC	TTTGTGTTG	CACACTGATT	TTCTCACCG	TCTGTCCC	2430
Ref	TGCTTACCTC	TTCTTTGAA	ACTTCATTC	GTATGATTTC	ATTTCTTC	TTTGTGTTG	CACACTGATT	TTCTCACCG	TCTGTCCC	2430
	2,440	2,460	2,480	2,500	2,520					
AU	CTCTTGCTT	GACAATATTTC	TCTTTGATTG	CAGCATCTT	TGTTCTTC	TTGAGTAGTT	CATTGTC	AGCCTTTCA	TGTATCTCCC	2520
Ref	CTCTTGCTT	GACAATATTTC	TCTTTGATTG	CAGCATCTT	TGTTCTTC	TTGAGTAGTT	CATTGTC	AGCCTTTCA	TGTATCTCCC	2520
	2,540	2,560	2,580	2,600						
AU	TTTTGAAATT	CGAACGCTCT	TCAACCTCTC	CTTTAGTTC	CGGAATCTTC	TCGTGACCCT	CATCTTCTT	AAATTCTGTT	TTATTTGTTT	2610
Ref	TTTTGAAATT	CGAACGCTCT	TCAACCTCTC	CTTTAGTTC	CGGAATCTTC	TCGTGACCCT	CATCTTCTT	AAATTCTGTT	TTATTTGTTT	2610
	2,620	2,640	2,660	2,680	2,700					
AU	CATCTCGTTG	GACCTCAGAA	GGCAATACAG	AGGTTGTGAT	ATGATGGTGG	TTAGTTTAC	CTCTCAAAT	GGCTTCGG	TCCCTTATCT	2700
Ref	CATCTCGTTG	GACCTCAGAA	GGCAATACAG	AGGTTGTGAT	ATGATGGTGG	TTAGTTTAC	CTCTCAAAT	GGCTTCGG	TCCCTTATCT	2700
	2,720	2,740	2,760	2,780						
AU	TGTCCTTTTC	CTGGCTTATC	TTTCTCTT	TCTTTACTCC	ATCAATGTG	CCATTTTCAG	TTCTTCTTTC	CTTCATATCC	TCCCTCCTCGG	2790
Ref	TGTCCTTTTC	CTGGCTTATC	TTTCTCTT	TCTTTACTCC	ATCAATGTG	CCATTTTCAG	TTCTTCTTTC	CTTCATATCC	TCCCTCCTCGG	2790
	2,800	2,820	2,840	2,860	2,880					
AU	TATTGCCAAC	TTCATCAACA	GTTTTGTGCT	TATACTCTC	AAATTGCAAT	GTGGCAACCA	AATTGAAAT	TTGCTCCTTG	GGTTTCTCGG	2880
Ref	TATTGCCAAC	TTCATCAACA	GTTTTGTGCT	TATACTCTC	AAATTGCAAT	GTGGCAACCA	AATTGAAAT	TTGCTCCTTG	GGTTTCTCGG	2880
	2,900	2,920	2,940	2,960						
AU	TTGCGGTAGT	CTGAGAAGTA	GCATTATCAT	CTACTGCAAG	AGCTTCATCC	TCCTTAAGCT	GTTCTTTTG	CAAGGTCACA	TCACTCTCAA	2970
Ref	TTGCGGTAGT	CTGAGAAGTA	GCATTATCAT	CTACTGCAAG	AGCTTCATCC	TCCTTAAGCT	GTTCTTTTG	CAAGGTCACA	TCACTCTCAA	2970
	2,980	3,000	3,020	3,040	3,060					
AU	TATCACTTCG	TAAGAATTTT	GCTTCTCCA	TTTCTTACA	TTCTATCTTC	CCGGCTGTAC	CAGCAGTCAT	TTCATCTTCT	TTGACAGCTT	3060
Ref	TATCACTTCG	TAAGAATTTT	GCTTCTCCA	TTTCTTACA	TTCTATCTTC	CCGGCTGTAC	CAGCAGTCAT	TTCATCTTCT	TTGACAGCTT	3060
	3,080	3,100	3,120	3,140						
AU	CTAGCCCCAAT	GGAACCTTCG	TTGTGTGAGC	TTTCTCTAC	TGTAATCTGT	GCATCAGAAC	CATCTTCTC	TGCTTGTCT	TTTATGATT	3150
Ref	CTAGCCCCAAT	GGAACCTTCG	TTGTGTGAGC	TTTCTCTAC	TGTAATCTGT	GCATCAGAAC	CATCTTCTC	TGCTTGTCT	TTTATGATT	3150
	3,160	3,180	3,200	3,220	3,240					
AU	CCCCCTCGCT	TGTTTCTCA	TCTTCTGTCT	CCTTAGTTTC	AGGCTACCA	GGTCCTTCCT	TTTCAAGCAT	TCAGGTTCC	TTCACCTTTT	3240
Ref	CCCCCTCGCT	TGTTTCTCA	TCTTCTGTCT	CCTTAGTTTC	AGGCTACCA	GGTCCTTCCT	TTTCAAGCAT	TCAGGTTCC	TTCACCTTTT	3240
	3,260	3,280	3,300	3,320						
AU	CTTCTTCTTT	GCTGGCATCA	GCTTCATGCT	TTGGGTTCTG	AAACAGGAAT	TCTAAATACA	AAGATTGTTTC	ATTGATTCC	TCCATTGAA	3330
Ref	CTTCTTCTTT	GCTGGCATCA	GCTTCATGCT	TTGGGTTCTG	AAACAGGAAT	TCTAAATACA	AAGATTGTTTC	ATTGATTCC	TCCATTGAA	3330
	3,340	3,360	3,380	3,400	3,420					
AU	GAGCTTGAGC	AGCAGCAGTT	TCATCGTTG	TTGCATCATT	AAGCTCTGA	AACTCTTGT	TTTCTAAGCA	TGTCTCAGCT	TTTTCTTCTG	3420
Ref	GAGCTTGAGC	AGCAGCAGTT	TCATCGTTG	TTGCATCATT	AAGCTCTGA	AACTCTTGT	TTTCTAAGCA	TGTCTCAGCT	TTTTCTTCTG	3420
	3,440	3,460	3,480	3,500						
AU	CATCAGAAGG	ATCTTCTGAA	ATCTTGTCCG	GACTCATGTC	TGCTTCATTT	TTTTCTCTTT	CTTCCATCTT	TCAGCTTTG	CTCACATT	3510
Ref	CATCAGAAGG	ATCTTCTGAA	ATCTTGTCCG	GACTCATGTC	TGCTTCATTT	TTTTCTCTTT	CTTCCATCTT	TCAGCTTTG	CTCACATT	3510
	3,520	3,540	3,560	3,580	3,600					
AU	CTTCTTCGTC	CGAAACATAAT	GGAGAGCTCG	TAGTCTTCTC	CTCGGGTTTA	TTAGTCTCAA	GAGTCTGCTC	ACCAGTTATT	CCATCTCCG	3600
Ref	CTTCTTCGTC	CGAAACATAAT	GGAGAGCTCG	TAGTCTTCTC	CTCGGGTTTA	TTAGTCTCAA	GAGTCTGCTC	ACCAGTTATT	CCATCTCCG	3600
	3,620	3,640	3,660	3,680						
AU	CAACAGCAGTC	TAGCTTGTG	GGTATTTCTT	CCTTTGTTT	GCCATATTCT	ATCTCCTCAT	TTATGGCTCC	AGTTCCATGT	TCGCTTCTC	3690
Ref	CAACAGCAGTC	TAGCTTGTG	GGTATTTCTT	CCTTTGTTT	GCCATATTCT	ATCTCCTCAT	TTATGGCTCC	AGTTCCATGT	TCGCTTCTC	3690
	3,700	3,720	3,740	3,760	3,780					
AU	TGGAAGCCAA	TGTTGAAAAT	GTAGTTGAA	TTTCTCTTC	CGGTTCCACA	TTAGGAAGAC	CCTCATCAGC	AATTGTC	ACTGCTTAA	3780
Ref	TGGAAGCCAA	TGTTGAAAAT	GTAGTTGAA	TTTCTCTTC	CGGTTCCACA	TTAGGAAGAC	CCTCATCAGC	AATTGTC	ACTGCTTAA	3780
	3,800	3,820	3,840	3,860						
AU	GCTCTCCTTG	TTCTTCATTT	GGCAAGCGTA	CCTCTTCTAT	TCTCTTGTG	TCAGAAGAAT	CTTCTCGGCT	CTCGTGTGTTT	ATTCAGCAT	3870
Ref	GCTCTCCTTG	TTCTTCATTT	GGCAAGCGTA	CCTCTTCTAT	TCTCTTGTG	TCAGAAGAAT	CTTCTCGGCT	CTCGTGTGTTT	ATTCAGCAT	3870
	3,880	3,900	3,920	3,940	3,960					
AU	CCTTCTGGAT	CTCTTCTTC	ATATTCTCAA	TCGGGTTTG	AATTCTGATA	TTTTCACCC	CAGAAAGCAA	TGTGCAAGCA	GAACATTGAA	3960
Ref	CCTTCTGGAT	CTCTTCTTC	ATATTCTCAA	TCGGGTTTG	AATTCTGATA	TTTTCACCC	CAGAAAGCAA	TGTGCAAGCA	GAACATTGAA	3960
	3,980	4,000	4,020	4,040						
AU	GTTGCTCATC	TGGTTCTCT	TCTGGAAAGAG	TATTGGTTGA	ACCAATATCA	TCGATGAGAA	TAGTTTCAAC	TCTTTGAGAG	ATGTAATCAG	4050
Ref	GTTGCTCATC	TGGTTCTCT	TCTGGAAAGAG	TATTGGTTGA	ACCAATATCA	TCGATGAGAA	TAGTTTCAAC	TCTTTGAGAG	ATGTAATCAG	4050
	4,060	4,080	4,100	4,120	4,140					
AU	ATGATGTTTC	AGGTACTCCT	TCAAATGTC	CAGTTTTATT	GTCTTCCAGT	TCTGCTGTTG	GGCCTTTCTC	AGGTTCAGCA	ACCTGATCTT	4140
Ref	ATGATGTTTC	AGGTACTCCT	TCAAATGTC	CAGTTTTATT	GTCTTCCAGT	TCTGCTGTTG	GGCCTTTCTC	AGGTTCAGCA	ACCTGATCTT	4140
	4,160	4,180	4,200	4,220						
AU	TGTGCATGTA	AAAGTGCATCT	GCATCTCTCT	TGATTACCTC	CACTTCATG	TTGTCATTAA	GTTGGACTTC	AAGTGCAG	TCGGAGGATA	4230
Ref	TGTGCATGTA	AAAGTGCATCT	GCATCTCTCT	TGATTACCTC	CACTTCATG	TTGTCATTAA	GTTGGACTTC	AAGTGCAG	TCGGAGGATA	4230
	4,240	4,260	4,280	4,300	4,320					
AU	TATCTTCAAG	CTGCCCCATC	TCTACTTCTT	AGTCTCTTG	ATTGGCATCA	TCTTAATGG	TTTCTAGACC	CATCTTGACT	GAGCTGATAT	4320
Ref	TATCTTCAAG	CTGCCCCATC	TCTACTTCTT	AGTCTCTTG	ATTGGCATCA	TCTTAATGG	TTTCTAGACC	CATCTTGACT	GAGCTGATAT	4320
	4,340	4,360	4,380	4,400						
AU	TCAAATGTTATT	TATGAGCTCC	TCATTCACAG	CATCGTTAA	GCTCATCTGT	GCTCTTAAGT	CTGTCGCAAC	TTCAATATGA	CTCTCTTCAC	4410
Ref	TCAAATGTTATT	TATGAGCTCC	TCATTCACAG	CATCGTTAA	GCTCATCTGT	GCTCTTAAGT	CTGTCGCAAC	TTCAATATGA	CTCTCTTCAC	4410
	4,420	4,440	4,460	4,480	4,500					
AU	GGCAGAGTTT	CACGTCACTG	ATAATCTCA	CTTTTCTTC	TGCTTCTCA	TCTTCACGG	ACACTTTTG	TGGATCTG	GACTCTTCAC	4500
Ref	GGCAGAGTTT	CACGTCACTG	ATAATCTCA	CTTTTCTTC	TGCTTCTCA	TCTTCACGG	ACACTTTTG	TGGATCTG	GACTCTTCAC	4500

AU CCTGTCTCCT TTCTTCAAT TCTTCTATAA TGTTGGATCC TTCTTGTCA TGGGTTGATT TTCTTCAAC CTCTGTTGT TCTGCTTCA 4590  
 Ref CCTGTCTCCT TTCTTCAAT TCTTCTATAA TGTTGGATCC TTCTTGTCA TGGGTTGATT TTCTTCAAC CTCTGTTGT TCTGCTTCA 4590  
 AU 4,600 4,620 4,640 4,660 4,680  
 AU TGTGCTCT GGACTCAGCT TCCTTGATAC ATTCAGTTGA GACTGAATTG TAGTCTTTG CCTGTTAAC AATAGTTAA ATGTTGAGA 4680  
 Ref TGTGCTCT GGACTCAGCT TCCTTGATAC ATTCAGTTGA GACTGAATTG TAGTCTTTG CCTGTTAAC AATAGTTAA ATGTTGAGA 4680  
 AU 4,700 4,720 4,740 4,760  
 AU TTTTGACACA TAAAAAGAAC TATTTTAATA GAAGTATATA ACTATTTATT AGAAATTACT GAGATGACCT TATTTGCTTC ATCTCCTCTT 4770  
 Ref TTTTGACACA TAAAAAGAAC TATTTTAATA GAAGTATATA ACTATTTATT AGAAATTACT GAGATGACCT TATTTGCTTC ATCTCCTCTT 4770  
 AU 4,780 4,800 4,820 4,840 4,860  
 AU GATACTGGAG GAATTTCATT TGACTACTT TCTTCCTCCA TGACATGTT TTTGATGGCT TCACTAACAT CAGCTTCTC AACCTCCTTA 4860  
 Ref GATACTGGAG GAATTTCATT TGACTACTT TCTTCCTCCA TGACATGTT TTTGATGGCT TCACTAACAT CAGCTTCTC AACCTCCTTA 4860  
 AU 4,880 4,900 4,920 4,940  
 AU TCAACCTGCC AACAAATGATA ATGTAACTGA AATTCAACCTT GAATAAACAT GAAGACAAAA TATCTTACAA GCTCTTACCC TTTTCAGACA 4950  
 Ref TCAACCTGCC AACAAATGATA ATGTAACTGA AATTCAACCTT GAATAAACAT GAAGACAAAA TATCTTACAA GCTCTTACCC TTTTCAGACA 4950  
 AU 4,960 4,980 5,000 5,020 5,040  
 AU AAACGACAAAC GTTGGTAACC TTGTTTTCAT TTTCTTCTT GTAAATATAG TCTTCATCCT TCAGTTTGT TTTCAGTTCC TCTGTAATAT 5040  
 Ref AAACGACAAAC GTTGGTAACC TTGTTTTCAT TTTCTTCTT GTAAATATAG TCTTCATCCT TCAGTTTGT TTTCAGTTCC TCTGTAATAT 5040  
 AU 5,060 5,080 5,100 5,120  
 AU GTTCCCTTGAT TTCTTGACTG GTTTCATCAT CTTGGCTCTT GGTGCTTCA GGGATTGTTT CATTACTTTG GTCTTCAGGCC ACGGGTTCCG 5130  
 Ref GTTCCCTTGAT TTCTTGACTG GTTTCATCAT CTTGGCTCTT GGTGCTTCA GGGATTGTTT CATTACTTTG GTCTTCAGGCC ACGGGTTCCG 5130  
 AU 5,140 5,160 5,180 5,200 5,220  
 AU AGTCGGATTG GGGCTTTCT TCAAAACCTG CTAGTGCTGG CTCAACATTC TGTTCGAAAT TAATACATTA GTCGAGGGCT TGAGAAATCT 5220  
 Ref AGTCGGATTG GGGCTTTCT TCAAAACCTG CTAGTGCTGG CTCAACATTC TGTTCGAAAT TAATACATTA GTCGAGGGCT TGAGAAATCT 5220  
 AU 5,240 5,260 5,280 5,300 5,400  
 AU GCTACTTAAA TGACCGAGAAG CAATTGAATG GTGATAACTG ATATGACAAA CGAACAAAGAT TAGACATACT ATACAACATAT GTTTTCCAT 5310  
 Ref GCTACTTAAA TGACCGAGAAG CAATTGAATG GTGATAACTG ATATGACAAA CGAACAAAGAT TAGACATACT ATACAACATAT GTTTTCCAT 5310  
 AU 5,320 5,340 5,360 5,380 5,400  
 AU TATCGTATAC ATATGAAGTT TTGGTTTTG TTTCTATTG ATAATTGAT TTGACGAGGT TCTTCATTGA GTTAACCATT TTCTATACTA 5400  
 Ref TATCGTATAC ATATGAAGTT TTGGTTTTG TTTCTATTG ATAATTGAT TTGACGAGGT TCTTCATTGA GTTAACCATT TTCTATACTA 5400  
 AU 5,420 5,440 5,460 5,480  
 AU AAACCTGAAT CCAGTTGTC ATTGGAACGT GAATTCAACA AGTCTGTTGC AGTTCCTATT ACATTTACAT TTGCATAAAA TATTATGCTA 5490  
 Ref AAACCTGAAT CCAGTTGTC ATTGGAACGT GAATTCAACA AGTCTGTTGC AGTTCCTATT ACATTTACAT TTGCATAAAA TATTATGCTA 5490  
 AU 5,500 5,520 5,540 5,560 5,580  
 AU ACTTTGAAAG GAGGGTTGAA AGTAGTCTGA CCTCTTCTGT TACTGTTATA GTTCAATCC TGGAAACTGGA AGCATGGCTT GGATCCTCAA 5580  
 Ref ACTTTGAAAG GAGGGTTGAA AGTAGTCTGA CCTCTTCTGT TACTGTTATA GTTCAATCC TGGAAACTGGA AGCATGGCTT GGATCCTCAA 5580  
 AU 5,600 5,620 5,640 5,660  
 AU AAGTTTATC TTCAATCTGG TTTCAATTAA TCTCAATTTC ACTGGTCTTT TGAGCTTCCT GAACGAGCTG TTATAGATTT AATATTCAAT 5670  
 Ref AAGTTTATC TTCAATCTGG TTTCAATTAA TCTCAATTTC ACTGGTCTTT TGAGCTTCCT GAACGAGCTG TTATAGATTT AATATTCAAT 5670  
 AU 5,680 5,700 5,720 5,740 5,760  
 AU ATGTATTTAT ATAAGAGAAA AAAATGGAAG ATGAACAAAG TTATAGTTTC TGTTTCTTA CCTCTCCATA TGTCCTGCCT TCAGTCGGGT 5760  
 Ref ATGTATTTAT ATAAGAGAAA AAAATGGAAG ATGAACAAAG TTATAGTTTC TGTTTCTTA CCTCTCCATA TGTCCTGCCT TCAGTCGGGT 5760  
 AU 5,780 5,800 5,820 5,840 5,940  
 AU TTTCTTTAC CTCATCTTC CCAGAGCCAT CTTTGATCAG CTCTTCCGAA CAGGAAGTTG AATAAGTTGT TTCACTAAT TTCTTGTCT 5850  
 Ref TTTCTTTAC CTCATCTTC CCAGAGCCAT CTTTGATCAG CTCTTCCGAA CAGGAAGTTG AATAAGTTGT TTCACTAAT TTCTTGTCT 5850  
 AU 5,860 5,880 5,900 5,920 5,940  
 AU TTTTGACCTG TCAAGTTGA CCCTTAGTTA ATAGTCATCT AGATAACAAAT AGAAACGAGA ATACATAGAA AATAGATGAA AAATTATGAA 5940  
 Ref TTTTGACCTG TCAAGTTGA CCCTTAGTTA ATAGTCATCT AGATAACAAAT AGAAACGAGA ATACATAGAA AATAGATGAA AAATTATGAA 5940  
 AU 5,960 5,980 6,000 6,020  
 AU TTTTGAGCT AGTATTGGTT ACCTGTTCTA TCGAAACTTC ATTCTTGGG ATTCTATTTC CTTCATTTGT TTCTCACAC TGGTTTACTT 6030  
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AU	TGTCCTTTT	GGACTCCTTG	AGCTAAAGCG	ACAAACTACA	GACATGGTCT	TCTTCACATT	CTATTGGTGT	TGTTTCTTCT	TTTGACGCCT	13860
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AU	AAACAATTGG	ATATATATAT	TGATGGAAGA	CTGAGTTCA	AAACATTGAA	TAAGTGGAAA	AGTTAAGATG	TTAGTGAGTC	ATACTATCTA	14040
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AU	TTGTTATCTT	CCTCATCTAG	AATTGTTTTT	TCTGTATTCT	CAACGGTGT	GTCTGGTTCC	AGATTTTCAG	AGACCTGAAT	AAAAAGTAAT	14220
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AU	TGGAAACAATG	GATTCTCTGT	CCTCTCTCAT	TTCAAGACAT	CCCTTTGTTA	TGGCCAGCT	CTTGTCTGTC	ACCTGTATGA	CTCTCACATC	15210
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AU	CACCGCAACG	TTTCTGAGT	GCTTGTGTTG	ATCTTGATCT	TCTTCGAA	TGTCTCGGC	TATTTTCTTA	TCTTCATGA	GAGTCGTATC	15300
Ref	CACCGCAACG	TTTCTGAGT	GCTTGTGTTG	ATCTTGATCT	TCTTCGAA	TGTCTCGGC	TATTTTCTTA	TCTTCATGA	GAGTCGTATC	15300
AU	ATATTTTCC	AAAGGCATAG	AAGAATGTTC	CTTAACATCA	TCTCCATAA	CCTCTTCAA	CAGAATCTCC	TTGGAGGTCA	TTTCCATTG	15390
Ref	ATATTTTCC	AAAGGCATAG	AAGAATGTTC	CTTAACATCA	TCTCCATAA	CCTCTTCAA	CAGAATCTCC	TTGGAGGTCA	TTTCCATTG	15390
AU	ATTTTCAC	CCTGTGTTA	GAAATTATC	CTGCACTCCA	TCTTCTTCTC	GGGAAATCTC	GATGTCTCTC	TCACTTGTC	CATATGATT	15480
Ref	ATTTTCAC	CCTGTGTTA	GAAATTATC	CTGCACTCCA	TCTTCTTCTC	GGGAAATCTC	GATGTCTCTC	TCACTTGTC	CATATGATT	15480
AU	TACTGATT	ACCTCAATT	CTGCACTTC	TAAACCAATC	CCTTCATTT	CACCATCAAC	AACTTATT	ATTCATTCA	TTTCCCTTTC	15570
Ref	TACTGATT	ACCTCAATT	CTGCACTTC	TAAACCAATC	CCTTCATTT	CACCATCAAC	AACTTATT	ATTCATTCA	TTTCCCTTTC	15570
AU	TTCTGGTTCA	GCTT TAGCTT	CCTTTAAGTT	TGTTT TAC	GTTTCCGCTG	TTTGTGATGT	GCAGTCAA	TCAGCAGTAT	TGATTGTT	15660
Ref	TTCTGGTTCA	GCTT TAGCTT	CCTTTAAGTT	TGTTT TAC	GTTTCCGCTG	TTTGTGATGT	GCAGTCAA	TCAGCAGTAT	TGATTGTT	15660
AU	TTTACCCCT	GTTTCTTCTT	CAGTATCCAT	TTTGGACTCC	TCGAGCTAA	ATGAAGAGCT	GCAGACAAGT	TCTTCTTGA	ATTGATTAC	15750
Ref	TTTACCCCT	GTTTCTTCTT	CAGTATCCAT	TTTGGACTCC	TCGAGCTAA	ATGAAGAGCT	GCAGACAAGT	TCTTCTTGA	ATTGATTAC	15750

AU AGTAATTTCT TTTCTGCTG TACCTGAAGC TGCAGTTCT GGTCTCTAC ACAGCTGCAG GATATAAAA ATTCATAT AACTCAGACA 15840  
 Ref AGTAATTTCT TTTCTGCTG TACCTGAAGC TGCAGTTCT GGTCTCTAC ACAGCTGCAG GATATAAAA ATTCATAT AACTCAGACA 15840  
 AU AAAGTCTGGA TGGTTATGGA CTTATAAGAC AAATTCCAAT GGGCCATGGG TCGCTTTGT GCTAAGTTCA TGAACTGAAC TTTTATTCT 15930  
 Ref AAAGTCTGGA TGGTTATGGA CTTATAAGAC AAATTCCAAT GGGCCATGGG TCGCTTTGT GCTAAGTTCA TGAACTGAAC TTTTATTCT 15930  
 AU TTGAAATCCA AGTTATCGTG AGGTTTGCTA CTAGCATCAG TTGCCATAG ACACCTCTCT TATTAAGGAT CTCTATTTT GGCTTATCT 16020  
 Ref TTGAAATCCA AGTTATCGTG AGGTTTGCTA CTAGCATCAG TTGCCATAG ACACCTCTCT TATTAAGGAT CTCTATTTT GGCTTATCT 16020  
 AU TTTCTGGTT TGATCAGAAC TCACTCAGCT CTATATTTC ACACGTGTA ATGGATAGCC CATCACTTC GATTGATTG TTTGCATTG 16110  
 Ref TTTCTGGTT TGATCAGAAC TCACTCAGCT CTATATTTC ACACGTGTA ATGGATAGCC CATCACTTC GATTGATTG TTTGCATTG 16110  
 AU TACGCATGTG TTTGCCAATA ATAATGGTT TTTGTTCCAG TTGTAATTCT ACACCTCTTC AATTGTAATC TGATTATTGT TGGTGTCTC 16200  
 Ref TACGCATGTG TTTGCCAATA ATAATGGTT TTTGTTCCAG TTGTAATTCT ACACCTCTTC AATTGTAATC TGATTATTGT TGGTGTCTC 16200  
 AU CTCCTCTAGA ATTTGTTTC CATCATTCTC AACTGTGTT TCCGGTTCTA GCTTTCAGC GACCTGAATG AAAATATATT CATAGATTGG 16290  
 Ref CTCCTCTAGA ATTTGTTTC CATCATTCTC AACTGTGTT TCCGGTTCTA GCTTTCAGC GACCTGAATG AAAATATATT CATAGATTGG 16290  
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 Ref TTGAAACATGG AGGTTTCAGA TTTTAGCAA TAGATTGAAA TTGAGTTAGT ATTCAATGATA GGAACCTTT TGATTACCTC AGCTTCATCC 16380  
 AU TTGTCACAG TTTCATATT TAGATTCTGC TTTGCACTT CATCATTTC ATCAGTTCGG ACATCAGGAT CCGATTCAAGA GGATATCTCT 16470  
 Ref TTGTCACAG TTTCATATT TAGATTCTGC TTTGCACTT CATCATTTC ATCAGTTCGG ACATCAGGAT CCGATTCAAGA GGATATCTCT 16470  
 AU TCAAGCTGCA CAATCTCCAC TTCTGTCCT TGAGCATCAC TCTTAATATT TTCTACAGTC TTCTCGGCTG AGCTTATATT TGATGCATT 16560  
 Ref TCAAGCTGCA CAATCTCCAC TTCTGTCCT TGAGCATCAC TCTTAATATT TTCTACAGTC TTCTCGGCTG AGCTTATATT TGATGCATT 16560  
 AU ACAATCTTGT CGGTGTCGTT CAAGCTCGTC TCTGCTGTTA CTTCTAGTAC AGCTTCAAGA CGACTCTCT CATGACACGG TTTGAGTC 16650  
 Ref ACAATCTTGT CGGTGTCGTT CAAGCTCGTC TCTGCTGTTA CTTCTAGTAC AGCTTCAAGA CGACTCTCT CATGACACGG TTTGAGTC 16650  
 AU TCAATAATCT CCTTGGTTT CTCTTCTGCT TCTCGATCTT TGCTACGCA ACTTACAGGA TCTCCAGACT CTTCAGCCTG ACTCATTCT 16740  
 Ref TCAATAATCT CCTTGGTTT CTCTTCTGCT TCTCGATCTT TGCTACGCA ACTTACAGGA TCTCCAGACT CTTCAGCCTG ACTCATTCT 16740  
 AU TCTAAATTCT TGGTGTGTT GGATGATTCT ACATCATTGG TTGAAATGTT TTTGACCTCC AAGTTTCTG CAACTATCTG CTCTTGGAC 16830  
 Ref TCTAAATTCT TGGTGTGTT GGATGATTCT ACATCATTGG TTGAAATGTT TTTGACCTCC AAGTTTCTG CAACTATCTG CTCTTGGAC 16830  
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 Ref TCTGCTCCA TGAAACATTC AGTTGAGACA GAACCTGTAGT CTTCCACCTG CGCATAACAC AGATAAAATG TTTAACAAATA TAACATCCGA 16920  
 AU AAATCTGTT GAGTATCTAT TCATTTGAA AGTAAAAAGT GTAATTATTC ACTAACGTT GATAACCTT TTCAATATCTT CAATTGTTT 17010  
 Ref AAATCTGTT GAGTATCTAT TCATTTGAA AGTAAAAAGT GTAATTATTC ACTAACGTT GATAACCTT TTCAATATCTT CAATTGTTT 17010  
 AU TTGTTTTTA ACAAGAAAAA GAGCAGAATA AGTATTCACT AACCTTGCA TTATAAGTC TGTAACCAT CTCAGGAATC TGATTTCTT 17100  
 Ref TTGTTTTTA ACAAGAAAAA GAGCAGAATA AGTATTCACT AACCTTGCA TTATAAGTC TGTAACCAT CTCAGGAATC TGATTTCTT 17100  
 AU TTTCTATTTC AGTATCTTCC TTCAAGCTT CCGGTTCTC CATGTTCT GCTAATTCT CTTCTTCAGT AGCCTTCTCA TCATCCTGT 17190  
 Ref TTTCTATTTC AGTATCTTCC TTCAAGCTT CCGGTTCTC CATGTTCT GCTAATTCT CTTCTTCAGT AGCCTTCTCA TCATCCTGT 17190  
 AU GAGTACCCAA GGTTGCTTC AAATCCACAG CTTCATCCAA TGGTTTCCA TCTTCATT TTGATCTATC CAGTCCTACA TCTGATGGAT 17280  
 Ref GAGTACCCAA GGTTGCTTC AAATCCACAG CTTCATCCAA TGGTTTCCA TCTTCATT TTGATCTATC CAGTCCTACA TCTGATGGAT 17280  
 AU TTTCAACTGG CAGTGCTTT GCTTCGCAAG CATTGGTTG ACTCTCGAGT TCTCTATGTT CTTCAACAGT AGAGACATCG GTGACCTCTA 17370  
 Ref TTTCAACTGG CAGTGCTTT GCTTCGCAAG CATTGGTTG ACTCTCGAGT TCTCTATGTT CTTCAACAGT AGAGACATCG GTGACCTCTA 17370  
 AU GATTAGATTC CTTGGCCAACT TGTTCCTCGA CTACTTGTC ACCTCCGTT CCCGTATCTT TAGACACAGC TTTAGGGACA TCTACTAGCG 17460  
 Ref GATTAGATTC CTTGGCCAACT TGTTCCTCGA CTACTTGTC ACCTCCGTT CCCGTATCTT TAGACACAGC TTTAGGGACA TCTACTAGCG 17460  
 AU GATTATCTTC ACTTTGTATT AATTTCTCAT CCGTTATTG TGGCATAGGA GTTTCTGAGT GCTCACGCTG ATCCTGATCT TTGTCGAAG 17550  
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 AU TTTCGTCAGT TATTTCCCT TCTTCAGAGA ATGTTGCCCT AGATTCCTTC ACTTCATCTC CTAATGTTT TTTCAAGGGAA ATCTCCTTGG 17640  
 Ref TTTCGTCAGT TATTTCCCT TCTTCAGAGA ATGTTGCCCT AGATTCCTTC ACTTCATCTC CTAATGTTT TTTCAAGGGAA ATCTCCTTGG 17640  
 AU TGGTCATCTC CAATTGATCT TCACCTCCTG CATATGGAAC TTTATCCTGC ACTGTATCTT CTTGTCGGA ATCTCCATGT CCTCTCGTGT 17730  
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 AU TGTTACGTAC GAGTCCCCGT ATTTCACATC AATTTCAGCA CTGCATACAC CACTCTCTTC ATGTTCAAGT CTTGCTGAGC AGTTCTCTT 17820  
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 AU GATGACAACA ACATTATAGA TTTTGTCTT TGTCTTTCTT CTTTCAGCTT CAGCT----- TCAGCAT CAGCATCAGC 17892  
 Ref GATGACAACA ACATTATAGA TTTTGTCTT TGTCTTTCTT CTTTCAGCTT CAGCTTCAGC ATCAGCAT CAGCATCAGC 17910  
 AU TTCACTTCA GCTTCAGCAT CAGCATCAGC TTTCAGCTTC GCTTCCTTTA TGTCTGTTTC TTCTGTTCT CAGCATCAGC 17982  
 Ref TTCACTTCA GCTTCAGCAT CAGCATCAGC TTTCAGCTTC GCTTCCTTTA TGTCTGTTTC TTCTGTTCT CAGCATCAGC 18000

AU TTCTTCTTTG GTGTCGTTTG TGGACTCCTT GAGCTCAAAC TCCGAACATAC GAGCCTGGTC TTCTTCACT TCTATTGGTG TTATTCCTTC 18072  
 Ref TTCTTCTTTG GTGTCGTTTG TGGACTCCTT GAGCTCAAAC TCCGAACATAC GAGCCTGGTC TTCTTCACT TCTATTGGTG TTATTCCTTC 18090  
 AU TTTTGACTCC TGTACAATAT CATACAGACT AGAATTGTTA TTTTGCTTTG CTGTTCTCT TATGGACACT GTTTCTCAA TGATTGTGTG 18162  
 Ref TTTTGACTCC TGTACAATAT CATACAGACT AGAATTGTTA TTTTGCTTTG CTGTTCTCT TATGGACACT GTTTCTCAA TGATTGTGTG 18180  
 AU GCTCGTCTGT AAAACATTGA ATATATAAGT TGATGAAAGA TTGAATTCAAG AGCATTGAA TAAGTGGAAA AGTTAACATG TTAGTGTATTG 18252  
 Ref GCTCGTCTGT AAAACATTGA ATATATAAGT TGATGAAAGA TTGAATTCAAG AGCATTGAA TAAGTGGAAA AGTTAACATG TTAGTGTATTG 18270  
 AU ATACAATCTA CTTACCACTG TTATCGACAC CACTGCTGTA GTATTCTT CCTTAGCTTT TTCTTCTTC GTCTCAGTCA TTAGTGTGTTG 18342  
 Ref ATACAATCTA CTTACCACTG TTATCGACAC CACTGCTGTA GTATTCTT CCTTAGCTTT TTCTTCTTC GTCTCAGTCA TTAGTGTGTTG 18360  
 AU GTGATCCTTC TTGTTGCTTT CCTCCTCTAG AAATTGTTT TCAGCATTG CTACTGTGTT GTCTGGTCC AGCTTTCAC AGACCTGAAC 18432  
 Ref GTGATCCTTC TTGTTGCTTT CCTCCTCTAG AAATTGTTT TCAGCATTG CTACTGTGTT GTCTGGTCC AGCTTTCAC AGACCTGAAC 18450  
 AU AAAAATAGTA TCAAAAGATT GGTAACAT GGAGGTTTC AGCTTCAAC TTGATCTTC GACTTCAAA ACAAACAAAA AAAATTCTT 18522  
 Ref AAAAATAGTA TCAAAAGATT GGTAACAT GGAGGTTTC AGCTTCAAC TTGATCTTC GACTTCAAA ACAAACAAAA AAAATTCTT 18540  
 AU AAGAACTAGT CTCTGCAAA TTTTAAAGT AACATGACTG TCACCTCTAC TTCAGGTTGAT GCTTCATTTT CATTCTTC TGAGCTTTTC 18612  
 Ref AAGAACTAGT CTCTGCAAA TTTTAAAGT AACATGACTG TCACCTCTAC TTCAGGTTGAT GCTTCATTTT CATTCTTC TGAGCTTTTC 18630  
 AU AGTGAGTCAT CAATGATAGT ACATTCTTC TTCACAATCG ACATGGACC TTCATGAACC TGCAAGAATAAGGTGTA CTTTATTAA 18702  
 Ref AGTGAGTCAT CAATGATAGT ACATTCTTC TTCACAATCG ACATGGACC TTCATGAACC TGCAAGAATAAGGTGTA CTTTATTAA 18720  
 AU TAATTACAAT GATAGTATCT GAAAGAGATT AAAATTGTT TAAAAAAAGT TTTTAAAC TGAGTTAGTA TTCAATTATG CCCTAATTAA 18792  
 Ref TAATTACAAT GATAGTATCT GAAAGAGATT AAAATTGTT TAAAAAAAGT TTTTAAAC TGAGTTAGTA TTCAATTATG CCCTAATTAA 18810  
 AU ACGCCTCTTT GGTTCTTC GGTTTTAAGT TCTCCATGGC ATCCTTGGT TCATGTTCT GTGCACTGCA GTTCTCCGAC TGCTGCAGTA 18882  
 Ref ACGCCTCTTT GGTTCTTC GGTTTTAAGT TCTCCATGGC ATCCTTGGT TCATGTTCT GTGCACTGCA GTTCTCCGAC TGCTGCAGTA 18900  
 AU TACTCTCTTC TTTGTTCT TAAAGCTTCAT TTTCCACTGG TCGAACACTG AGGAGCTCAT ACTGCTTTTC GTGCTCATTTG TCCACTAGTT 18972  
 Ref TACTCTCTTC TTTGTTCT TAAAGCTTCAT TTTCCACTGG TCGAACACTG AGGAGCTCAT ACTGCTTTTC GTGCTCATTTG TCCACTAGTT 18990  
 AU GTCCAACATA TGAATTCTATA ACATGTAAT CATCAGCCTT TCGTTTGAATCTCAACTT CCACGGTTT ATTAAATGTCA CTTGCTTTAG 19062  
 Ref GTCCAACATA TGAATTCTATA ACATGTAAT CATCAGCCTT TCGTTTGAATCTCAACTT CCACGGTTT ATTAAATGTCA CTTGCTTTAG 19080  
 AU TTTGTAGGTG GTCAGTTGGT GAGTCTTCG TTGTTCTCC CTCAGATCTC TCGCATGGT CTTTGACTGA TGTTGCTTCT TCGATGCTTG 19152  
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 AU CATTCTTTG CTTGAAGCTC TCCTCCTGGA TTTCGTCCCT GTGGGAATTTC TCCCTCTGAT TCTTCTGTAT GAAACTGATA TTTCTACTT 19332  
 Ref CATTCTTTG CTTGAAGCTC TCCTCCTGGA TTTCGTCCCT GTGGGAATTTC TCCCTCTGAT TCTTCTGTAT GAAACTGATA TTTCTACTT 19350  
 AU CATTAGTTT TACTTCAGGC ATTGGAACAA TGGATTCTGC GTCCCTTCTC ATTCCTGCTG ATTCTCTTGT TATAGCCAGC TCCTCATCTG 19422  
 Ref CATTAGTTT TACTTCAGGC ATTGGAACAA TGGATTCTGC GTCCCTTCTC ATTCCTGCTG ATTCTCTTGT TATAGCCAGC TCCTCATCTG 19440  
 AU TCTCCTGTAC AACTTTGCG TCCACCATAT TTTCAGGGTG TTTGTATAGA GCTTGATCTT CTTCTGAAGT TTCTCAACT ATTCTCTTCT 19512  
 Ref TCTCCTGTAC AACTTTGCG TCCACCATAT TTTCAGGGTG TTTGTATAGA GCTTGATCTT CTTCTGAAGT TTCTCAACT ATTCTCTTCT 19530  
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 Ref CTTCAATGAG AGCTGCATCA TTTCTTCTG AAGGCATAGT CGAACCTTC TTAACATCAT CTCCTGATAC CTCTTCAAT GGAATCTCCT 19620  
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 Ref CATTCTTTTC CTCTTCTGGT TCAGCTTTAG CTTCTTTAA GTTTGTTCT TCCGTTCTG CTGTTGTA TGTCAGTCC AACTCAGCAT 19890  
 AU TACTGATATA TTCTTTACT GTGGCGGCTT CTTCAGTATC CATCTTGGAC TGCTCGAGCT CAGATGAAGA GCTACAGACA AGTTCTTCTT 19962  
 Ref TACTGATATA TTCTTTACT GTGGCGGCTT CTTCAGTATC CATCTTGGAC TGCTCGAGCT CAGATGAAGA GCTACAGACA AGTTCTTCTT 19980  
 AU TGACTTCAAT TGGAGTTATC TCTTCTTTG CTGTACCTGA AGCTGCAGTT TCTGATCTT TACACATCTG TGTGGTAAAC AAAATTCAA 20052  
 Ref TGACTTCAAT TGGAGTTATC TCTTCTTTG CTGTACCTGA AGCTGCAGTT TCTGATCTT TACACATCTG TGTGGTAAAC AAAATTCAA 20070  
 AU CATAAACATG ATAAATTCT GGACGATCAC GGACTATATT ATAAAAACAA TTCCAACAAAG TCATGAGCCA TCTTTGTGCT AAGTTCATGA 20142  
 Ref CATAAACATG ATAAATTCT GGACGATCAC GGACTATATT ATAAAAACAA TTCCAACAAAG TCATGAGCCA TCTTTGTGCT AAGTTCATGA 20160  
 AU ACGBAATTTC TTATTCTTG AAATCCAAGT TATCTTGAGG TTTGCTAGCC GGCACCAAGTT GCCTCATGAC ACCTCCCTTA TTAATGAAC 20232  
 Ref ACGBAATTTC TTATTCTTG AAATCCAAGT TATCTTGAGG TTTGCTAGCC GGCACCAAGTT GCCTCATGAC ACCTCCCTTA TTAATGAAC 20250

20,260                    20,280                    20,300                    20,320                    20,340  
 AU CTGTTTTGG CCTTTCTGA TTTTGATCAG ATAACCTTT GATGTCGCAT AAGACTTCTG GTTGTGCTAT TTTGCATTGT ACGCATGTTG 20322  
 Ref CTGTTTTGG CCTTTCTGA TTTTGATCAG ATAACCTTT GATGTCGCAT AAGACTTCTG GTTGTGCTAT TTTGCATTGT ACGCATGTTG 20340  
 20,360                    20,380                    20,400                    20,420  
 AU TTGCCAATAA TGGTTATTTG TCCATGTC ATTAAATGAA TAGTTCTCGA ATTATATTAT AAGTTTGAAG TTGAAAGATT AGAGAGTTGT 20412  
 Ref TTGCCAATAA TGGTTATTTG TCCATGTC ATTAAATGAA TAGTTCTCGA ATTATATTAT AAGTTTGAAG TTGAAAGATT AGAGAGTTGT 20430  
 20,440                    20,460                    20,480                    20,500                    20,520  
 AU ATAATTTCT TACCTCTTC ATTGGAAATCT GATTATTGTA GGTTGCTTCC TTCTCTAGAA TTTGTTTCC ATCATTCTCA GCTATGTTG 20502  
 Ref ATAATTTCT TACCTCTTC ATTGGAAATCT GATTATTGTA GGTTGCTTCC TTCTCTAGAA TTTGTTTCC ATCATTCTCA GCTATGTTG 20520  
 20,540                    20,560                    20,580                    20,600  
 AU CCGGTTCTAG CTTTCAGTG ACCTGAATGG AAATATATTG GAAGATTGGT TAAGCATGGA GGTTCTGCT TTTAGCAAAT AGATTGAAAT 20592  
 Ref CCGGTTCTAG CTTTCAGTG ACCTGAATGG AAATATATTG GAAGATTGGT TAAGCATGGA GGTTCTGCT TTTAGCAAAT AGATTGAAAT 20610  
 20,620                    20,640                    20,660                    20,680                    20,700  
 AU TGGGTTAGTA TTCATGATAG GAACTGTTT GATTACCTCA GCTTCATCCT TGTCACAGT TTCCATATT AGGTTCTGCT TTTGCACTTC 20682  
 Ref TGGGTTAGTA TTCATGATAG GAACTGTTT GATTACCTCA GCTTCATCCT TGTCACAGT TTCCATATT AGGTTCTGCT TTTGCACTTC 20700  
 20,720                    20,740                    20,760                    20,780  
 AU ATCATTGTC TTAGTCGGA CATCAGGATC CAATTCCGAA GATATCTT CAAGCTGCAC ATTCTCTGCT TCTGTCCTT GAGCATCACT 20772  
 Ref ATCATTGTC TTAGTCGGA CATCAGGATC CAATTCCGAA GATATCTT CAAGCTGCAC ATTCTCTGCT TCTGTCCTT GAGCATCACT 20790  
 20,800                    20,820                    20,840                    20,860                    20,880  
 AU CTTAATATTT CCTACAGTCG TCTCGGCTGA GCTGATATT GATGCATTTA CAATCTTGT GGTGTCGTT AAGCTCGTCT CTGCTGTTAC 20862  
 Ref CTTAATATTT CCTACAGTCG TCTCGGCTGA GCTGATATT GATGCATTTA CAATCTTGT GGTGTCGTT AAGCTCGTCT CTGCTGTTAC 20880  
 20,900                    20,920                    20,940                    20,960  
 AU TTCTGTTACA GCTTCAATAC AACTCTCTC ATGACACGGT TTCGAGTCAT CAATAATCTC CTCGGTTTCT TCTTCTGCTT CTCGATCTTT 20952  
 Ref TTCTGTTACA GCTTCAATAC AACTCTCTC ATGACACGGT TTCGAGTCAT CAATAATCTC CTCGGTTTCT TCTTCTGCTT CTCGATCTTT 20970  
 20,980                    21,000                    21,020                    21,040                    21,060  
 AU GCTACACAGA CTTACAGGAT CTCGGACTC TTCAACCTGT CCCATTCTT CTAAATTCTT GGTGATTTGG GATGCTTCTC CATCATTGGT 21042  
 Ref GCTACACAGA CTTACAGGAT CTCGGACTC TTCAACCTGT CCCATTCTT CTAAATTCTT GGTGATTTGG GATGCTTCTC CATCATTGGT 21060  
 21,080                    21,100                    21,120                    21,140  
 AU TGAATGTTCT TTGACCTCCA AGTTTCTGC AACATCTGC TCTTGGAAAT CGGCTCCAT GGAAACATTCA GTTGAGACTG AACTATAAGTC 21132  
 Ref TGAATGTTCT TTGACCTCCA AGTTTCTGC AACATCTGC TCTTGGAAAT CGGCTCCAT GGAAACATTCA GTTGAGACTG AACTATAAGTC 21150  
 21,160                    21,180                    21,200                    21,220                    21,240  
 AU TGCCACCTGC GCATAAACATA GTTACAATGT TAAACAATAT AACATCCAAA ATCTGTTTG AGTATCTATT CATTGAAAAA ATGAAAAGTG 21222  
 Ref TGCCACCTGC GCATAAACATA GTTACAATGT TAAACAATAT AACATCCAAA ATCTGTTTG AGTATCTATT CATTGAAAAA ATGAAAAGTG 21240  
 21,260                    21,280                    21,300                    21,320  
 AU TAATTATTCA GTAACGCTTG ATACCCCTTT TTATATCTT AATTGATT TGTTATTTA GAAGAAAAAG AGTGAAAAAGA GTAGAATATC 21312  
 Ref TAATTATTCA GTAACGCTTG ATACCCCTTT TTATATCTT AATTGATT TGTTATTTA GAAGAAAAAG AGTGAAAAAGA GTAGAATATC 21330  
 21,340                    21,360                    21,380                    21,400                    21,420  
 AU TATTCACTAA CCTTGCATT ATAAGTCATG GTAAACCAC TCAAGAACATG GTTTCTTT TCTTTCTTCA CATCTTCTTT CAAGCTTTCA 21402  
 Ref TATTCACTAA CCTTGCATT ATAAGTCATG GTAAACCAC TCAAGAACATG GTTTCTTT TCTTTCTTCA CATCTTCTTT CAAGCTTTCA 21420  
 21,440                    21,460                    21,480                    21,500  
 AU GTTTTCTA TGTTCTTAGC TGAATTCTCT TCTTCAGTAG CCTTTGCC ATCCTTGCAA GTACCAAGG TTGCTTCCAA ATCCATGGTT 21492  
 Ref GTTTTCTA TGTTCTTAGC TGAATTCTCT TCTTCAGTAG CCTTTGCC ATCCTTGCAA GTACCAAGG TTGCTTCCAA ATCCATGGTT 21510  
 21,520                    21,540                    21,560                    21,580                    21,600  
 AU TCATCCAGCG GCTTCCATC TCAAATTTC GACCTTCTA GTCCTACATC TGATGAATT TCTGCTGGCA GTGTTTTTC TTGCAAGCA 21582  
 Ref TCATCCAGCG GCTTCCATC TCAAATTTC GACCTTCTA GTCCTACATC TGATGAATT TCTGCTGGCA GTGTTTTTC TTGCAAGCA 21600  
 21,620                    21,640                    21,660                    21,680  
 AU TCGGTTTGAT TCTCTAGTC TCTATGTTCT TCACCACTGAG AGACATTGGT GACATCTAGA TTAGATTCT CAGCCAACGT TTCTTGTACT 21672  
 Ref TCGGTTTGAT TCTCTAGTC TCTATGTTCT TCACCACTGAG AGACATTGGT GACATCTAGA TTAGATTCT CAGCCAACGT TTCTTGTACT 21690  
 21,700                    21,720                    21,740                    21,760                    21,780  
 AU TCATGTCTAC TCTCGTTCC TGTATCTCA GGCTCAGCTT TAGGGACATC TGCTAGCGGA TTATCTTCAT TTAGAATTAA TTCTCATCC 21762  
 Ref TCATGTCTAC TCTCGTTCC TGTATCTCA GGCTCAGCTT TAGGGACATC TGCTAGCGGA TTATCTTCAT TTAGAATTAA TTCTCATCC 21780  
 21,800                    21,820                    21,840                    21,860  
 AU GTTACTTG TGCACTGGAGT TTCTGAGTGC TCACACAAAT CATGATCTT TTCTGAAGTT TCATCAGTTA TTTTCTTTT TTCAATGAGT 21852  
 Ref GTTACTTG TGCACTGGAGT TTCTGAGTGC TCACACAAAT CATGATCTT TTCTGAAGTT TCATCAGTTA TTTTCTTTT TTCAATGAGT 21870  
 21,880                    21,900                    21,920                    21,940                    21,960  
 AU GTGGCCTCAT ATTCTTCAC TTCACTCCT AGTATTTCTT TCAAGGAAT CTCCCTGGTG GTCATCTCCA ATTGATCTTC ATTCTTCATCA 21942  
 Ref GTGGCCTCAT ATTCTTCAC TTCACTCCT AGTATTTCTT TCAAGGAAT CTCCCTGGTG GTCATCTCCA ATTGATCTTC ATTCTTCATCA 21960  
 21,980                    22,000                    22,020                    22,040  
 AU TTGCGAACCT TATCTGCAT TGCACTCTCT TCTTGGGAA TCTCCATGTC CTTCTCATGT GTTACGATG AGTCCCCCTGA TTTCACATCA 22032  
 Ref TTGCGAACCT TATCTGCAT TGCACTCTCT TCTTGGGAA TCTCCATGTC CTTCTCATGT GTTACGATG AGTCCCCCTGA TTTCACATCA 22050  
 22,060                    22,080                    22,100                    22,120                    22,140  
 AU ATTTCACTAC TGCTTACACC AATCTGTTCA TGTTCTGTC TTGCTGAGCT GTTCTCTTCG ATGACAACAA CATTACATT TTGGTTCTTT 22122  
 Ref ATTTCACTAC TGCTTACACC AATCTGTTCA TGTTCTGTC TTGCTGAGCT GTTCTCTTCG ATGACAACAA CATTACATT TTGGTTCTTT 22140  
 22,160                    22,180                    22,200                    22,220  
 AU TCCTTTCTT GTTCAGCGTC AGCTTCTTT ATATTGTTT CTTCGGTTTC TGCTGTATAT GATGCTGAGT TTCTTCTTT GGTATCGTTT 22212  
 Ref TCCTTTCTT GTTCAGCGTC AGCTTCTTT ATATTGTTT CTTCGGTTTC TGCTGTATAT GATGCTGAGT TTCTTCTTT GGTATCGTTT 22230  
 22,240                    22,260                    22,280                    22,300                    22,320  
 AU TTGGACTCCC CAAGCTCAAC TGAGGAACTA GGGACATGAT CTTCTTCAC TTCTATTGGT GCTATTTCTT CTTTGTACT CTGTATAATA 22302  
 Ref TTGGACTCCC CAAGCTCAAC TGAGGAACTA GGGACATGAT CTTCTTCAC TTCTATTGGT GCTATTTCTT CTTTGTACT CTGTATAATA 22320  
 22,340                    22,360                    22,380                    22,400                    22,500  
 AU TCAAAACAGAC TAGAATTATT ATTGGTTCT CTTGTTCTCT AGCTATTAGT GAGTCATACA ATCTACTTAC CTCTGTTATC GAAACCACTG 22392  
 Ref TCAAAACAGAC TAGAATTATT ATTGGTTCT CTTGTTCTCT AGCTATTAGT GAGTCATACA ATCTACTTAC CTCTGTTATC GAAACCACTG 22410  
 22,420                    22,440                    22,460                    22,480                    22,500  
 AU GAATATATAT GTTGATGGAA GATTGAGTC AGAACAGTT AGCTATTAGT GAGTCATACA ATCTACTTAC CTCTGTTATC GAAACCACTG 22482  
 Ref GAATATATAT GTTGATGGAA GATTGAGTC AGAACAGTT AGCTATTAGT GAGTCATACA ATCTACTTAC CTCTGTTATC GAAACCACTG 22500

AU CTGCACTATT CTCTTCTTTA GCTTTTCAT CCATCGTC GGCCATTAAT GTGGTCTGAT CCTTCTTGT ATCTTCTCC TCTAGAAATT 22572  
 Ref CTGCACTATT CTCTTCTTTA GCTTTTCAT CCATCGTC GGCCATTAAT GTGGTCTGAT CCTTCTTGT ATCTTCTCC TCTAGAAATT 22590  
 AU GTTTTTCGGG ATTCTCAACT GTGTTGTCG GTTCCAGCTT TTACACAGACC TGAAACAAAA ATAATATCAA AAGTTCGGTG AAACATGGAG 22662  
 Ref GTTTTCGGG ATTCTCAACT GTGTTGTCG GTTCCAGCTT TTACACAGACC TGAAACAAAA ATAATATCAA AAGTTCGGTG AAACATGGAG 22680  
 AU GTTTTCAGCT TTCAACTTGA TTGTTGAAAAAA AAATAAAAAC TTACATCTTAA GAACTAGGCT CTTCAAAATT TTAGAAGTAA CATGAATTTC 22752  
 Ref GTTTTCAGCT TTCAACTTGA TTGTTGAAAAAA AAATAAAAAC TTACATCTTAA GAACTAGGCT CTTCAAAATT TTAGAAGTAA CATGAATTTC 22770  
 AU ACCTCTACTT CAGGGTGGC TTCACTTCA TTTCTTCTG TAGCTTCAG TGAGTCATTG ATGATAGCAC ATTCTTCTT CACAATGAAT 22842  
 Ref ACCTCTACTT CAGGGTGGC TTCACTTCA TTTCTTCTG TAGCTTCAG TGAGTCATTG ATGATAGCAC ATTCTTCTT CACAATGAAT 22860  
 AU GTTGAGCCTT CATGAACCTG CATGAAGAAC ATGGTGTACT TTTTTAAACA ATTACAGTGC TAATCTTGGA GAAGTGTGTTA AAATTGTTT 22932  
 Ref GTTGAGCCTT CATGAACCTG CATGAAGAAC ATGGTGTACT TTTTTAAACA ATTACAGTGC TAATCTTGGA GAAGTGTGTTA AAATTGTTT 22950  
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 Ref AAAAGAACGT TGAAACTGAG TTAGTATTCA TTAGTGCTT AAATCCAGAC CTCTTGGTT TCTTCCTGAT TTAAGTTATC TTCTACGTT 23040  
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 Ref TCCTTGGCAT CCTTTGGTAG TGCACTTCTG TTCTCTGACT CCTGCTGTAT ACTCTCTTCT TTGTTCTA AAGCTTCATT TTCCACTGGT 23130  
 AU TCAACATCGG AGAACCTGTA CTGCTTTTCG TGCTCATTGT CCATTAGTTG TCCAACGAAT GAGTCAGTA CATGTAATC ATCAGCATTC 23202  
 Ref TCAACATCGG AGAACCTGTA CTGCTTTTCG TGCTCATTGT CCATTAGTTG TCCAACGAAT GAGTCAGTA CATGTAATC ATCAGCATTC 23220  
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 Ref TCGTTGAAA TCTCACTTC CACGGTTCA TTACTCTGAC TTGCTTTAGT TTGTTAGGTGG TCAGTTGGTG AGTCTTCCCT CGTTTGTCCC 23310  
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 Ref TCGGATCTCT TGTCACTGGTC TTGACTGGT GTGCTTCTT CAATGCCGT GCTGTTCT GCAGTTTCGG CACAGATGGC AGTATTGACA 23400  
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 Ref TCCTCAGAAA CTGCATTCCA AGGACATGCA CGGTTCTCT TTGCTGGTT ATTCTTTGC TTGAAGCTCT CCTCTAGGAT TTGTCCTTC 23490  
 AU TTGGAATTTC CCTCCTGATT CTTCTGTGCG AGACCAATAT TTTCTACTTC ACTAATTTC CTTTCAGGCA TTGAAACAAT GGATTCTTTG 23562  
 Ref TTGGAATTTC CCTCCTGATT CTTCTGTGCG AGACCAATAT TTTCTACTTC ACTAATTTC CTTTCAGGCA TTGAAACAAT GGATTCTTTG 23580  
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 Ref TGCTTTGTT GAGCTTGATC TTCTTCAGAA GTTTCATCAA CTAATTCCCT CACTTCTAAG AGAGTTGCA CGTTTCTTC TGAATGCATA 23760  
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 Ref GCAGAATGTT CCTAACATC ATCTCCTAAT ATATCTTCA ACAGAATCTT GTGGTTGTC ATTTCCTATT GATTTCACT GCCTGCATTC 23850  
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 Ref TCAGCACTTC CTAACCAATT CCCATGATCT TCACCATCAA CGACATTATT TATTCATTC TTTTCTCGT CCTCTTCTGC TTGGCTTTA 24030  
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 AU GCTGTACCA AAGCTCCGT TTCTGATCCT TCACATGCCT TCACGATAAA CAAATTTCA ATATAAACAT GATAAGTTTC CAGATGGTTA 24282  
 Ref GCTGTACCA AAGCTCCGT TTCTGATCCT TCACATGCCT TCACGATAAA CAAATTTCA ATATAAACAT GATAAGTTTC CAGATGGTTA 24300  
 AU CAGACTATTA TAAGACAAAT TCCAACAGGT CACGGGTCA CTTAGTGCTA AGTTAATGAA CTGAATTTC TATCCTTGA AATCCCAGTT 24372  
 Ref CAGACTATTA TAAGACAAAT TCCAACAGGT CACGGGTCA CTTAGTGCTA AGTTAATGAA CTGAATTTC TATCCTTGA AATCCCAGTT 24390  
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 Ref ATCTTGTGGT TTGCTTGCTG GCATCAGTT CTTCATGACA TCTCCTATTG TGGAACCTCT TTTTGGCCT TCTCTTTCT GAATTTGATC 24480  
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 Ref TGAATTACT CGGCTCTATT ACTTTACAC TGTAAATGG ATAGCTTTTC GATGGGGCAT AAGACTTCTG GTTGTGCTGT TTTGCAATTGT 24570  
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 Ref ATGCATGTT ATGCCAAAGA TTTTATTTG TTCCATGTC ATTGATGGAA GTTCGAGTTC AGAAATAT TATAAGTTTG AAGTTTGATC 24660  
 AU ATTAGAGAGT CATATAATT TCTTACCTCT TTAAGTGTAA TCTGATCATT GTGGTGTCT TCCTCTAGAA TTGTTTTCC ATCATTCTCA 24732  
 Ref ATTAGAGAGT CATATAATT TCTTACCTCT TTAAGTGTAA TCTGATCATT GTGGTGTCT TCCTCTAGAA TTGTTTTCC ATCATTCTCA 24750

AU ACTGTGTTGT CCAGTCCAG CTTTCAGTG ACCTGATTG AAAATATTG AGAGATTGGT TAAAAATGGA GCTTTCAGAT TTTAACAAAA 24822  
 Ref ACTGTGTTGT CCAGTCCAG CTTTCAGTG ACCTGATTG AAAATATTG AGAGATTGGT TAAAAATGGA GCTTTCAGAT TTTAACAAAA 24840  
 AU AAATTGAAAT TGAGTTAGTA TTATGATCTA GAATCAGACC TCTT TAGTT CTTCGTGTT CAAGTTATCT TCTATGTGAT CATTTAGCTC 24912  
 Ref AAATTGAAAT TGAGTTAGTA TTATGATCTA GAATCAGACC TCTT TAGTT CTTCGTGTT CAAGTTATCT TCTATGTGAT CATTTAGCTC 24930  
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 Ref TTGCTGTATC CCCTTGTTAA AGCTTCACTG CACTCTGAGC CATTCCGGA GTTCAAGTGA TATGCCCTCTT TTATAAACAT 25020  
 AU TGTGTTTCA TTGCCCATTT CACTTGTTTCA AGATTTAAA TCTGTTGCTG CTGTTCTGT GAATATGTC TCTGGAATTG TGTAACTACT 25092  
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 AU ATCGGAGAAC TCAAACGTCT TTTCATACCT GTTTCACTG GATTGTTGAA CTGATGAATC AGTAACATGT AAATCATCAG CATTGTCATT 25272  
 Ref ATCGGAGAAC TCAAACGTCT TTTCATACCT GTTTCACTG GATTGTTGAA CTGATGAATC AGTAACATGT AAATCATCAG CATTGTCATT 25290  
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 Ref AAGCTTTAT TTCATTTAC ATATGTGTTA TTTATTCTCT GGAAGAAATAA GCTCTTTGG TAACAAAAAA ATGGCATAC TAAATCTAT 27810  
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 Ref AGGTGGTTGT GCAAGCATCG TGAACACAAT ACAAACATG TCCTCTCGG GGGTCGACTA ACCCTCGAG GCTTTCATTC AATGGCGTGT 28170  
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 Ref CTCCCTGGAT TTGAAACCAA CACTTCTTAC AACACATATA GTAGTCTATC TTGAAAACAT AGTACCGTTT GATCTTGCA AAAAGACCG 28260  
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 Ref CTCAAAAGCA GAAGTGCTAT TAAAGATGA AACCTGACGC ATGATCTTTC TGGTTAATC TGTCATCAA GACATCTGAT CTTGATGAC 28800  
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 Ref GTAGAGTGTGTT CTAAGGTTTG CTTAAAGCAC AATACGTGTA AGATGAAAGA ACATTACAC TATTGAAATG GGCTCTTGAA TGATCACTGC 28890  
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 Ref TGAAGAAACT TTACCACTAT GAAAATGGGT TCTTGAATAA TCACTGATTC AGTTGCAATT TTTTCTGCA CAAGTGATCG AATTGAGTT 29070  
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 Ref GGTTCTGAA TGATCGCTGC TTCAAGTTGCC ATCTACTTT CCTGCAGATA TAATAGAATT CG 29222