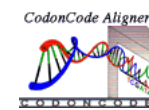


CodonCode Aligner: AE1176, AE1176_HVR

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<< AE1176 F1-2r	GAAGCAGATT	TGGGTACCAG	CCAAGTATTG	ACTCAGCCAT	CAACAACCGG	TATGTATTTC	GTACATTACT	GCCAGCCACC	ATGAATATTG	TACGGTACCA	
<< AE1176 B2r		T	TGGGTACCAG	CCAAGTATTG	ACTCAGCCAT	CAACAACCGG	TATGTATTTC	GTACATTACT	GCCAGCCACC	ATGAATATTG	TACGGTACCA
AE1176 B2f						GG	TATGTATTTC	GTACATTACT	GCCAGCCACC	ATGAATATTG	TACGGTACCA
AE1176 F1f						GG	TATGTATTTC	GTACATTACT	GCCAGCCACC	ATGAATATTG	TACGGTACCA
<< AE1176 C1-2r									CACC	ATGAATATTG	TACGGTACCA
<< AE1176 C1r									CACC	ATGAATATTG	TACGGTACCA
<< AE1176 C2r									CACC	ATGAATATTG	TACGGTACCA
NC_012920_2 (r...	gaagcagatt	tgggtaccag	ccaagtattg	actcagccat	caacaacccg	tatgtatttc	gtacattact	gccagccacc	atgaatattg	tacgggtacca	
	16036	16046	16056	16066	16076	16086	16096	16106	16116	16126	
AE1176_HVR	GAAGCAGATT	TGGGTACCAG	CCAAGTATTG	ACTCAGCCAT	CAACAACCGG	TATGTATTTC	GTACATTACT	GCCAGCCACC	ATGAATATTG	TACGGTACCA	
<< AE1176 F1-2r	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAGTACAGTA	ATCAACCCCG	AACATATACA	
<< AE1176 B2r	TAAATAC										
AE1176 B2f	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATC						
AE1176 F1f	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAGTACAGTA	ATCAACCCCG	AACATATACA	
<< AE1176 C1-2r	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AA			
<< AE1176 C1r	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GG				
<< AE1176 C2r	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAA				
AE1176 C1f			GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAGTACAGTA	ATCAACCCCG	AACATATACA	
AE1176 C2f			ACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAGTACAGTA	ATCAACCCCG	AACATATACA	
<< AE1176 D1-2r								AAGTACAGTA	ATCAACCCCG	AACATATACA	
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AE1176 G1-2f											A
NC_012920_2 (r...	taaatacttg	accacctgta	gtacataaaa	acccaatcca	catcaaaacc	ccctccccat	gttacaaagg	aagtacagta	atcaacccc	aactatcaca	
	16136	16146	16156	16166	16176	16186	16196	16206	16216	16226	
AE1176_HVR	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAGTACAGTA	ATCAACCCCG	AACATATACA	

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