

# CodonCode Aligner: A78, A78\_HVR

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Page 1 of 2



<< A78 F1r	GAAGCAGATT	TGGGTACCAG	CCAAGTATTG	ACTCAGCCAT	CAACAACCGG	TATGTATTTC	GTACATTACT	GCCAGCCACC	ATGAATATTG	TACGGTACCA		
<< A78 B1r		TGGGTACCAG	CCAAGTATTG	ACTCAGCCAT	CAACAACCGG	TATGTATTTC	GTACATTACT	GCCAGCCACC	ATGAATATTG	TACGGTACCA		
<< A78 B2r		TGGGTACCAG	CCAAGTATTG	ACTCAGCCAT	CAACAACCGG	TATGTATTTC	GTACATTACT	GCCAGCCACC	ATGAATATTG	TACGGTACCA		
A78 B1f						CG	TATGTATTTC	GTACATTACT	GCCAGCCACC	ATGAATATTG	TACGGTACCA	
A78 B2f						CG	TATGTATTTC	GTACATTACT	GCCAGCCACC	ATGAATATTG	TACGGTACCA	
<< A78 C1r									CA	CG	ATGAATATTG	TACGGTACCA
<< A78 C2r									CA	CG	ATGAATATTG	TACGGTACCA
NC_012920_2 (r...	gaagcagatt	tggtaccag	ccaagtattg	actcagccat	caacaacggg	tatgtatttc	gtacattact	gccagccacc	atgaatattg	tacgggtacca		
	16036	16046	16056	16066	16076	16086	16096	16106	16116	16126		
A78_HVR	GAAGCAGATT	TGGGTACCAG	CCAAGTATTG	ACTCAGCCAT	CAACAACCGG	TATGTATTTC	GTACATTACT	GCCAGCCACC	ATGAATATTG	TACGGTACCA		

<< A78 F1r	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAGTACAGTA	ATCAACCCTC	AACTATCACA
<< A78 B1r	TAAATAC									
<< A78 B2r	TAAATAC									
A78 B1f	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATC					
A78 B2f	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATC					
<< A78 C1r	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAG		
<< A78 C2r	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAG		
A78 F1f	TAACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAGTACAGTA	ATCAACCCTC	AACTATCACA
A78 C1f			ATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAGTACAGTA	ATCAACCCTC	AACTATCACA
A78 C2f			ATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAGTACAGTA	ATCAACCCTC	AACTATCACA
<< A78 G1-2r						CCAT	GTTACAAGC	AAGTACAGTA	ATCAACCCTC	AACTATCACA
<< A78 D1r								AAGTACAGTA	ATCAACCCTC	AACTATCACA
<< A78 D2r								AAGTACAGTA	ATCAACCCTC	AACTATCACA
NC_012920_2 (r...	taaatacttg	accacctgta	gtacataaaa	acccaatcca	catcaaaacc	ccctccccat	gttacaagc	aagtacagta	atcaaccctc	aaactatcaca
	16136	16146	16156	16166	16176	16186	16196	16206	16216	16226
A78_HVR	TAAATACTTG	ACCACCTGTA	GTACATAAAA	ACCCAATCCA	CATCAAAACC	CCCTCCCCAT	GTTACAAGC	AAGTACAGTA	ATCAACCCTC	AACTATCACA

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Page 2 of 2

