

Project: Untitled Contig 1

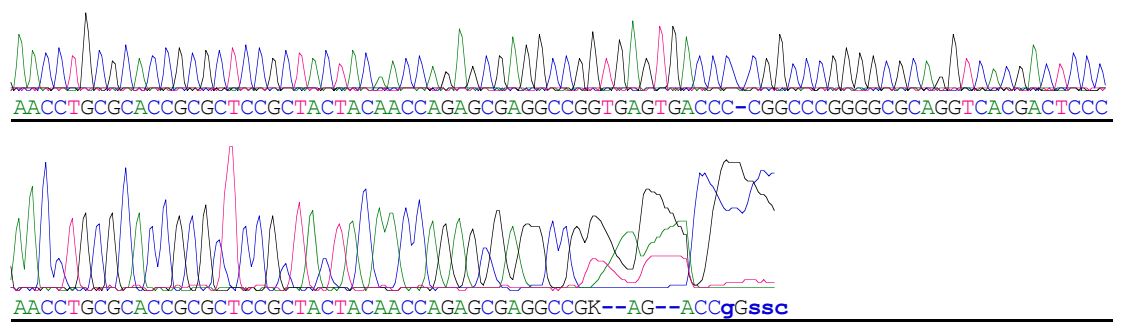
330 340 350 360 370 380 390 400 410

AACCTGCGCACCGCGCTCCGCTACTACAACCAGAGCGAGGCCGGT**tgAGTg**ACCC-**CGG**CCCCGGGGCGCAGGTCACGACTCCC

HLA-B 130101.SEQ (1>970) →
Sample-F1.ab1 (10>350) →

AACCTGCGCACCGCGCTCCGCTACTACAACCAGAGCGAGGCCGGT**tgAGTg**ACCC-**CGG**CCCCGGGGCGCAGGTCACGACTCCC

Sample-R1.ab1 (1>311) ←

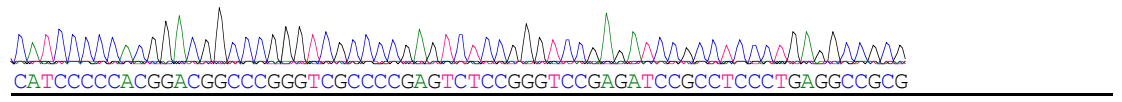


420 430 440 450 460 470 480 490

CATCCCCACGGACGGCCCGGGTTCGCCCGAGTCTCCGGTCCGAGATCCGCCTCCCTGAGGCCGCGGGACCCGCCAGACC

HLA-B 130101.SEQ (1>970) →
Sample-F1.ab1 (10>350) →

CATCCCCACGGACGGCCCGGGTTCGCCCGAGTCTCCGGTCCGAGATCCGCCTCCCTGAGGCCGCGGGACCCGCCAGACC



500 510 520 530 540 550 560 570

CTCGACCGGCGAGAGCCCCAGGCGGTTTTACCCGGTTTCATTTTCAGTTGAGGCCAAAATCCCCGCGGGTTGGTCGGGGCGG

HLA-B 130101.SEQ (1>970) →
Sample-R2.ab1 (1>370) ←

CTCGACCGGCGAGAGCCCCAGGCGGTTTTACCCGGTTTCATTTTCAGTTGAGGCCAAAATCCCCGCGGGTTGGTCGGGGCGG

△
G

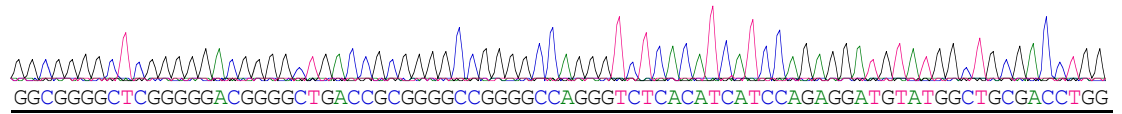
580 590 600 610 620 630 640 650

GGCGGGGCTCGGGGACGGGGCTGACCGCGGGCCGGGGCCAGGGTCTCACATCATCCAGAG**GgATgTAT**GGCTGCGACCTGG

HLA-B 130101.SEQ (1>970) →
Sample-F2.ab1 (2>397) →

GGCGGGGCTCGGGGACGGGGCTGACCGCGGGCCGGGGCCAGGGTCTCACATCATCCAGAGGATGTATGGCTGCGACCTGG

Sample-R2.ab1 (1>370) ←

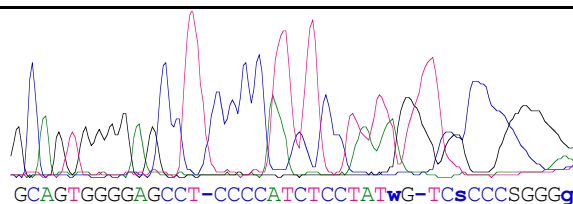




910 920 930 940 950 960 970 980

GCAGTGGGGAGCCTtCCCCATCTCCTATAGgTCGCC-GGGGATGGCCTCCCACGAGAAGAGGAGGAAAATGGGATCAGCGCT

Sample-R2.ab1 (1>370)

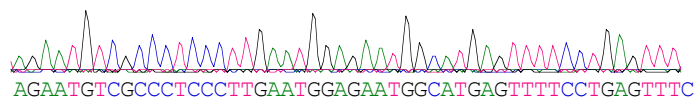


GCAGTGGGGAGCCT-CCCCATCTCCTATwG-TCsCCCSGGGgy

990 1000 1010 1020 1030

AGAATGTCGCCCTCCCTTGAATGGAGAATGGCATGAGTTTTCTGAGTTTC

Sample-F2.ab1 (2>397)



AGAATGTCGCCCTCCCTTGAATGGAGAATGGCATGAGTTTTCTGAGTTTC