## Sequencing results of 11. Mai 2017 (GroupB)

## GroupB-Sample 1

Your sample has been re-sequenced. We have been able to improve your initial result.

## GroupB-Sample 2

GCCTCCTTACGGTTAGCTCAACGCCTTCGAGTGAATCCAACTCCCATGGTGTGACGGGCGGTGTGTACAAGG CCTGGGAACGTATTCACCGCGGCATGCTGATCCGCGATTACTAGCGATTCCGCCTTCACGCTCTCGAGTTGC AGAGAACGATCCGAACTGAGACGACTTTTGGAGATTAGCTCCCTCTCGCGAGGTGGCTGCCCACTGTAGTCG CCATTGTAGCACGTGTGTAGCCCAACGCGTAAGGGCCATGAGGACTTGACGTCATCCCCACCTTCC

Your sample has been re-sequenced. We have been able to improve your initial result.

#### GroupB-Sample 3

GCTACCTACTTNTGGTAAAACCCGCTCCCRTGGTGTGACGGGCGGTGTGTACAAGACCCGGGAACGTATTCACCGCGACATGCTGATCCGCGATTACTAGCGATTCCAACTTCACGCAGTCGAGTTGNAGACTGNGATCCGGACTACGA

The signal strength of your sequencing reaction is very low. Therefore, your sequencing result needs careful evaluation.

## GroupB-Sample 4

TNRTTAGCTGCCTCCTTGCGGTTAGCGCACTACCTTCGGGTAGAACCAACTCCCATGGTGTGACGGGCGGTGTGTACAAGGCCCGGGAACGTTAGCGCACTTCATCGCACTTCATGCACTGCACTGCAGATTGCAAGGCCCGGGAACCGAACTGAGATGGCTTTTGGAGATTAGCTCGACCTCGCGGTCTCGCCCACTGTCACCACCATTGTAGCACGTGTGTAGCCCAGCCGTAAGGGCCATGAGGACTTGACGTCATCCCACCTTCCTCCGGCTTATCACCGGCAGTCCCCTTAGAGTGCCCAACTAAATGCTGGCAACTAAGGGCGAGGGTTGCGCTCGTTGCGGGGACTTAACCCAACATCTCACGACACGAGCTGACGACAGCCATGCAGCACCTGTCTCCGATCCAGCCGAACTGAAGGAATACATCTCTGTAATCCGCGATCGGGATGTCAAGGGCTGGTAAGGTTCTGCGCGTTGCTTCGAATTAAACCACATGCTCCACCGCTTGTGCGGGCCCCCGTCAATTCCTTTGAGTTTTAATCTTGCGACCGTACTCCCCAGGCGGAACTGCCCGACGGCTAACACTCATCGTTTACGGCGTGGACTACCACGGTTAGCTGCGCCCCCCCGCCACCGCTTTCGCACCTCAGCGTCACCACGTTAACCCCCACGCTTTCGCACCTCCACCGCTTAACCACGTTAATCCTGTTTTGCTCCCCCACGCTTTCGCACCTCAGCGTCACCACGGTAACACACATGAGCCACTCCCCACGCTTCGCCACCTGG

Quality control passed.

Moderate or poor sequencing result.

## GroupB-Sample 9

Your sample has been re-sequenced. We have been able to improve your initial result.

## GroupB-Sample 12

NNGGTTACCTCACCGACTTCGGGTGTTACAAACTCTCGTGGTGTGACGGGCGGTGTGTACAAGGCCCGGGA ACGTATTCACCGCGCGTGCTGATCCGCGATTACTAGCGATTCCGGCTTCATGCAGGCGAGTTGCAGCCTGC AATCCGAACTGAGAGAGCTTTAAGAGATTTGCATGACCTCGCGGTCTAGCGACTCGTTGTACTTCCCATTGT AGCACGTGTGTAGCCCAGGTCATAAGGGGCATGATGATTTGACGTCATCCCCACCTTCCTCCGGTTTGTCAC CGGCAGTCTCGCTAGAGTGCCCAACTAAATGATGGCAACTAACAATAAGGGTTGCGCTCGTTGCGGGACTTA ACCCAACATCTCACGACACGAGCTGACGACAACCATGCACCACCTGTCACTTTGTCCCCGAAGGGAAAGCTC TATCTCTAGAGTGGTCAAAGGATGTCAAGACCTGGTAAGGTTCTTCGCGTTGCTTCGAATTAAACCACATGCT CCACCGCTTGTGCGGGCCCCCGTCAATTCCTTTGAGTTTCAACCTTGCGGTCGTACTCCCCAGGCGGAGTGC TTAATGCGTTTGCTGCAGCACTGAAGGGCGGAAACCCTCCAACACTTAGCACTCATCGTTTACGGCGTGGAC TACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGAGCCTCAGCGTCAGTTACAGACCAGAGAGCCGC CTCAAGTCTCCCAGTTTCCAATGACCCTCCCGGTTGAGCCGGGGGCTTTCACATCAGACTTAAGAAACCGC CTGCGCTCGCTTTACGCCCAATAAATCCGGACAACGCTTGCCACCTACGTATTACCGCGGCTGCTGGCACGT ANTTAGCCGTGGCTTTCTGGTNAGATACCGTCAGGGGACGTTCAGTTACTAACGTCCTTGTTCTTCTAACA ACAGAGTTTTACGATCCGAAAACCTTCTTCACTCACGCGGCGTTGCTCGGTCAGACTTTCGTCCATTGCCAAA AATTCCCTACTGCTGCCTCCCGTAAGAAT

Your sample has been re-sequenced. We have been able to improve your initial result.

# GroupB-Sample 13

GCTACTTCTGGTGCAACCCACTCCCATGGTGTGACGGGCGGTGTGTACAAGGCCCGGGAACGTATTCACCGCGCACATTCTGATTCGCGATTACTAGCGATTCCGACTTCACGCAGTTGCAGACTGCGACTGCGACTACCGACCTTGTTTTGTGAGATTAGCTCCACCTCGCGGCTTTGGCAACCCTCTGTACCGACCATTGTAGCACGTGTGTAGCCCAGGCCGTAAGGGCCATGATGACTTGACGTCATCCCCACCTTCCTCCGGTTTGTCACCGGCAGTCTCCTTAGAGTGCCCACCATNACGTGCTGGTAACTAAGGACAAGGGTTGCGCTCGTTACGGGACTTAACCCAACATCTCACGACACGAGCACGAGCTGACNACAGCCATGCAGCACCTGTGTCANAGTTCCCGAAGGCACCAATCCATCTCTGGAAGGTTCTCTGCATGT

Moderate or poor sequencing result.

GCAACCCACTCCCATGGTGTGACGGGCGGTGTGTACAAGGCCCGGGAACGTATTCACCGTAGCATTCTGAT CTACGATTACTAGCGATTCCGACTTCATGGAGTCGAGTTGCAGACTCCAATCCGGACTACGACATACTTTATG AGGTCCGCTTGCTCTCGCGAGGTCGCTTCTCTTTGTATATGCCATTGTAGCACGTGTGTAGCCCTGGTCGTAA GGGCCATGATGACTTGACGTCATCCCCACCTTCCTCCAGTTTATCACTGGCAGTCTCCTTTGAGTTCCCGGCC TAACCGCTGGG

Moderate or poor sequencing result.

## GroupB-Sample 16

Your sample has been re-sequenced. We have been able to improve your initial result.

# GroupB-Sample 19

Quality control passed.

### GroupB-Sample 23

TCCATCTCTGGAAAGTTCTCGACATGTCAAGACCAGGTAAGGTTCTTCGCGTTGCATCGAATTAAACCACATA CTCCACCGCTTGTGCGGGCCCCGTCAATTCCTTTGAGTTTCAGTCTTGCGACCGTACTCCCCAGGCGGCGA ACTTAACGCGTTAGCTTCGATACTGCGTGCCAAATTGCACCCAACATCCAGTTCGCATCGTTTAGGGCGTGG ACTACCAGGGTATCTAAWCCYGTTTGCTCCCCACGCTTTCGTGCCTCAGTGTCAATGTTGGTCCASGTAGCT GCCTTCGCCAYGGATGTTCCTCCCGNTCTCTACGCATTTCACTGCTAYACCGGGAATTCCACTACCCYCYMCC ACMCYCTWGTCGCCA

Your sample has been re-sequenced. We have been able to improve your initial result.

#### GroupB-Sample 27

CCCTCCGAAGGTTAAGCTACCTACTTCTTTTGCAACCCACTCCCATGGTGTGACGGGCGGTGTGTACAAGG
CCCGGGAACGTATTCACCGTGGCATTCTGATCCACGATTACTAGCGATTCCGACTTCATGGAGTCGAGTTGC
AGACTCCAATCCGGACTACGACGCACTTTATGAGGTCCGCTTGCTCTCGCGAGGTCGCTTCTCTTTGTATGC
GCCATTGTAGCACGTGTGTAGCCCTGGTCGTAAGGGCCATGATGACTTGACGTCATCCCCACCTTCCTCCAG
TTTATCACTGGCAGTCTCCTTTGAGTTCCCGGCCGGACCGCTGGCAACAAAGGATAAGGGTTGCGCTCGTTG
CGGGACTTAACCCAACATTTCACAACACGAGCTGACGACAGCCATGCAGCACCTGTCTCACGGTTCCCGAAG
GCACATTCTCATCTCTGAAAACTTCCGTGGATGTCAAGACCAGGTAAGGTTCTTCGCGTTGCATCGAATTAAA
CCACATGCTCCACCGCTTGTGCGGGCCCCCGTCAATTCATTTGAGTTTTAACCTTGCGGCCGTACTCCCCAG
GCGGTCGACTTAACGCGTTAGCTCCGGAAGCCACGCCTCAAGGGCACAACCTCCAAGTCGACATCGTTTACG
GCGTGGACTACCAGGGTATCTAATCCTGTTTGCTCCCCACGCTTTCGCACCTGAGCGTCAGTCTTCGTCCAG
GGGGCCGCCTTCGCCACCGGTATTCCTCCAGATCTCTACGCATTTCAYCGCTACACCTGGAATTCTACCCCCC
TCTACGAGACTCAAGCYTGCCAGTATCAGATGCAGTTCCCCAGNYTGAGCCCGGSGATTTCACATCTGACTTAA
CAAACCGCCTGCSTGCGCTTTACGT

Your sample has been re-sequenced. We have been able to improve your initial result.

# GroupB-Sample 29

GNAGCAACCCACTCCCATGGTGTGACGGGCGGTGTGTACAAGGCCCGGGAACGTATTCACCGTGACGTTCT
GATTCACGATTACTAGCGATTCCGACTTCACGCAGTCGAGTTGCAGACTGCGATCCGGACTACGATCGGTTT
TATGGGATTAGCTCCACCTCGCGGCTTGGCAACCCTTTGTACCGACCATTGTAGCACGTGTGTAGCCCTGGC
CGTAAGGGCCATGATGACTTGACGTCATCCCCACCTTCCTCCGGTTTGTCACCGGCAGTCTCCTTAGAGTGC
CCACCATAACGTGCTGGTAACTAAGGACAAGGGTTGCGCTCGTTACGGGACTTAACCCAACATCTCACGACA
CGAGCTGACGACAGCCATGCAGCACCTGTGTCTGAGTTCCCGAAGGCACCAATCCATCTCTGGAAAGTTCTC
AGCATGTCAAGGCCAGGTAAGGTTCTTCGCGTTGCTTCGAATTAAACCACATGCTCCACCGCTTGTGCGGGC
CCCCGTCAATTCATTTGAGTTTTAACCTTGCGGCCGTACTCCCCAGGCGGTCAACTTAATGCGTTAGCTGCGC
CACTAAGATCTCAAGGATCCCAACGGCTAGTTGACATCGTTTACGGCGTGGACTACCAGGGTATCTAATCCT
GTTTGCTCCCCACGCTTTCGCACCTCAGTGTCAGTGTCAGTCCACGGTAGTCGCCCTTCGCCACTGGTGTTCCTT
CCAATATCTACGCATTTCACCGCTACACTGGAAATTCCACTCCACCTCTACCGCACCTCTAGCCAGACAGTTTTG
GATGCAGTTCCCAGGTTGAGCCCGGGGATTTCACATCCAACTTATCAAGCCACCTACSCGCGCTTTATCCC
AGTAATTCCGATTAACGCTTGCACCCTTCGTATTACCGCGGGCTGCTGGCACGAAGTTAGCCGGTGCTTATTCT
GTTGGTAACGTCAAAACTCACAGGTATTCGCTATGAACCCCTTCCCCCAACTTAAAGTGCTTTACGACCCN

Your sample has been re-sequenced. We have been able to improve your initial result.

# GroupB-Sample 31 TGGGCCACCGGCTTCGGGTGTTACCGACTTTCNNGACTTGACGG

The signal strength of your sequencing reaction is very low. Therefore, your sequencing result needs careful evaluation.

CTCKCGNGGTGTGACGGGNGGTGTGTASAGACCCGGGAACGTATTCACCGTAKCANGCTGATCTACRATWA CGAKCGATWCCTGCTKCATGTAGTCKAGTTRCACACTACAATCCKAACTGAGAACAAYTTTATGGRATTTSCT TGACCTCGCGRTTTAGCTGCCCTTTGTATTGTYCATTGTAGCANGTGTGTAGCCCMAATCATAASGGGCATGA TKATTTGMCKYCMTCCCCACCTTCCTCYKGTTTGTCACCGGCNNTCWACTTATASTGCCCWACTTAATGATG GCAACTANNCTWARGGKTTGCGCTCKTTGCGRNAYTTAMCCCAACMTCTCAMGACACGAKCTGAMSACCAC CRTGCMCCMCCTGTCWCTTTGTCCCCCGAARGKSAAGACTCTATCTCTRAAKCGSTCAAAGGATGTCMAGAT TTGGKAWSGTTCTTCGCSTTSCTTCTAATTAAACCRCATGCTCCMCCGCTTGTGCSSSTCCCCGTYMATTYCW TTGAKTTTCAWCCTTGCGGTCKYACTCCCCAGGCKNAGTGCTTAATGCGTTNNCTGCAKCACTANNGGGCGN AAACCCCCTAACACTTASCACTCWTC

Your sample has been re-sequenced. The result obtained in the initial trial is confirmed.

# GroupB-Sample 33

ACCGACTTCGGGTGTTACAAACTCTCGTGGTGTGACGGGCGGTGTGACAAGGCCCGGGAACGTATTCACC GCGCATGCTGATCCGCGATTACTAGCGATTCCAGCTTCATGTAGGCGAGTTGCAGCCTACAATCCGAACTG AGAACGGTTTTATGAGATTAGCTCCACCTCGCGGTCTTGCAGCTCTTTGTACCGTCCATTGTAGCACGTGTGT AGCCCAGGTCATAAGGGGCATGATGATTTGACGTCATCCCCACCTTCCTCCGGTTTGTCACCGGCAGTCANC NTAGAGTGCCCAACTNAATGNTGGCAACTAAGNTNAAGGGTTGCGCTCGTTGCGGGACTTAACCCAACATCT CACGACACNAGCTGACNACAACCATGCACCACCTGTCACTTTGTCCCCCN

Your sample has been re-sequenced. We have been able to improve your initial result.

# GroupB-Sample 35

Moderate or poor sequencing result.

# GroupB-Sample 36

Your sample has been re-sequenced. We have been able to improve your initial result.

ACCGGCTTCGGGTGTTACAAACTCTCGTGGTGTGACGGGCGGTGTGTACAAGACCCGGGAACGTATTCACC GTAGCATGCTGATCTACGATTACTANCGATTCCAGCTTCATGTAGTCNAGTTGCAGACTACAATCCNAACTGA GAACAACTTTATGGGATTTGCTTGACCTCGCGGTTTCGCTGCCCTTTGTATTGTCCATTGTAGCACGTGTGTA GCCCAAATCATAAGGGGCATGATGATTTGACGTCATCCCCACCTTCCTCCGGTTTGTCACC

The signal strength of your sequencing reaction is very low. Therefore, your sequencing result needs careful evaluation.

## GroupB-Sample 39

Your sample has been re-sequenced. We have been able to improve your initial result.

# GroupB-Sample 41

TATCNNGNNGGNNSCSGGNTCTCCNGGNTGNCSGWTCASNNCCCCCCWANACCYNTGGCTTGACGGGCG GTGTGTACAAGGTCCGGGAACRTATTCACCGCGCCATGGCTGATGCGCGATTACTAGCGATTCCAGCTTCAT GGGGTCGGGTTGCAGACCCCAATCCGAACTGTGACCGGCTTTACAAGATTGGCTCCGGGTTACCCCCTCGC TACCCGCTGTACCGACCATTGTAGCACGTGTGTCGCCCTGGACGTAAGGGCCATGATGACTTGACGTCGTCC CCCCCTTCCTCTGCTTGCGCAGGCAGTCTACTTAGAGTTCCCGACATTACTCGCTGGCAACTAAGTACAGG GGTTGCGCTCGTTGCGGGACTTAACCCAACACCTCACGGCACGAGCTGACGACAGCCATGCAGCACCTTGTT TTGTGTGTATTGCTACACGGACCCATTTCTGAACCCTTCACGCACATTCTAGCCCAGGTAAGGTTCCTCGCGT ATCATCGAATTAAACCACATGCTCCACCGCTTGTGCGGACCCCCGTCAATTCCTTTGAGTTTCACTGTTGCCA GCGTACTCCCCAGGTGGATTACTTAACGCTTTCGCTCAGCCACACATGTTAGAACACATACAGCCAGTAATCA TCGTTTACGGCATGGACTACCAGGGTATCTAATCCTGTTCGCTACCCATGCTCTCGTGCCTCAGTGTCAATCA AGTCGTAGTAGCCTGCCTTCGCAATCGGTGTTCCGGGTCATATCTATGCATTTCACCGCTACATGACCCATTC CGGCTACCGCCAACCCATTCAAGTCATTCAGTTTCCAGCCACATCAGACGGTTAAGCCGCCTGCTTTCAAACC AGACTTAAATGACCACCTACGCACCCTTTAAACCCAATAAATCCGGACAACGCTTGCACCCTCCGTATTACCG CGGCTGCTGGCACGGAATTAGCCGGTGCTTATTCCTTGGCTACCGTCAMACAATCCCGCAGGACTGCTGTTC TTCACCAATAAAAGCAGTTTACAACGCTGAACGCCTTCMTCCTGGMCGCGGCATGGYTGGGTCAAACTTCCG TCCATTGCCCAAAATTCCYAATGGTGCCTCCCGKAGGAATTGGGTCCGTATCTCANTACCCNNGGTGGGGGC CAATCTNNTCCAAAACCCTANNNGAHCHTNGG

Your sample has been re-sequenced. We have been able to improve your initial result.

# GroupB-Sample 50

YRGGTGTTACCRNCTTTCGTGACTTGACGGGCGGTGTGTACAAGGCCCGGGAACGTATTCACCGCAGCGTTGCTGATCTGCGATTACTAGCGACTCCGACTTCATGGGGTCGAGTTGCAGACCCCAATCCGAACTGAGACCGGCCATTGTAGCATGCGTGAAGCCCAACCCATTGTACCGGCCATTGTAGCATGCGTGAAGCCCAA

GACATAAGGGGCATGATTTGACGTCGTCCTCACCTTCCTCCGAGTTGACCCCGGCAGTCTCCCATGAGT CCCCACCACKACGTGCTGGCAACATGGAACGAGGGTTGCGCTCGTTGCGGGACTTAACCCAACATCTCACG ACACGAGCTGACGACAACCATGCACCACCTGTGAACCCGCCCCAAAGGGGAAACCGTATCTCTACGGCGAT CGAGAACATGTCAAGCCTTGGTAAGGTTCTTCGCGTTGCATCGAATTAATCCGCATGCTCCGCCGCTTGTGC GGGCCCCCGTCAATTCCTTTGAGTTTTAGCCTTGCGGCCGTACTCCCCAGGCGGGGCACTTAATGCGTTAGC TGCGGCGCGGGAAACCGTGGAATGGTCCCCACACCTAGTGCCCAACGTTTACGGCATGGACTACCAGGGTAT CTAATCCTGTTCGCTCCCCATGCTTTCGCTCCTCAGCGTCAGTTACAGCCCAGAGACCTGCCTTCGCCATCGG TGTTCCTCCTGATATCTGCGCATTCCACCGCTACACCAGGAATTCCAGTCTCCCCTACTGCACTCTAGTCTGC CCGTACCCACCGCAGATCCGGGGTTAAGCCCCGGACTTTCACGACAGACGCGACAAACCGCCTACSAGCTC TTTACGCCCAATAATTCCGGATAACGCTCGCACCCTACGTATTACCGCGGCTGCTGGCACGTAATTAGCCGG TGCTTCTTCTCGCACGTCATTTCCCTACTGAAAAAAAGGTTTTACAACCCGAAGGCCGTCATY CCT

Your sample has been re-sequenced. We have been able to improve your initial result.

## GroupB-Sample 51

GGTTACTCCACCGGCTTCGGGTGTTACAAACTCTCGTGGTGTGACGGGCGGTGTGTACAAGACCCGGGAAC GTATTCACCGTAGCATGCTGATCTACGATTACTAGCGATTCCAGCTTCATGTAGTCGAGTTGCAGACTACAAT CCGAACTGAGAACAACTTTATGGGATTTGCTTGACCTCGCGGTTTAGCTGCCCTTTGTATTGTCCATTGTAGC ACGTGTGANCCCAAATCATAAGGGGCATGATGATTTGACGTCNTCCCCACCTTCCTCCGGTTTGTCACCGG CNGTCAACTTA

Your sample has been re-sequenced. We have been able to improve your initial result.

GroupB-Sample 52 NNNNN

Moderate or poor sequencing result.

# GroupB-Sample 53

GCTCCATAAAGGTTACCTCACCGACTTCRGGTGTTGCAAACTCTCGTGGTGTGACGGGCGGTGTGTACAAGG CCCGGGAACGTATTCACCGCGGCATGCTGATCCGCGATTACNAGCGATTCCNGCTTCACGCAGTCGAGTTG CAGACTGCGATCCGAACTGAGAACAGATTTATGGGATTGGCTAAACCTTGCGGTCTCGCAGCCCTTTGTTCT GTCCATTGTAGCACGTGTGTAGCCCAGGTCATAAGGGGCATGATGATTTGACGTCATCCCCACCTTCCTCCG GTTTGTCACCGGCAGTCACCTTAGAGTGCCCAACTAAATGCTGGCAACTAAGATCAAGGGTTGCGCTCGTTG CGGGACTTAACCCAACATCTCACGACACGAGCTGACGACAACCATGCACCACCTGTCACTCTGTCCCCGAAG GGAAAGCCCTATCTCTAGGGTTGTCAGAGGATGTCAAGACCTGGTAAGGTTCTTCGCGTTGCTTCGAATTAA ACCACATGCTCCACCGCTTGTGCGGGCCCCCGTCAATTCCTTTGAGTTTCAGTCTTGCGACCGTACTCCCCA GGCGGAGTGCTTAATGCGTTAGCTGCAGCACTAAGGGGCGGAAACCCCCTAACACTTAGCACTCATCGTTTA CGGCGTGGACTACCAGGGTATCTAATCCTGTTCGCTCCCCACGCTTTCGCTCCTCAGCGTCAGTTACAGACC AGAGAGTCGCCTTCGCCACTGGTGTTCCTCCACATCTCTACGCATTTCACCGCTACACGTGGAATTCCACTCT CCTCTTCTGCACTCAAGTTTCCCAGTTTCCAATGACCCTCCCCGGTTGAGCCGGGGNCTTTCACATCAGACTT AAGAAACCGCCTGCGAGCCCTTTACGCCCAATAATTCCGGACAACGCTTGCCACCTACGTATTACCGCGGCT GCTGGCACGTANTTANCCGTGGCTTTCTGGTNAGNTACCGTCNAGGTGCGAGCAGTTACTCTCGCACTTGTT CTTCCCTAACAACAGAACTTTACGATCCSAAAANCYTCNYCAYTCACGCSGCGTTSCTCCGTCAAACTTTCTTC CATTGCGGAAAATTCCCTACWGCTGCCTCCCCTAAGAAATC

Your sample has been re-sequenced. We have been able to improve your initial result.

GroupB-Sample 54

ANGGGGTTAGGCCACCGGCTTCGGGTGTTACCAACTTTCGTGACTTGACGGGCGGTGTGTACAAGGCCCGG

Quality control passed.

## GroupB-Sample 55

CAAGGGGTTAGGCCACCGGCTTCGGGTGTTACCAACTTTCGTGACTTGACGGGCGGTGTGTACAAGGCCCGGGAACGTATTCACCGCAGCGTTGCTGATCTGCGATTACTAGCGACTCCAACTTCATGAGGTCGAGTTGCAGACCCCAATCCGAACTGAGACCGGCTTTTTGGGATTAGCTCCACCTCACAGTATCGCAACCCTTTGTACCGGCCATTGTAGCATGCGTGAAGCCCAAGACATAAGGGGCATGATGATTTGACGTCGTCCCCACCTTCCTCCGAGTTGACCCCGGCAGTCTCCCCATGAGTCCCCGGCATAACCCGCTGGCAACATGGAACGAGGGTTGCGCTCGTTGCGGG

Your sample has been re-sequenced. We have been able to improve your initial result.

# GroupB-Sample 57

Your sample has been re-sequenced. The result obtained in the initial trial is confirmed.

## GroupB-Sample 60

AGGTTAGACTACTTCTGGTGCAACCCACTCCCATGGTGTACCGGGCGGTGTGTACAAGGCCCGGGAA CGTATTCACCGCGACATTCTGATTCGCGATTACTAGCGATTCCGACTTCACGCAGTCGAGTTGCAGACTGCG ATCCGGACTACGATCGGATTGTGAGATTAGCTCCACCTCGCGGCTTGGCAACCCTCTGTACCGACCATTGT ANCACGTGTGAGCCCANGCCGTAAGGGCCATGATGACTTGACGTCATCCCCACCTTCCTCCGGTTTGTCAN CGGNA

The signal strength of your sequencing reaction is very low. Therefore, your sequencing result needs careful evaluation.

CCTCCCGAAGGTTAAGCTACCTACTTCTTTTGCAACCCACTCCCATGGTGTGACGGGCGGTGTGTACAAGGC CCGGGAACGTATTCACCGTGGCATTCTGATCCACGATTACTAGCGATTCCGACTTCATGGAGTCGAGTTGCA GACTCCAATCCGGACTACGACGCACTTTATGAGGTCCGCTTGCTCTCGCGAGGTCGCTTCTCTTTGTATGCG CCATTGTAGCACGTGTGAGCCCTGGTCGTAAGGGCCATGATGACTTGACGTCATCCCCACCTTCCTCCAGT TTATCACTGGNNGTCTCCTTTGAGTTCCCGGCCGGACCGCTGGCAACAAAGGATAAGGGTTGCGCTCGTTGC GGGACTTAACCCAACATTTCACAACACGAGCTGACGACAGCCATGCAGCA

The signal strength of your sequencing reaction is very low. Therefore, your sequencing result needs careful evaluation.

## GroupB-Sample 63

TTACTCCACCGGCTTCGGGTGTTACAGACTCTCGTGGTGTGACGGGCGGTGTGTACAAGACCCGGGAACGT ATTCACCGTAGCATGCTGATCTACGATTACTAGCGATTCCNGCTTCATGTAGTCGAGTTGCAGACTACAATCC GAACTGAGAACAACTTTATGGGATTTGCATGACCTCGCGGTTTAGCTGCCCTTTGTATTGTCCATTGTAGCAC GTGTGTAGCCCAAATCATAAGGGGCATGATGATTTGACGTCATCCCCACCTTCCTCCGGTTTGTCACCGGCA GTCAACCTAGAGTGCCCAACTAAATGCTGGCAACTAAGTTTAAGGGTTGCGCTCGTTGCGGGACTTAACCCA ACATCTCACGACACGAGCTGACGACAACCATGCACCACCTGTCACTTTGTCCCCCGAAGGGRAAGGCTCTAT CTCTAGAGTTTTCAAAGGATGTCAAGATTTGGTAAGGTTCTTCGCGTTGCTTCGAATTAAACCACATGCTCCA CCGCTTGTGCGGGTCCCCGTCAATTCCTTTGAGTTTCAACCTTGCGGTCGTACTCCCCAGGCGGAGTGCTTA ATGCGTTAGCTGCAGCACTAAGGGGCGGAAACCCCCTAACACTTAGCACTCATCGTTTACGGCGTGGACTAC CAGGGTATCTAATCCTGTTTGATCCCCACGCTTTCGCACATCAGCGTCAGTTACAGACCAGAAAGTCGCCTTC GCCACTGGTGTTCCTCCATATCTCTGCGCATTTCACCGCTACACATGGAATTCCACTTTCCTCTTCTGTACTCA AGTTTCCCAGTTTCCAATGACCCTCCACGGTTGANCCGTGGGCTTTCACATCAGACTTAAGAAACCGCCTACG CGCGCTTTACGCCCAATAATTCCGGATAACGCTTGCCACCTACGTATTACCGCGGCTGCTGGCACGTANTTA GCCGTGGCTTTCTGATTANGTACCGTCAAGATGTGCACAGTTACTTACACATTTGTTCTTCCCTAATAACAAAA TTTTAMGATCCGAAAACCTTCATCATTCNGGCGGCGTTGCTCCGTCMGGCTTTNNCCCATTGCGGAAAAATT **CCCTACTGCTGC** 

Your sample has been re-sequenced. We have been able to improve your initial result.

## GroupB-Sample 64

Quality control passed.