## **Supporting Information**

## Multidimensional Data Organization and Random Access in Large-Scale DNA Storage Systems

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## Results of In Silico PCR Experiments of Sixteen Random-Access Patterns

## (1) UFP+URP: all blocks.

#### Data retrieval target:

All blocks (entry #1, 3, 5, 7, 9, 11, 13, 15, 2, 4, 6, 8, 10, 12, 14, 16)

#### Input primer list:

>UFP

AAGGCAAGTTGTTACCAGCA

>URF

TGCGACCGTAATCAAACCAA

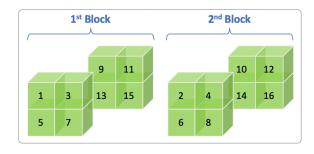
#### In silico PCR result:

In silico Primer(s) search for: ////seq1:

>ufp 1->20 5'-aaggcaagttgttaccagca >urp 106<-125 5'-tgcgaccgtaatcaaaccaa

>1-125 Amplicon size: 125bp Ta=40°C

In silico Primer(s) search for: ////seq3:



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>ufp 1->20 5'-aaggcaagttgttaccagca 116<-135 >urp 5'-tgcgaccgtaatcaaaccaa >1-135 Amplicon size: 135bp Ta=40°C a agg caagt t g trace ag caag c t caa accat g g c t cat t t caca a t c g g ta aa caa accat g g c t c g t t t a caa a t t a g g c g a c g c g a t t t g g t t g a t t a c g g t c g c g c g a t t g g t t g a t a c g g c g a c gIn silico Primer(s) search for: ////seq5: >ufp 1->20 5'-aaggcaagttgttaccagca 126<-145 5'-tgcgaccgtaatcaaaccaa >1-145 Amplicon size: 145bp gattacggtcgc In silico Primer(s) search for: ////seq7: >ufp 1->20 5'-aaggcaagttgttaccagca 136<-155 >urp 5'-tgcgaccgtaatcaaaccaa >1-155 Amplicon size: 155bp Ta=41°C a agg caag t t g t a caag caag c caa a caatttggtttgattacggtcgc In silico Primer(s) search for: ////seq9: >ufp 1->20 5'-aaggcaagttgttaccagca >urp 146<-165 5'-tgcgaccgtaatcaaaccaa >1-165 Amplicon size: 165bp Ta=41°C  ${\tt gccgcctccagtttttggtttgattacggtcgc}$ In silico Primer(s) search for: ////seq11: >ufp 1->20 5'-aaggcaagttgttaccagca >urp 156<-175 5'-tgcgaccgtaatcaaaccaa >1-175 Amplicon size: 175bp Ta=41°C gttttaatttgccgcctccagtttttggtttgattacggtcgc In silico Primer(s) search for: ////seq13: >ufp 1->20 5'-aaggcaagttgttaccagca >urp 166<-185 5'-tgcgaccgtaatcaaaccaa

>1-185 Amplicon size: 185bp Ta=41°C

aatcaacgacgaacgaataatttgccgcctccagtttttggtttgattacggtcgc In silico Primer(s) search for: ////seq15: >ufp 1->20 5'-aaggcaagttgttaccagca >urp 176<-195 5'-tgcgaccgtaatcaaaccaa >1-195 Amplicon size: 195bp Ta=41°C aaaaaaaacaaact catggctccgttttaatttgccgcctccagtttttggtttgattacggtcgcIn silico Primer(s) search for: ////seq2: >ufp 1->20 5'-aaggcaagttgttaccagca >urp 111<-130 5'-tgcgaccgtaatcaaaccaa >1-130 Amplicon size: 130bp Ta=40°C a agg caagt t g trace a cag cag trace a attgcg t g cac attg g creat trace a cag cag access a case can calculate the case can be a calculated and can be a calculated as a case can be a calculated as a caIn silico Primer(s) search for: ////seq4: 1->20 >ufp 5'-aaggcaagttgttaccagca 121<-140 5'-tgcgaccgtaatcaaaccaa >1-140 Amplicon size: 140bp Ta=40°C In silico Primer(s) search for: ////seq6: 1->20 5'-aaggcaagttgttaccagca >urp 131<-150 5'-tgcgaccgtaatcaaaccaa >1-150 Amplicon size: 150bp Ta=41°C ggtttgattacggtcgc In silico Primer(s) search for: ////seq8: >ufp 1->20 5'-aaggcaagttgttaccagca >urp 141<-160 5'-tgcgaccgtaatcaaaccaa >1-160 Amplicon size: 160bp Ta=41°C cgcgaatttggtttgattacggtcgc

In silico Primer(s) search for: ///seq10:

1->20

>ufp

S3

5'-aaggcaagttgttaccagca 151<-170

5'-tgcgaccgtaatcaaaccaa

>1-170 Amplicon size: 170bp Ta=41°C

a atttgccgcctccagtttttggtttgattacggtcgc

In silico Primer(s) search for: ////seq12:

>ufp 5'-aaggcaagttgttaccagca 161<-180 >urp 5'-tgcgaccgtaatcaaaccaa

>1-180 Amplicon size: 180bp Ta=41°C

gctccgttttaatttgccgcctccagtttttggtttgattacggtcgc

In silico Primer(s) search for: ////seq14:

>ufp 1->20 5'-aaggcaagttgttaccagca >urp 171<-190 5'-tgcgaccgtaatcaaaccaa

>1-190 Amplicon size: 190bp Ta=41°C

aaccaatcaacgacgaacgaataatttgccgcctccagtttttggtttgattacggtcgc

In silico Primer(s) search for: ////seq16:

>ufp 1->20 5'-aaggcaagttgttaccagca 181<-200 5'-tgcgaccgtaatcaaaccaa

>1-200 Amplicon size: 200bp

aaaaaaaaaaaaaaacaaactcatggctccgttttaatttgccgcctccagtttttggtttgattacggtcgc

## (2) UFP+TP: specific table from all blocks.

#### Data retrieval target:

The 1<sup>st</sup> table from all blocks (entry #1, 3, 5, 7, 2, 4, 6, 8)

#### Input primer list:

>UFP

AAGGCAAGTTGTTACCAGCA

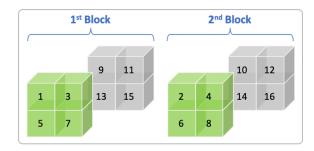
>TP1

ATTCGCGTCGCCTAATTTGT

#### In silico PCR result:

In silico Primer(s) search for: ////seq1:

>ufp 1->20



5'-aaggcaagttgttaccagca	
>tp1 86<-105 5'-attcgcgtcgcctaatttgt	
>1-105 Amplicon size: 105bp Ta=40°C	
aaggcaagttgttaccagcaagccgacaagttcaaacacatggctcatttcacaatcggtaaaa	aaccaatcaacgacgaacgaaacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq3:	
>ufp 1->20	
5'-aaggcaagttgttaccagca	
>tp1 96<-115 5'-attcgcgtcgcctaatttgt	
>1-115 Amplicon size: 115bp Ta=40°C aaggcaagttgttaccagcaagccgacaagttcaaacacatggctcatttcacaatcggtaaaa	aaaaaaaaaaaaacaaactcatggctccgtttacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq5:	
>ufp 1->20	
5'-aaggcaagttgttaccagca	
>tp1 106<-125 5'-attcgcgtcgcctaatttgt	
>1-125 Amplicon size: 125bp Ta=40°C aaggcaagttgttaccagcaagccgacaagttcaaacacaataaatgacctgccgtgcaaaaa	aaaaaaaaaaaaaaaaaaaaaaaaccaatcaacgacgaacga
In silico Primer(s) search for: ////seq7:	
>ufp 1->20	
5'-aaggcaagttgttaccagca	
>tp1 116<-135 5'-attcgcgtcgcctaatttgt	
>1-135 Amplicon size: 135bp Ta=40°C	aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaacaaactcatggctccgtttacaaattaggcgacgcg
aa	
In silico Primer(s) search for: ////seq9:	
In silico Primer(s) search for: ////seq11:	
In silico Primer(s) search for: ////seq13:	
In silico Primer(s) search for: ////seq15:	
In silico Primer(s) search for: ////seq2:	
>ufp 1->20	
5'-aaggcaagttgttaccagca	
>tp1 91<-110 5'-attcgcgtcgcctaatttgt	
>1-110 Amplicon size: 110bp Ta=40°C aaggcaagttgttaccagcagttcaaattgcgtgcgacattggctcatttcacaatcggtaaaaa	aaaaaaccaatcaacgacgaacgaaacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq4:	

>ufp 1->20 5'-aaggcaagttgttaccagca	
>tp1 101<-120	
5'-attcgcgtcgcctaatttgt	
>1-120 Amplicon size: 120bp Ta=40°C aaggcaagttgttaccagcagttcaaattgcgtgcgacattggctcatttcacaatcggtaa	aaaaaaaaaaaaaaaaaaaaaacaaactcatggctccgtttacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq6:	
>ufp 1->20	
5'-aaggcaagttgttaccagca	
>tp1 111<-130	
5'-attcgcgtcgcctaatttgt	
>1-130 Amplicon size: 130bp Ta=40°C	
	ааааааааааааааааааааааааааааааааааааа
In silico Primer(s) search for: ////seq8:	
>ufp 1->20	
5'-aaggcaagttgttaccagca	
>tp1 121<-140	
5'-attcgcgtcgcctaatttgt	
>1-140 Amplicon size: 140bp Ta=40°C	
a agg caagt t g taccag cag t tcaa at t g c g t g c g a catataa at g acct g c c g t g caa a c g c g a a c g c g a a c g c g	aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq10:	
In silico Primer(s) search for: ////seq12:	
In silico Primer(s) search for: ////seq14:	
In silico Primer(s) search for: ////seq16:	

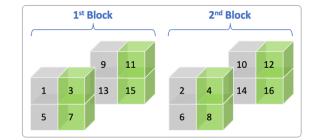
## (3) UFP+CP: specific column from all tables in all blocks.

#### Data retrieval target:

In silico PCR result:

The 2<sup>nd</sup> column from all tables in all blocks (entry #3, 7, 11, 15, 4, 8, 12, 16)

# Input primer list: >UFP AAGGCAAGTTGTTACCAGCA >CP2 AAACGGAGCCATGAGTTTGT



In silico Primer(s) search for: ////seq1:

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In silico Primer(s) search for: ////seq3:
>ufp 1->20
5'-aaggcaagttgttaccagca
>cp2 76<-95
5'-aaacggagccatgagtttgt
>1-95 Amplicon size: 95bp Ta=40°C aaggcaagttgttaccagcaagctgacaagttcaaacacatggctcatttcacaatcggtaaaaaaaa
In silico Primer(s) search for: ////seq5:
In silico Primer(s) search for: ////seq7:
>ufp 1->20
5'-aaggcaagttgttaccagca
>cp2 96<-115
5'-aaacggagccatgagtttgt
>1-115 Amplicon size: 115bp Ta=40°C aaggcaagttgttaccagcaagccgacaagttcaaacacaataaatgacctgccgtgcaaaaaaaa
In silico Primer(s) search for: ////seq9:
In silico Primer(s) search for: ////seq11:
>ufp 1->20
5'-aaggcaagttgttaccagca
>cp2 116<-135
5'-aaacggagccatgagtttgt
>1-135 Amplicon size: 135bp Ta=40°C
aaggcaagttgttaccagcaagccgacaagttcaaacacatggctcatttcacaatcggtaaaaaaaa
gtt
In silico Primer(s) search for: ////seq13:
In silico Primer(s) search for: ////seq15:
>ufp 1->20
5'-aaggcaagttgttaccagca
>cp2 136<-155
5'-aaacggagccatgagtttgt
>1-155 Amplicon size: 155bp Ta=41°C
aaggcaagttgttaccagcaagccgacaagttcaaacacaataaatgacctgccgtgcaaaaaaaa
aaaaaaaacaaactcatggctccgtt
In silico Primer(s) search for: ////seq2:
550
In silico Primer(s) search for: ////seq4:
>ufp 1->20
5'-aaggcaagttgttaccagca
>cp2 81<-100
5'-aaacggagccatgagtttgt

>1-100 Amplicon size: 100bp Ta=40°C

In silico Primer(s) search for: ////seq6: In silico Primer(s) search for: ////seq8: >ufp 1->20 5'-aaggcaagttgttaccagca >cp2 101<-120 5'-aaacggagccatgagtttgt >1-120 Amplicon size: 120bp Ta=40°C In silico Primer(s) search for: ////seq10: In silico Primer(s) search for: ////seq12: >ufp 1->20 5'-aaggcaagttgttaccagca >cp2 121<-140 5'-aaacggagccatgagtttgt >1-140 Amplicon size: 140bp gctccgtt In silico Primer(s) search for: ////seq14: In silico Primer(s) search for: ////seq16: >ufp 1->20 5'-aaggcaagttgttaccagca >cp2 141<-160 5'-aaacggagccatgagtttgt >1-160 Amplicon size: 160bp Ta=41°C aaaaaaaaaaaacaaactcatggctccgtt (4) RP+URP: specific row from all tables in all blocks.

#### Data retrieval target:

The 1<sup>st</sup> row from all tables in all blocks (entry #1, 3, 9, 11, 2, 4, 10, 12)

#### Input primer list:

>RP1

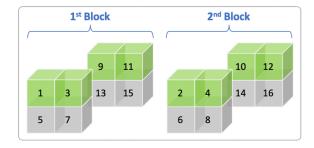
**TGGCTCATTTCACAATCGGT** 

>URP

TGCGACCGTAATCAAACCAA

#### In silico PCR result:

In silico Primer(s) search for: ////seq1:



>rp1 41->60
5'-tggctcatttcacaatcggt
>urp 106<-125 5'-tgcgaccgtaatcaaaccaa
J. ISCENICE FUNCTION
>41-125 Amplicon size: 85bp Ta=40°C tggctcatttcacaatcggtaaaaaaccaatcaacgacgaacga
In silico Primer(s) search for: ////seq3:
>rp1 41->60
5'-tggctcatttcacaatcggt
>urp 116<-135
5'-tgcgaccgtaatcaaaccaa
>41-135 Amplicon size: 95bp Ta=40°C tggctcatttcacaatcggtaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq5:
In silico Primer(s) search for: ////seq7:
In silico Primer(s) search for: ////seq9:
>rp1 41->60
5'-tggctcatttcacaatcggt
>urp 146<-165 5'-tgcgaccgtaatcaaaccaa
3 1,00,000,000,000
>41-165 Amplicon size: 125bp Ta=40°C
tggctcatttcacaatcggtaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq11:
111 3111co 1 11111c1(3) 3 carett 101. ////3 cq11.
>rp1 41->60
5'-tggctcatttcacaatcggt
>urp 156<-175 5'-tgcgaccgtaatcaaaccaa
>41-175 Amplicon size: 135bp Ta=40°C tggctcatttcacaatcggtaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq13:
In silico Primer(s) search for: ////seq15:
In silico Primer(s) search for: ////seq2:
>rp1 41->60
5'-tggctcatttcacaatcggt
>urp 111<-130
5'-tgcgaccgtaatcaaaccaa
>41-130 Amplicon size: 90bp Ta=40°C tggctcatttcacaatcggtaaaaaaaaaccaatcaacgacgaacga

In silico Primer(s) search for: ////seq4:
>rp1 41->60 5'-tggctcatttcacaatcggt >urp 121<-140 5'-tgcgaccgtaatcaaaccaa
>41-140 Amplicon size: 100bp Ta=40°C tggctcatttcacaatcggtaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq6:
In silico Primer(s) search for: ////seq8:
In silico Primer(s) search for: ////seq10:
>rp1 41->60 5'-tggctcatttcacaatcggt >urp 151<-170 5'-tgcgaccgtaatcaaaccaa
>41-170 Amplicon size: 130bp Ta=40°C tggctcatttcacaatcggtaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq12:
>rp1 41->60 5'-tggctcatttcacaatcggt >urp 161<-180 5'-tgcgaccgtaatcaaaccaa
>41-180 Amplicon size: 140bp Ta=40°C tggctcatttcacaatcggtaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq14:
In silico Primer(s) search for: ////seq16:

## (5) RP+TP: specific row from specific table in all blocks.

#### Data retrieval target:

The 2<sup>nd</sup> row from the 1<sup>st</sup> table in all blocks (entry #5, 7, 6, 8)

## Input primer list:

>RP2

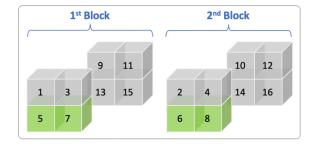
ATAAATGACCTGCCGTGCAA

>TP1

ATTCGCGTCGCCTAATTTGT

### In silico PCR result:

In silico Primer(s) search for: ////seq1:



In silico Primer(s) search for: ////seq3:	
In silico Primer(s) search for: ////seq5:	
>rp2 41->60 5'-ataaatgacctgccgtgcaa >tp1 106<-125	
5'-attcgcgtcgcctaatttgt	
>41-125 Amplicon size: 85bp Ta=40°C ataaatgacctgccgtgcaaaaaaaaaaaaaaaaaaaaa	gacgaacgaaacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq7:	
>rp2 41->60 5'-ataaatgacctgccgtgcaa >tp1 116<-135 5'-attcgcgtcgcctaatttgt	
>41-135 Amplicon size: 95bp Ta=40°C ataaatgacctgccgtgcaaaaaaaaaaaaaaaaaaaaa	aacaaactcatggctccgtttacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq13:	
In silico Primer(s) search for: ////seq15:	
In silico Primer(s) search for: ////seq2:	
In silico Primer(s) search for: ////seq4:	
In silico Primer(s) search for: ////seq6:	
>rp2 41->60	
5'-ataaatgacctgccgtgcaa >tp1 111<-130	
5'-attcgcgtcgcctaatttgt	
>41-130 Amplicon size: 90bp Ta=40°C ataaatgacctgccgtgcaaaaaaaaaaaaaaaaaaaaa	atcaacgacgaacgaaacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq8:	
>rp2 41->60 5'-ataaatgacctgccgtgcaa >tp1 121<-140	
5'-attegegtegectaatttgt	
>41-140 Amplicon size: 100bp Ta=40°C ataaatgacctgccgtgcaaaaaaaaaaaaaaaaaaaaa	aaaaaaacaaactcatggctccgtttacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq14:	

In silico Primer(s) search for: ////seq16:

S11

# (6) UFP+TP then UFP+CP: specific column from specific table in all blocks.

## Data retrieval target:

gttttaatttgccgcctccagtt

In silico Primer(s) search for: ////seq13:

The 1 <sup>st</sup> column from the 2 <sup>nd</sup> table in all blocks (en	try #9, 13, 10, 14)	
Input primer list: >UFP	1 <sup>st</sup> Block	2 <sup>nd</sup> Block
AAGGCAAGTTGTTACCAGCA	9 11	10 12
>TP2 AAACTGGAGGCGGCAAATTA	1 3 13 15	2 4 14 16
Then, >UFP	5 7	6 8
AAGGCAAGTTGTTACCAGCA		
>CP1 TTCGTTCGTCGTTGATTGGT		
In silico PCR result:		
>>> 1st PCR round >>>	_	
In silico Primer(s) search for: ////seq1:		
In silico Primer(s) search for: ////seq3:	-	
In silico Primer(s) search for: ////seq5:	-	
In silico Primer(s) search for: ////seq7:	-	
In silico Primer(s) search for: ////seq9:	-	
>ufp 1->20 5'-aaggcaagttgttaccagca >tp2 126<-145		
5'-aaactggaggcggcaaatta		
>1-145 Amplicon size: 145bp Ta=40°C aaggcaagttgttaccagcaagccgacaagttcaaacacatggctcatttcacaatcggccgcctccagtt	gtaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa	aaaaaaaccaatcaacgacgaacgaataatti
In silico Primer(s) search for: ////seq11:	-	
>ufp 1->20 5'-aaggcaagttgttaccagca >tp2 136<-155 5'-aaactggaggcggcaaatta		
>1 155 Amplican ciza: 155hp T2-41°C		

S12

>ufp 1->20 5'-aaggcaagttgttaccagca >tp2 146<-165 5'-aaactggaggcggcaaatta >1-165 Amplicon size: 165bp Ta=41°C aatcaacgacgaacgaataatttgccgcctccagtt In silico Primer(s) search for: ////seq15: >ufp 1->20 5'-aaggcaagttgttaccagca 156<-175 5'-aaactggaggcggcaaatta >1-175 Amplicon size: 175bp Ta=41°C aaaaaaaaaaaactcat ggctccgttttaatttgccgcctccagttIn silico Primer(s) search for: ////seq2: In silico Primer(s) search for: ////seq4: In silico Primer(s) search for: ////seq6: In silico Primer(s) search for: ////seq8: In silico Primer(s) search for: ////seq10: >ufp 1->20 5'-aaggcaagttgttaccagca >tp2 131<-150 5'-aaactggaggcggcaaatta >1-150 Amplicon size: 150bp Ta=41°C aatttgccgcctccagtt In silico Primer(s) search for: ////seq12: >ufp 1->20 5'-aaggcaagttgttaccagca >tp2 141<-160 5'-aaactggaggcggcaaatta >1-160 Amplicon size: 160bp Ta=41°C  ${\tt gctccgttttaatttgccgcctccagtt}$ In silico Primer(s) search for: ////seq14: >ufp 1->20 5'-aaggcaagttgttaccagca >tp2 151<-170 5'-aaactggaggcggcaaatta

>1-170 Amplicon size: 170bp

Ta=41°C

a agg caa gtt gtt accag cag tt caa att gc gt gc gac at at aaat gac ct gc c gt gc aaccaat caa c gac gaa c gaa ta att tt gc gc ct cca gt t	aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq16:	
>ufp 1->20 5'-aaggcaagttgttaccagca >tp2 161<-180 5'-aaactggaggcggcaaatta	
>1-180 Amplicon size: 180bp Ta=41°C aaggcaagttgttaccagcagttcaaattgcgtgcgacatataaatgacctgccgtgcaaaaaaaa	aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa
Then,	
>>> 2 <sup>nd</sup> PCR round >>>	
In silico Primer(s) search for: ////seq9:	
>ufp 1->20 5'-aaggcaagttgttaccagca >cp1 106<-125 5'-ttcgttcgtcgttgattggt	
>1-125 Amplicon size: 125bp Ta=40°C aaggcaagttgttaccagcaagccgacaagttcaaacacatggctcatttcacaatcgg	gtaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq11:	
In silico Primer(s) search for: ////seq13:	
>ufp 1->20 5'-aaggcaagttgttaccagca >cp1 126<-145 5'-ttcgttcgtcgttgattggt	
>1-145 Amplicon size: 145bp Ta=40°C aaggcaagttgttaccagcaagccgacaagttcaaacacaataaatgacctgccgtgcaatcaacgacgaacga	ааааааааааааааааааааааааааааааааааааааа
In silico Primer(s) search for: ////seq15:	
In silico Primer(s) search for: ////seq10:	
>ufp 1->20 5'-aaggcaagttgttaccagca >cp1 111<-130 5'-ttcgttcgtcgttgattggt	
>1-130 Amplicon size: 130bp Ta=40°C	ааааааааааааааааааааааааааааааааааааааа
In silico Primer(s) search for: ////seq12:	
In silico Primer(s) search for: ////seq14:	
>ufp 1->20	

5'-aaggcaagttgttaccagca 131<-150 5'-ttcgttcgtcgttgattggt >1-150 Amplicon size: 150bp aaccaatcaacgacgaacga In silico Primer(s) search for: ////seq16: (7) RP+CP: specific entry from all tables in all blocks. Data retrieval target: Entry at the 1<sup>st</sup> row 2<sup>nd</sup> column from all tables in all blocks (entry #3, 11, 4, 12) Input primer list: 1<sup>st</sup> Block 2<sup>nd</sup> Block >RP1 TGGCTCATTTCACAATCGGT 9 11 10 12 >CP2 AAACGGAGCCATGAGTTTGT 15 2 5 6 8 In silico PCR result: In silico Primer(s) search for: ////seq1: In silico Primer(s) search for: ////seq3: 41->60 5'-tggctcatttcacaatcggt 76<-95 >cp2 5'-aaacggagccatgagtttgt >41-95 Amplicon size: 55bp Ta=40°C In silico Primer(s) search for: ////seq7: In silico Primer(s) search for: ////seq9: In silico Primer(s) search for: ////seq11: >rp1 41->60 5'-tggctcatttcacaatcggt >cp2 116<-135 5'-aaacggagccatgagtttgt >41-135 Amplicon size: 95bp Ta=40°C In silico Primer(s) search for: ////seq15:

In silico Primer(s) search for: ////seq2:

In silico Primer(s) search for: ////seq4: 41->60 >rp1 5'-tggctcatttcacaatcggt >cp2 81<-100 5'-aaacggagccatgagtttgt >41-100 Amplicon size: 60bp Ta=40°C In silico Primer(s) search for: ////seq8: In silico Primer(s) search for: ////seq10: In silico Primer(s) search for: ////seq12: 41->60 >rp1 5'-tggctcatttcacaatcggt

>cp2 121<-140 5'-aaacggagccatgagtttgt

>41-140 Amplicon size: 100bp Ta=40°C

In silico Primer(s) search for: ////seq16:

## (8) UFP+TP then RP+CP: specific entry from specific table in all blocks.

#### Data retrieval target:

Entry at the 2<sup>nd</sup> row 2<sup>nd</sup> column from the 1<sup>st</sup> table in all blocks (entry #7, 8)

#### Input primer list:

>UFP

AAGGCAAGTTGTTACCAGCA

>TP1

ATTCGCGTCGCCTAATTTGT

Then,

>RP2

ATAAATGACCTGCCGTGCAA

>CP2

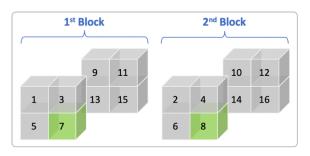
AAACGGAGCCATGAGTTTGT

#### In silico PCR result:

#### >>> 1st PCR round >>>

In silico Primer(s) search for: ////seq1:

>ufp 1->20 5'-aaggcaagttgttaccagca >tp1 86<-105



5'-attcgcgtcgcctaatttgt	
>1-105 Amplicon size: 105bp Ta=40°C aaggcaagttgttaccagcaagccgacaagttcaaacacatggctcatttcacaatcggtaaa	aaaccaatcaacgacgaacgaaacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq3:	
>ufp 1->20 5'-aaggcaagttgttaccagca >tp1 96<-115 5'-attcgcgtcgcctaatttgt	
>1-115 Amplicon size: 115bp Ta=40°C aaggcaagttgttaccagcaagccgacaagttcaaacacatggctcatttcacaatcggtaaa	aaaaaaaaaaaaaactcatggctccgtttacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq5:	
>ufp 1->20 5'-aaggcaagttgttaccagca >tp1 106<-125 5'-attcgcgtcgcctaatttgt	
>1-125 Amplicon size: 125bp Ta=40°C aaggcaagttgttaccagcaagccgacaagttcaaacacaataaatgacctgccgtgcaaaa	ааааааааааааааааааааааааааассааtcaacgacgaacgaaacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq7:	
>ufp 1->20 5'-aaggcaagttgttaccagca >tp1 116<-135 5'-attcgcgtcgcctaatttgt	
>1-135 Amplicon size: 135bp Ta=40°C aaggcaagttgttaccagcaagccgacaagttcaaacacaataaatgacctgccgtgcaaaa aa	aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq9:	
In silico Primer(s) search for: ////seq11:	
In silico Primer(s) search for: ////seq13:	
In silico Primer(s) search for: ////seq15:	
In silico Primer(s) search for: ////seq2:	
<pre>&gt;ufp 1-&gt;20 5'-aaggcaagttgttaccagca &gt;tp1 91&lt;-110 5'-attcgcgtcgcctaatttgt</pre>	
>1-110 Amplicon size: 110bp Ta=40°C aaggcaagttgttaccagcagttcaaattgcgtgcgacattggctcatttcacaatcggtaaaa	aaaaaaaccaatcaacgacgaacgaaacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq4:	

>ufp

1->20

5'-aaggcaagttgttaccagca	
>tp1 101<-120 5'-attcgcgtcgcctaatttgt	
>1-120 Amplicon size: 120bp Ta=40°C	
aaggcaagttgttaccagcagttcaaattgcgtgcgacattggctcatttcacaatcggtaa	aaaaaaaaaaaaaaaaaaaaacaaactcatggctccgtttacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq6:	
>ufp 1->20	
5'-aaggcaagttgttaccagca	
>tp1 111<-130	
5'-attcgcgtcgcctaatttgt	
>1-130 Amplicon size: 130bp Ta=40°C aaggcaagttgttaccagcagttcaaattgcgtgcgacatataaatgacctgccgtgcaaa	aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaccaatcaacgacgaacga
In silico Primer(s) search for: ////seq8:	
>ufp 1->20	
5'-aaggcaagttgttaccagca	
>tp1 121<-140	
5'-attcgcgtcgcctaatttgt	
>1-140 Amplicon size: 140bp Ta=40°C aaggcaagttgttaccagcagttcaaattgcgtgcgacatataaatgacctgccgtgcaaa cgcgaa	aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq10:	
In silico Primer(s) search for: ////seq12:	
In silico Primer(s) search for: ////seq14:	
In silico Primer(s) search for: ////seq16:	
Then,	
>>> 2 <sup>nd</sup> PCR round >>>	
In silico Primer(s) search for: ////seq3:	
In silico Primer(s) search for: ////seq5:	
In silico Primer(s) search for: ////seq7:	
>rp2 41->60	
5'-ataaatgacctgccgtgcaa	
>cp2 96<-115 5'-aaacggagccatgagtttgt	
J uuu-oppagelaigagiiigi	
>41-115 Amplicon size: 75bp Ta=40°C ataaatgacctgccgtgcaaaaaaaaaaaaaaaaaaaaa	aaactcatggctccgtt
In silico Primer(s) search for: ////seq4:	

In silico Primer(s) search for: ////seq6:
In silico Primer(s) search for: ////seq8:

>rp2 41->60
5'-ataaatgacctgccgtgcaa
>cp2 101<-120
5'-aaacggagccatgagtttgt</pre>

>41-120 Amplicon size: 80bp Ta=40°C

## (9) BP+URP: specific block.

#### Data retrieval target:

The 2<sup>nd</sup> block (entry #2, 4, 6, 8, 10, 12, 14, 16)

#### Input primer list:

>BP2

GTTCAAATTGCGTGCGACAT

>URP

TGCGACCGTAATCAAACCAA

#### In silico PCR result:

In silico Primer(s) search for: ////seq1:

In silico Primer(s) search for: ////seq3:

In silico Primer(s) search for: ////seq5:

In silico Primer(s) search for: ////seq7:

In silico Primer(s) search for: ////seq9:

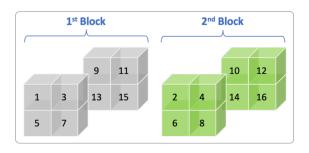
In silico Primer(s) search for: ////seq11:

In silico Primer(s) search for: ////seq13:

In silico Primer(s) search for: ////seq15:

>bp2 21->40 5'-gttcaaattgcgtgcgacat >urp 111<-130 5'-tgcgaccgtaatcaaaccaa

>21-130 Amplicon size: 110bp Ta=40°C



In silico Primer(s) search for: ////seq4: >bp2 21->40 5'-gttcaaattgcgtgcgacat >urp 121<-140 5'-tgcgaccgtaatcaaaccaa >21-140 Amplicon size: 120bp Ta=40°C In silico Primer(s) search for: ////seq6: >bp2 21->40 5'-gttcaaattgcgtgcgacat >urp 131<-150 5'-tgcgaccgtaatcaaaccaa >21-150 Amplicon size: 130bp Ta=40°C In silico Primer(s) search for: ////seq8: 21->40 >bp2 5'-gttcaaattgcgtgcgacat 141<-160 5'-tgcgaccgtaatcaaaccaa >21-160 Amplicon size: 140bp Ta=40°C gtcgc In silico Primer(s) search for: ////seq10: >bp2 21->40 5'-gttcaaattgcgtgcgacat 151<-170 >urp 5'-tgcgaccgtaatcaaaccaa >21-170 Amplicon size: 150bp Ta=41°C gtttgattacggtcgc In silico Primer(s) search for: ////seq12: >bp2 21->40 5'-gttcaaattgcgtgcgacat 161<-180 >urp 5'-tgcgaccgtaatcaaaccaa >21-180 Amplicon size: 160bp Ta=41°C ccagtttttggtttgattacggtcgc In silico Primer(s) search for: ////seq14: >bp2 21->40 5'-gttcaaattgcgtgcgacat 171<-190

5'-tgcgaccgtaatcaaaccaa

>21-190 Amplicon size: 170bp Ta=41°C

at a atttgccgcctccagtttttggtttgattacggtcgc

In silico Primer(s) search for: ////seq16:

>bp2 21->40  $5'\mbox{-} gttcaaattgcgtgcgacat$ >urp 181<-200 5'-tgcgaccgtaatcaaaccaa

>21-200 Amplicon size: 180bp Ta=41°C

atggctccgttttaatttgccgcctccagtttttggtttgattacggtcgc

### (10) BP+TP: specific table from specific block.

#### Data retrieval target:

The 1<sup>st</sup> table from the 2<sup>nd</sup> block (entry #2, 4, 6, 8)

#### Input primer list:

>BP2

GTTCAAATTGCGTGCGACAT

#### >TP1

ATTCGCGTCGCCTAATTTGT

#### In silico PCR result:

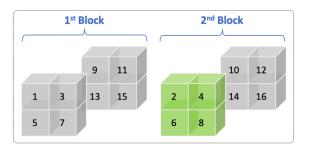
In silico Primer(s) search for: ////seq1: In silico Primer(s) search for: ////seq3: In silico Primer(s) search for: ////seq5: In silico Primer(s) search for: ////seq7: In silico Primer(s) search for: ////seq2:

>bp2 21->40 5'-gttcaaattgcgtgcgacat >tp1 91<-110 5'-attcgcgtcgcctaatttgt

>21-110 Amplicon size: 90bp Ta=41°C

In silico Primer(s) search for: ////seq4:

21->40 5'-gttcaaattgcgtgcgacat >tp1 101<-120 5'-attcgcgtcgcctaatttgt



>21-120 Amplicon size: 100bp Ta=41°C gttcaaattgcgtgcgacattggctcatttcacaatcggtaaaaaaaa	acaaactcatggctccgtttacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq6:	
>bp2 21->40 5'-gttcaaattgcgtgcgacat >tp1 111<-130 5'-attcgcgtcgcctaatttgt	
>21-130 Amplicon size: 110bp Ta=41°C gttcaaattgcgtgcgacatataaatgacctgccgtgcaaaaaaaa	naaaaaaaaaaaccaatcaacgacgaacgaaacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq8:	
>bp2 21->40 5'-gttcaaattgcgtgcgacat >tp1 121<-140 5'-attcgcgtcgcctaatttgt	
>21-140 Amplicon size: 120bp Ta=41°C gttcaaattgcgtgcgacatataaatgacctgccgtgcaaaaaaaa	naaaaaaaaaaaaaaaaaaaaaaaaaaaactcatggctccgtttacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq10:	
In silico Primer(s) search for: ////seq12:	
In silico Primer(s) search for: ////seq14:	
In silico Primer(s) search for: ////seq16:	

## (11) BP+CP: specific column from all tables in specific block.

#### Data retrieval target:

The 2<sup>nd</sup> column from all tables in the 1<sup>st</sup> block (entry #3, 7, 11, 15)

#### Input primer list:

>BP1

AGCCGACAAGTTCAAACACA

>CP2

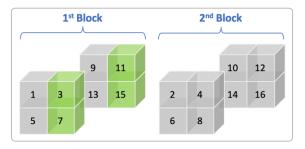
AAACGGAGCCATGAGTTTGT

#### In silico PCR result:

In silico Primer(s) search for: ////seq1:

In silico Primer(s) search for: ////seq3:

21->40 5'-agccgacaagttcaaacaca >cp2 76<-95 5'-aaacggagccatgagtttgt



>21-95 Amplicon size: 75bp Ta=40°C agccgacaagttcaaacacatggctcatttcacaatcggtaaaaaaaa	caaactcatggctccgtt
In silico Primer(s) search for: ////seq5:	-
In silico Primer(s) search for: ////seq7:	-
>bp1 21->40	
5'-agccgacaagttcaaacaca >cp2 96<-115	
>cp2 96<-115 5'-aaacggagccatgagtttgt	
>21-115 Amplicon size: 95bp Ta=40°C agccgacaagttcaaacacaataaatgacctgccgtgcaaaaaaaa	aaaaaaaaaaaaaaaaaaaaaacaaactcatggctccgtt
In silico Primer(s) search for: ////seq9:	-
In silico Primer(s) search for: ////seq11:	-
>bp1 21->40	
5'-agccgacaagttcaaacaca	
>cp2 116<-135 5'-aaacggagccatgagtttgt	
J-aaalggagccatgagtttgt	
>21-135 Amplicon size: 115bp Ta=40°C agccgacaagttcaaacacatggctcatttcacaatcggtaaaaaaaa	aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq13:	-
In silico Primer(s) search for: ////seq15:	-
>bp1 21->40	
5'-agccgacaagttcaaacaca	
>cp2 136<-155 5'-aaacggagccatgagtttgt	
>21-155 Amplicon size: 135bp Ta=40°C	
agccgacaagttcaaacacaataaatgacctgccgtgcaaaaaaaa	аааааааааааааааааааааааааааааааааааааа
In silico Primer(s) search for: ////seq4:	-
In silico Primer(s) search for: ////seq8:	-
In silico Primer(s) search for: ////seq12:	-
In silico Primer(s) search for: ////seq16:	-

## (12) BP+URP then RP+URP: specific row from all tables in specific block.

Data retrieval target: The  $1^{st}$  row from all tables in the  $2^{nd}$  block (entry #2, 4, 10, 12)

1st Block 2<sup>nd</sup> Block >BP2 GTTCAAATTGCGTGCGACAT 9 10 11 12 >URP TGCGACCGTAATCAAACCAA 13 15 14 16 5 7 6 8 Then, >RP1 TGGCTCATTTCACAATCGGT >URP TGCGACCGTAATCAAACCAA In silico PCR result: >>> 1st PCR round >>> In silico Primer(s) search for: ////seq1: In silico Primer(s) search for: ////seq3: In silico Primer(s) search for: ////seq5: In silico Primer(s) search for: ////seq7: In silico Primer(s) search for: ////seq9: In silico Primer(s) search for: ////seq11: In silico Primer(s) search for: ////seq13: In silico Primer(s) search for: ////seq15: In silico Primer(s) search for: ////seq2: >bp2 21->40 5'-gttcaaattgcgtgcgacat 111<-130 >urp 5'-tgcgaccgtaatcaaaccaa >21-130 Amplicon size: 110bp Ta=40°C In silico Primer(s) search for: ////seq4: >bp2 21->40 5'-gttcaaattgcgtgcgacat 121<-140 >urp 5'-tgcgaccgtaatcaaaccaa >21-140 Amplicon size: 120bp Ta=40°C

Input primer list:

In silico Primer(s) search for: ////seq6: >bp2 21->40 5'-gttcaaattgcgtgcgacat >urp 131<-150 5'-tgcgaccgtaatcaaaccaa >21-150 Amplicon size: 130bp Ta=40°C In silico Primer(s) search for: ////seq8: >bp2 21->40 5'-gttcaaattgcgtgcgacat >urp 141<-160 5'-tgcgaccgtaatcaaaccaa >21-160 Amplicon size: 140bp Ta=40°C gtcgc In silico Primer(s) search for: ////seq10: >bp2 21->40 5'-gttcaaattgcgtgcgacat >urp 151<-170 5'-tgcgaccgtaatcaaaccaa >21-170 Amplicon size: 150bp Ta=41°C gtttgattacggtcgc In silico Primer(s) search for: ////seq12: >bp2 21->40 5'-gttcaaattgcgtgcgacat 161<-180 5'-tgcgaccgtaatcaaaccaa >21-180 Amplicon size: 160bp Ta=41°C  ${\tt ccagtttttggtttgattacggtcgc}$ In silico Primer(s) search for: ////seq14: >bp2 21->40 5'-gttcaaattgcgtgcgacat 171<-190 >urp 5'-tgcgaccgtaatcaaaccaa >21-190 Amplicon size: 170bp Ta=41°C at a atttgccgcctccagtttttggtttgattacggtcgcIn silico Primer(s) search for: ////seq16: >bp2 21->40 5'-gttcaaattgcgtgcgacat 181<-200 >urp

5'-tgcgaccgtaatcaaaccaa	
>21-200 Amplicon size: 180bp Ta=41°C gttcaaattgcgtgcgacatataaatgacctgccgtgcaaaaaaaa	наазаазаазаазаазаазаазаазаазаазаазаазааз
Then,	
>>> 2 <sup>nd</sup> PCR round >>>	
In silico Primer(s) search for: ////seq2:	
>rp1 21->40 5'-tggctcatttcacaatcggt >urp 91<-109 5'-tgcgaccgtaatcaaaccaa	
>21-110 Amplicon size: 90bp Ta=40°C tggctcatttcacaatcggtaaaaaaaaaaaaccaatcaacgacgaacga	ggcgacgcgaatttggtttgattacggtcgc
In silico Primer(s) search for: ////seq4:	
>rp1 21->40 5'-tggctcatttcacaatcggt >urp 101<-119 5'-tgcgaccgtaatcaaaccaa	
>21-120 Amplicon size: 100bp Ta=40°C tggctcatttcacaatcggtaaaaaaaaaaaaaaaaaaa	ttacaaattaggcgacgcgaatttggtttgattacggtcgc
In silico Primer(s) search for: ////seq6:	
In silico Primer(s) search for: ////seq8:	
In silico Primer(s) search for: ////seq10:	
>rp1 21->40 5'-tggctcatttcacaatcggt >urp 131<-149 5'-tgcgaccgtaatcaaaccaa	
>21-150 Amplicon size: 130bp Ta=40°C tggctcatttcacaatcggtaaaaaaaaaaaaaaaaaaa	aaaaaaaaaaaaaaccaatcaacgacgaacgaataatttgccgcctccagtttttggtttgattacggtcgc
In silico Primer(s) search for: ////seq12:	
>rp1 21->40 5'-tggctcatttcacaatcggt >urp 141<-159 5'-tgcgaccgtaatcaaaccaa	
>21-160 Amplicon size: 140bp Ta=40°C tggctcatttcacaatcggtaaaaaaaaaaaaaaaaaaa	aaaaaaaaaaaaaaaaaaaaaaaaaaaactcatggctccgttttaatttgccgcctccagtttttggtttgattacg
In silico Primer(s) search for: ////seq14:	

In silico Primer(s) search for: ////seq16:

## (13) BP+TP then RP+TP: specific row from specific table in specific block.

#### Data retrieval target:

The 2<sup>nd</sup> row from the 2<sup>nd</sup> table in the 1<sup>st</sup> block (entry #13, 15)

#### Input primer list:

>BP1

AGCCGACAAGTTCAAACACA

>TP2

AAACTGGAGGCGGCAAATTA

Then,

>RP2

ATAAATGACCTGCCGTGCAA

AAACTGGAGGCGGCAAATTA

#### In silico PCR result:

#### >>> 1st PCR round >>>

In silico Primer(s) search for: ////seq1:
In silico Primer(s) search for: ////seq3:
In silico Primer(s) search for: ////seq5:
In silico Primer(s) search for: ////seq7:
In silico Primer(s) search for: ////seq9:
>bp1 21->40
5'-agccgacaagttcaaacaca

126<-145 >tp2

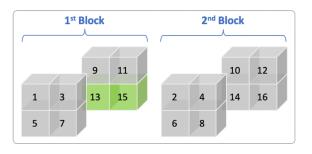
5'-aaactggaggcggcaaatta

>21-145 Amplicon size: 125bp Ta=40°C

In silico Primer(s) search for: ////seq11:

>bp1 21->40 5'-agccgacaagttcaaacaca 136<-155 >tp2 5'-aaactggaggcggcaaatta

>21-155 Amplicon size: 135bp Ta=40°C



In silico Primer(s) search for: ////seq13: >bp1 21->40 5'-agccgacaagttcaaacaca >tp2 146<-165 5'-aaactggaggcggcaaatta >21-165 Amplicon size: 145bp Ta=40°C tttgccgcctccagtt In silico Primer(s) search for: ////seq15: >bp1 21->40 5'-agccgacaagttcaaacaca 156<-175 >tp2 5'-aaactggaggcggcaaatta >21-175 Amplicon size: 155bp Ta=41°C tccgttttaatttgccgcctccagtt In silico Primer(s) search for: ////seq10: In silico Primer(s) search for: ////seq12: In silico Primer(s) search for: ////seq14: In silico Primer(s) search for: ////seq16: Then, >>> 2nd PCR round >>> In silico Primer(s) search for: ////seq9: In silico Primer(s) search for: ////seq11: In silico Primer(s) search for: ////seq13: >rp2 21->40 5'-ataaatgacctgccgtgcaa 126<-144 5'-aaactggaggcggcaaatta >21-145 Amplicon size: 125bp Ta=40°C In silico Primer(s) search for: ////seq15: 21->40 >rp2 5'-ataaatgacctgccgtgcaa 136<-154 >tp2 5'-aaactggaggcggcaaatta

>21-155 Amplicon size: 135bp Ta=40°C

## (14) BP+TP then BP+CP: specific column from specific table in specific block.

#### Data retrieval target:

The 1<sup>st</sup> column from the 1<sup>st</sup> table in the 2<sup>nd</sup> block (entry #2, 6)

### Input primer list:

>BP2

GTTCAAATTGCGTGCGACAT

>TP1

ATTCGCGTCGCCTAATTTGT

Then,

>BP2

GTTCAAATTGCGTGCGACAT

>CP1

>tp1

101<-120 5'-attcgcgtcgcctaatttgt

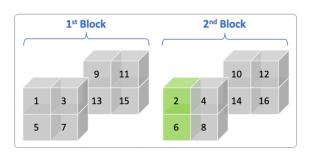
>21-120 Amplicon size: 100bp Ta=41°C

TTCGTTCGTCGTTGATTGGT

#### In silico PCR result:

#### >>> 1st DCP round >>>

>>> 1st PCR round >>>	
In silico Primer(s) search for: ////seq1:	
In silico Primer(s) search for: ////seq3:	
In silico Primer(s) search for: ////seq5:	
In silico Primer(s) search for: ////seq7:	
In silico Primer(s) search for: ////seq2:	
>bp2 21->40 5'-gttcaaattgcgtgcgacat >tp1 91<-110 5'-attcgcgtcgcctaatttgt	
>21-110 Amplicon size: 90bp Ta=41°C gttcaaattgcgtgcgacattggctcatttcacaatcggtaaaaaaaa	acgaaacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq4:	
>bp2 21->40 5'-gttcaaattgcgtgcgacat	



S29

In silico Primer(s) search for: ////seq6:	
>bp2 21->40	
5'-gttcaaattgcgtgcgacat	
>tp1 111<-130	
5'-attcgcgtcgcctaatttgt	
>21-130 Amplicon size: 110bp Ta=41°C gttcaaattgcgtgcgacatataaatgacctgccgtgcaaaaaaaa	aaaaaaaaaaaaaaccaatcaacgacgaacgaaacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq8:	
>bp2 21->40	
5'-gttcaaattgcgtgcgacat	
>tp1 121<-140	
5'-attcgcgtcgcctaatttgt	
>21-140 Amplicon size: 120bp Ta=41°C gttcaaattgcgtgcgacatataaatgacctgccgtgcaaaaaaaa	aaaaaaaaaaaaaaaaaaaaaaaaaaactcatggctccgtttacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq10:	
In silico Primer(s) search for: ////seq12:	
In silico Primer(s) search for: ////seq14:	
In silico Primer(s) search for: ////seq16:	
Then,	
>>> 2 <sup>nd</sup> PCR round >>>	
In silico Primer(s) search for: ////seq2:	
>bp2 1->20	
5'-gttcaaattgcgtgcgacat	
>cp1 51<-70	
5'-ttcgttcgtcgttgattggt	
>1-70 Amplicon size: 70bp Ta=40°C	
gttcaaattgcgtgcgacattggctcatttcacaatcggtaaaaaaaa	ocgacgaacga
In silico Primer(s) search for: ////seq4:	
In silico Primer(s) search for: ////seq6:	
>bp2 1->20	
5'-gttcaaattgcgtgcgacat	
>cp1 71<-90	
5'-ttcgttcgtcgttgattggt	
>1-90 Amplicon size: 90bp Ta=40°C gttcaaattgcgtgcgacatataaatgacctgccgtgcaaaaaaaa	aaaaaaaaaaaaaccaatcaacgacgaacga

In silico Primer(s) search for: ////seq8:

## (15) BP+URP then RP+CP: specific entry from all tables in specific block.

>21-130 Amplicon size: 110bp Ta=40°C

In silico Primer(s) search for: ////seq4:

Data retrieval target: Entry at the 2 <sup>nd</sup> row 2 <sup>nd</sup> column from all tables in	the 2 <sup>nd</sup> block (entry	y #8, 16)				
Input primer list: >BP2		1 <sup>st</sup> Block			2 <sup>nd</sup> Block	
GTTCAAATTGCGTGCGACAT						
>URP TGCGACCGTAATCAAACCAA	1	3 13	15	2	10	12
Then,	5	7		6	8	
>RP2 ATAAATGACCTGCCGTGCAA						
>CP2 AAACGGAGCCATGAGTTTGT						
In silico PCR result:						
>>> 1st PCR round >>>						
In silico Primer(s) search for: ////seq1:	_					
In silico Primer(s) search for: ////seq3:	_					
In silico Primer(s) search for: ////seq5:	_					
In silico Primer(s) search for: ////seq7:	_					
In silico Primer(s) search for: ////seq9:	_					
In silico Primer(s) search for: ////seq11:	_					
In silico Primer(s) search for: ////seq13:	_					
In silico Primer(s) search for: ////seq15:	_					
In silico Primer(s) search for: ////seq2:	_					
>bp2 21->40 5'-gttcaaattgcgtgcgacat >urp 111<-130						

21->40 5'-gttcaaattgcgtgcgacat 121<-140 >urp 5'-tgcgaccgtaatcaaaccaa >21-140 Amplicon size: 120bp Ta=40°C In silico Primer(s) search for: ////seq6: >bp2 21->40 5'-gttcaaattgcgtgcgacat 131<-150 5'-tgcgaccgtaatcaaaccaa >21-150 Amplicon size: 130bp Ta=40°C In silico Primer(s) search for: ////seq8: >bp2 21->40 5'-gttcaaattgcgtgcgacat >urp 141<-160 5'-tgcgaccgtaatcaaaccaa >21-160 Amplicon size: 140bp Ta=40°C In silico Primer(s) search for: ////seq10: >bp2 21->40 5'-gttcaaattgcgtgcgacat >urp 151<-170 5'-tgcgaccgtaatcaaaccaa >21-170 Amplicon size: 150bp Ta=41°C gtttgattacggtcgc In silico Primer(s) search for: ////seq12: 21->40 >bp2 5'-gttcaaattgcgtgcgacat 161<-180 >urp 5'-tgcgaccgtaatcaaaccaa >21-180 Amplicon size: 160bp Ta=41°C ccagtttttggtttgattacggtcgc In silico Primer(s) search for: ////seq14: >bp2 21->40 5'-gttcaaattgcgtgcgacat 171<-190 5'-tgcgaccgtaatcaaaccaa

>21-190 Amplicon size: 170bp Ta=41°C

gttcaaattgcgtgcgacatataaatgacctgccgtgcaaaaaaaa	aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa
In silico Primer(s) search for: ////seq16:	
>bp2 21->40 5'-gttcaaattgcgtgcgacat >urp 181<-200 5'-tgcgaccgtaatcaaaccaa	
>21-200 Amplicon size: 180bp Ta=41°C gttcaaattgcgtgcgacatataaatgacctgccgtgcaaaaaaaa	паавааваавааваавааваавааваавааваавааваав
Then,	
>>> 2 <sup>nd</sup> PCR round >>>	
In silico Primer(s) search for: ////seq4:	
In silico Primer(s) search for: ////seq6:	
In silico Primer(s) search for: ////seq8:	
>rp2 21->40 5'-ataaatgacctgccgtgcaa >cp2 81<-100 5'-aaacggagccatgagtttgt	
>21-100 Amplicon size: 80bp Ta=40°C ataaatgacctgccgtgcaaaaaaaaaaaaaaaaaaaaa	aacaaactcatggctccgtt
In silico Primer(s) search for: ////seq12:	
In silico Primer(s) search for: ////seq14:	
In silico Primer(s) search for: ////seq16:	
>rp2 21->40 5'-ataaatgacctgccgtgcaa >cp2 121<-140 5'-aaacggagccatgagtttgt	
>21-140 Amplicon size: 120bp Ta=40°C ataaatgacctgccgtgcaaaaaaaaaaaaaaaaaaaaa	aaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaaa
(16) BP+TP then RP+CP: specific entry f	rom specific table in specific block.
Data retrieval target: Entry at the $2^{nd}$ row $1^{st}$ column from the $1^{st}$ table in th	e 2 <sup>nd</sup> block (entry #6)
Input primer list: >BP2	
GTTCAAATTGCGTGCGACAT	

>TP1

#### ATTCGCGTCGCCTAATTTGT 2<sup>nd</sup> Block 1st Block Then, >RP2 9 11 10 12 ATAAATGACCTGCCGTGCAA 13 15 14 16 >CP1 5 7 6 8 TTCGTTCGTCGTTGATTGGT In silico PCR result: >>> 1st PCR round >>> In silico Primer(s) search for: ////seq1: In silico Primer(s) search for: ////seq3: In silico Primer(s) search for: ////seq5: In silico Primer(s) search for: ////seq7: In silico Primer(s) search for: ////seq2: >bp2 21->40 5'-gttcaaattgcgtgcgacat >tp1 91<-110 5'-attcgcgtcgcctaatttgt >21-110 Amplicon size: 90bp Ta=41°C In silico Primer(s) search for: ////seq4: >bp2 21->40 5'-gttcaaattgcgtgcgacat 101<-120 5'-attcgcgtcgcctaatttgt >21-120 Amplicon size: 100bp Ta=41°C In silico Primer(s) search for: ////seq6: 21->40 >bp2 5'-gttcaaattgcgtgcgacat 111<-130 >tp1 5'-attcgcgtcgcctaatttgt >21-130 Amplicon size: 110bp Ta=41°C In silico Primer(s) search for: ////seq8:

>bp2 21->40 5'-gttcaaattgcgtgcgacat

121<-140

>tp1

S34

5'-attcgcgtcgcctaatttgt	
>21-140 Amplicon size: 120bp Ta=41°C gttcaaattgcgtgcgacatataaatgacctgccgtgcaaaaaaaa	aaaaaaaaaaaaaaaaaaaaaaaaaaacaaactcatggctccgtttacaaattaggcgacgcgaa
In silico Primer(s) search for: ////seq10:	
In silico Primer(s) search for: ////seq12:	
In silico Primer(s) search for: ////seq14:	
In silico Primer(s) search for: ////seq16:	
Then,	
>>> 2 <sup>nd</sup> PCR round >>>	
In silico Primer(s) search for: ////seq2:	
In silico Primer(s) search for: ////seq6:	
>rp2 21->40 5'-ataaatgacctgccgtgcaa >cp1 71<-90 5'-ttcgttcgtcgttgattggt >21-90 Amplicon size: 70bp Ta=40°C	
ataaatgacctgccgtgcaaaaaaaaaaaaaaaaaaaaa	tcaacgacgaacga

In silico Primer(s) search for: ////seq8: