

Genomic mapping of transformed DNA fragments

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Working

Protist Research to Optimize Tools in Genetics (PROT-G)



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ABSTRACT

Mapping of transformed DNA within mutated diatoms. A stepwise specific PCR reaction that allow for identification of genome location of transformed fragments such as antibiotic resistance genes or GFP constructs.

Degenerate primer 1: 5'-GCCGGACTTAGCGAGCA TGTGTAGTACCTTGCCG ATCCTNASATANSATANT TC - 3' Degenerate primer 2: 5'-GCCGGACTTAGCGAGCA TGTGTAGTACCTTGCCG ATCCCNTSABGNACYTN CTG - 3' Degenerate primer 3: 5' - GCCGGACTTAGCGAGCA TGTGTAGTACCTTGCCG ATCCNGACGARWGANA WGAC - 3' Degenerate primer 4: 5'-GCCGGACTTAGCGAGCA TGTGTAGTACCTTGCCG ATCCTAHATGDAGKACN TAC - 3'

Specific primer 1: 5' - GCCGGACTTAGCGAGCA TGTGTAG - 3' Specific primer 2: 5' - CATGTGTAGTACCTTGC CGATCC - 3'

Internal primers must be designed with regard to each construct that are to be mapped. They should not overlap but instead cover consecutive stretches of known DNA sequence of the transformed construct.

PROTOCOL STATUS

Working

We use this protocol in our group and it is working

MATERIALS

CATALOG # **VENDOR** NAME

Gel electrophoresis equipment

View Thermo Fisher Scientific SuperFi Polymerase

SAFETY WARNINGS

BEFORE STARTING

Make sure to have good quality DNA

Primary, 25uL PCR reaction

Vol. Component

5uL 5x SuperFi Buffer dNTP - Mix (10mM Each) 0.8uL 3uL Internal Primer (10uL)

0.2uL SuperFi 12.5uL Water

2.5uL Degenerate Primer (40uM)



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Primary PCR program

Step	Temp	Time		
1	96	5 min		
2	98	10 sec		
3	70	15 sec		
4	72	3 min		
5 Go	5 Goto 2 - 30 more times			
6	98	10 sec		
7	25	1 min		
8	8 Ramp to 72 0.5C per sec			
9	72	3 min		
10 G	oto 6 - 1	more time		
11	98	10 sec		
12	70	15 sec		
13	72	3 min		
14	98	10 sec		
15	70	15 sec		
16	72	3 min		
17	98	10 sec		
18	70	15 sec		
19	72	3 min		
20	98	10 sec		
21	40	45 sec		
22	72	3 min		
23 G	oto 11 -	10 More times		
24	72	5min		
25	4	Forever		

2 Dilute Primary PCR reaction 40x times (5uL + 195uL water)

3 Secondary, 25uL PCR reaction

Vol.Component5uL5x SuperFi Buffer0.5uLdNTP - Mix (10mM Each)1.5uLInternal Primer 2 (10uM)0.2uLSuperFi1.5uLSpecific primer 1 (10uM)15.3uLWater1uLDiluted primary PCR reaction

Secondary PCR program

Step	Temp	Time
1	98	30 sec
2	98	10 sec
3	70	15 sec
4	72	3 min
5	98	10 sec
6	70	15 sec
7	72	3 min
8	98	10 sec
9	57	30 sec
10	72	3 min
11	Goto 2	- 12 More times
12	72	5min
13	4	Forever

4 Dilute Secondary PCR reaction 40x times (5uL + 195uL water)

5 Tertiary, 50uL PCR reaction

Vol. Component

10uL 5x SuperFi Buffer

0.8uL dNTP - Mix (10mM Each)

2uL Internal Primer 3 (10uM)

0.3uL SuperFi

2uL Specific primer 2 (10uM)

25.4uL Water

2uL Diluted Secondary reaction

Tertiary PCR program

Ste	p Temp	Time
1	98	30 sec
2	98	10 sec
3	57	30 sec
4	72	3 min
5	Goto 2 - 14	More times
6	72	5min
7	4	Forever

- Run a 1% agarose gel, excise bands and cleanup using a gel cleanup kit (Ex. Promega Wizard)
- 7 Clone 2uL of purified bands into Blunt TOPO vector (Thermo Fisher Scientific) and transform competent E.coli dH5a (Thermo Fisher Scientific) by 30 sec heat shock at 42C. Grow 5mL overnight cultures of picked colonies and extract vectors.
- 8 Sanger sequence using the TOPO vectors internal primers.

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