## **Primer3 Output**

```
PRIMER PICKING RESULTS FOR Puccinia recondita consensus consensus group 0
Template masking not selected
No mispriming library specified
No internal oligo mishyb library specified
Using 1-based sequence positions
OLIG0
                start len
                               tm
                                     qc%
                                          any th 3' th hairpin seq
                   80
                       23
                            57.97
                                    43.48
                                            0.00
LEFT PRIMER
                                                   0.00
                                                           0.00 tctqqcatcaaaaqatactcctc
RIGHT PRIMER
                  273
                       20
                            57.91
                                    50.00
                                           10.34
                                                   0.00
                                                           0.00 gcaattgactccgtaccact
INTERNAL OLIGO
                  163
                       21
                            58.93
                                    52.38
                                            0.00
                                                   0.00
                                                          46.23 ggtctcaaccaggaagaatcc
SEQUENCE SIZE: 349
INCLUDED REGION SIZE: 349
PRODUCT SIZE: 194, PAIR ANY TH COMPL: 0.00, PAIR 3' TH COMPL: 0.00
   1 ttgctaacgattttccggccaccttatggttcttcaaggatccttttatacagtatgtta
  61 gatatcaaggaaaatcgattctggcatcaaaagatactcctcttctgatgaataagtgga
                       >>>>>>>>>>>>
 ^^^^^
  181 tccatataaaccaattatccaagcattcctttgatttttttgggttatctttcaagtatac
 241 gaccgaatatttcagtggtacggagtcaattgctagaaaattcgtttttaatggataatg
                  <<<<<<<<
  301 ctatgaagaagcttgataNattatttccaattaNtcNaatgataggatc
KEYS (in order of precedence):
>>>>> left primer
<<<<< right primer
^^^^^ internal oligo
ADDITIONAL OLIGOS
                   start len _____tm ___gc% _any th _3' th hairpin seq
 1 LEFT PRIMER
                          22
                               58.11
                                       40.91
                     68
                                              11.83
                                                      0.00
                                                              0.00 aggaaaatcgattctggcatca
  RIGHT PRIMER
                     266
                          23
                               58.32
                                       43.48
                                               0.00
                                                      0.00
                                                              0.00 actccgtaccactgaaatattcg
                               58.93
                                       52.38
  INTERNAL OLIGO
                     163
                          21
                                               0.00
                                                      0.00
                                                             46.23 ggtctcaaccaggaagaatcc
  PRODUCT SIZE: 199, PAIR ANY_TH COMPL: 0.00, PAIR 3'_TH COMPL: 0.00
 2 LEFT PRIMER
                                               0.00
                      75
                          23
                               57.01
                                       39.13
                                                      0.00
                                                              0.00 tcgattctggcatcaaaagatac
```

	RIGHT PRIMER		272	22	58.34	58.34 4		0.00	0.00	0.0	0 caa	ttgacto	cgtacc	actgaa		
	INTERNAL OLIGO		163	21	58.93	3 5	2.38	0.00	0.00	46.2	3 ggt	ctcaaco	aggaag	aatcc		
	PR0DUCT	SIZ	ZE: 198	, PAII	R ANY_	_TH COME	PL: 0	.00,	PAIR 3'	_TH CON	1PL: 0.0	0				
3	3 LEFT PRIMER			72	23	57.10	3	4.78	11.83	0.00	0.0	0 aaa	tcgattc	tggcat	caaaaga	а
	RIGHT PRIMER			269	23	57.33	3	9.13	0.00	0.00	0.0	00 ttga	) ttgactccgtaccactgaaatat			
	INTERNAL OLIGO			163	21	58.93	3 5	2.38	0.00	0.00	46.2	3 ggt	3 ggtctcaaccaggaagaatcc			
	PRODUCT SIZE: 198, PAIR ANY_TH COMPL: 0.00, PAIR 3'_TH COMPL: 0.00															
Statistics																
	C	on	too	in	in	not		n	o tm	tm	high	high	high		high	
	S	id	many	tar	excl	ok	bad	G	C too	too	any_th	3'_th	hair-	poly	end	
	er	ed	Ns	get	reg	reg	GC%	clam	p low	high	compl	compl	pin	Χ	stab	ok
Lef	t 10	48	0	0	0	0	314		0 575	16	0	0	0	29	0	114
Rig	ht 9:	11	36	0	0	0	286		0 498	0	0	0	32	29	0	30
Int	l 28	43	31	0	0	0	20		0 1822	123	0	0	6	22	0	819
Pai	r Stats	:														

considered 2347, unacceptable product size 2341, primer in pair overlaps a primer in a better pair 131, ok 6 libprimer3 release 2.4.0

(primer3\_results.cgi release 4.1.0)