## **Primer3 Output**

```
PRIMER PICKING RESULTS FOR Puccinia recondita elongation factor 1 alpha elongation factor 1 consensus group 4
Template masking not selected
No mispriming library specified
No internal oligo mishyb library specified
Using 1-based sequence positions
0LIG0
                 start len
                                 tm
                                        qc%
                                            any th 3' th hairpin seq
                    25
                         20
                              59.24
                                      50.00
                                               0.00
                                                      0.00
                                                              0.00 tgtctccgttgaagccagaa
LEFT PRIMER
RIGHT PRIMER
                   206
                         20
                              58.11
                                      45.00
                                               0.00
                                                      0.00
                                                              0.00 tgcttgcgttcacattaggt
INTERNAL OLIGO
                   138
                         20
                              59.74
                                      50.00
                                               0.00
                                                      0.00
                                                             36.37 catttggccgtgtccatctt
SEQUENCE SIZE: 312
INCLUDED REGION SIZE: 312
PRODUCT SIZE: 182, PAIR ANY TH COMPL: 0.00, PAIR 3' TH COMPL: 1.53
   1 agttNNcggaattatcaatcatattgtctccgttgaagccagaaattggcacaaatggaa
                              >>>>>>>>>
  61 caaccttqqqqttaaatccgactttctttatNaagttggtcgtctccttgatNatttcaa
  121 ggtatctNgcttcagcccatttggccgtgtccatcttNttaatggcaacaatcaattgct
                       ^^^^^^
  181 ttNgacacctaatgtgaacgcaagcaNtgcatgttctctggtttgaccatcctttgaaat
            <<<<<<<<
  241 acctgcctcNaactcaccggtaccggccgcgatgatgagaatggcacagtcagcttgaga
  301 ggttccNgtaat
KEYS (in order of precedence):
>>>>> left primer
<<<<< right primer
^^^^^ internal oligo
ADDITIONAL OLIGOS
                         <u>len</u> <u>tm</u> <u>gc% any th</u> 3' th hairpin seq
                    <u>start</u>
 1 LEFT PRIMER
                      136
                           18
                                 60.28
                                         61.11
                                                  0.00
                                                         0.00
                                                                 0.00 cccatttqqccqtqtcca
   RIGHT PRIMER
                      295
                            20
                                 59.31
                                         50.00
                                                  0.00
                                                         0.00
                                                                 0.00 agctgactgtgccattctca
                      187
                                 59.02
                                         45.00
   INTERNAL OLIGO
                            20
                                                  0.00
                                                         0.00
                                                                45.62 acctaatgtgaacgcaagca
  PRODUCT SIZE: 160, PAIR ANY_TH COMPL: 0.00, PAIR 3'_TH COMPL: 0.00
 2 LEFT PRIMER
                       64
                            20
                                 57.20
                                         50.00
                                                  0.00
                                                         0.00
                                                                 0.00 ccttggggttaaatccgact
```

RIGHT PRIMER	247	22	58.63	45.45	0.00	0.00	0.00	ggcaggtatttcaaaggatggt
INTERNAL OLIGO	138	20	59.74	50.00	0.00	0.00	36.37	catttggccgtgtccatctt
PRODUCT SIZE: 184,	PAIR	ANY	TH COMPL:	0.00,	PAIR 3'	TH COMPL	: 0.00	

## Statistics

	con	too	in	in	not		no	tm	tm	high	high	high		high	
	sid	many	tar	excl	ok	bad	GC	too	too	any_th	3'_th	hair-	poly	end	
	ered	Ns	get	reg	reg	GC%	clamp	low	high	compl	compl	pin	Χ	stab	ok
Left	544	88	0	0	0	32	0	213	42	0	0	80	0	0	89
Right	503	101	0	0	0	26	0	178	81	0	0	31	0	0	86
Intl	1502	187	0	0	0	1	0	367	425	0	0	9	0	0	513
Dain Chaha.															

Pair Stats:

considered 5190, unacceptable product size 5185, primer in pair overlaps a primer in a better pair 2431, ok 5 libprimer3 release 2.4.0

(primer3\_results.cgi release 4.1.0)