Primer3 Output

PRIMER PICKING RESULTS FOR Puccinia_recondita_consensus_consensus_group_1

Template masking not selected
No mispriming library specified
No internal oligo mishyb library specified
Using 1-based sequence positions

OLIGO	start	len	tm	gc%	any_th	3' th	hairpin	seq
LEFT PRIMER	101	20	59.04	50.00	7.63	7.63	0.00	aagtgcttcagttgccatgg
RIGHT PRIMER	296	21	59.06	47.62	0.00	0.00	0.00	caacgccttacatattcgcca
INTERNAL OLIGO	253	21	58.72	47.62	23.47	0.00	45.52	gaagggtacaagtttgaccca

SEQUENCE SIZE: 310
INCLUDED REGION SIZE: 310

PRODUCT SIZE: 196, PAIR ANY_TH COMPL: 0.00, PAIR 3'_TH COMPL: 0.00

```
    1 cggactggatNtcctNtggtcgatgccgggatgagagagttatgggctacNggttggttg


   61 cacgatcgNataagagtagttgtttcaagcttctNtgttaaagtgcttcagttgccatgg
                                            >>>>>>>>>>>>>>>>>>>>


  121 cgatgggggatgaagtaNttNtgggacacccttNtcgatgcagatttaNaaagtgatgct


  181 NttggttggcaatacNttaNtggtactNtcccNgatagccgggagtttgatNgcatagat


  241 aacccNcNgtttgaagggtacaagtttgacccaaatggcgaatatgtaaggcgttggctt
               ^^^^^^^^^^^^^^^^^^^^^  <<<<<<<<<<<<<<<<<<<<<


  301 ccagaNctct
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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	96	22	58.98	40.91	0.00	0.00	0.00	tgttaaagtgcttcagttgcca
RIGHT PRIMER	294	22	58.79	40.91	0.00	0.00	0.00	acgccttacatattcgccattt
INTERNAL OLIGO	119	19	59.79	57.89	0.00	0.00	0.00	ggcgatgggggatgaagta

PRODUCT SIZE: 199, PAIR ANY_TH COMPL: 0.00, PAIR 3'_TH COMPL: 0.00

Statistics

	con sid ered	too many Ns	in tar get	in excl reg	not ok reg	bad GC%	no GC clamp	tm too low	tm too high	high any_th compl	high 3'_th compl	high hair- pin	poly X	high end stab	ok
Left	331	121	0	0	0	3	0	74	48	0	0	18	9	0	58
Right	347	121	0	0	0	0	0	115	5	0	0	10	0	0	96
Intl	960	238	0	0	0	0	0	169	269	0	0	0	0	0	284

Pair Stats:
considered 3385, unacceptable product size 3383, primer in pair overlaps a primer in a better pair 1102, ok 2
libprimer3 release 2.4.0

(primer3_results.cgi release 4.1.0)