

# Primer3 Output

PRIMER PICKING RESULTS FOR Puccinia\_recondita\_consensus\_consensus\_group\_3

Template masking not selected  
No mispriming library specified  
No internal oligo mishyb library specified  
Using 1-based sequence positions  
NO PRIMERS FOUND - [Help](#)

SEQUENCE SIZE: 256  
INCLUDED REGION SIZE: 256

1 attgctggcaataNatggccNcggNcNcggccNNNacgctacctgtccttgctgccaccNg  
61 tNctccctcgtcgtcccagcgaaccctcctcgtcccNNNaaacccNNNcNcaagggcNcNcg  
121 aNgccaggagNNNNNNaaNNcccNccgtcccagNNctgccNNtcaNaaNcacNNgggcca  
181 cNgtggcNagNNacggactgcNacaNaaagcagNcccNaNtgNnaaggctNtgccNcNac  
241 ctgNggcNaaagccca

## Statistics

	con sid ered	too many Ns	in tar get	in excl reg	not ok reg	bad GC% clamp	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	192	90	0	0	0	62	0	0	27	0	0	0	0	0	13
Right	102	102	0	0	0	0	0	0	0	0	0	0	0	0	0
Intl	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0

Pair Stats:  
considered 0, ok 0  
libprimer3 release 2.4.0

(primer3\_results.cgi release 4.1.0)