

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
NO PRIMERS FOUND - [Help](#)

SEQUENCE SIZE: 211
INCLUDED REGION SIZE: 211

```
1 tagatttagcttctgttgaagctccttctataaatggataatcttttgataaaatagatt
61 ctcaattgcagcaaatgccgaatctNNNNNNagattcggcatttgtaaaaaaaaaaNa
121 atacttttttttNaagataaaaaccNttttaaaaataggggcggacgtaNccaagtggat
181 taagggaatggNttgtgNaNccccNtgcg
```

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	312	0	0	0	0	173	0	124	0	0	0	15	0	0	0
Right	73	52	0	0	0	0	0	13	0	0	0	0	5	0	3
Intl	814	120	0	0	0	125	0	370	24	0	0	0	6	0	169

Pair Stats:
considered 0, ok 0
libprimer3 release 2.4.0

(primer3_results.cgi release 4.1.0)