## **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: 330** 

INCLUDED REGION SIZE: 330

301 accgtgagggttcaaNtccctctaNcccca

## **Statistics**

	con	too	in	in	not		no	tm	tm					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		Χ	stab	ok
Left	686	80	0	0	0	111	0	358	11	0	0	0	0	0	126
Right	688	60	0	0	0	406	0	199	0	0	0	3	20	0	0
Intl	1802	144	0	0	0	134	0	970	84	0	0	0	19	0	451

Pair Stats:

considered 3, unacceptable product size 3, ok 0 libprimer3 release 2.4.0  $\,$ 

(primer3\_results.cgi release 4.1.0)