

# Development of a Real-Time Polymerase Chain Reaction Method to Measure Ligation Efficiency

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## SUPPLEMENTAL MATERIAL

**SUPPLEMENTARY TABLE 1 Raw Data from replicates used in the initial qPCR with oligonucleotide cohesive ends of 0%, 50%, and 100% GC content and controls.**

Sample	Replicate Number	Quantification Cycle (Cq) Value	Mean Cq Value for Sample	Standard Error for Mean Cq Value	Melt Temperature (°C)
Control A1	1	*NS	**N/A	N/A	None
	2	NS			None
	3	NS			None
Control B1	1	NS	N/A	N/A	None
	2	NS			None
	3	NS			None
Control C1	1	NS	N/A	N/A	None
	2	NS			None
	3	NS			None
Control D	1	18.5	19.4	1.0	76.0
	2	18.2			76.0
	3	21.4			75.5
0% GC Content	1	NS	N/A	N/A	None
	2	NS			None
	3	NS			None
50% GC Content	1	NS	N/A	N/A	None
	2	NS			None
	3	NS			None
100% GC Content	1	NS	N/A	N/A	None
	2	NS			None
	3	NS			None

\*NS: No Signal

\*\*N/A: Not Applicable as some replicates contained no signal

**SUPPLEMENTARY TABLE 2 Raw Data from replicates used in qPCR on modified oligonucleotide with controls.**

Sample	Replicate Number	Quantification Cycle (Cq) Value	Mean Cq Value for Sample	Standard Error for Mean Cq Value	Melt Temperature (°C)
Control A2	1	*NS	**N/A	N/A	None
	2	NS			None
	3	NS			None
Control B2	1	17.6	N/A	N/A	81.0
	2	NS			None
	3	NS			None
Control B3	1	23.8	N/A	N/A	80.5
	2	29.1			None
	3	NS			None
Control C2	1	23.0	N/A	N/A	80.5
	2	NS			None
	3	28.3			80.0
Control C3	1	NS	N/A	N/A	None
	2	NS			None
	3	26.5			80.0
Control D	1	21.8	22.0	1.0	76.0
	2	23.8			75.5
	3	20.4			75.5
1 Hour Ligated DNA Sample	1	14.2	14.3	0.0	80.5
	2	14.4			80.5
	3	14.4			80.5

\*NS: No Signal

\*\*N/A: Not Applicable as some replicates contained no signal