Benchmark Analysis NZYSupreme qPCR Green Master Mix Product No MB419







services

Stratech



HIGHLY OPTIMIZED KITS AND ENZYMES



MKT-010_V01/2022

Two Real-Time PCR Experiments

NZYSupreme qPCR Green Master Mix was benchmarked against a total of 9 market-leading green master mixes considered to be the gold-standard in qPCR Master Mixes.

Detection of b2m from mouse cDNA



A 5-fold serial dilution of cDNA reverse transcribed from total mouse liver was used as template for a real time qPCR experiment to detect the b2m housekeeping gene.

Detection of Large1 from human gDNA



A 5-fold serial dilution of human genomic DNA was used as template for a real time qPCR experiment to detect a region from the LargeI gene, which encodes a member of the N acetylglucosaminyltransferase family and is ubiquitous expressed in different human tissues.

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Detection of b2m from mouse cDNA - Amplification Plot

Excellent sensitivity and linearity in the amplification using a 5-fold serial dilution of cDNA reverse transcribed from total mouse liver.



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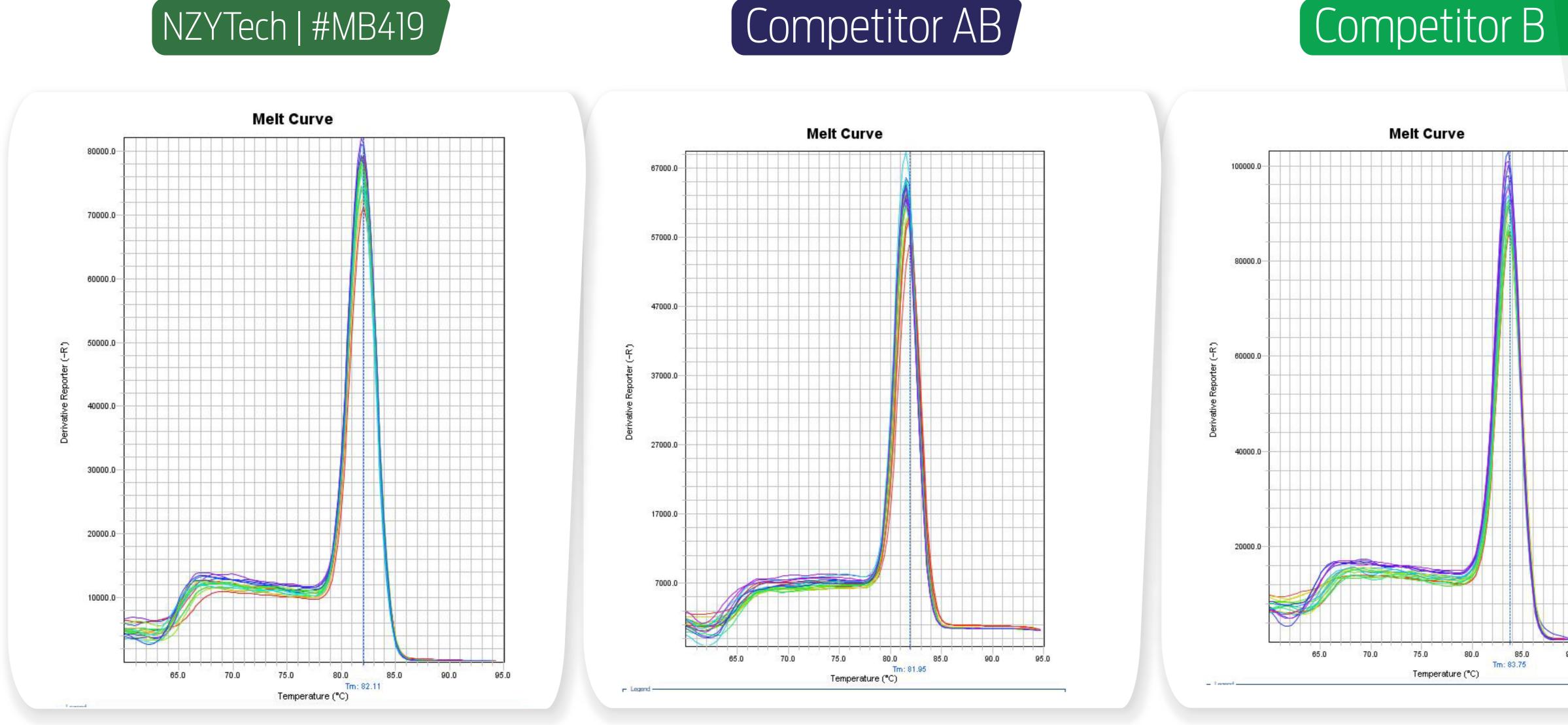
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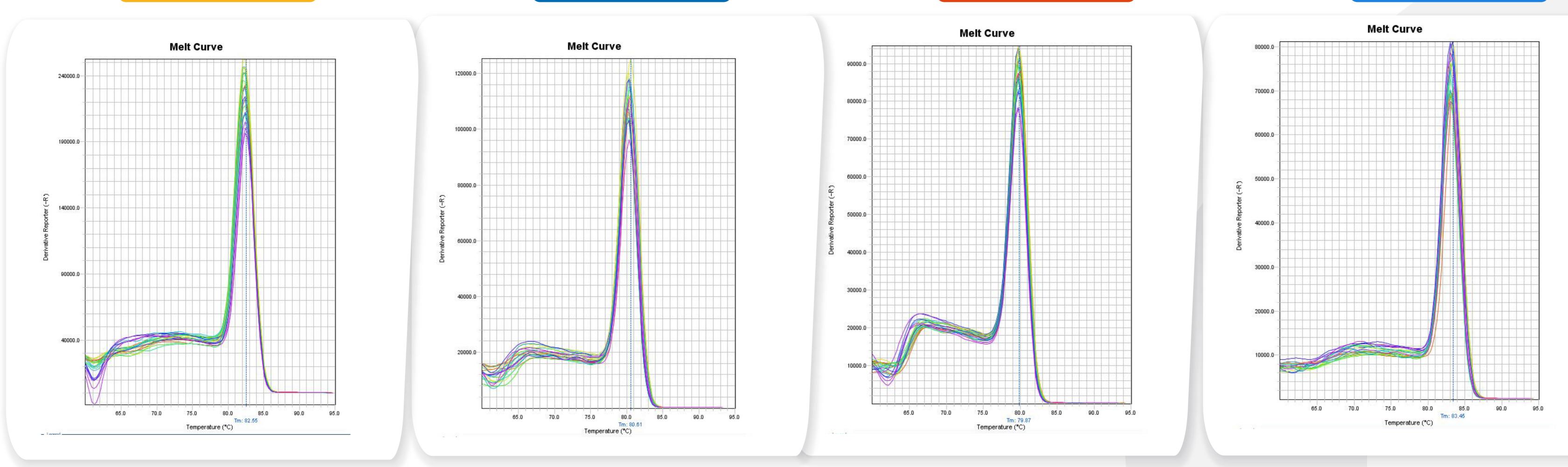


Detection of b2m from mouse cDNA - Melt Curve

The exceptional specificity of NZYSupreme qPCR Green Master Mix (MB419) is evidenced by the single specific melting peak observed for all dilution series.





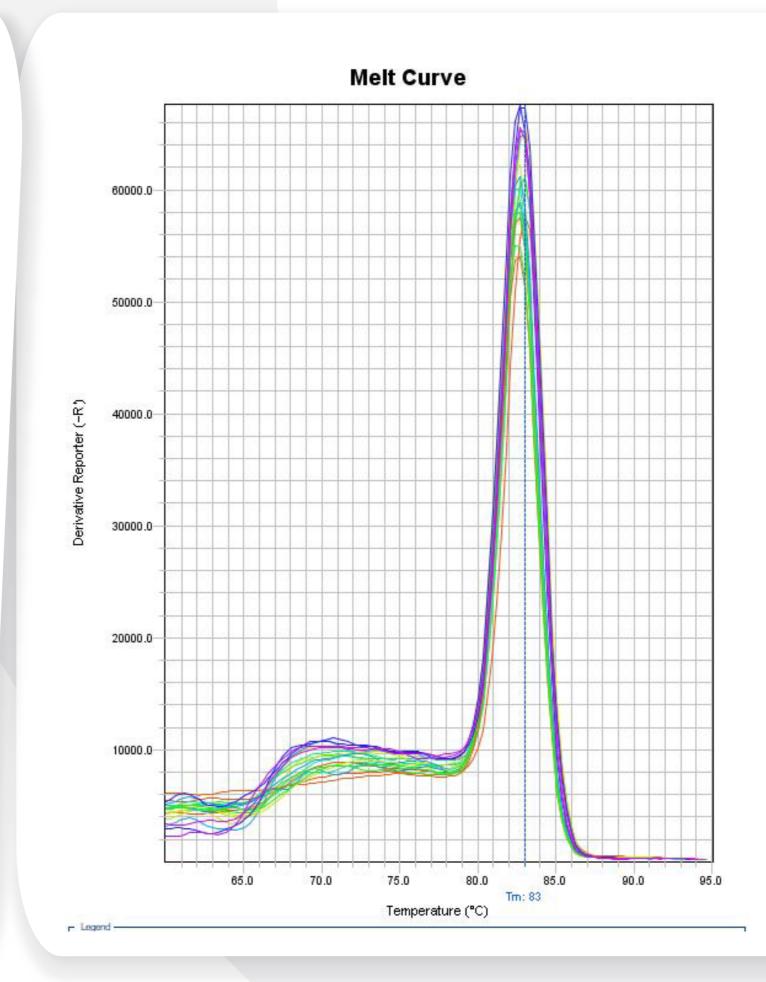


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Competitor Q



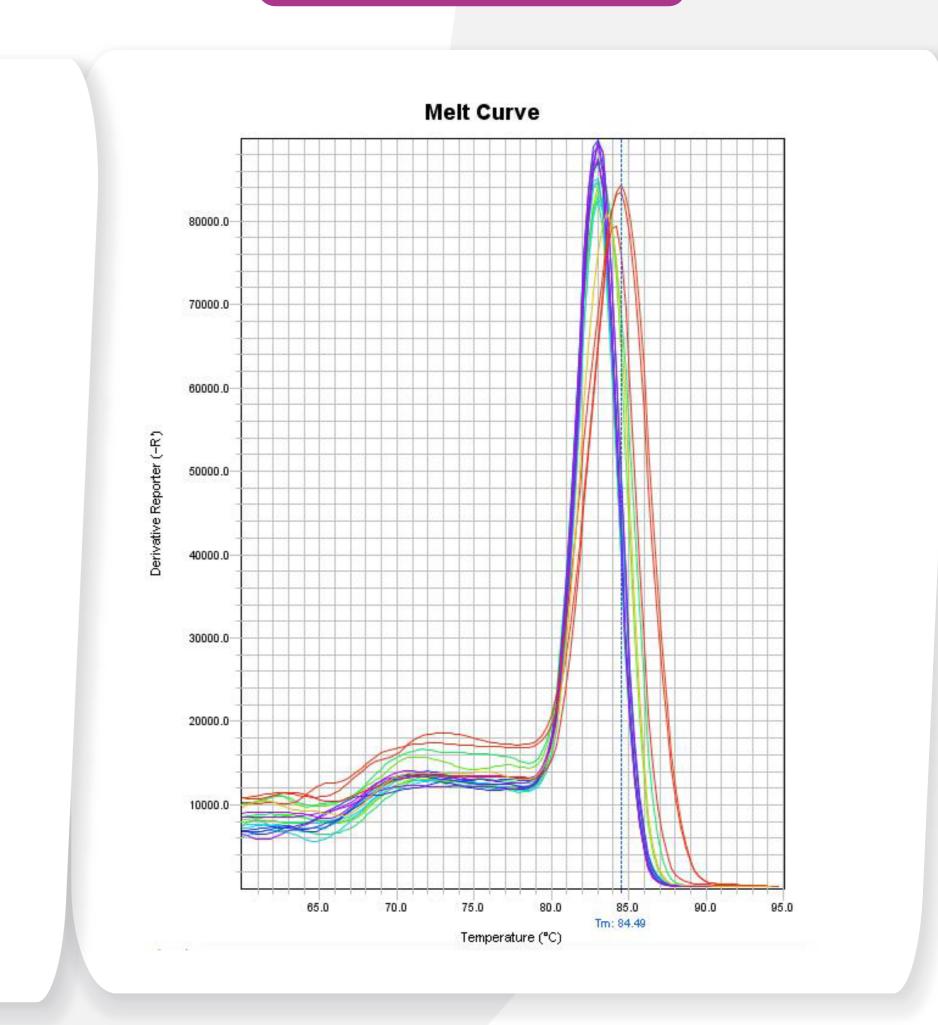
85.0 90.0 95.0 Tm: 83.75



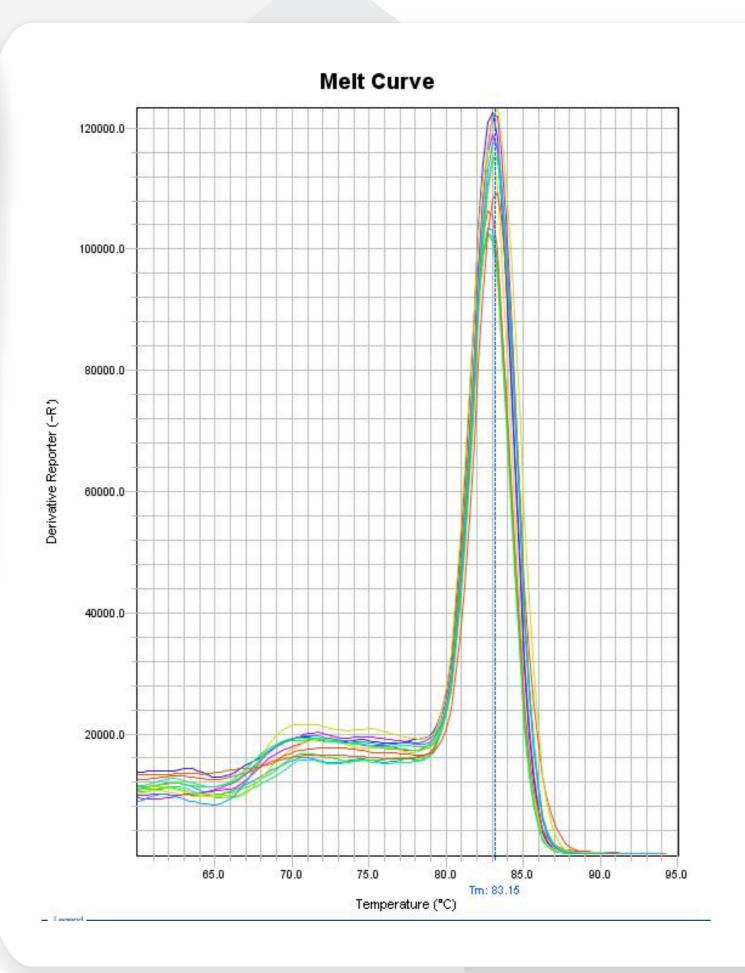
Competitor T

Competitor N

Competitor QB

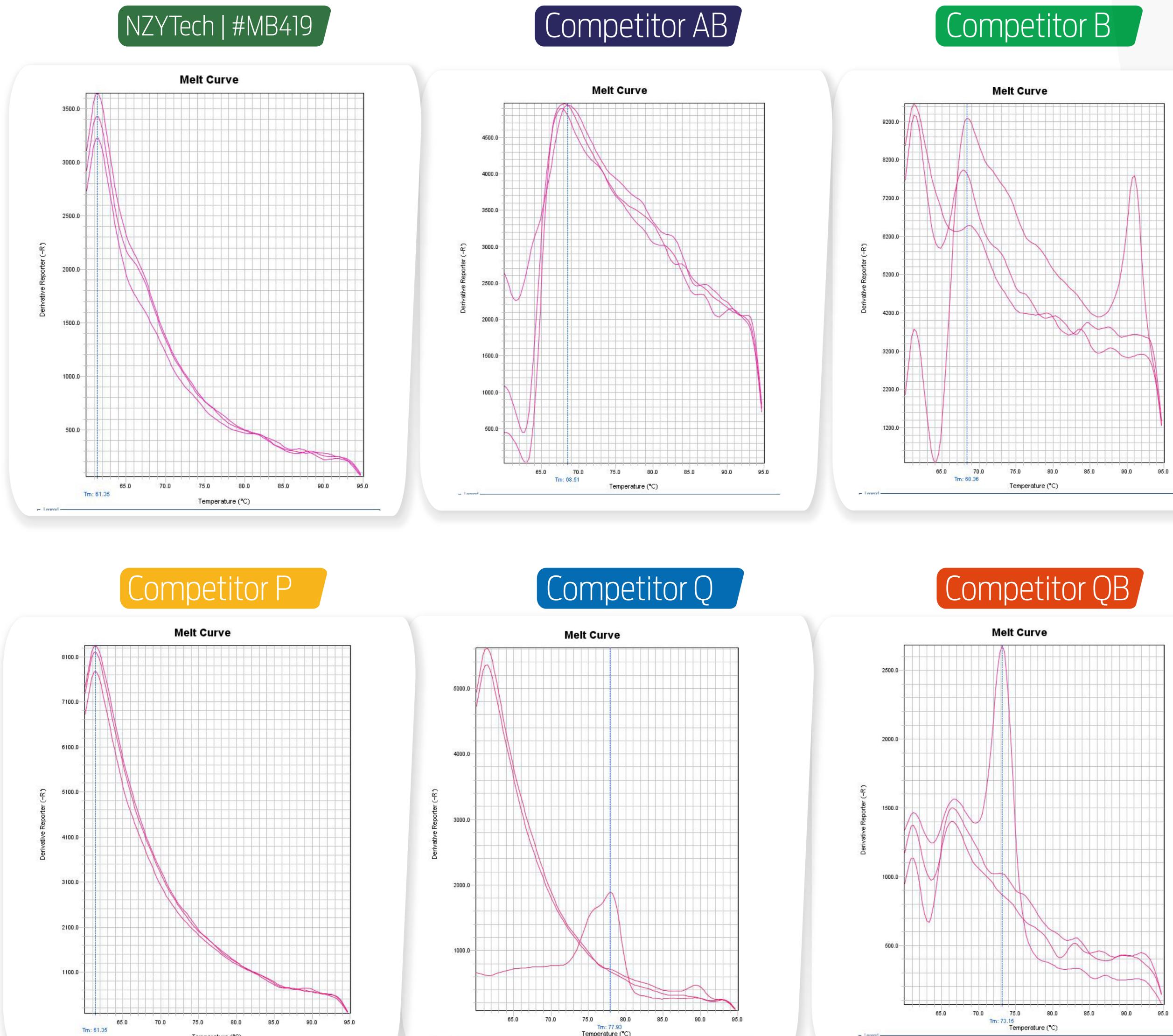


Competitor PB

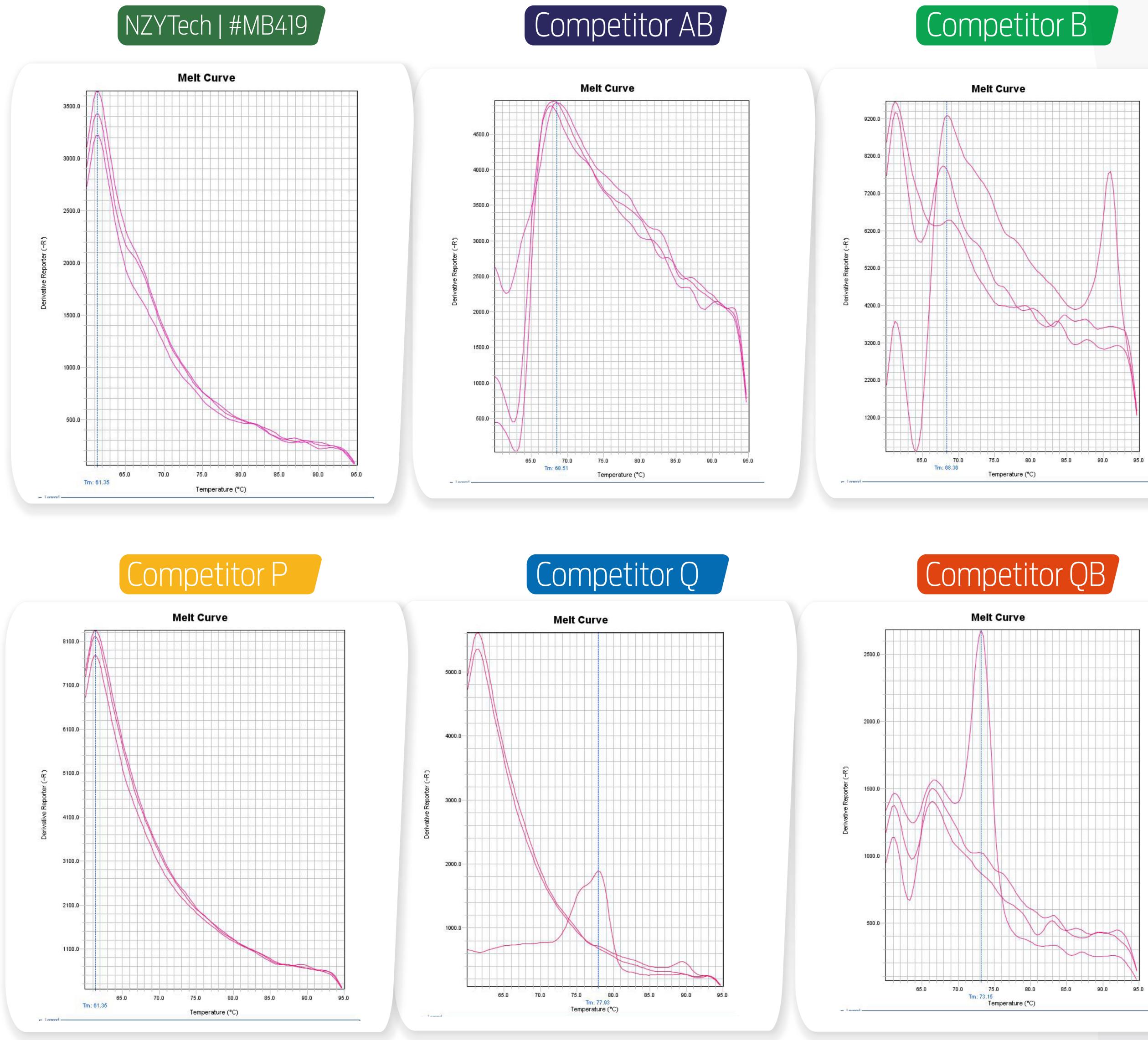




Detection of b2m from mouse cDNA - Melt Curve Analysis of NTCs Clean melting curves plot for NTC's, including absence of primer-dimer formation (low Tm), reinforce specificity of MB419, a reliable hot-start master mix.





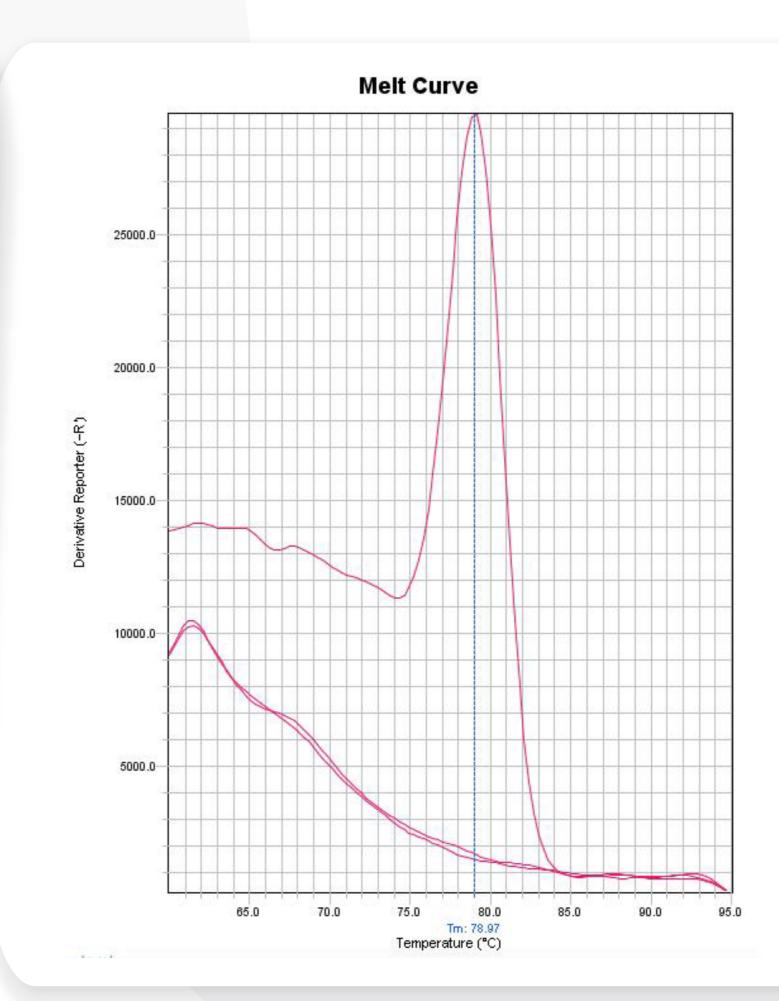


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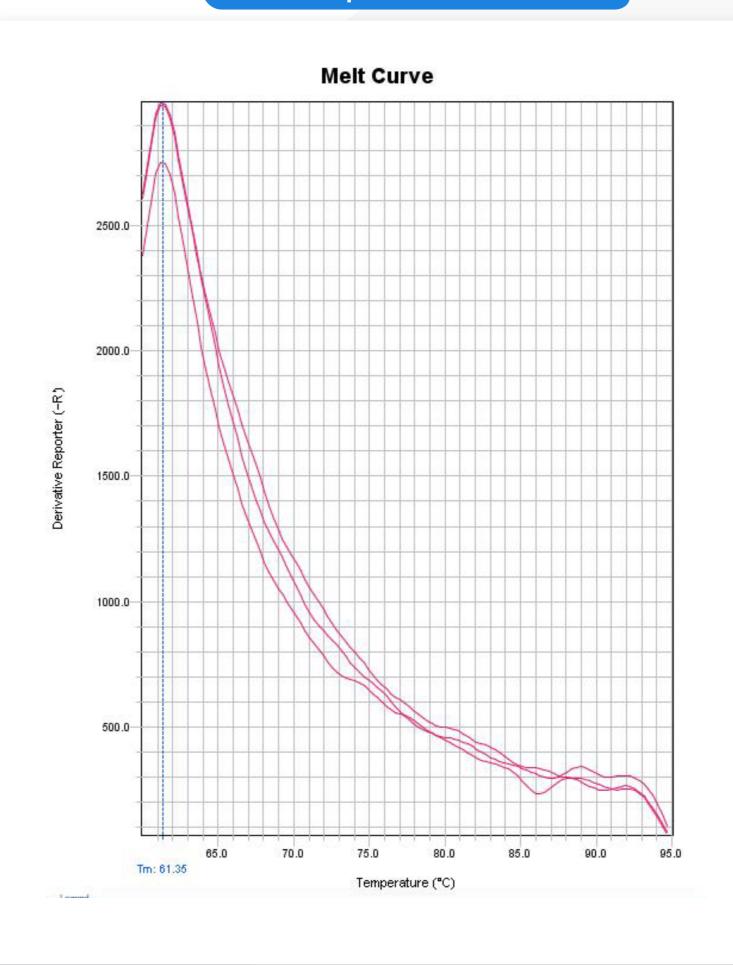
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Competitor N

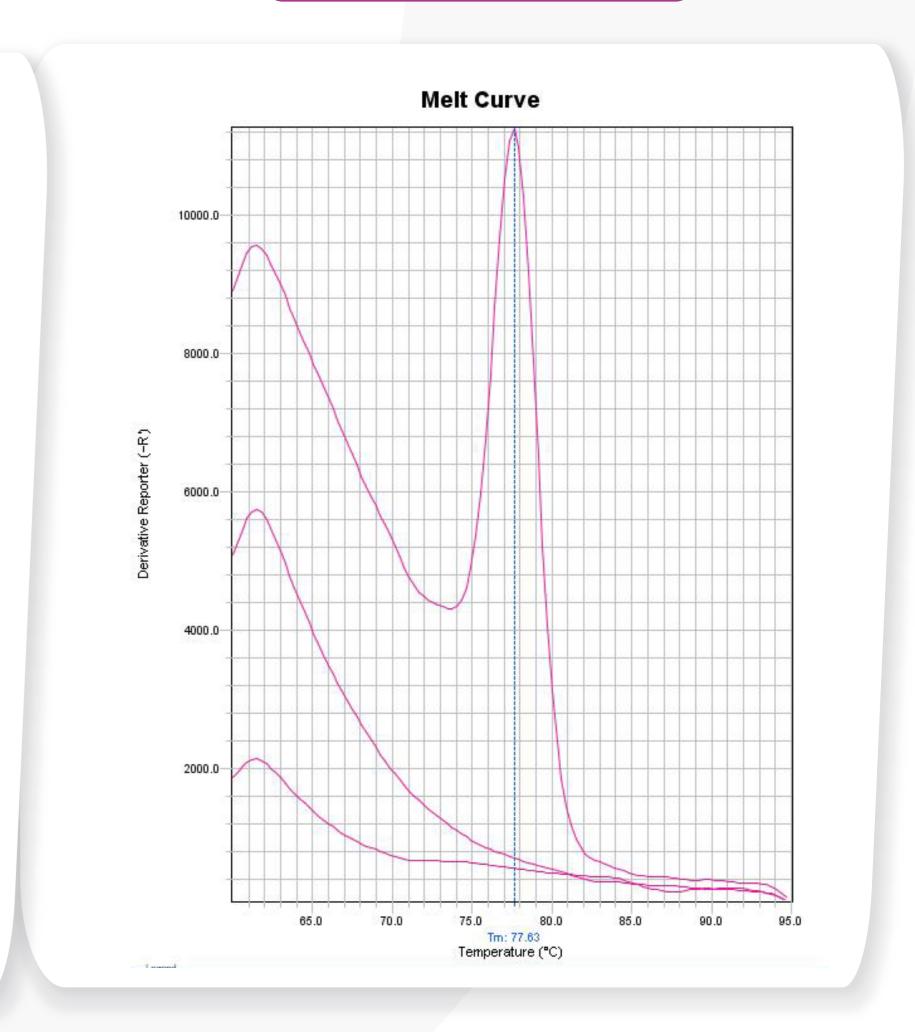


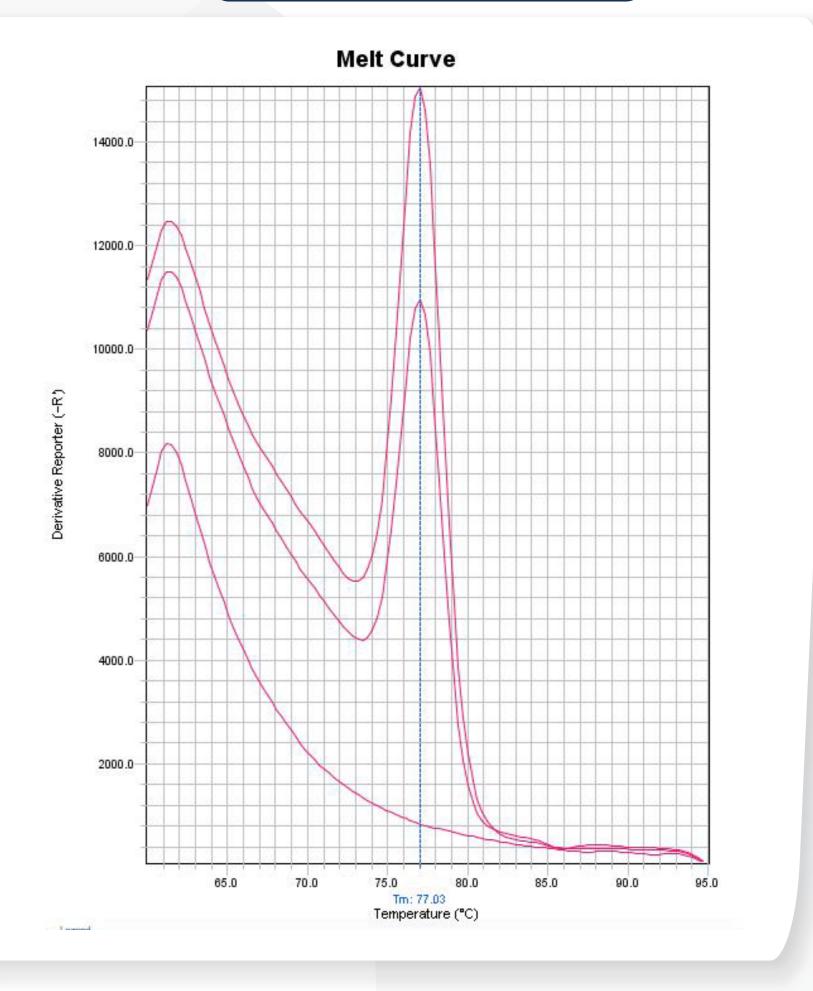
Competitor T





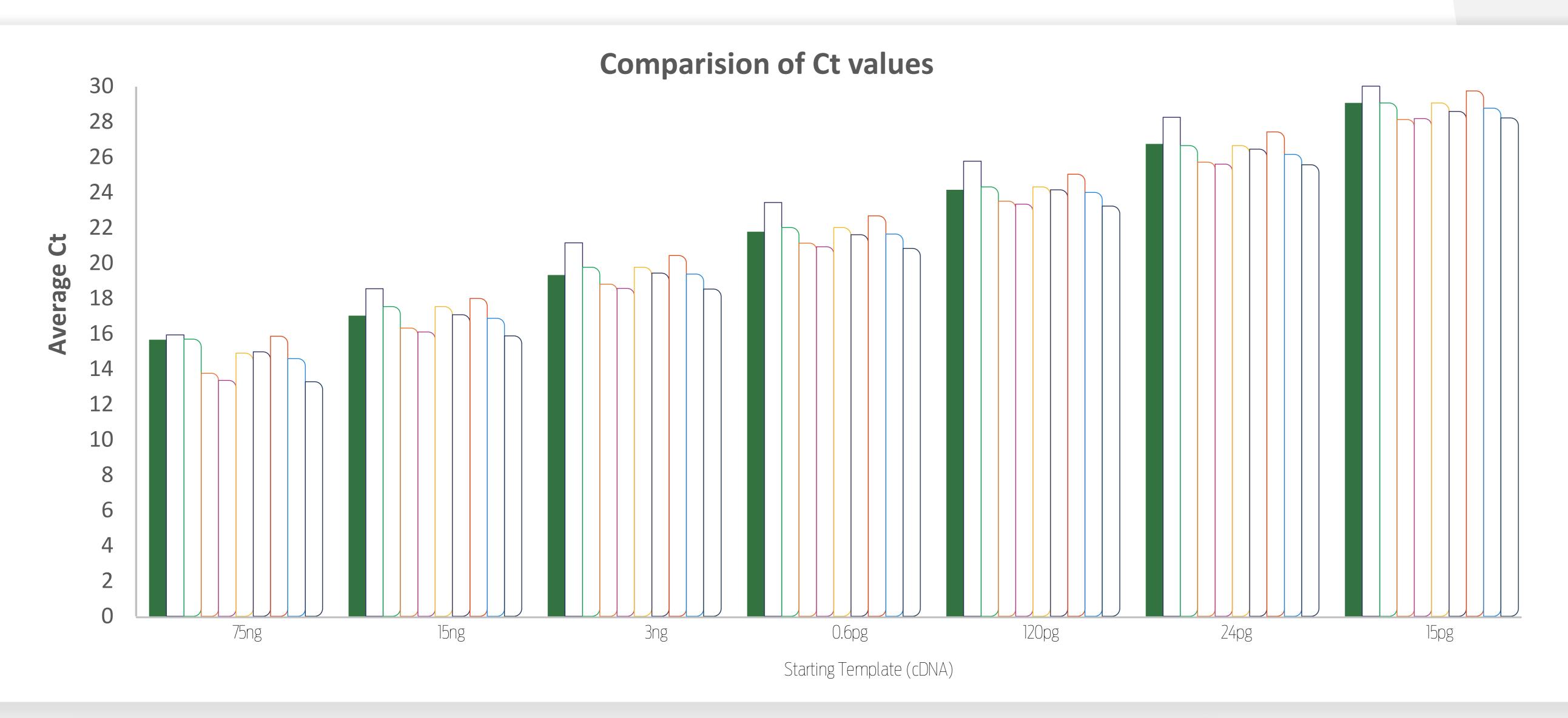
Competitor PB



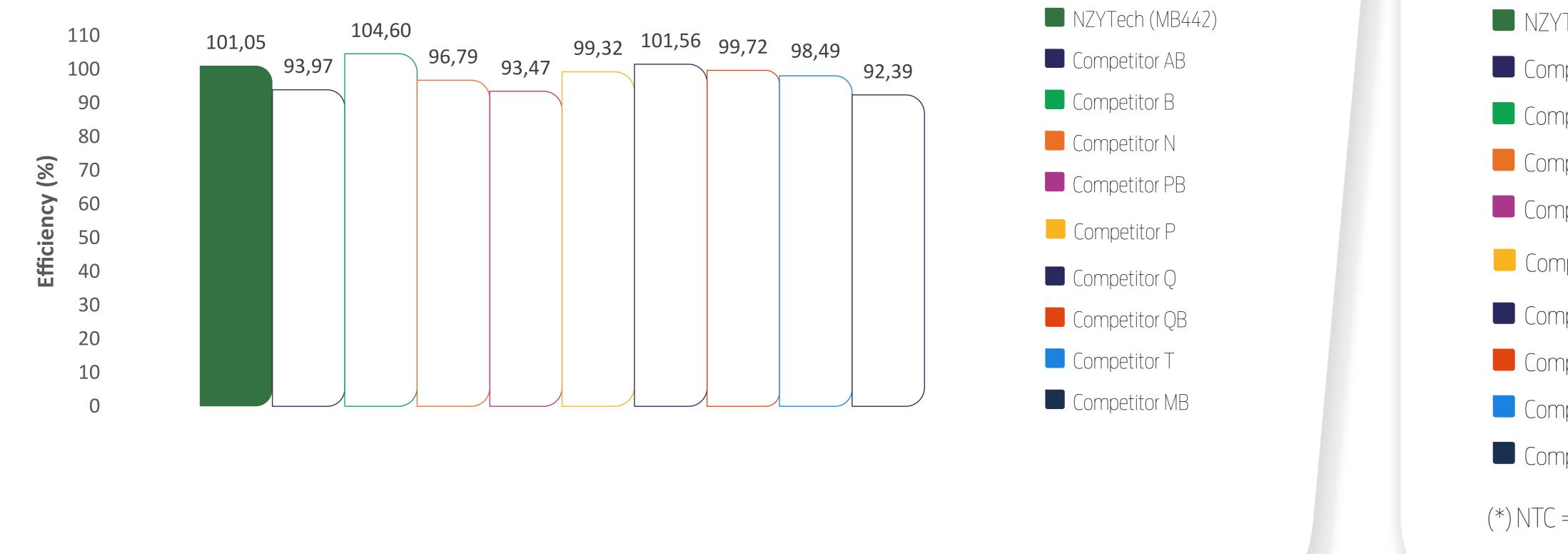




Global Analysis Comparison of Ct Values - Detection of b2m from mouse cDNA



Comparison of Efficiencies



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	Starting remptate (CDINA)						
	75 ng	15 ng	3 ng	0.6 pg	120 pg	24 pg	15 pg
■ NZYTech (MB442)	15,7	17,0	19,3	21,8	24,1	26,8	29,1
 Competitor AB Competitor B Competitor N Competitor PB Competitor Q Competitor QB Competitor T Competitor MB 	15,9	18,6	21,1	23,4	25,8	28,3	30,6
	15,7	17,5	19,8	22,0	24,3	26,7	29,1
	13,8	16,3	18,8	21,1	23,5	25,7	28,1
	13,4	16,1	18,6	20,9	23,3	25,6	28,2
	14,9	17,5	19,8	22,0	24,3	26,7	29,1
	15,0	17,1	19,4	21,6	24,2	26,5	28,6
	15,9	18,0	20,4	22,7	25,0	27,4	29,8
	14,6	16,9	19,4	21,6	24,0	26,2	28,8
	13,3	15,9	18,5	20,8	23,2	25,б	28,2

		Ct values for NTC*				
	Replicates					
]	2	3			
ZYTech (MB442)	Undetermined	Undetermined	Undetermined			
ompetitor AB	Undetermined	Undetermined	Undetermined			
ompetitor B	Undetermined	Undetermined	Undetermined			
ompetitor N	Undetermined	Undetermined	38,5			
ompetitor PB	Undetermined	Undetermined	Undetermined			
ompetitor P	Undetermined	Undetermined	Undetermined			
ompetitor Q	Undetermined	Undetermined	Undetermined			
ompetitor QB	Undetermined	Undetermined	Undetermined			
ompetitor T	Undetermined	Undetermined	Undetermined			
ompetitor MB	Undetermined	Undetermined	40,0			

(*) NTC = No template control

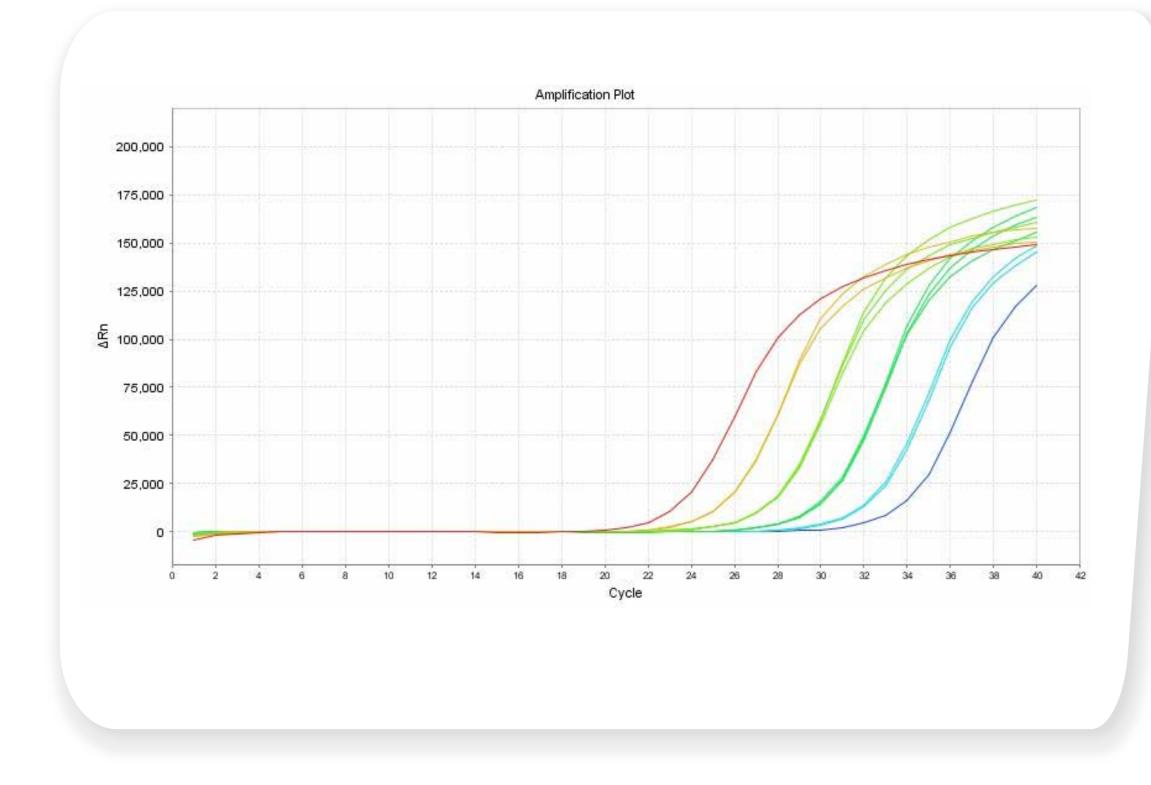


Starting Template (cDNA)



Detection of Large1 from human gDNA - Amplification

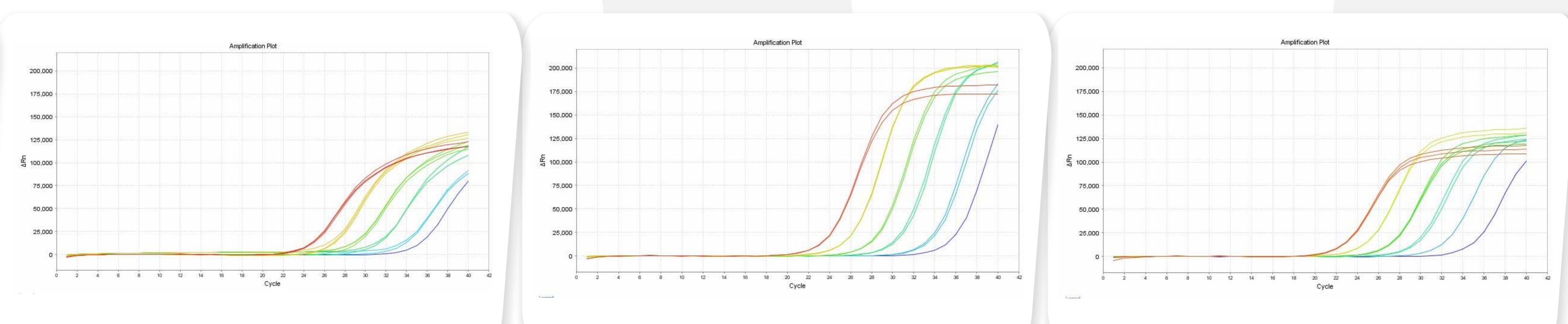
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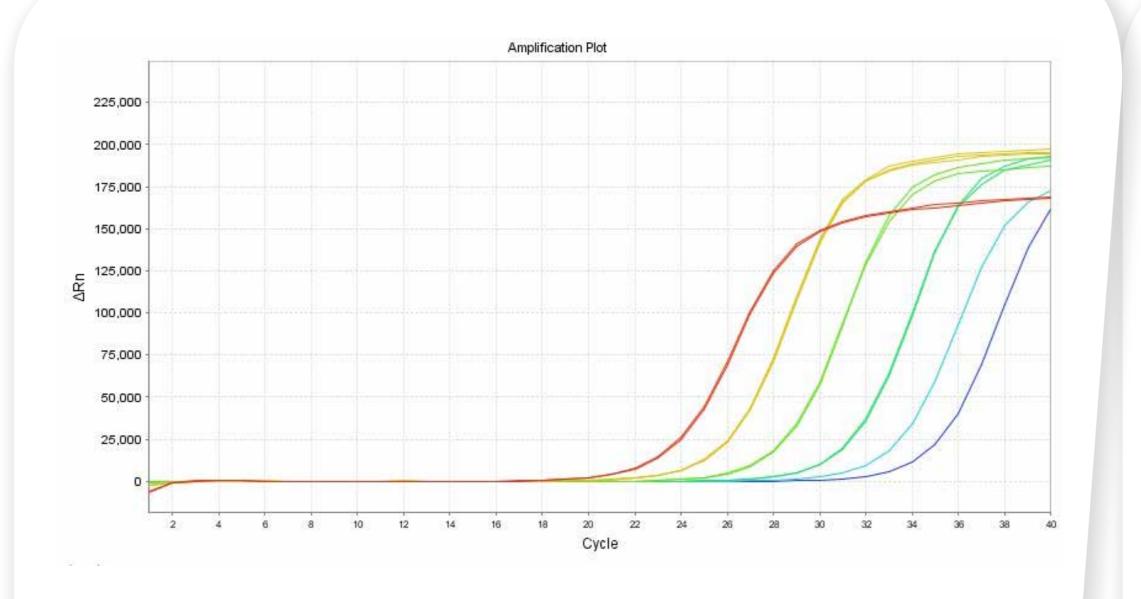
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Excellent sensitivity and linearity in the amplification using a 5-fold serial dilution of human genomic DNA.

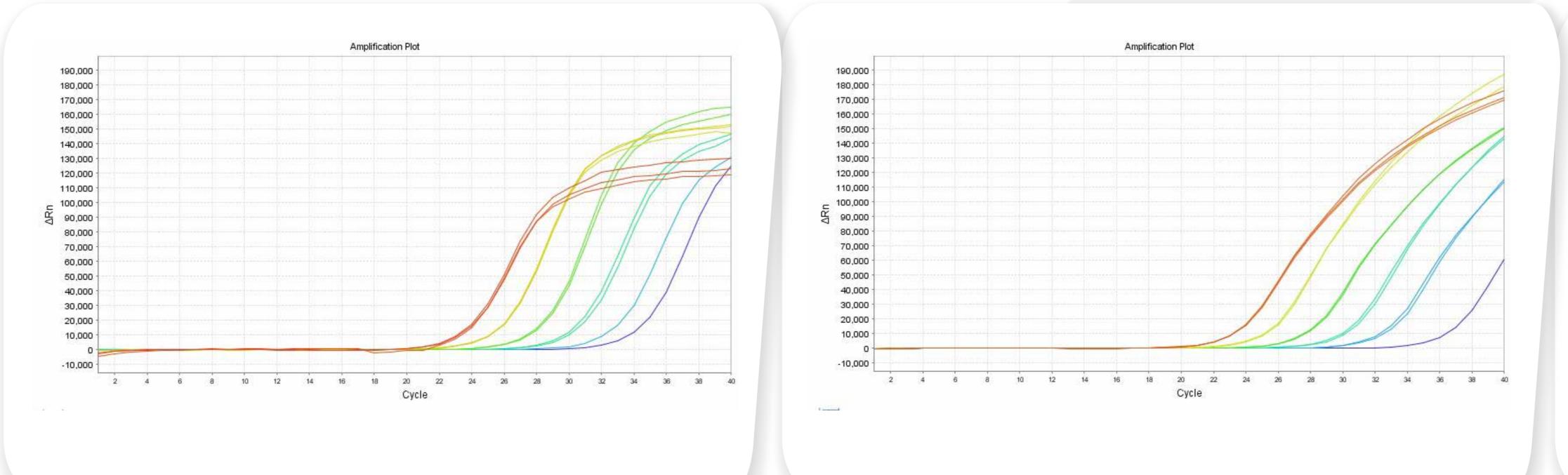




Competitor PB



Competitor QB

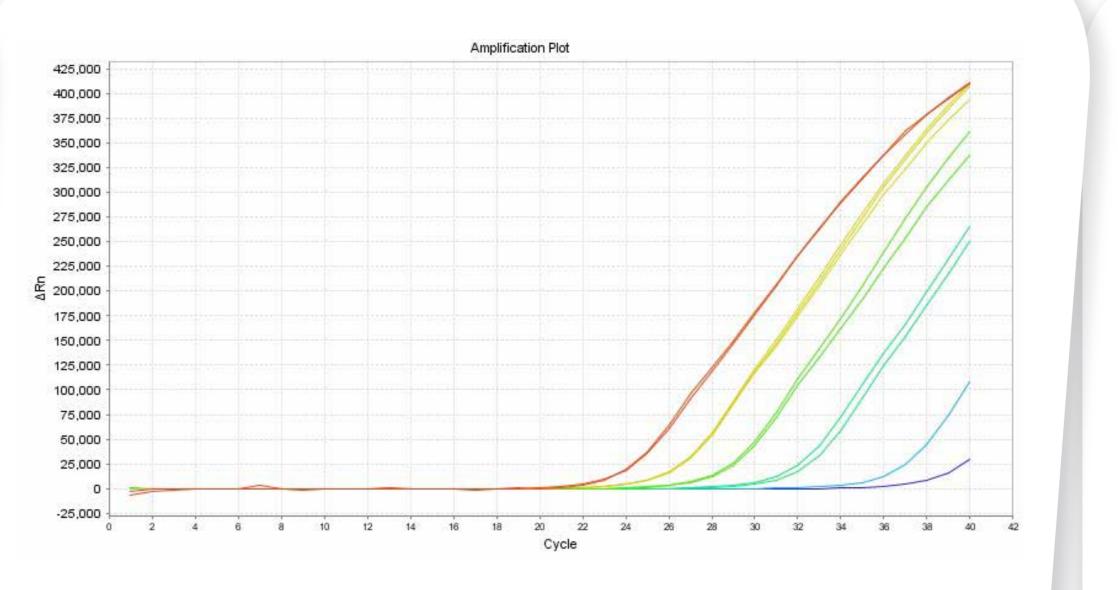


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Competitor B

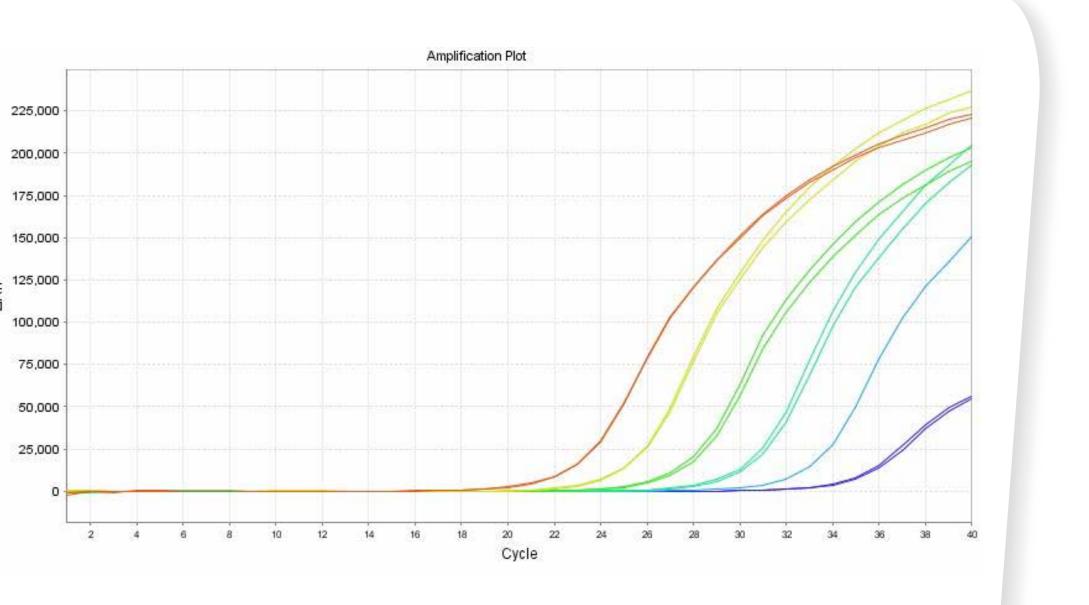
Competitor P

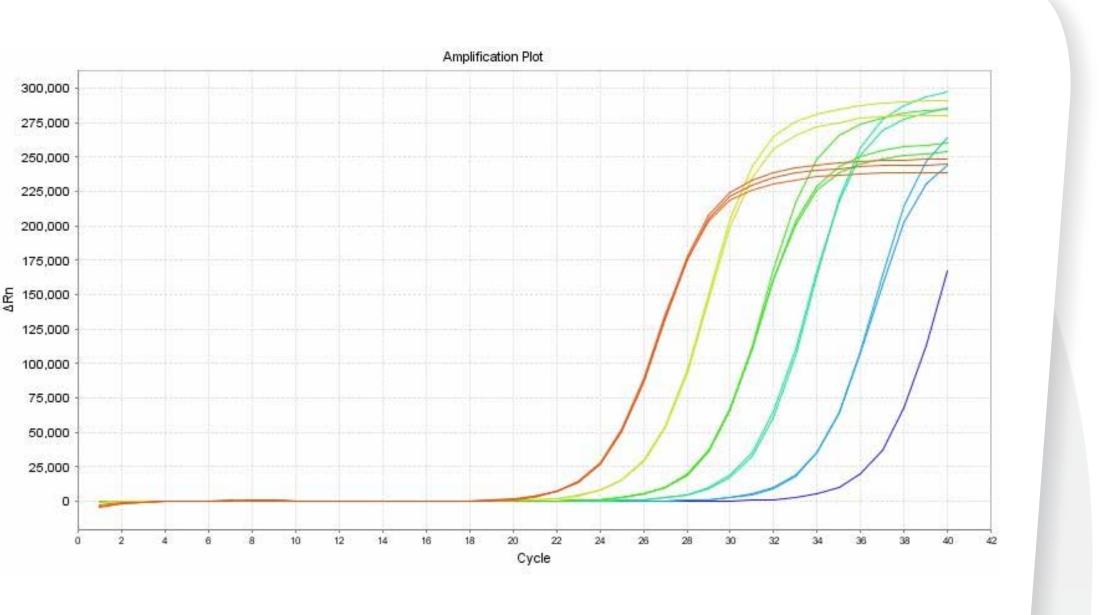


Competitor T

Competitor N

Competitor Q

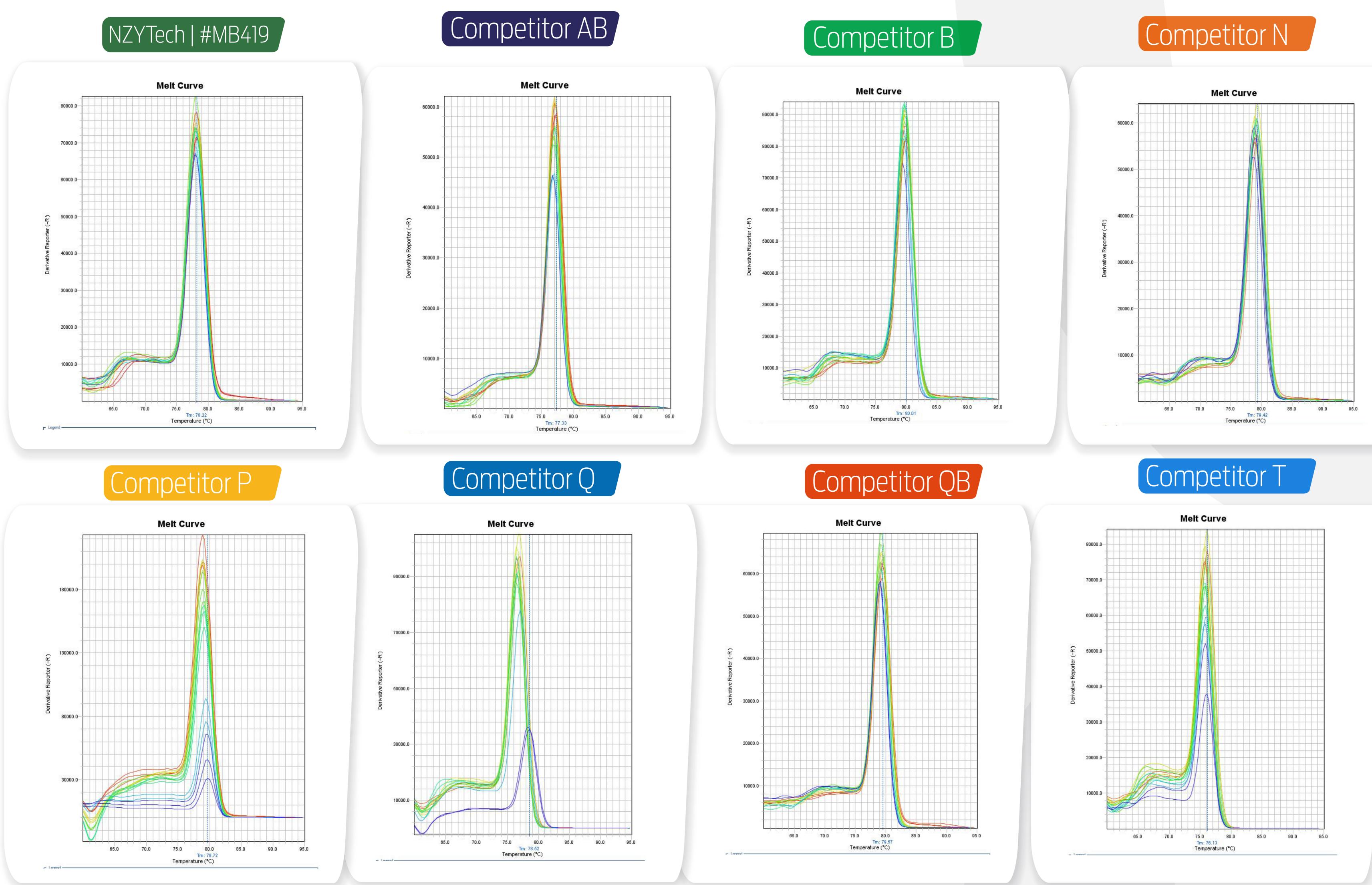


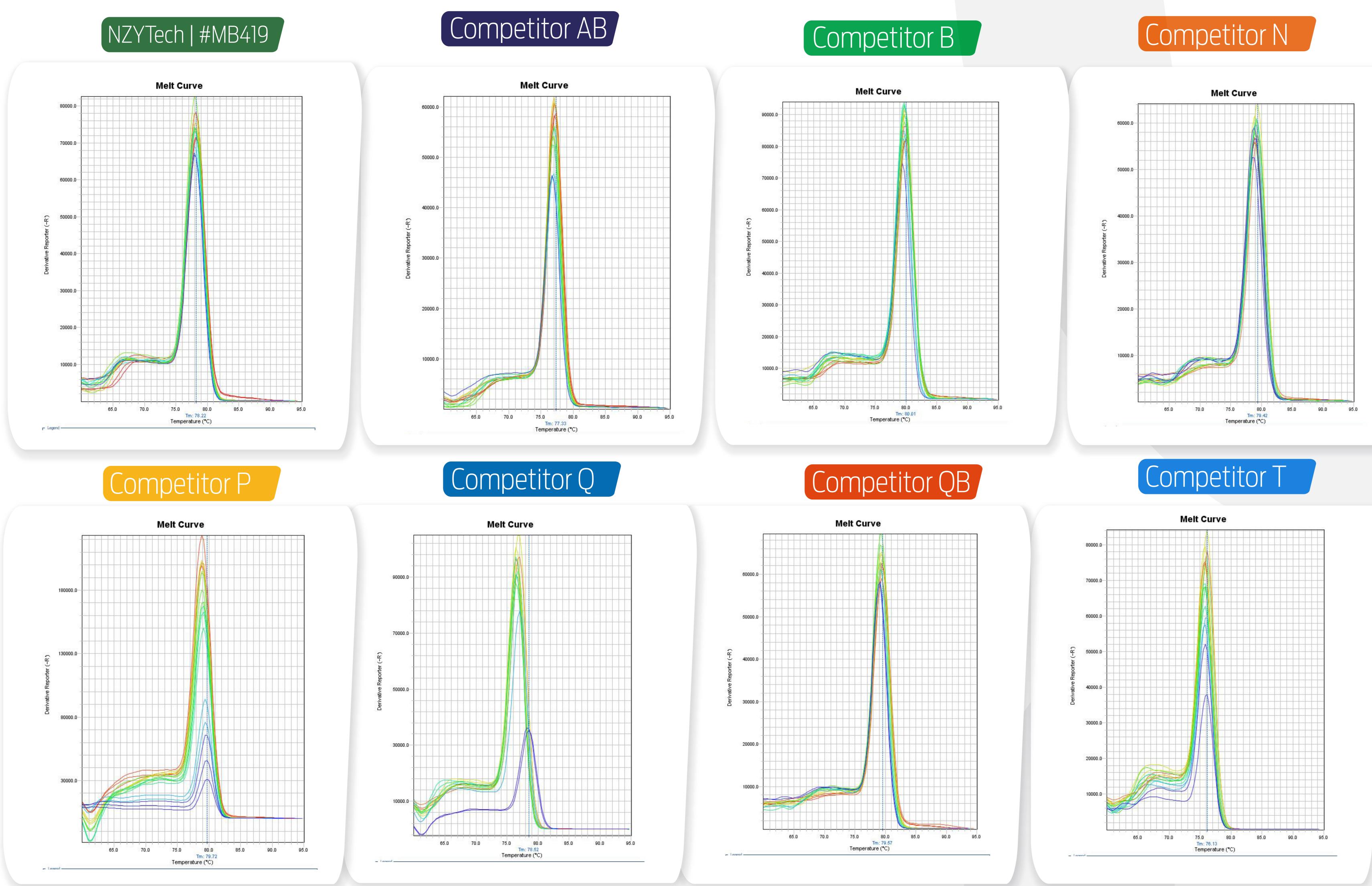




Detection of Large1 from human gDNA - Melt Curve

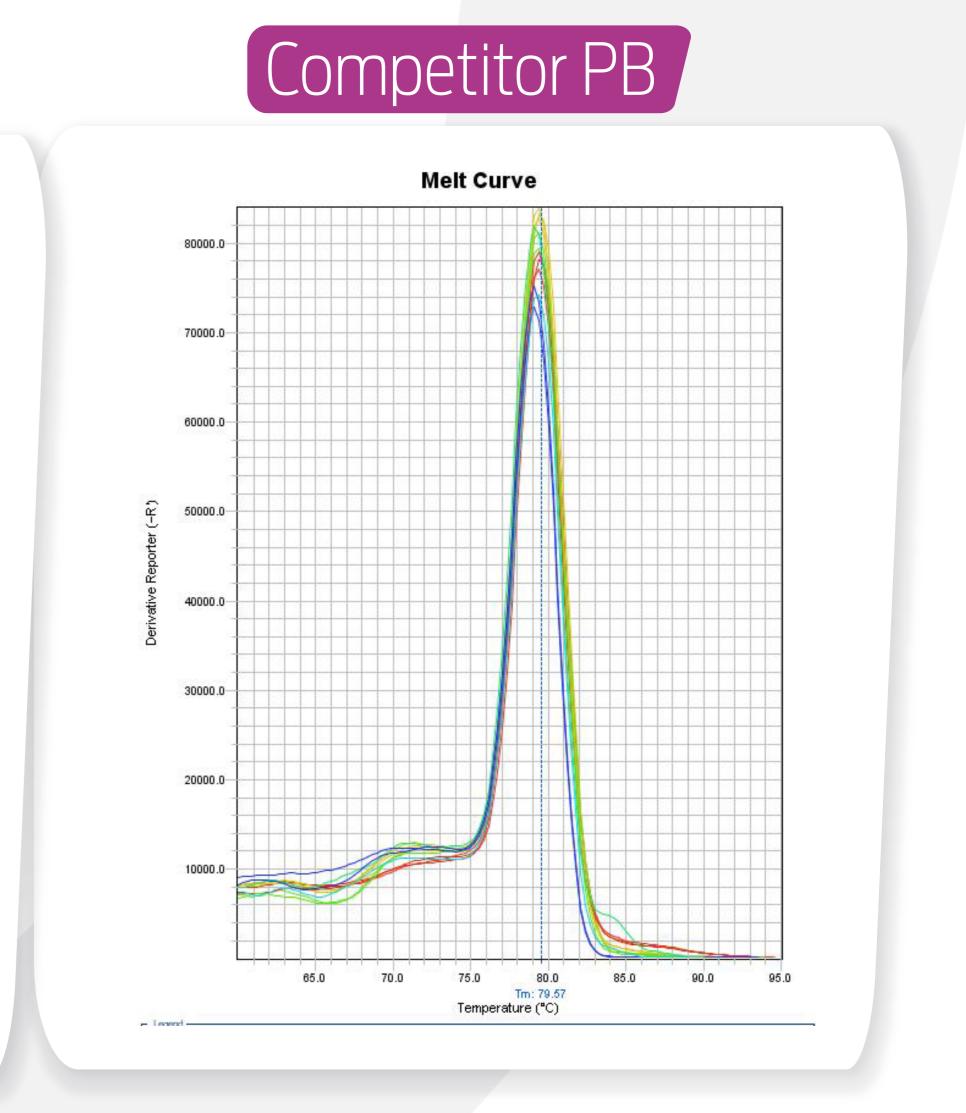
Specific detection of human target is observed for all dilutions of template when using the NZYTech's master mix, as shown by the unique peak corresponding to the desired amplicon.

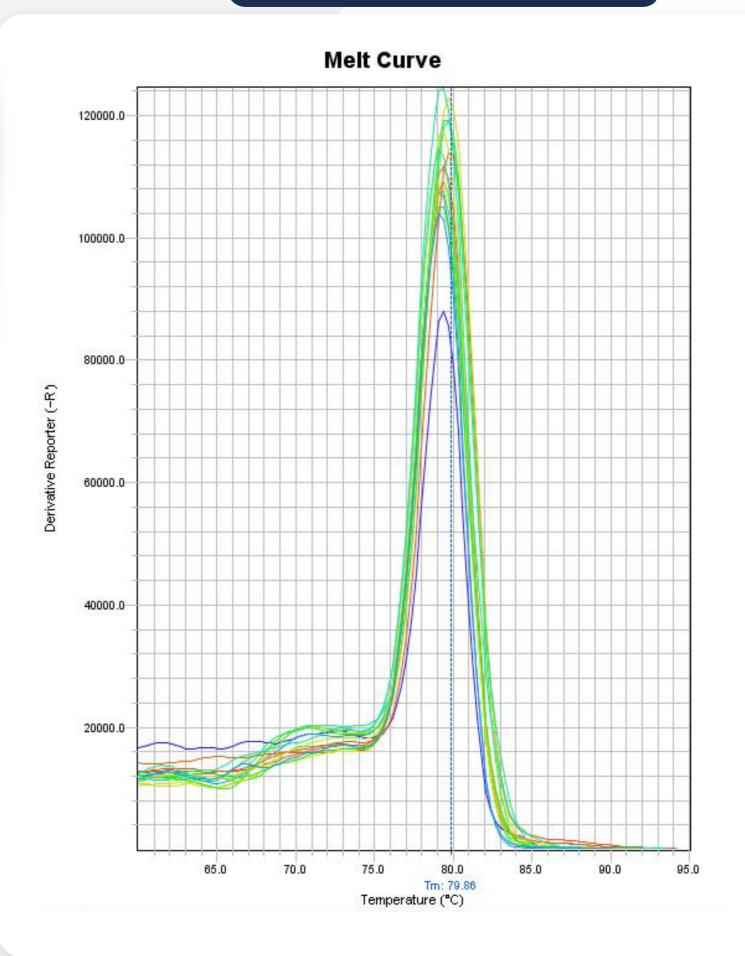




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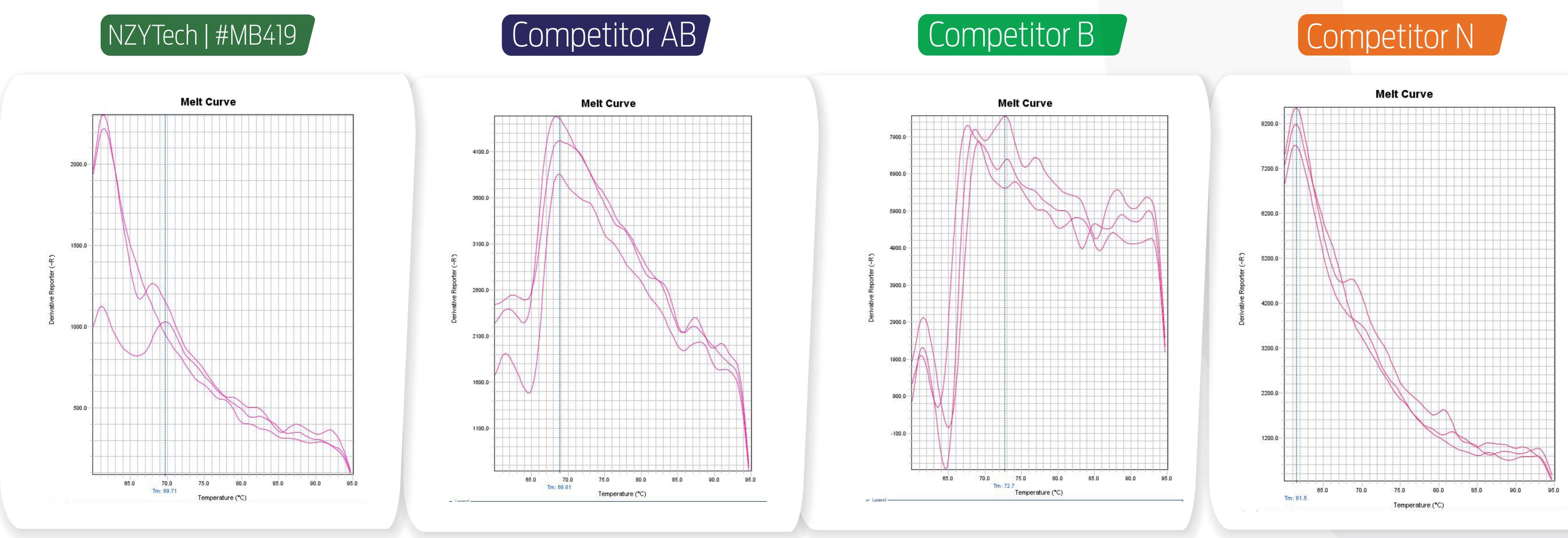




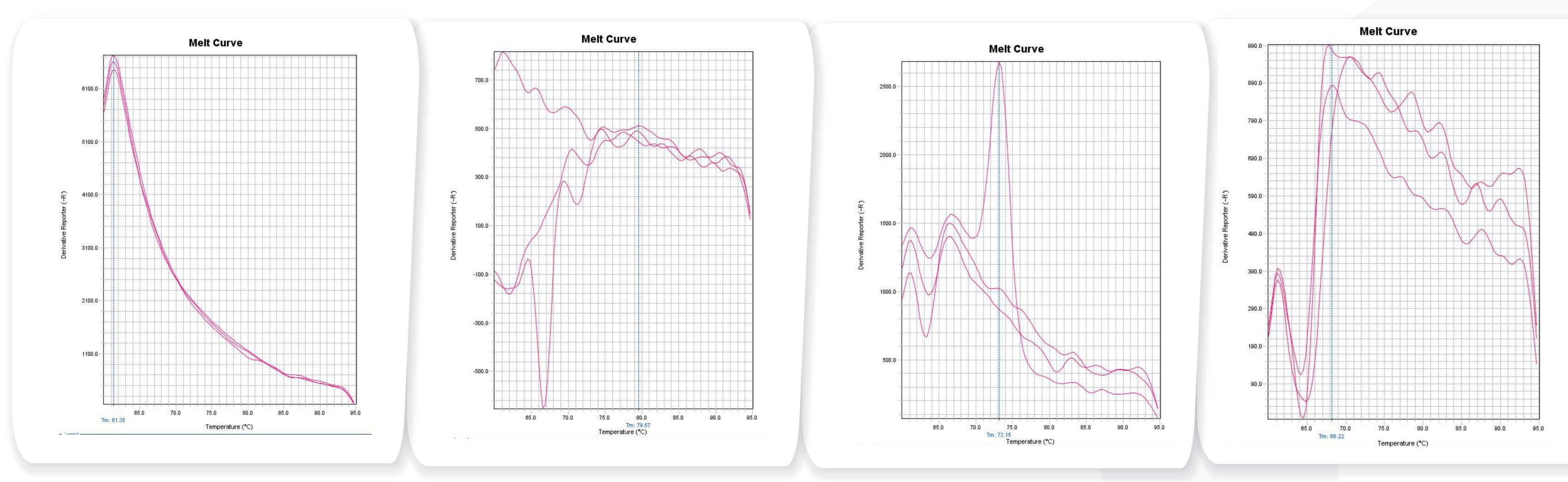


Detection of Large1 from human gDNA - Melt Curve Analysis of NTCs

NZYTech's master mix was designed to increase confidence in all qPCR experiments. Absence of primer-dimers nor non-specific amplifications corroborate the specificity of NZYSupreme qPCR Green Master Mix (MB419).



Competitor P



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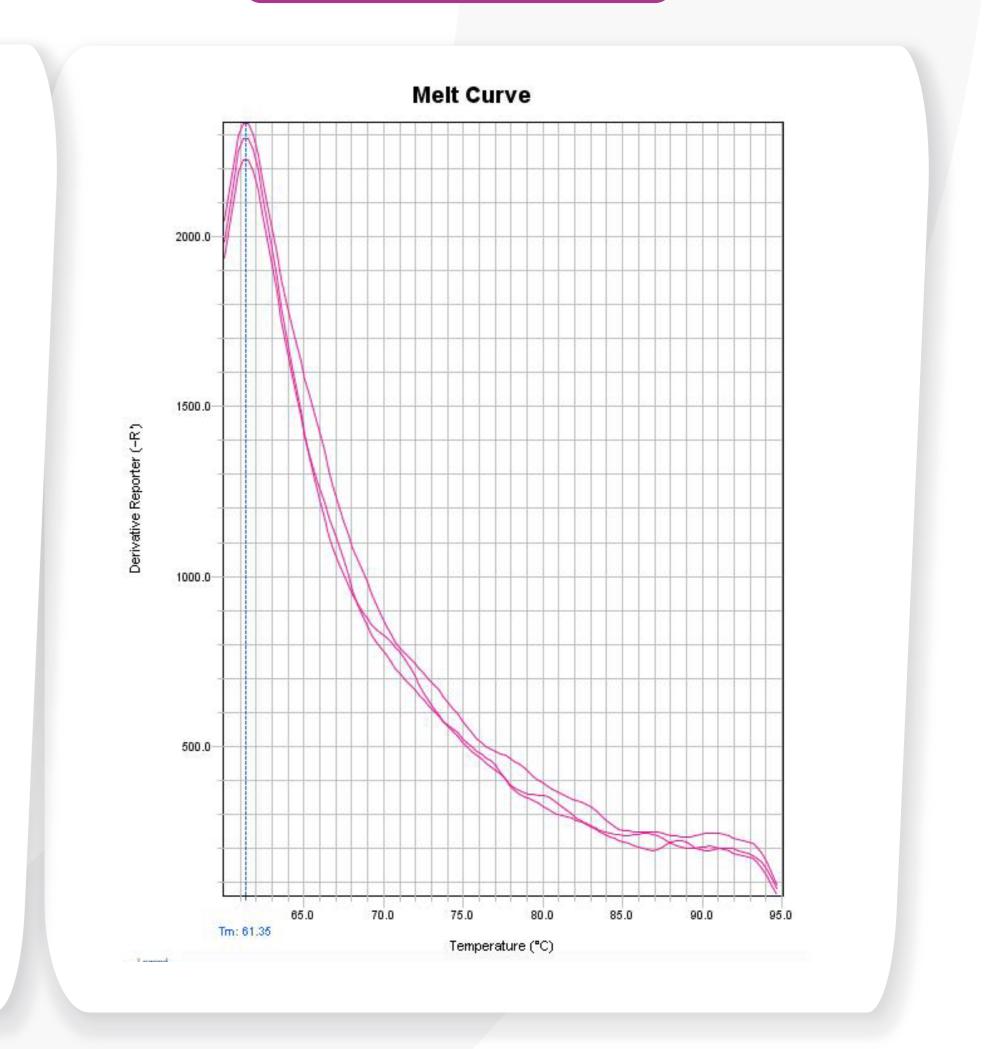
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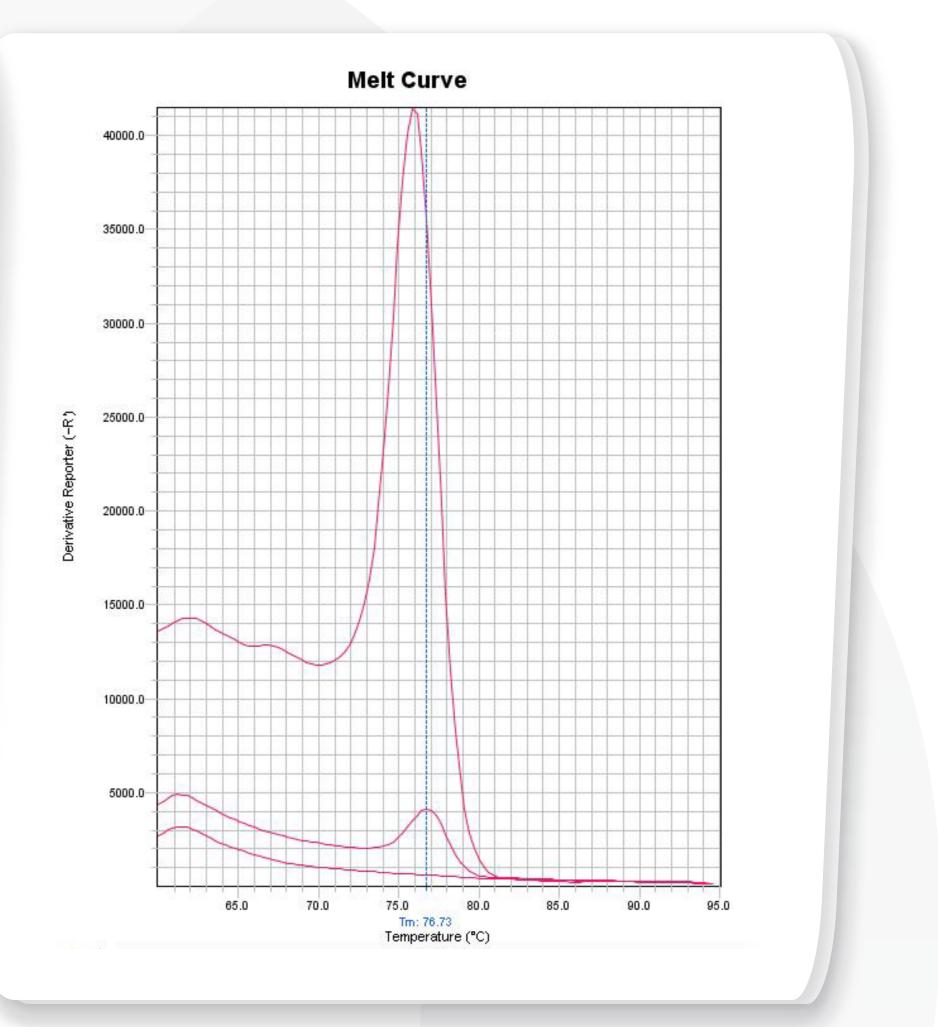






Competitor PB

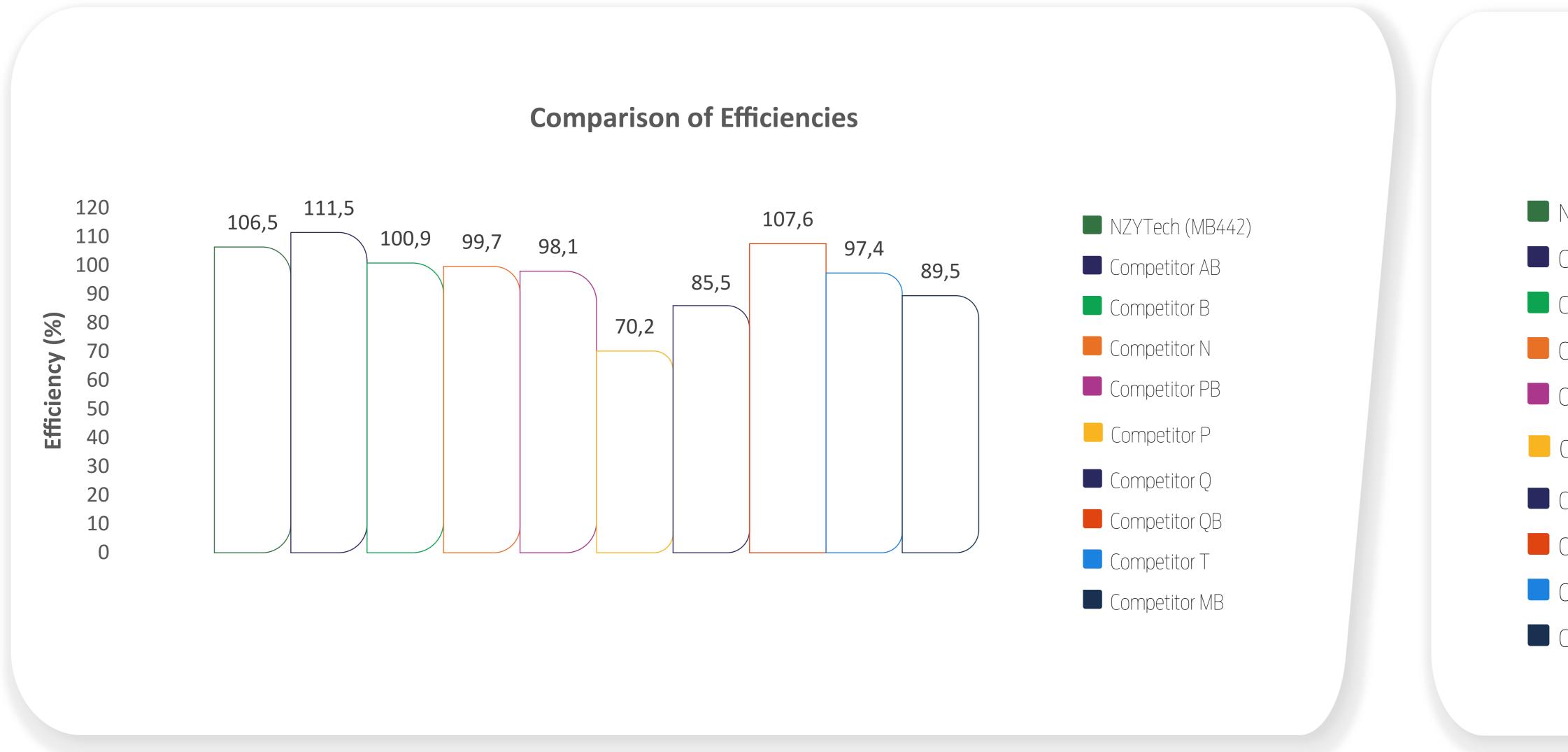






Global Analysis **Comparison of Ct Values -** Detection of Largel from human gDNA





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	Starting Template (gDNA)						
	30 ng	бng	1.2 ng	240 pg	48 pg	9.6 pg	
NZYTech (MB442)	2) 23,3	25,2	27,4	29,8	32,2	34,2	
Competitor AB	24,7	26,6	28,8	31,0	33,5	35,2	
Competitor B	23,0	25,0	27,6	29,7	32,3	34,4	
Competitor N	23,6	25,6	27,9	30,2	33,5	34,7	
Competitor PB	22,7	24,9	27,4	30,2	32,3	34,2	
Competitor P	23,2	25,5	27,9	31,1	36,2	37,3	
Competitor Q	22,6	24,7	27,2	29,8	32,6	35,6	
Competitor QB	23,4	25,3	27,7	30,0	32,9	33,9	
Competitor T	23,4	25,4	27,8	30,1	32,6	35,2	
Competitor MB	22,7	24,6	27,3	29,5	32,4	35,2	

Ct values for NTC*

NZYTech (MB442) Competitor AB Competitor B Competitor N Competitor PB Competitor P Competitor Q Competitor QB Competitor T Competitor MB

Undetermined Undetermined Undetermined Undetermined Undetermined Undetermined Undetermined Undetermined Undetermined Undetermined

Replicates

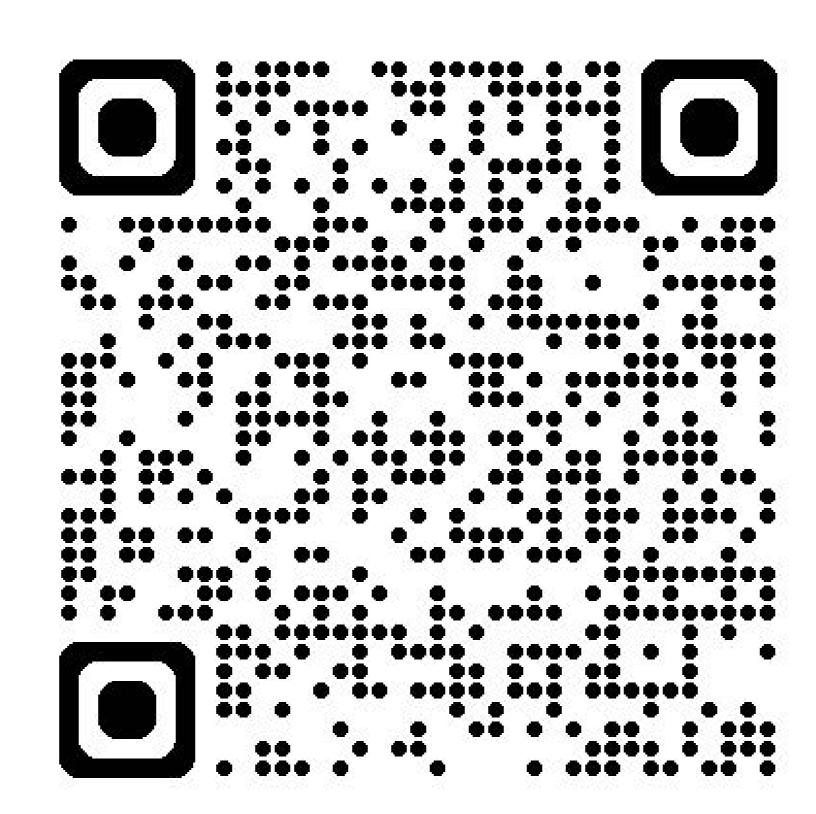
Undetermined 3

Undetermined Undetermined Undetermined Undetermined Undetermined Undetermined Undetermined Undetermined Undetermined 37.5

(*) NTC = No template control







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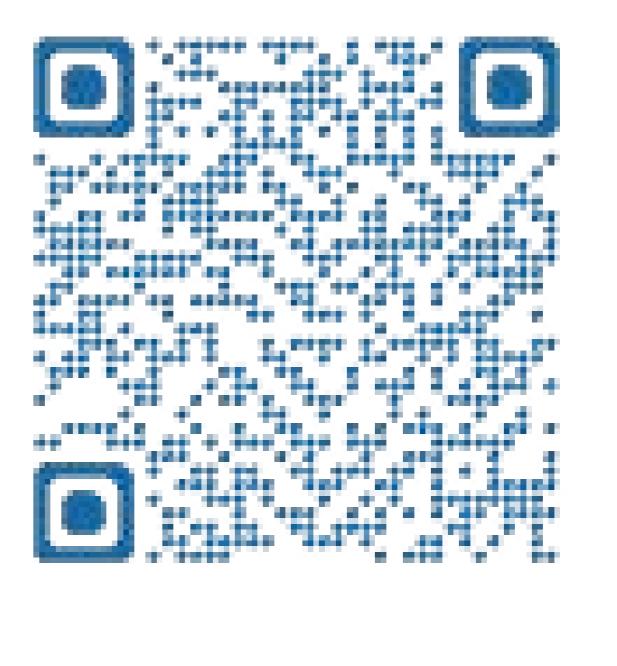
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