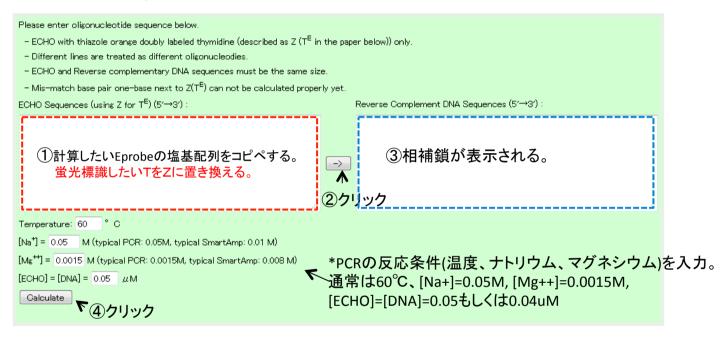
ECHO DNA Thermodynamics (https://www.dnaform.com/devel/echo/thermodynamics)にアクセスする。

ECHO/DNA Thermodynamics



Users of the ECHO/DNA Thermodynamics page are requested to cite:

Kimura, Y., Hanami, T., Tanaka, Y., de Hoon, M.J.L., Soma, T., Harbers, M., Lezhava, A., Hayashizaki, Y. and Usui, K. Effect of Thiazole Orange Doubly Labeled Thymidine on DNA Duplex Formation. Biochemistry, 51, 6056-6067 (2012).

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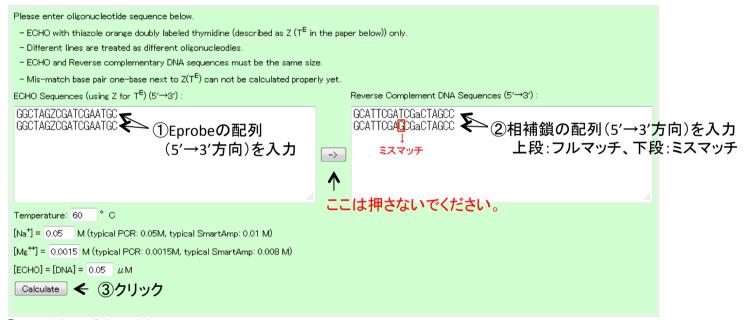
Tm計算例(5'-GGCTAGZCGATCGAATGC-3'のTm予測)

ECHO/DNA Thermodynamics																
Please enter oligonucleotide se																
- ECHO with thiazole orange of - Different lines are treated a																
- EOHO and Reverse complem																
- Mis-match base pair one-ba																
ECHO Sequences (using Z for T	for T ^E) (5'→3') :				Reverse Complement DNA Sequences (5'→3') :											
GGCTAGZCGATCGAATGC GCATTCGATCGaCTAGCC ->																
				at												
Temperature: 60 ° C																
[Na ⁺] = 0.05 M (typical PCR: 0.05M, typical SmartAmp: 0.01 M)																
[Mg ⁺⁺] = 0.0015 M (typical PCR: 0.0015M, typical SmartAmp: 0.008 M)																
[ECHO] = [DNA] = 0.05 μM																
DNA/DNA = チアゾールピンク標識の場合の予測Tm値 (一)																
⑤予測結果が表示される。 ECO/DNA (by fluorescence) = チアゾールオレンジ標識の場合の予測Tm値 ()																
○ 1 以前未り、ないこれの。																
ΔH[kcal/mol], ΔS[cal/mol	I/K], ∆G ₆₀	[kcal/mol],	α(the fra	action of c	ouble-stra	nded DNA),	T ^M [。 C]									
	DNA/DNA					ECHO/DNA						ECHO/DNA (by fluorescence)				
Sequence	ΔH	Δ8	∆G ₆₀	α	T _M	ΔН	Δ8	∆G ₆₀	α	T _M	ΔH	Δ8	∆G ₆₀	а	T _M	
5'-GGCTAGZCGATCGAATGC-3' 3'-CCGATCaGCTAGCTTACG-5'	-143.0	-400.0	-10.1	0.093	55.6	-126.0	-341.0	-12.5	0.61	61.2	-156.0	-428.0	-13.6	0.8	63.3	
Users of the ECHO/DNA Thermo Kimura,Y., Hanami,T., Tanaka,Y., o				zhava,A., Haya	shizaki,Y. an	d Usui,K. <u>Effe</u>	ct of Thiazok	e Orange Dou	bly Labeled 1	Γhymidine on	DNA Duplex F	Formation. Bio	ochemistry, 5	1, 6056-6067	7 (2012).	

フルマッチとミスマッチのTm値予測例

フルマッチ(上側がEprobe配列、下側が相補鎖) 5' GGCTAGZCGATCGAATGC 3' 3' CCGATCaGCTAGCTTACG 5' ミスマッチ(上側がEprobe配列、下側がミスマッチを含む相補鎖) 5' GGCTAGZCGATCGAATGC 3' 3' CCGATCaGCGAGCTTACG 5'

ECHO/DNA Thermodynamics



④予測結果が表示される(上段:フルマッチ、下段:ミスマッチ)。

 $\Delta H[kcal/mol], \Delta S[cal/mol/K], \Delta G_{sn}[kcal/mol], \alpha$ (the fraction of double-stranded DNA), $T_{u}[^{\circ}]$ C] DNA/DNA ECHO/DNA ECHO/DNA (by fluorescence) Sequence ΔG_{eo} T_M ΔG_{co} T_M ΔG_{60} T_M ΔH ΔS ΔH ΔS ΔH ΔS 5'-GGCTAGZCGATCGAATGC-3' -143.0 -400.0 -10.1 0.093 -126.0 -341.0 -12.5 0.61 61.2 -156.0 -428.0 -13.6 63.3 55.6 0.8 3'-CCGATCaGCTAGCTTACG-5' 5'-GGCTAGZCGATCGAATGC-3 -129.0 -9.32 -301.0 58.9 -388.0 -12.7 0.66 61.7 -360.0 0.031 53.1 -112.0 -11.70.41 -142.0 3'-CCGATCaGCGAGCTTACG-5

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