Certificate of Analysis & Product Manual



PCR Additives, DNA & RNA Precipitation Solutions
Fluorescent Probes, siRNA, Custom Primers and Probes
Hybridization and Detection Reagents

PCR Additives & Enhancers

Catalog Numbers: See Material Supplied List

Storage Condition: -20°C

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Material Supplied

Storage Condition: Store at -20°C

PCR Additives				
Item	Catalog No.	Product	Size	
	40-3020-05	dNTP 10mM	0.5 mL	
	40-3021-11	dNTP 2mM (10X)	1.1 mL	
	40-3022-16	MgCl ₂ ; 25 mM	1.6 mL	
	40-3023-16	MgCl ₂ ; 50 mM	1.6 mL	
	40-3059-10	KCI; 300 mM	1 mL	
	40-3053-10	TMAC (Tetramethyl ammonium chloride) 100 mM	1 mL	
	40-3032-10	Betaine; 5M	1 mL	
	40-3031-10	DMSO	1 mL	

Certificate of Analysis & Product Specifications

PCR additive solutions are a basic requirement for all molecular biology laboratories. Gene Link prepared a collection of common and popular solutions using nuclease free water and all are molecular biology grade.

Appropriate molarity solutions of each PCR additive are prepared and are validated for performance in PCR. All solutions are certified to be free of nucleases and nucleic acids and validated for use PCR as additives. Appropriate nuclease free handling, dispensing and storage conditions required.

Manufacturing lot numbers are stated on the label of each product and accompanying packing slip.



Product Description & Application

PCR Additives

DNA polymerases need to elongate rapidly and accurately to function effectively *in vivo* and *in vitro*, yet certain DNA regions appear to interfere with their progress. One common problem is pause sites, at which DNA polymerase molecules cease elongation for varying lengths of time. Many strong DNA polymerase pauses are at the beginnings of regions of strong secondary structure such as template hairpins (1). Taq polymerase use in PCR suffers the same fate and GC-rich DNA sequences often require laborious work to optimize the amplification assay. The GC-rich sequences possess high thermal and structural stability, presumably because the high duplex melting temperature that permits stable secondary structures to form, thus preventing completion of a faithful replication (2).

Nucleotide analog 7-deaza dGTP is effective in reducing the secondary structure associated with GC rich region by reducing the duplex stability(4). Betaine, DMSO and formamide reduces the Tm and the complex secondary structure thus the duplex stability (1-5). Tetramethyl ammonium chloride (TMAC) actually increases the specificity of hybridization and increases the Tm. The use of TMAC is recommended in PCR conditions using degenerate primers and in combination with the use of DMSO. These PCR additives and enhancing agents have been used to increase the yield, specificity and consistency of PCR reactions. These additives may have beneficial effects on some amplification and it is impossible to predict which agents will be useful in a particular context and therefore they must be empirically tested for each combination of template and primers.

Betaine, DMSO & Formamide

Betaine, DMSO and formamide helps in reducing the secondary structure of GC rich templates and assists amplification of these templates. Betaine is used at 3.5M to 0.1M, DMSO should be used between 2-8%, however 10% DMSO can reduce *Taq* polymerase activity by up to 50%. Formamide is generally used at 1-5%. It has been our experience that the use of DMSO in combination with betaine is superior to using formamide.

TMAC

TMAC is generally used at a final concentration of 15-100 mM to eliminate non-specific priming. TMAC increases the specificity of hybridization and increases the Tm. It should be used in combination with the use of DMSO and/or Betaine.

BSA & Gelatin

BSA and gelatin stabilizes the Taq polymerase and is particularly effective when amplifying old/ancient DNA contain PCR inhibitors such as melanin. These are also helpful in reducing the loss of enzyme by non-selective adsorption to tube walls.



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PCR Additives & Enhancers					
Additive	Purpose & Function	Concentration			
7-deaza-2'- deoxyguanosine; 7-deaza dGTP	GC rich region amplification. Reduce the stability of duplex DNA	Totally replace dGTP with 7-deaza dGTP; or use 7-deaza dGTP: dGTP at 3:1			
Betaine (N,N,N-trimethylglycine = [carboxymethyl]trimethyla mmonium)	Reduces Tm facilitating GC rich region amplification. Reduces duplex stability	Use 3.5M to 0.1M betaine. Be sure to use Betaine or Betaine (mono)hydrate and not Betaine HCl.			
BSA (bovine serum albumin)	BSA has proven particularly useful when attempting to amplify ancient DNA or templates which contain PCR inhibitors such as melanin.	BSA concentration of 0.01µg/µl to 0.1µg/ µl can be used.			
DMSO (dimethyl sulfoxide)	DMSO is thought to reduce secondary structure and is particularly useful for GC rich templates.	DMSO at 2-10% may be necessary for amplification of some templates, however 10% DMSO can reduce <i>Taq</i> polymerase activity by up to 50% so it should not be used routinely.			
Formamide	Reduces secondary structure and is particularly useful for GC rich templates.	Formamide is generally used at 1-5%. Do not exceed 10%.			
Non-ionic detergents e.g. Triton X-100, Tween 20 or Nonidet P-40 (NP-40)	Non-ionic detergents stabilise <i>Taq</i> polymerase and may also supress the formation of secondary structure.	0.1-1% Triton X-100, Tween 20 or NP-40 may increase yield but may also increase non-specific amplification. As little as 0.01% SDS contamination of the template DNA (left-over from the extraction procedure) can inhibit PCR by reducing <i>Taq</i> polymerase activity to as low as 10%, however, inclusion of 0.5% Tween-20 or -40 will effectively neutralize this effect.			
TMAC (tetramethylammonium chloride)	TMAC is used to reduce potential DNA-RNA mismatch and improve the stringency of hybridization reactions. It increases Tm and minimizes mis-pairing.	TMAC is generally used at a final concentration of 15-100mM to eliminate non-specific priming.			



Appendix

PCR Components and Analysis

Buffer Condition

PCR buffer conditions vary and it is imperative to optimize buffer conditions for each amplification reaction. At Gene Link most amplification reactions have been optimized to work with the following standard buffer condition, unless indicated.

	Standard Gene Link PCR
	<u>Buffer</u>
10 X PCR buffer	1 X PCR buffer
100 mM Tris-HCl pH 8.3	10 mM
500 mM KCI	50 mM
15 mM MgCl ₂	1.5 mM
0.01% Gelatin	0.001%

dNTP Concentration

Standard dNTP concentration of 0.2mM of each base is used. See section on PCR additives when dNTP concentration is changed.

F	Recipe		
2.0 mM dNTP Stock Solution Preparation*			
100 mM dGTP	100 μl		
100 mM dATP	100 μl		
100 mM dTTP	100 μl		
100 mM dCTP	100 μl		
Water	4.6 ml		
Total Volume	5 ml		
*Aliquot and freeze			

MqCl₂ Concentration

The concentration of Mg^{++} will vary from 1-5 mM, depending upon primers and substrate. Since Mg^{2+} ions form complexes with dNTPs, primers and DNA templates, the optimal concentration of $MgCl_2$ has to be selected for each experiment. Low Mg^{2+} ions result in a low yield of PCR product, and high concentrations increase the yield of non-specific products and promote mis-incorporation. Lower Mg^{2+} concentrations are desirable when fidelity of DNA synthesis is critical. The recommended range of $MgCl_2$ concentration is 1-4 mM, under the standard reaction conditions specified. At Gene Link, using the standard PCR buffer with KCl, a final dNTP concentration of 0.2 mM, a $MgCl_2$ concentration of 1.5 is used in most cases. If the DNA samples contain EDTA or other chelators, the $MgCl_2$ concentration in the reaction mixture should be raised proportionally. Given below is an $MgCl_2$ concentration calculation and addition table using a stock solution of 25 mM $MgCl_2$.

MgCl₂ Concentration & Addition Table								
Final concentration of MgCl₂ in 50µl reaction mix, (mM) 1.0 1.25 1.5 1.75 2.0 2.5 3.0 4.0					4.0			
Volume of 25mM MgCl ₂ , µl	2	2.5	3	3.5	4	5	6	8



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Primer Concentration

The final concentration of primers in a PCR reaction is usually 0.5 to 1 μ M (micromolar). This is equivalent to 0.5 to 1 μ M (micromolar). This is equivalent to 0.5 to 1 μ M (micromolar). This is equivalent to 0.5 to 1 μ M (micromolar). This is equivalent to 0.5 to 1 μ M (micromolar). This is equivalent to 0.5 to 1 μ M (micromolar). This is equivalent to 0.5 to 1 μ M (micromolar). This is equivalent to 0.5 to 1 μ M (micromolar). This is equivalent to 0.5 to 1 μ M (micromolar). This is equivalent to 0.5 to 1 μ M (micromolar). This is equivalent to 0.5 to 1 μ M (micromolar).

Primer Reconstitution

Stock Primer Mix: Dissolve the supplied 10 nmols of lyophilized Genemer™ in 100 µl sterile TE. The 10 nmols of primers when dissolved in 100 µl will give a solution of 100 µM i.e. 100 pmols/µl.

Primer Mix: Prepare a 10 pmols/µl Primer Mix solution by a ten fold dilution of the stock primer mix. Example: Add 180 µl sterile TE to a new tube, to this tube add 20 µl of primer stock solution. Label this tube as Primer Mix 10 pmols/µl.

Amplification Thermal Cycling

Hot Start: It is essential to have a 'Hot Start' profile for amplification of any fragment from a complex template like human genomic DNA. Taq polymerase has low activity at room temperature and it is essential to minimize any mispriming in the first cycle of amplification. A typical hot start profile is given below. Various enzyme preparations are available which are activated by heat in the first cycle. A simple hot start protocol is given below that can be used with regular Taq polymerase. See the section on PCR additives for amplification of products from high GC content templates.

Hot Start					
Step Time & Temperature Cycles					
Initial Denaturation	95°C for 5 minutes	1			
Annealing	60°C Hold Infinity	Hold			
Comments: Add Tag premix while on hold.					

Amplification File

The initial denaturation step at 94 °C for 30 seconds is sufficient for all templates. The number of cycles is usually set to 30 and is sufficient to amplify 1-10 µg of product depending on the initial concentration of template. A higher number of cycles from 35-45 cycles may be used, but internal priming on the product and over amplification of unwanted bands often result from over-cycling. Generally, it is better to focus on optimizing reaction conditions than to go beyond 35 cycles.

Typical Amplification File					
Step	Cycles				
Denaturation	94 °C	30 sec.			
Annealing	*	30 sec.	30		
Elongation	72 °C	30 sec.			
Fill in	72 °C	7	1		
Extension	72°C	minutes			
Hold 4°C		Infinity	Hold		
*Based on the Tm of the primers. Usually varies from					
50°C to 65°C					



Typical Reaction Premix

Typical PCR Premix (/50µl)				
Component	Volume			
10 x PCR Buffer	5 μΙ			
2.0 mM dNTP mix (each)	5 μΙ			
Primer Mix (10 pmol/µl each) or 2.5µl of	2.5 μl			
10 pmol/μl of individual primer				
(final 25 pmol of each primer/50µl)				
H ₂ O	37.5 μΙ			
Total Volume	50 µl			

Typical PCR Reaction Mix

PCR reaction (/50µl)				
Component	Volume			
PCR premix	45 μΙ			
100ng/μl diluted DNA	1 μΙ			
Hot start and then add				
Taq premix	5 μΙ			

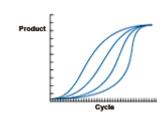
Taq Premix Preparation

Taq Premix (/50μl)				
Component	Volume			
PCR Premix	6 μΙ			
Taq polymerase (5 u/μl)	0.25µl			
Add 5µl/50 µl rxn. After initial denaturation				
Use 2.5 units of Taq for 100 µl reactions. Taq is usually supplied at a concentration of 5 units/µl				

Yield and Kinetics

The target will be amplified by up to 10^6 fold in a successful reaction, but the amplification will usually plateau at 1-10 μ g. Thus, 1 pg of target sequence in the reaction is a good place to begin.

PCR reactions produce product in a nonlinear pattern. Amplification follows a typical exponential curve until some saturation point is reached. Generally products will not be further amplified once 1-5 µg has been generated. Saturation by one product of a reaction does not always prevent further amplification of other generally unwanted products. Overcycling may decrease the quality of an otherwise good reaction. When first optimizing a reaction, it is advisable to take samples every 5 or 10 cycles to determine the number of cycles actually needed.





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References:

- 1. Kovarova, M; and Draber, P; (2000) New Specificity and yield enhancer for polymerase chain reactions (2000) Nucl. Acids. Res. 28: e70.
- 2. Henke, W., Herdel, K., Jung, K., Schnorr, D. and Stefan A. Loening, S. (1997) Betaine improves the PCR amplification of GC-rich DNA sequences. Nucl. Acids Res. 25: 3957-3958.
- 3. Daniel S. Mytelka, D.S., and Chamberlin, M.J., (1996) Analysis and suppression of DNA polymerasepauses associated with a trinucleotide consensus. Nuc. Acids Res., 24:2774–278.
- 4. Keith, J. M., Cochran, D.A.E., Lala, G.H., Adams, P., Bryant, D.and Mitchelson, K.R. (2004) Unlocking hidden genomic sequence. Nucl. Acids Res. 32: e35.
- 5. Owczarzy, R., Dunietz, I., Behlke, M.A., Klotz, I.M. and Joseph A. Walder. (2003) Thermodynamic treatment of oligonucleotide duplex-simplex equilibria. PNAS, 100:14840-14845.



Ordering Information

PCR Additives & Reagents					
Product	Catalog No.	Unit Size			
Taq DNA Polymerase 300 units; 5 μ/μL; 60 μL	40-5200-30	300 units			
PCR Buffer Standard (10 X); 1.6 mL	40-3060-16	1.6 mL			
PCR Buffer Mg Free (10 X); 1.6 mL	40-3061-16	1.6 mL			
Taq Polymerase Dilution Buffer; 1 mL	40-3070-10	1 mL			
dNTP 2mM (10X); 1.1 mL	40-3021-11	1.1 mL			
MgCl ₂ ; 25 mM; 1.6 mL	40-3022-16	1.6 mL			
Omni-Marker™ Universal Unlabeled; 1 mL	40-3005-10	1 mL			
Primer and Template Mix; 500 bp; 40 reactions	40-2026-60PT	100 μL			
Nuclease Free Water, 10 X 1.6 mL	40-3001-16	10 X 1.6 mL			
DMSO, 1 mL	40-3031-10	1 mL			
TMAC (Tetramethyl ammonium chloride) 100 mM; 1 mL	40-3053-10	1 mL			
KCI 300 mM; 1 mL	40-3059-10	1 mL			
Betaine 5M; 1 mL	40-3032-10	1 mL			

Related Products Ordering Information

DNA & RNA Precipitation Solutions		
Product	Catalog No.	Unit Size
DNA & RNA Precipitation Solutions Pack (contains the following; Glycogen Solution 10 mg/mL; 1 mL [40-5112-01]; Linear Acrylamide Solution 5mg/mL; 1 mL [40-5113-01] LiCI RNA Precipitation Solution [40-5131-05]; Sodium Acetate DNA & RNA Precipitation Solution [40-5132-05]; Sodium Chloride DNA & RNA Precipitation [40-5134-05] and Ammonium Acetate 7.5M DNA & RNA Precipitation Solution [40-4135-05])	40-5130-00	1 Pack
Glycogen Solution 10 mg/mL; 1 mL	40-5112-01	1 mL
Linear Acrylamide Solution (Linear polyacrylamide, LPA; 5mg/mL); 1 mL	40-5113-01	1 mL
LiCI RNA Precipitation Solution (7.5M LiCI, 50 mM EDTA pH 8.0); 50 mL	40-5131-05	50 mL
Sodium Acetate 3M pH 5.5; DNA & RNA Precipitation Solution; 50 mL	40-5132-05	50 mL
Potassium Acetate 3M pH 5.5; DNA & RNA Precipitation Solution; 50 mL	40-5133-05	50 mL
Sodium Chloride 5M DNA & RNA Precipitation; 50 mL	40-5134-05	50 mL
Ammonium Acetate 7.5M DNA & RNA Precipitation Solution; 50 mL	40-5135-05	50 mL
Ammonium Acetate 5 M DNA & RNA Precipitation Solution; 50 mL	40-5136-05	50 mL



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Related Products Ordering Information

DNA & RNA Reconstitution Solutions				
Product	Catalog No.	Unit Size		
DNA & RNA Reconstitution Solutions Pack (contains 50 mL each of DEPC Treated Water [40-3000-05], Nuclease Free Water (DEPC Free) [40-3001-05], TE pH 7.0 [40-5011-05] and RNA Reconstitution Solution[40-5014-05)	40-3000-00	1 Pack		
RNA Reconstitution & Storage Solution (1 mM Sodium Citrate pH 6.4;) 10 X 1.6 mL	40-5014-16	10 X 1.6 mL		
RNA Reconstitution & Storage Solution (1 mM Sodium Citrate pH 6.4); 50 mL	40-5014-05	50 mL		
TE Buffer 1X solution pH 7.0; 50 mL	40-5011-05	50 mL		
TE Buffer 1X solution pH 7.5; 50 mL	40-5012-05	50 mL		
TE Buffer 1X solution pH 8.0; 50 mL	40-5013-05	50 mL		
Nuclease Free Water (DEPC Free); 10 X 1.6 mL	40-3001-16	10 X 1.6 mL		
Nuclease Free Water (DEPC Free;) 50 mL	40-3001-05	50 mL		
Nuclease Free Water (DEPC Free); 500 mL	40-3001-50	500 mL		
Nuclease Free Water (DEPC Free); 1L	40-3001-01	1 L		
DEPC Treated Water; 10 X 1.6 mL	40-3000-16	10 X 1.6 mL		
DEPC Treated Water; 50 mL	40-3000-05	50 mL		
DEPC Treated Water; 500 mL	40-3000-50	500 mL		
DEPC Treated Water; 1L	40-3000-01	1 L		

$Omni-Marker^{TM}$; Molecular Weight Size Standards for Gel Electrophoresis

Product	Catalog No.	Unit Size
Omni- Marker™ DNA 1 kb mw Universal unlabeled; 500 µL	40-3005-05	500 μL
Omni-Marker™ DNA 1 kb mw Universal unlabeled; 1 mL	40-3005-10	1 mL
Omni-Marker™ DNA 100 bp mw Low unlabeled; 500 µL	40-3006-05	500 μL
Omni- Marker™ DNA 100 bp mw Low unlabeled; 1 mL	40-3006-10	1 mL

Loading Buffers; DNA non-denaturing and denaturing buffers

Product	Catalog No.	Unit Size
Loading Buffer 5X BPB/XC non-denaturing; 1 mL	40-3002-10	1 mL
Loading Buffer 5X BPB/XC non-denaturing; 15 mL	40-3002-15	15 mL
Loading Buffer 5X Orange G/XC non-denaturing; 1 mL	40-3004-10	1 mL
Loading Buffer 5X Orange G/XC non-denaturing; 15 mL	40-3004-15	15 mL
Loading Buffer 2X BPB/XC Denaturing for Sequencing; 1 mL	40-5027-10	1 mL
Loading Buffer 2X BPB/XC Denaturing for Sequencing; 15 mL	40-5027-15	15 mL



Electrophoresis Buffers & Hybridization Reagents **Product** Catalog No. **Unit Size** Agarose LE Molecular Biology Grade; 100 g 40-3010-10 100 g Agarose LE Molecular Biology Grade; 500 g 40-3010-50 500 g Hybwash A, Hybridization Wash Solution; 200 mL 40-5020-20 200 mL Hybwash B, Hybridization Wash Solution; 200 mL 40-5021-10 100 mL 100 mL TAE Buffer; 50X Concentrate; 100 mL 40-3007-01 TAE Buffer; 50X Concentrate; 1 L 40-3007-10 1 L TBE Buffer; 5X Concentrate; 1 L 40-3008-10 1 L 200 mL 40-5025-20 10x Washing buffer; 200 mL 100 mL 10% Blocking solution; 100 mL 40-5026-10 100 mL 10x AP Detection buffer; 100 mL 40-5031-10 Lumisol™ I Hybridization Solution; contains formamide; 200 mL 40-5022-20 200 mL Lumisol™ II Hybridization Solution; for non-toxic hybridizations; 200 40-5023-20 200 mL Lumisol™ III Hybridization Solution; for oligo probes; 200 mL 40-5024-20 200 mL



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