Date 3/15-/2013

Objective PCR amplification of triplet repeat region of FMR1 with a modified program (as below):

- Without hot start

- Three step amplification process

- Denaturation temp set according to denat temp grad expt

(see report denat 75-95 comments)

- Annealing temp set according to anng temp grad expt

(see report anng 50-70 comments)

- Extension temp grad 55-75deg/ 2:30mins

Description T

Template used: 100ng genomic DNA NA20230, (53/54 rpt sample from Coriell)

Primers: FP2-RP2

Polymerase: Native Taq Polymerase (Life Technologies)

Solvents: No Solvt rxns

Rxns in 1M NMP

Rxns in 1M Formyl Morpholine

Rxns in 0.5M TMSO Rxns in 1M TMSO

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Reaction	Com	position:

	No Solvt Rxns	1M NMP	1M For. Mor	0.5M TMSO	1M TMSO
Component	Final Conc/ amt				
100					
Taq Buffer	1 X	1 X	1 X	1 X	1 X
MgCl2	1.5 mM				
dNTPs	0.4 mM				
FP2	0.4 uM				
RP2	0.4 uM				
gDNA NA203230	100 ng				
Taq Polymerase	2.5 U				
Water					
NMP		1 M			
For. Mor			1 M		
TMSO				0.5 M	1 M
Betaine					
total	25ul	25ul	25ul	25ul	25ul
denat temp	91.7	76.4	82.9	79.1	76.4
anng temp	51.3/70	68.8	68.8	70.0	70

Cycling Conditions:

76.4degC/ 30secs
70degC/ 1:30mins
55-75degC/ 2:30mins
GOTO 2 40 times

Gradient steps: 75, 73.9, 71.6, 67.6, 62.8, 59, 56.3, 55degC

Gel Electrophoresis: 10% precast polyacrylamide gels (Life Technologies)

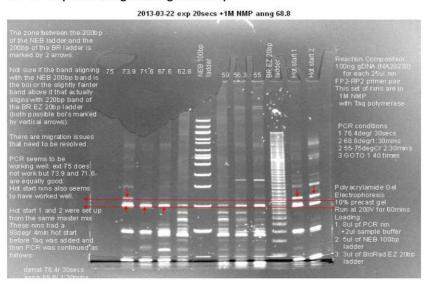
8ul of PCR rxn + 2ul of (5X) sample buffer loaded

Molecular Ladders: NEB 100bp ladder (5ul) BioRad EZ 20bp ladder (3ul)

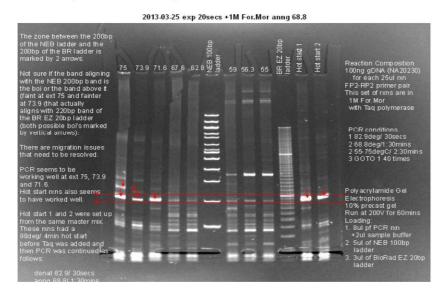
Gel Pictures

Lanes are marked according to the annealing temp of the particular rxn. Expected product size is 224bp

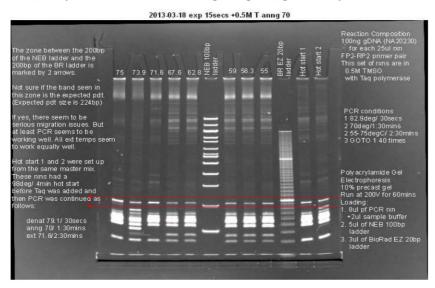
FP-RP-Taq-NMP anng 68.8deg 20sec exp



FP-RP-Taq-For.Mor 20sec exp



FP-RP-Taq-0.5M TMSO denat 79.1deg- anng 70deg 15sec exp



FP-RP-Taq-1M TMSO anng 70deg 20sec exp



Comments

As mentioned previously,

even tho' the 1000bp band migrate similarly for the NEB 100bp ladder and the BioRad EZ 20bp ladder, there is a marked difference in the 200bp band for both ladders

Because of this and the slight differences in the sizes of bands obtained in the 200bp range, a box has been drawn keeping the two ladders as boundaries.

It is possible that for the test rxns, the bands inside this box are the expected band.

If yes, there seem to be serious migrations differences for the same expected pdt under different conditions.

If all these ARE the expected band, then the PCR seems to be working quite well.

Conditions selected from ext grad are as follows:

	denat	anng	ext temp			
	temp	temp	1 st	2 nd	3 rd	
1 Taq + 0.5M T	79.1	70.0	75.0	73.9	71.6	
2 Taq + 1M T	76.4	70.0	72.0			
3 Taq + For. Mor	82.9	68.8	75.0	73.9	71.6	
4 Taq + NMP	76.4	68.8	73.9	71.6		
5 Taq - No solvt	91.7	70/51.3	no data yet*			

^{*:} the PCRs did not work, presumably, Taq expired. Both PCRs need to be rpt'd with new Taq.