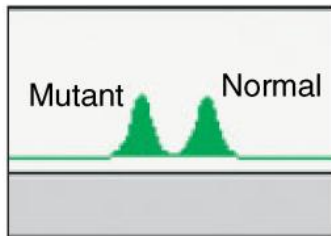


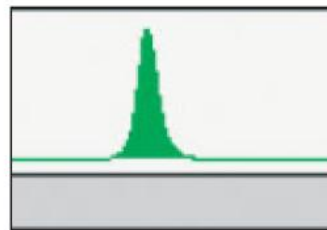
SINGLE CELL SEQUENCING

beta-Thalassemia codon 41-42

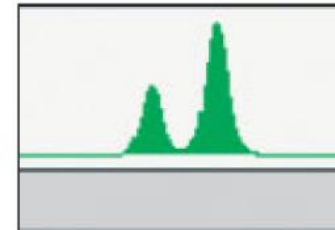
EA



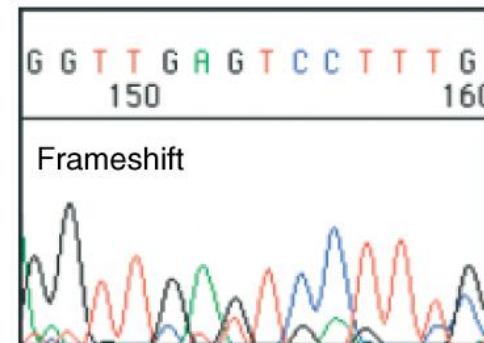
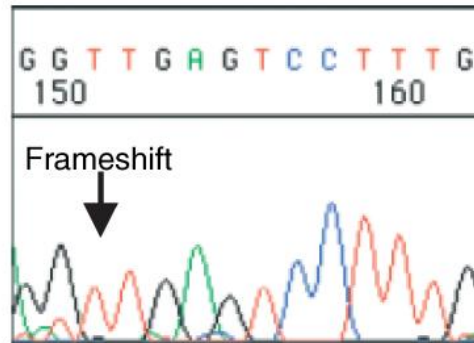
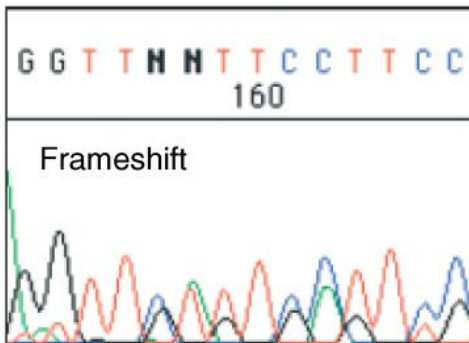
ADON



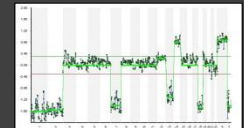
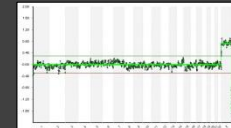
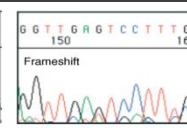
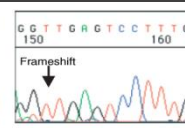
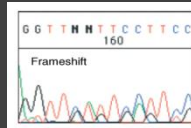
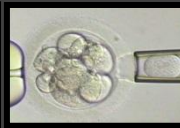
PAM



Fluorescent
PCR



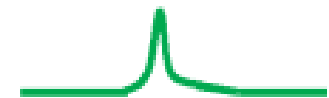
Sequencing



MINI-SEQUENCING

3' - A T A G T C T T G A T C -5'

T C A G A **A**



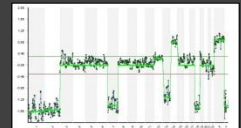
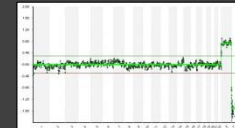
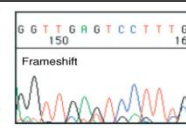
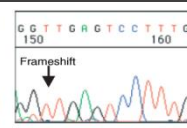
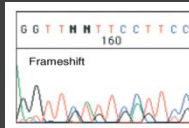
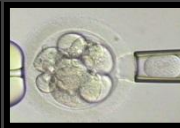
A

3' - A T A G T C T A G A T C -5'

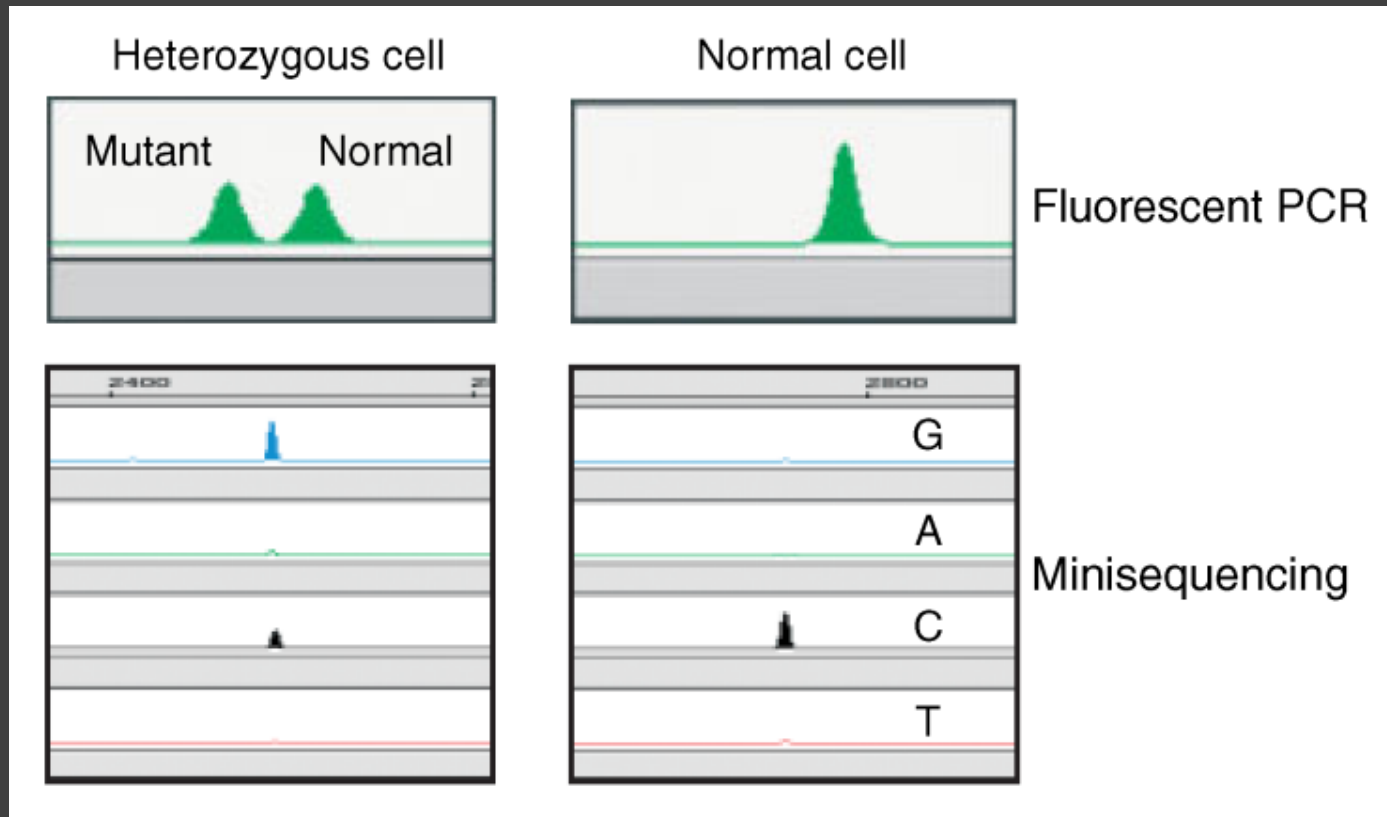
T C A G A **T**

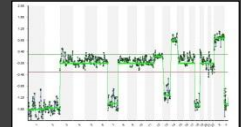
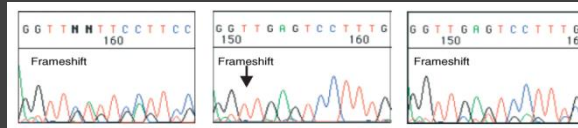


T



SINGLE CELL MINI-SEQUENCING beta-Thalassemia codon 41-42





CONTAMINATION

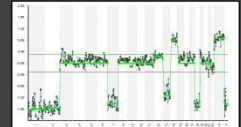
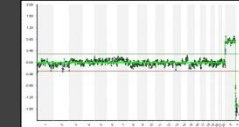
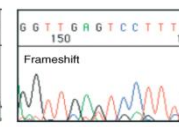
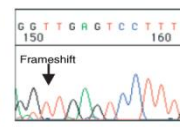
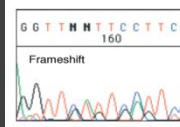
- Dominant disorders

- interpret normal as affected*



- interpret affected as affected





CONTAMINATION

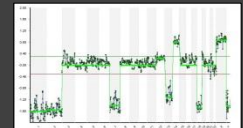
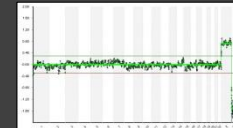
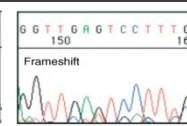
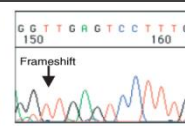
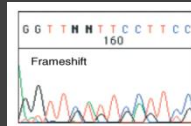
■ Recessive disorders

- interpret normal as heterozygous (carrier)



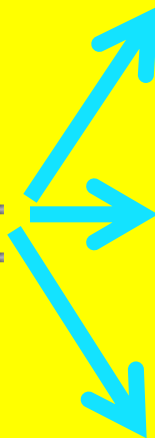
- interpret affected as heterozygous (carrier)*





Allele Dropout

Heterozygous cell



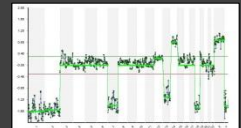
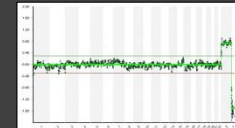
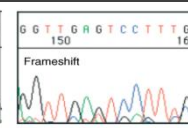
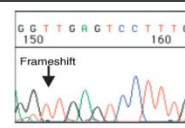
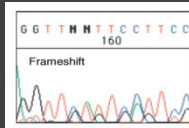
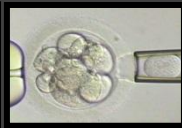
Amplification of both alleles
Heterozygous genotype predicted



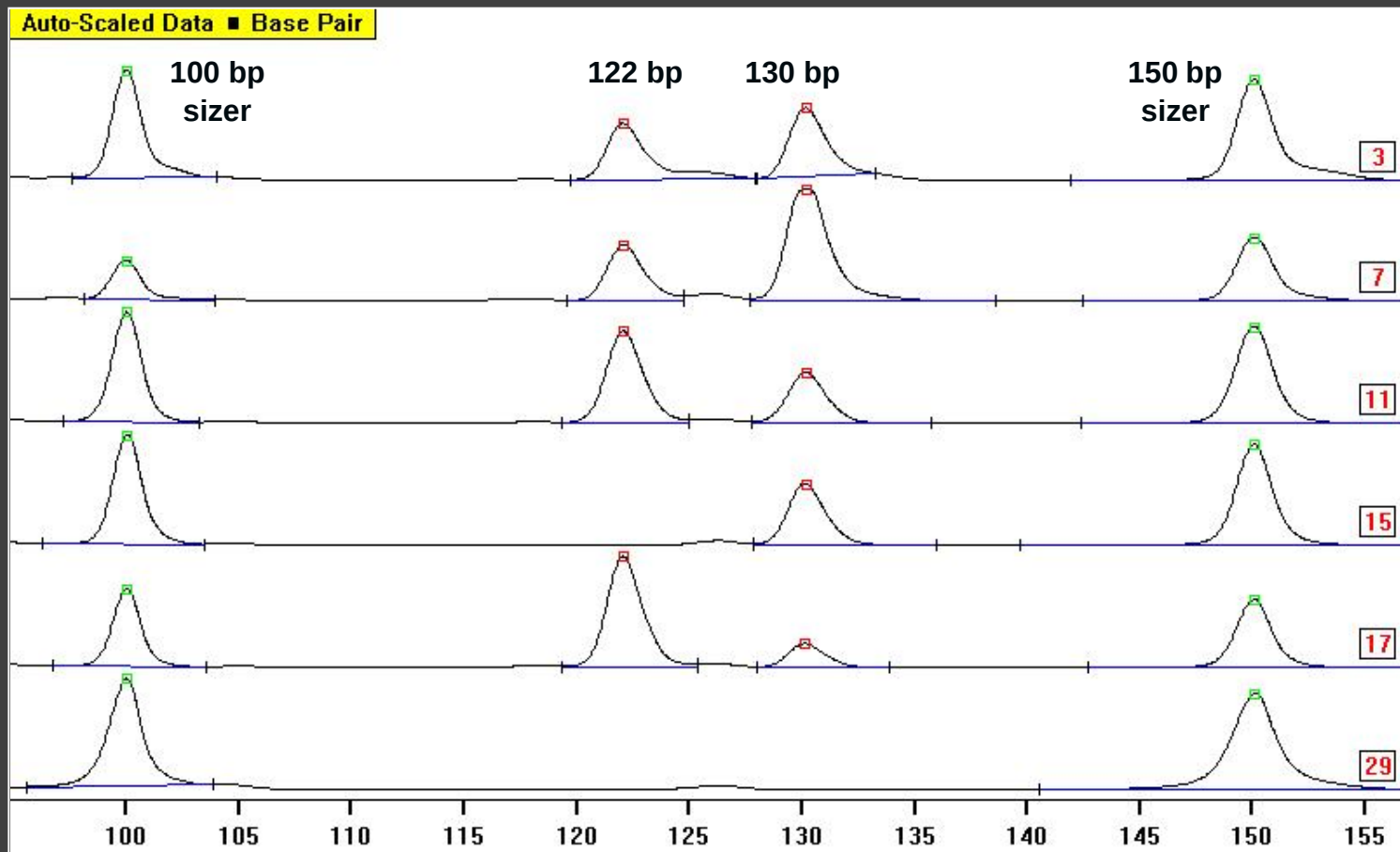
ADO of mutant allele
Amplification of normal allele only
Homozygous normal genotype predicted

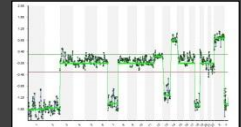
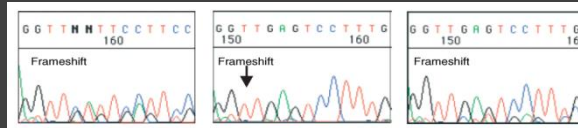


ADO of normal allele
Amplification of mutant allele only
Homozygous affected genotype predicted



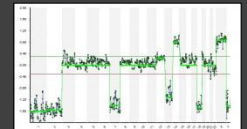
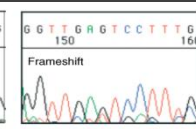
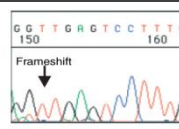
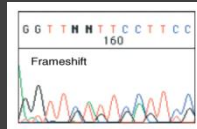
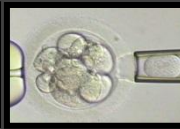
ALLELE DROP-OUT





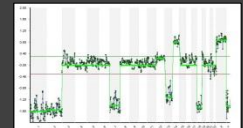
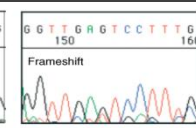
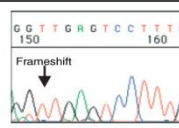
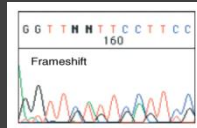
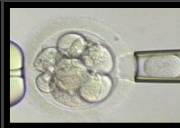
PGD for Single Gene Disorders

- ❑ Duchenne muscular dystrophy(Liu et al., 1995)
- ❑ Fragile X syndrome (Sermon et al., 1999)
- ❑ Tay Sachs disease (Gibbons et al., 1995)
- ❑ Marfan's syndrome (Harton et al., 1996)
- ❑ Myotonic dystrophy (Piyamongkol et al., 2001b)
- ❑ Charcot Marie Tooth type 1A (De Vos et al., 1998)
- ❑ Familial adenomatous polyposis coli (FAPC) (Ao et al., 1998)
- ❑ Huntington's chorea (Sermon et al., 1998)



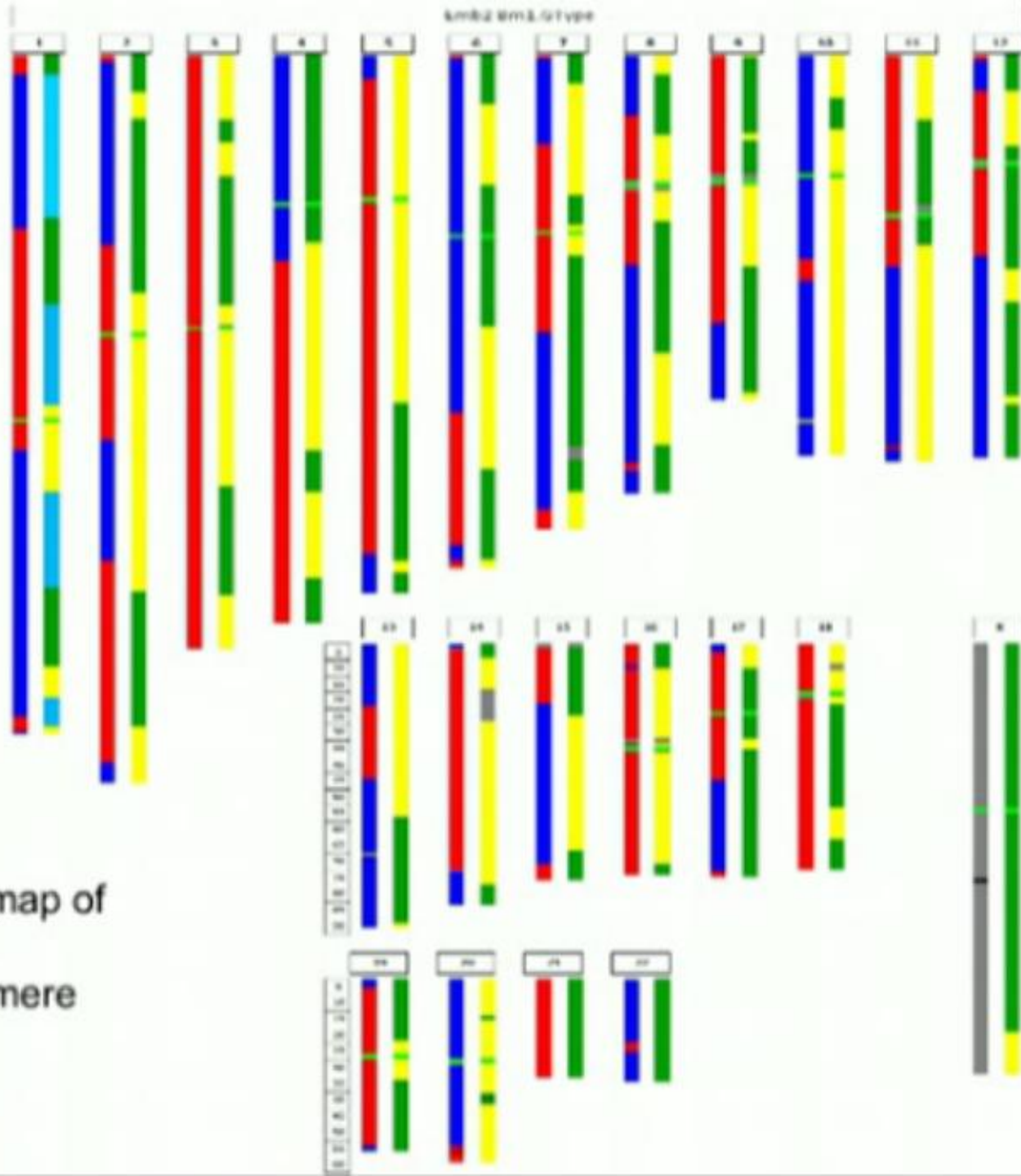
PGD for Single Gene Disorders

- ❑ Severe inherited skin diseases (McGrath and Handyside, 1998)
- ❑ Sickle cell anaemia (Rechitsky et al., 1998)
- ❑ Spinal muscular atrophy (Dreesen et al., 1998)
- ❑ β -thalassaemia (Kanavakis et al., 1999; Kuliev et al., 1999; Piyamongkol et al., 2006)
- ❑ Congenital adrenal hyperplasia (Van de Velde et al., 1999)
- ❑ Lesch Nyhan syndrome (Ray et al., 1999),
- ❑ Medium chain acyl CoA dehydrogenase (MCAD) deficiency (Ioulianos et al., 2000)



PGD of beta-Thalassaemia

- ❑ Sexing - Handyside, 1989
- ❑ beta-Thalassaemia: RFLP - Kuliev, 1998
- ❑ beta-Thalassaemia: DGGE - Kanavakis, 1999
- ❑ beta-Thalassaemia: RFLP - De Rycke, 2001
- ❑ beta-Thalassaemia: RFLP & reverse dot-blot - Chamayou, 2002
- ❑ beta-Thalassaemia: sequencing - Hussey, 2002
- ❑ beta-Thalassaemia: WGA + reverse dot-blot - Jiao, 2003
- ❑ beta-Thalassaemia: F-PCR - Piyamongkol, 2006



Karyomap of
single
blastomere

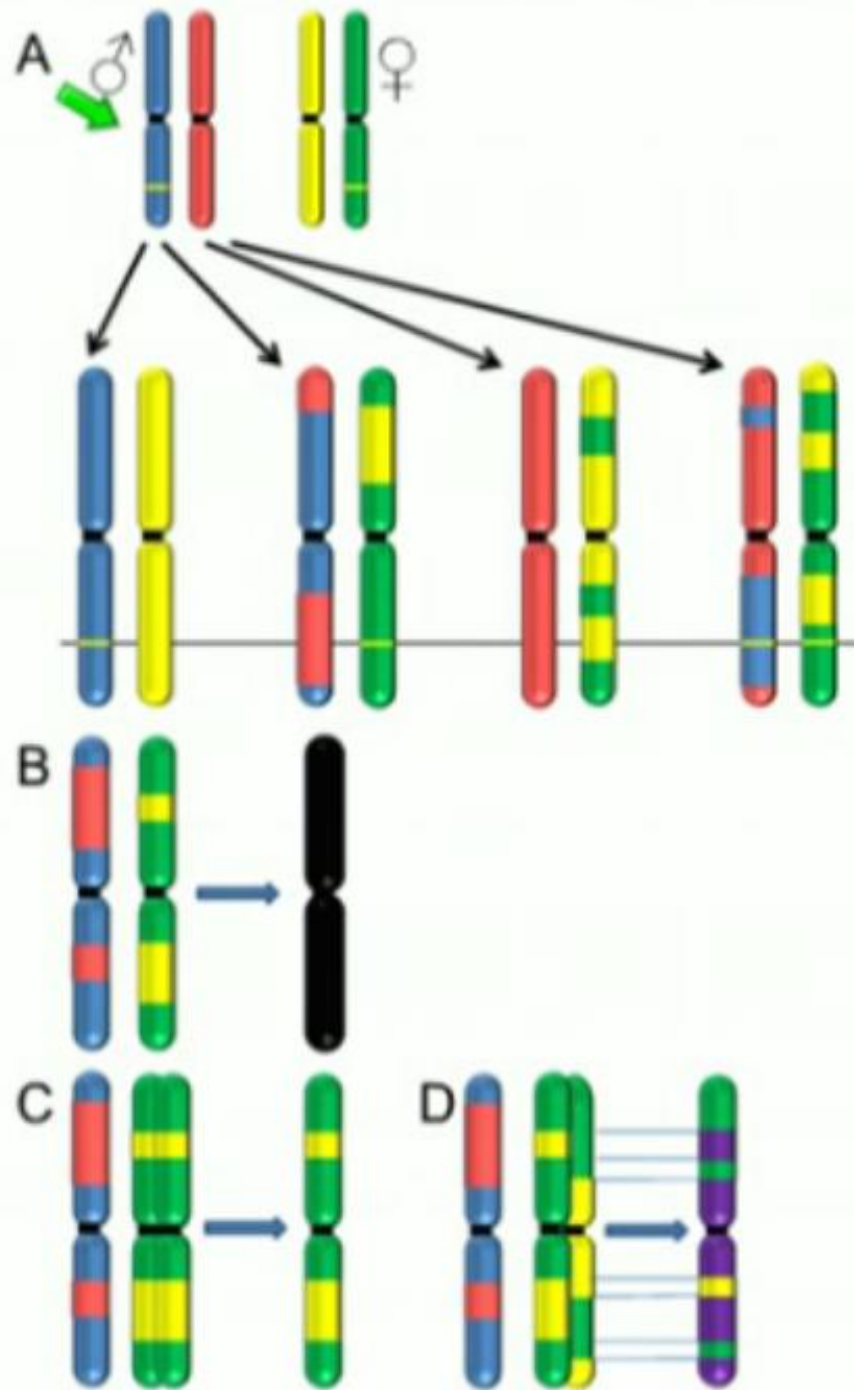
SNP ID & POSITION

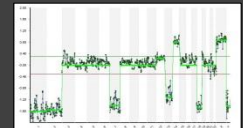
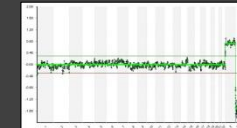
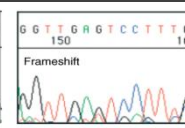
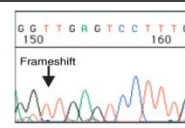
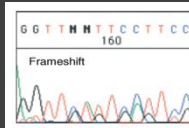
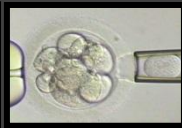
SNP GENOTYPES

KARYOMAP

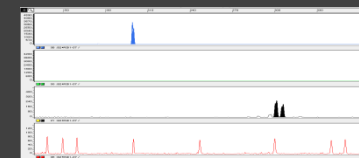
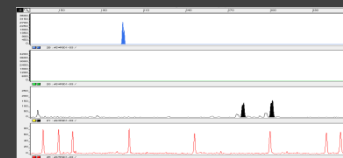
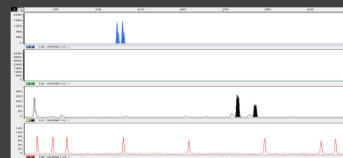
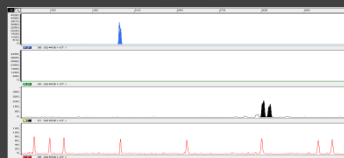
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						P1	M1		P1	P2	M1	M2
rs3094315	1	742429	AB	X	AB	AA		AA				
rs2073813	1	743404	BB		BB	BB		BB				
rs2905040	1	760079	AB	→	AA	AA		AB	→	B		
rs12124819	1	766409	BB		BB	BB		BB				
rs2980314	1	771121	BB	→	AB	BB		BB				
rs6684487	1	781716	BB	→	AB	AB		AB			B	
rs4245756	1	789326	BB		BB	BB		NC				
rs12086311	1	798632	BB	→	AB	BB		BB				
rs13303077	1	806480	BB		BB	BB		BB				
rs28625089	1	830190	AA	→	AB	AA		AA				
rs4475691	1	836671	AB	→	BB	AB		BB				
rs28587382	1	842635	BB		BB	BB		BB				
rs13303029	1	848359	BB		BB	BB		NC				
rs2340589	1	854618	BB	→	AB	BB		AB		B		
rs28576697	1	860508	BB		BB	BB		NC				
rs1110052	1	863421	AA	→	AB	AB		AB			B	
rs7523549	1	869180	BB		BB	BB		BB				
rs2272756	1	871896	AB	→	BB	AB		AB		A		
rs28711536	1	878083	AB	→	AA	AB		AB		B		
rs13303010	1	884436	BB		BB	BB		BB				
rs3935086	1	890593	AB	→	AA	AA		AA				
rs13302925	1	893896	BB		BB	BB		BB				
rs2340594	1	899418	BB	→	AB	AA		AA				
rs13303355	1	905090	AA		AA	AA		AA				
rs6693747	1	911872	AA	→	AB	AB		AB			A	
rs9777703	1	918699	BB		BB	BB		NC				

Karyomapping combines genome wide linkage based detection of single gene defects (A) with chromosomal aneuploidy including monosomy/deletions (B) and trisomies involving inheritance of two different meiotic chromosomes from one parent (D). Chromosome duplication is not detected (C).





PGD of beta-thalassaemia codon41-42

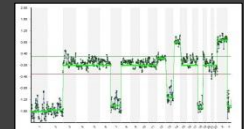
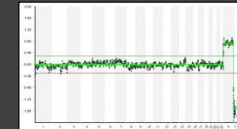
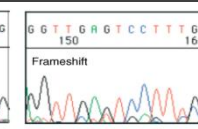
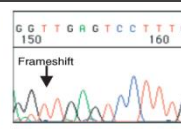
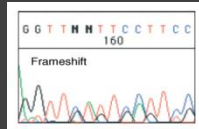
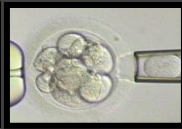


normal
ET

heterozygous
ET

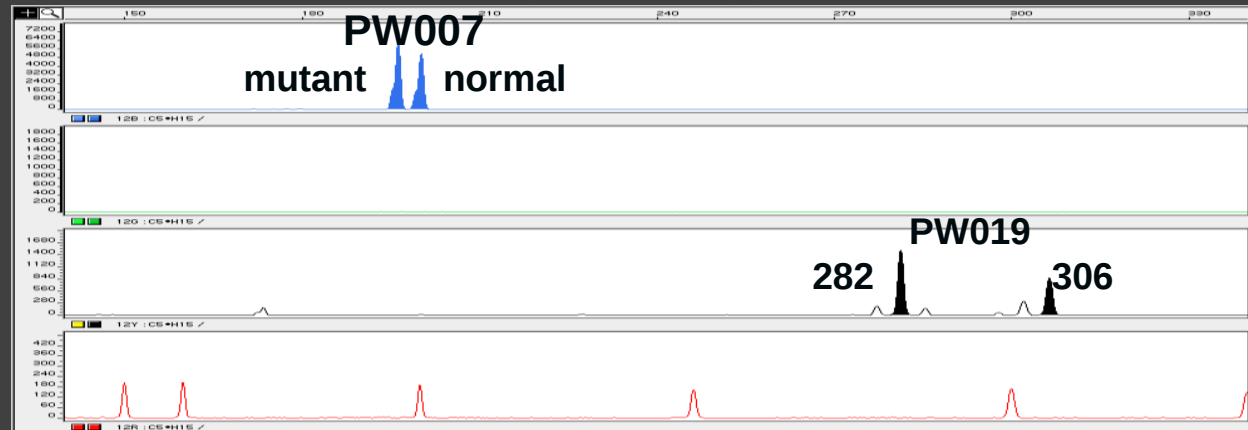
affected

normal
ET

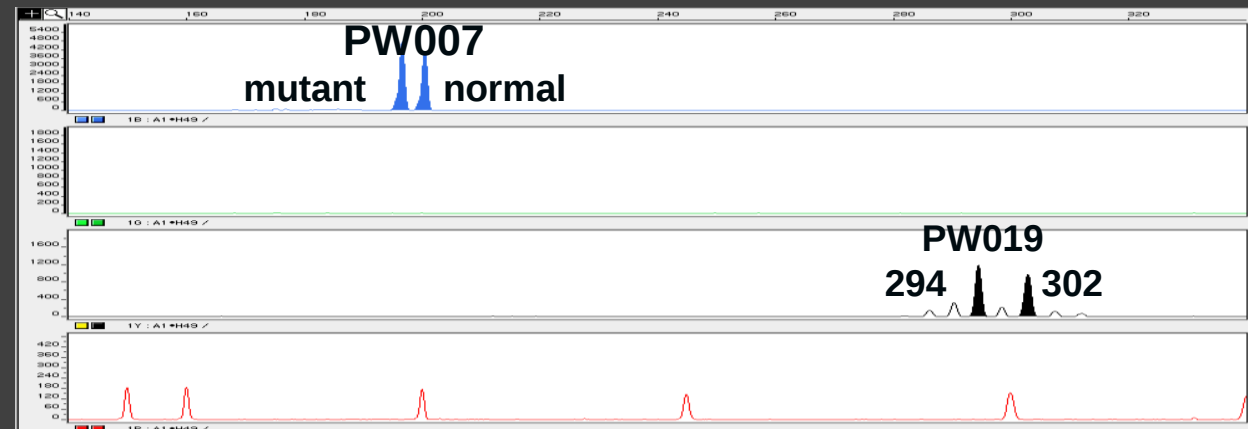


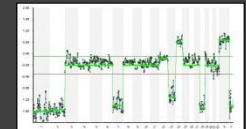
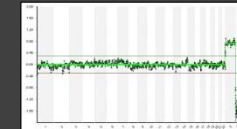
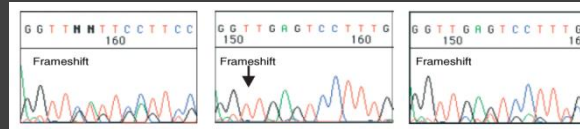
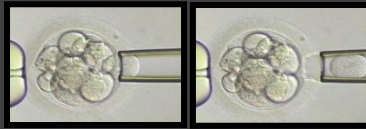
Work-up of PGD Protocol for beta-Thalassaemia codon 41-42

Father



Mother



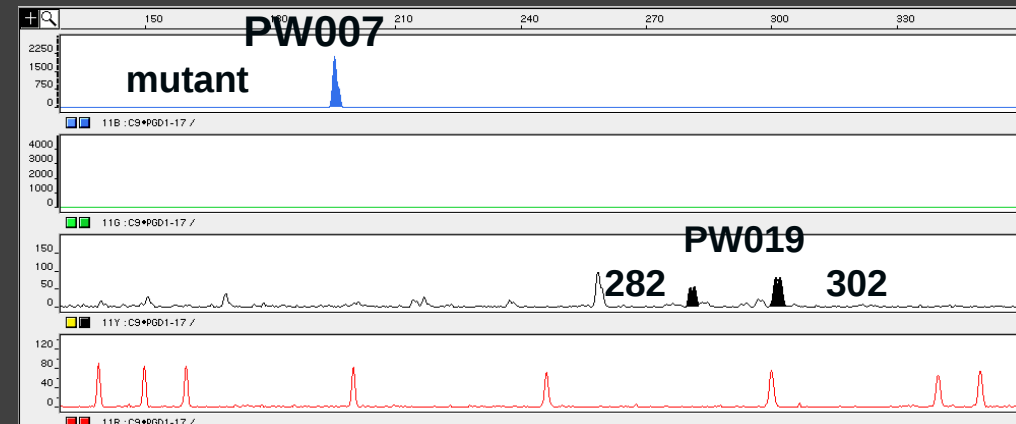
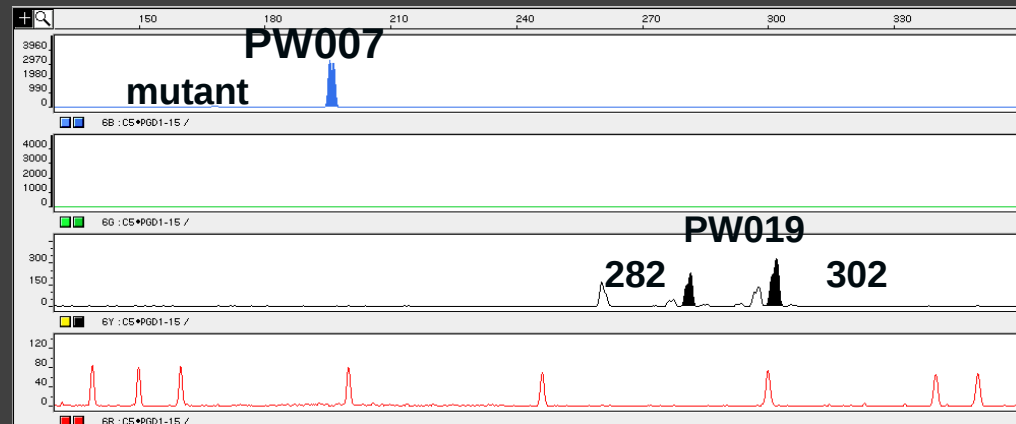


PGD of beta-Thalassaemia codon 41-42

Embryo 1

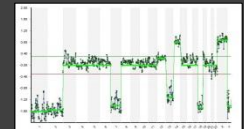
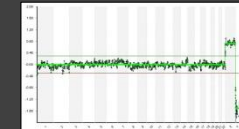
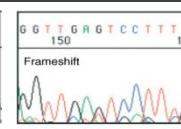
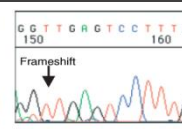
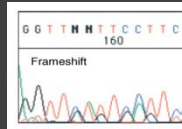
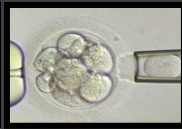
8-cell

Affected



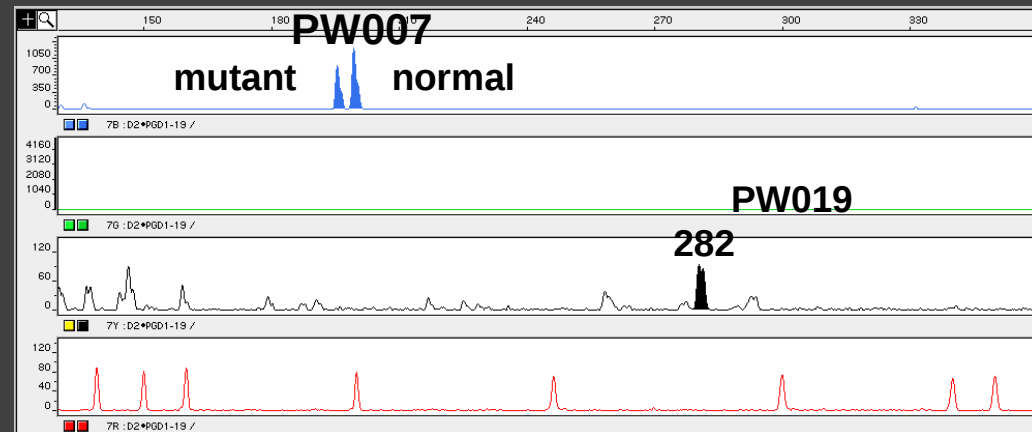
Confirmation -

Affected

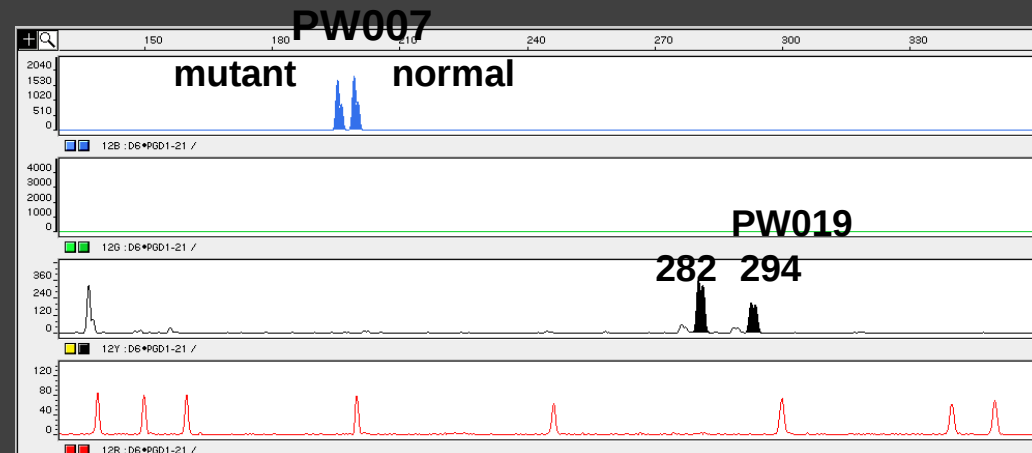


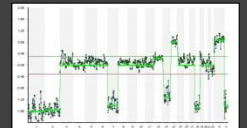
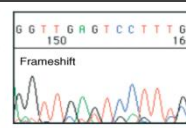
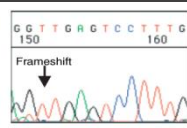
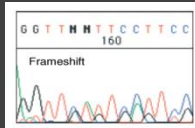
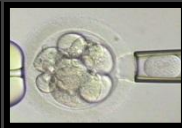
PGD of beta-Thalassaemia codon 41-42

Embryo 8
8-cell
Heterozygous



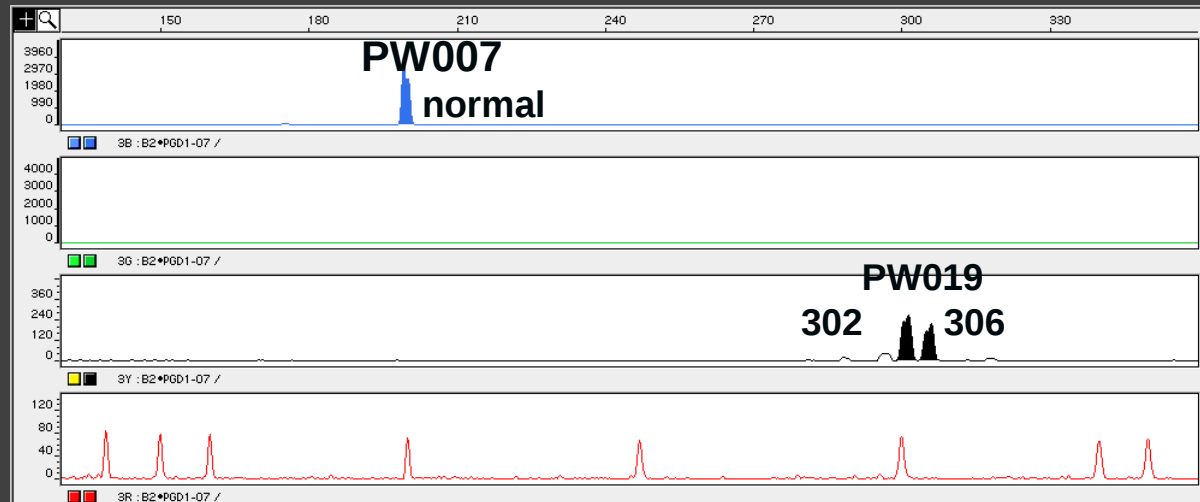
Confirmation -
Heterozygous



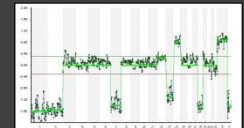
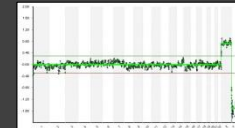
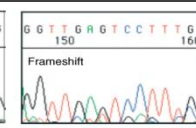
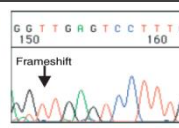
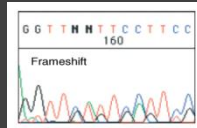
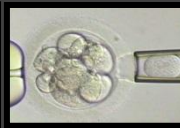


PGD of beta-Thalassaemia codon 41-42

Embryo 4
8-cell
Normal



ET



Work-up of PGD Protocol for alpha-Thalassemia-SEA

