# Principles of Genetic Testing (DMD)

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# Duchenne / Becker Muscular Dystrophy

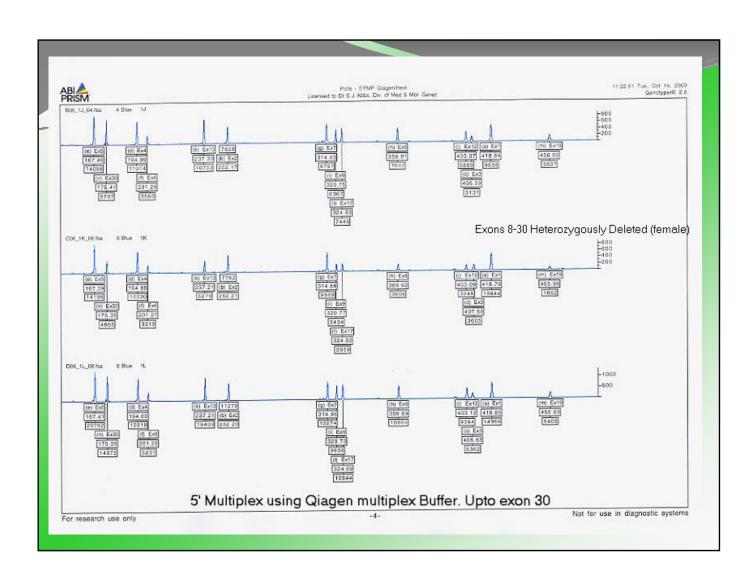
- >Highly elevated CK in young children
- >X-linked Dystrophin Gene
  - 2.4Mb 79 Exons
- >Females are carriers, Boys are affected
- ➤ Vast mutation spectrum

#### DMD/BMD

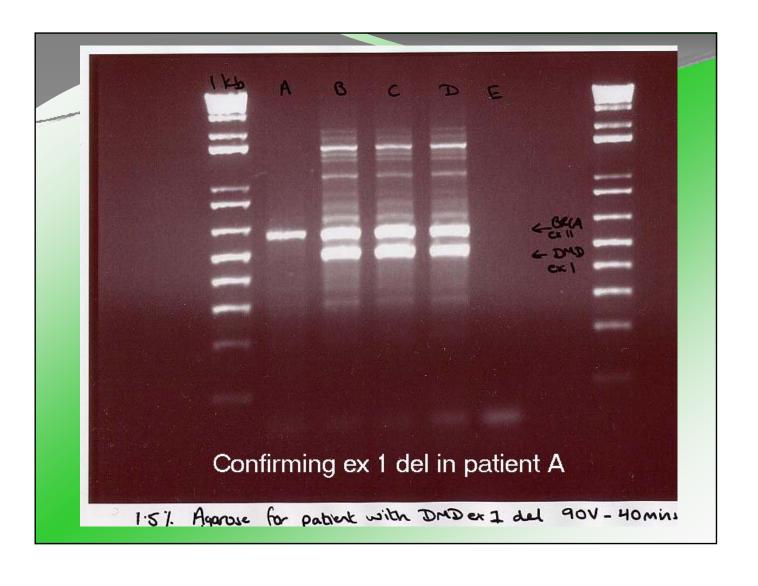
- > Mutations
  - Point mutations
  - Whole exon deletions/duplications
  - Small insertions/deletion
  - Non sense, frameshift (truncating protein)
- >Upto 10% Mosaicism
  - No mutation detected in Mother of boys with known mutations

# Screening - QF PCR

- DNA Quality Test
- 2 Multiplexes of 14 Exons
- ABI 3100 Genetic Analyser
- Analysed on Software and Excel sheet
- Detect female carrier status, duplicated males/ females
- ➤ Detects 99% del, 96% Dup & >1bp ins/del
- ➤ However, only accounts for ~ 65-70% dystrophin mutations



							5'1	FMP QI	AGEN2	STATS	S						
							- 551			SALEATA							
AME:	2K 40						RESULT										
	CONTROL SAMPLE							EXO	NS								
EXONS			Pm	2	3	4	5	6	7	8	9	12	13	17	19	30	MEAN
Pm	13332	12495		1.12	1.03	1.19	1.13	1.22	1.07	2.33	2.1	2	2.24	2	2.02	2.26	1.67
2	10520	8818	0.89		0.92	1.06	1.01	1.09	0.96	2.09	1.88	1.79	2.01	1.79	1.81	2.03	1.49
3	5174.5	4725	0.97	1.09		1.16	1.1	1.19	1.04	2.27	2.05	1.95	2.18	1.95	1.97	2,21	1.62
4	16270	12838	0.84	0.94	0.86		0.95	1.02	0.9	1.96	1.77	1.68	1.89	1.68	1.7	1.91	1.39
5	18532.5	15403	0.89	0.99	0.91	1.05		1.08	0.95	2.07	1.86	1.77	1.99	1.77	1.79	2,01	1.47
6	5381	4145	0.82	0.92	0.84	0.98	0.93		0.88	1.92	1.73	1.64	1.84	1.64	1.66	1.86	1.36
7	12427.5	10872	0.93	1.04	0.96	1.11	1.05	1.14		2.18	1.96	1.87	2.09	1.87	1.88	2.11	1.55
8	10267.5	4125	0.43	0.48	0.44	0.51	0.48	0.52	0.46		0.9	0.86	0.96	0.86	0.87	0.97	0.67
9	9109	4061	0.48	0.53	0.49	0.57	0.54	0.58	0.51	1.11		0.95	1.07	0.95	0.96	1.08	0.75
12	8271	3879	0.5	0.56	0.51	0.59	0.56	0.61	0.54	1.17	1.05		1.12	1	1.01	1.13	0.80
13	15860	6630	0.45	0.5	0.46	0.53	0.5	0.54	0.48	1.04	0.94	0.89		0.89	0.9	1.01	0.70
17	9678	4538	0.5	0.56	0.51	0.59	0.56	0.61	0.54	1.17	1.05	1	1.12		1.01	1.13	0.80
19	5497.5	2552	0.5	0.55	0.51	0.59	0.56	0.6	0.53	1.16	1.04	0.99	1.11	0.99		1.12	0.79
30	14348	5939	0.44	0.49	0.45	0.52	0.5	0.54	0.47	1.03	0.93	0.88	0.99	0.88	0.89		0.69
		MEAN	0.66	0.75	0.68	0.8	0.76	0.83	0.72	1.65	1.48	1.41	1.59	1.41	1.42	1.6	



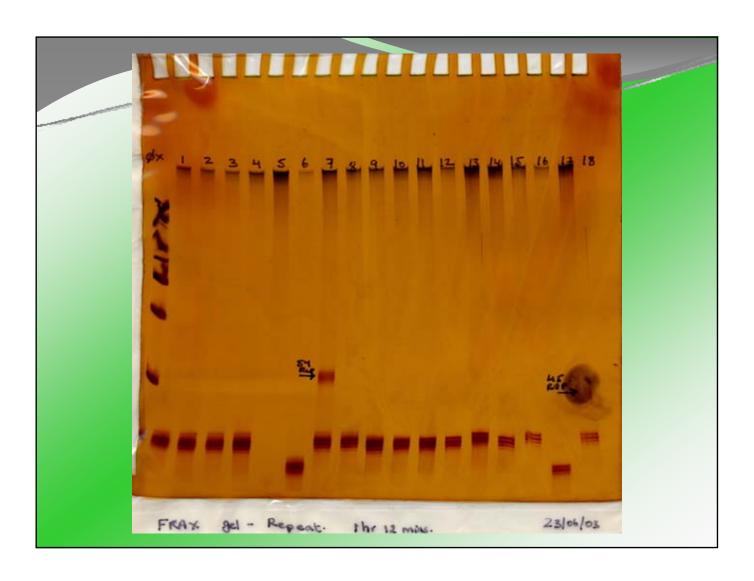
#### Full Screen

- >Full Screen
  - Full DMD gene screening (12 multiplexes,
    7 exons each)
  - Heteroduplex analysis
  - QF PCR all exons
  - Both done on ABI 3100 Genetic Analyser
- ➤ Gene Sequencing
- >PTT (Muscle biopsy mosaicism)

# Fragile X Syndrome

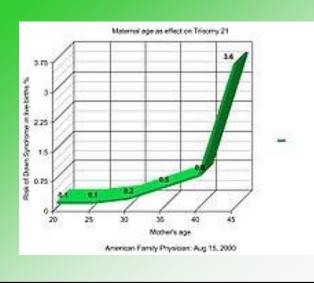
- ➤ Mutation on FMR1
  - CGG Repeat > 200
- ➤ Elongated face, large or protruding ears, flat feet, MR, speech problems
- ➤ Affects 1 in 3600 males and 1 in 4000-6000 Females
- ➤ PCR for up to 200 repeats and Southern Blot





### Down's Syndrome

- >1 in 800 to 1000 births worldwide
- ➤Trisomy 21 (full or part)
  - Meiotic Non-dysjunction
  - Full Trisomy 21 95% of Down's
  - 88% maternal
- ➤ Maternal Age Risk
- >Amniocentesis, CVS
  - Karyotyping, FISH
  - QF-PCR
- ➤ Triple Test
  - bHCG AFP, E2



# PCR –GOLD STANDARD BY WHO AND CDC

- ➤ Rapid, Ultra Sensitive and Specific
- ➤ CHIK, H1N1, MTB, Chlamydia etc.,
- ➤ Detects DNA unique to pathogen
- Accurately pick-up virus even in mild infections
- ➤ Used for Viral Loads (Hep B, C, HIV)
- >Human Papilloma Virus cervical cancer
- **≻**Septicaemia

