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Constitutional postnatal aCGH & DD/MR/CA cohort

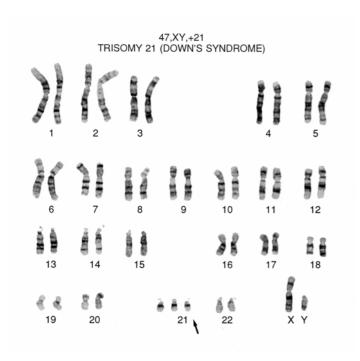
- Potential impact of aCGH on NHS Service Delivery Models*
- Risks associated with too slow an implementation of aCGH as a first lie test



POST-NATAL APPLICATIONS OF ARRAY-CGH



WHAT IS THE POTENTIAL FOR ARRAY-CGH?



KARYOTYPE ~ 5 Mb



ARRAY-CGH ≤50 Kb*

Resolution* increases from the chromosome band to the DNA base pair





Category	%
Abnormal	26
Normal	74
Total	100

^{* 4}x44k customised oligo array



DD/MR/CA - % abnormalities detected*

Category	No	% abnormals
Confirmed de novo	27	22.1%
Segregating abnormalities**	10	8.2%
CNV* ◆	37	30.3%
Abnormal in progress	48	39.3%
Sub-total abnormals	122	100%
** Same abnormality found in parent with s	imilar pheno	type or
penetrance of abnormality known to be vai	riable e.g. de	l(1)(q21.1)
* Same abnormality found in apparently ph	enotypically	normal parent



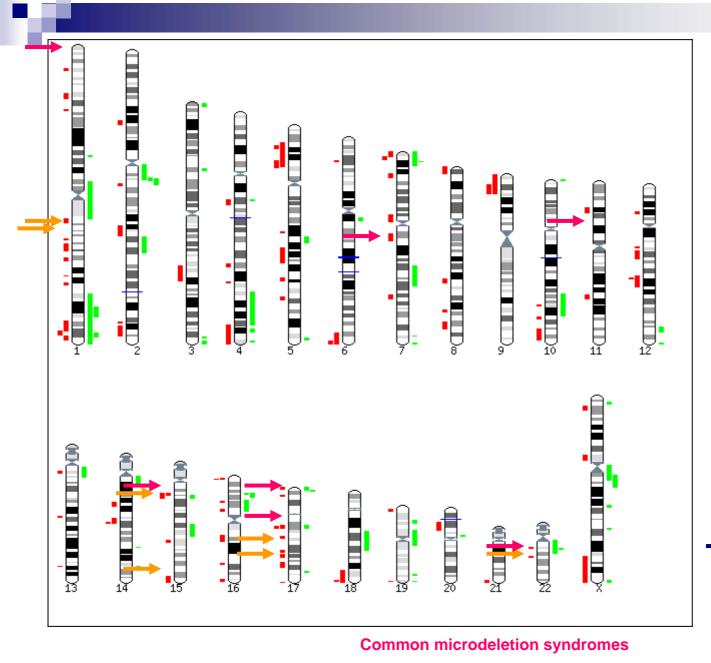
^{*12} month audit 2008; all retrospective cases



Resolution of aCGH (DD/MR/CA)

- 44k oligo aCGH gives ≥20% diagnostic enrichment of "pathogenic imbalances" over conventional karyotyping
- Will increasing this resolution provide significant improvements in diagnoses?
- NHS implementation could/should provide flexibility for platform variation





28th Jan 2009

161 patients registered with DECIPHER

deletions

amplifications

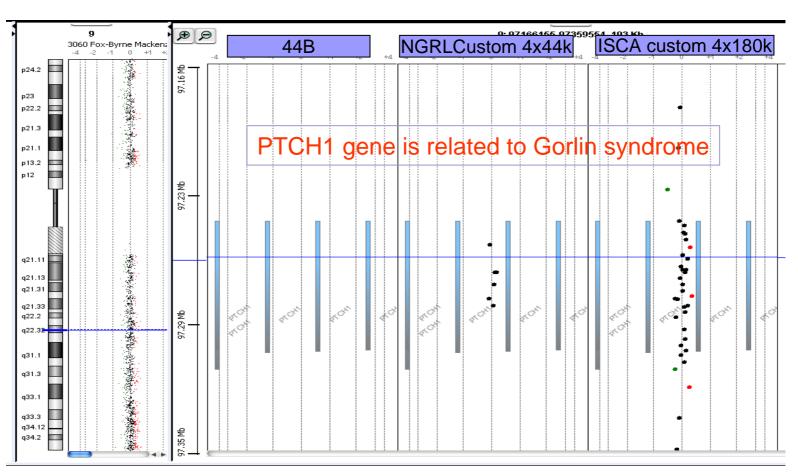
____ Translocation breakpoint

Recurring array-cgh copy number changes



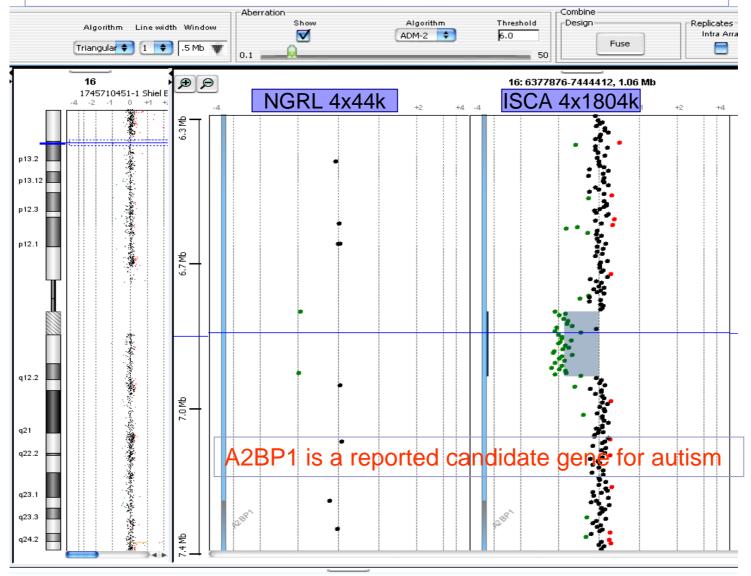
Array design – why is it important?

Example 1: some regions are not covered by catalogue array.





Higher resolution increases confidence of calls but increases CNV "noise".







- Has a proven diagnostic enhancement utility in a retrospective <u>post-natal</u> setting
- Will detect <u>all</u> chromosome <u>imbalances</u> currently diagnosed by karyotyping

Will not detect balanced chromosome rearrangents



De novo CCRs and translocations **Array detected abnormalities (+abnormal phenotype) CCR** Study **Deletions** Normal Total DeGregori 16 18 Present 6 24 (96%) **Total** 3 25 **ABSCR**

Deletions

11

4

6

21 (39%)



Total

27

16

10

53

Normal

16

12

32

Study

DeGregori

Baptista

Gribble

Total

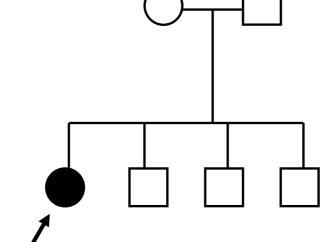


Karyotyping v array-cgh in the DD/MR/CA cohort

■ A thought.....



Pedigree

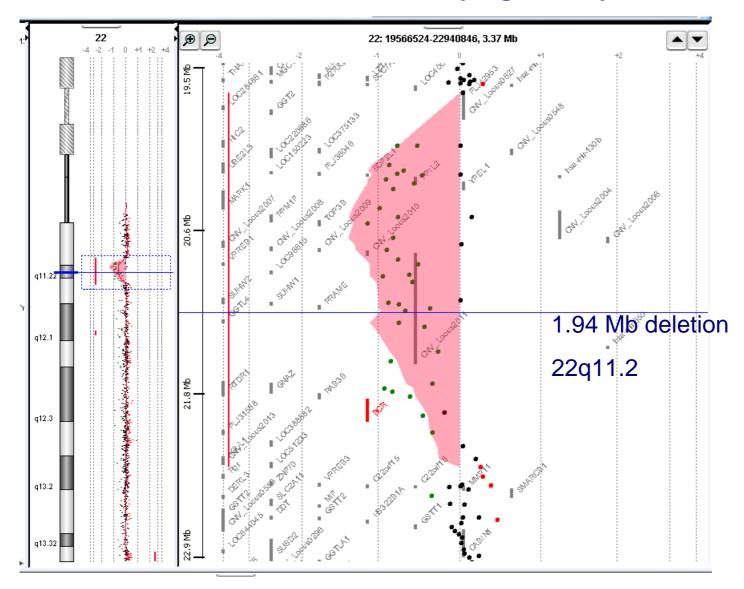


- 12 yr old female
- developmental delay
- dysmorphic features

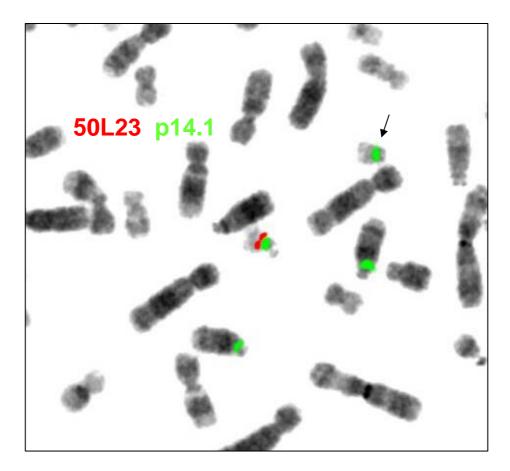
Karyotype January 2008: 46,XX (Q7) – no parental follow-up



Proband referred for array-cgh – July 2008



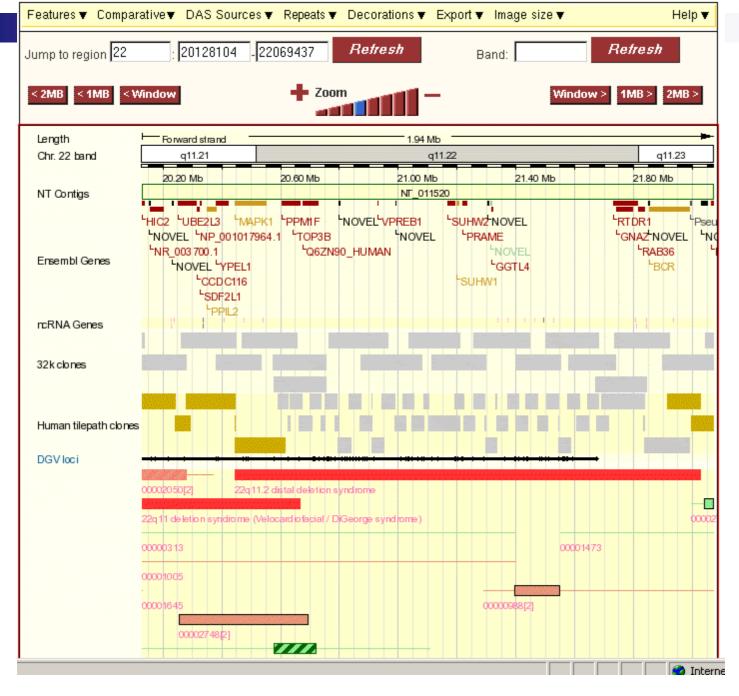




22q11.2 Distal Deletion: A Recurrent Genomic Disorder Distinct from DiGeorge Syndrome and Velocardiofacial Syndrome

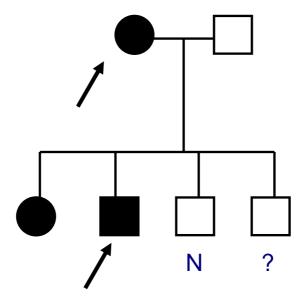
Shay Ben-Shachar et al







Pedigree



Karyotype January 2008: 46,XX (Q7)

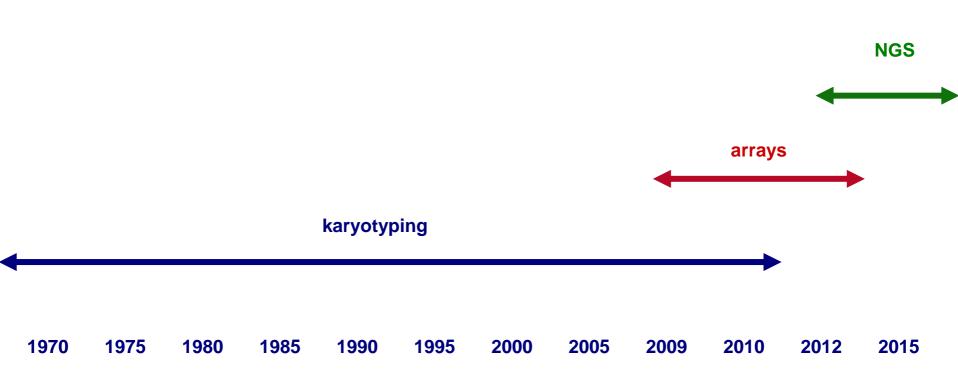


Recurrence risk ~ 50%

- 44k oligo array-cgh currently detecting ≥20% enrichment of *de novo* abnormalities not detectable by conventional karyotyping
- Will soon be actionable if an array had not been performed and a second affected child is born?
- In Wessex, this amounts to ~200 prospective "missed" diagnoses per annum if we continue with karyotyping as the front line test.



aCGH - implementation timeframe







Recommendation

- DD/MR/CA ascertainment groups must have array-cgh at minimum of 44k oligo as first line test
- Low resolution (? automated) karyotyping introduced for all other referral categories
- Retraining needed for bioinformatics and data analyses.
- U.K. NHS consortium approach to purchasing, platform and analytical stages of array-cgh





What is the right balance between array-CGH and karyotyping?

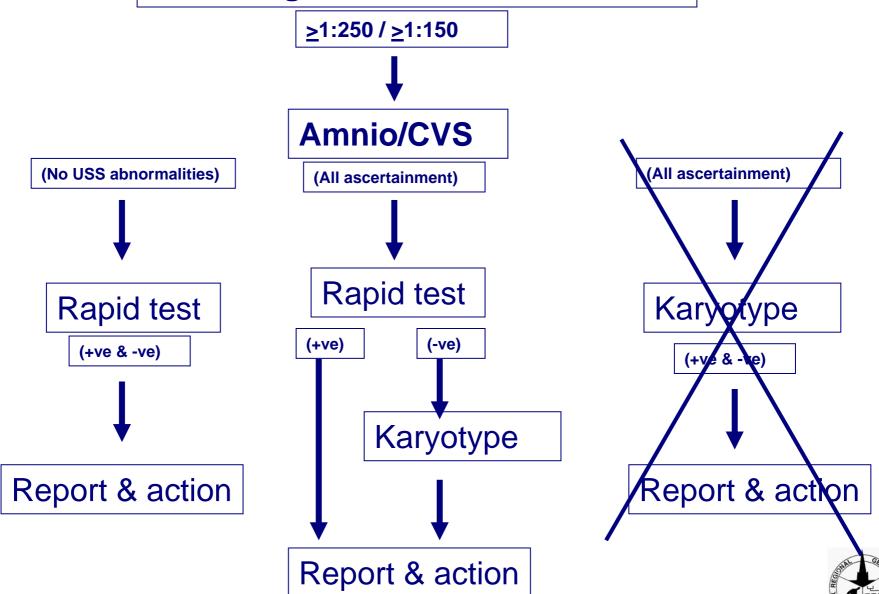
Difficult choices will have to be made about current service configurations.



Pre-natal application of array-cgh



Screening and DS risk assessment



North East North West 5,035,849 Yorkshire and The Humber **Regions providing** rapid testing as stand alone test East Midlands West Midlands East of England South Central 7,428,590 South East Coast South West



Requirements for routine prenatal array-cgh - 1

- Specifically designed array required
- Fast turn around times
- Fast throughput and bioinformatic tools
- Ability to distinguish between CNVs and pathogenic changes





Requirements for routine prenatal array-cgh - 2

- Rapid trio (family studies) required in relatively high proportion of cases
- Interpretation for microdeletions with know variable penetrance (e.g. 22q11.2, 1q21.2)
- Standardised protocols and platforms so that data can be shared rapidly
- Laboratory follow up studies conducted as rapidly as the array itself



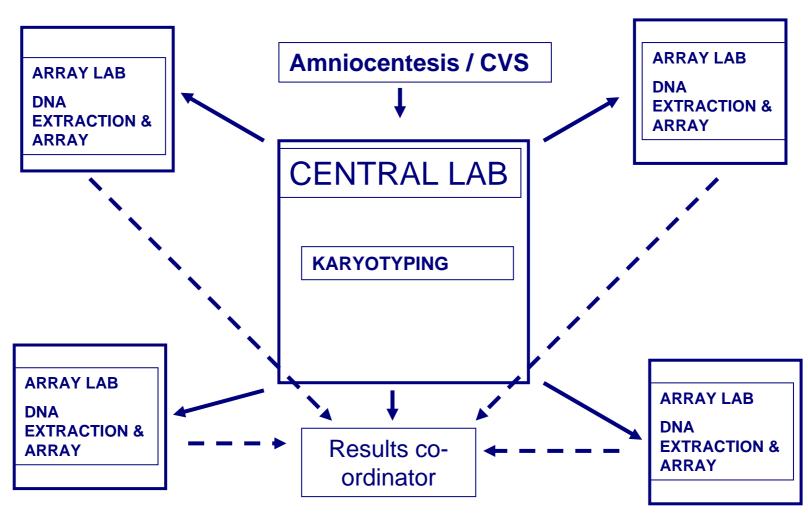
USA PROSPECTIVE ARRAY-CGH PRENATAL TRIAL

- 4 X 44K OLIGO "CUSTOMISED" ARRAY
- PERICENTRIC AND TELOMERIC DENSE COVERAGE
- NIMBLEGEN AND AFFY PLATFORMS BEING EVALUATED

- •4,000 PREGNANCIES
 - ~2,000 "ROUTINE" (MATERNAL AGE)
 - ~2,000 WITH AT LEAST 2 USS ABNORMALITIES



USA PROSPECTIVE ARRAY PRENATAL TRIAL





Screening and DS risk assessment

(No USS abnormalities)



Rapid test

(+ve & -ve)

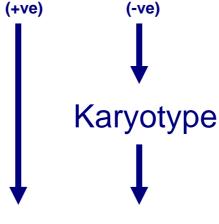


Report & action



Amnio/CVS

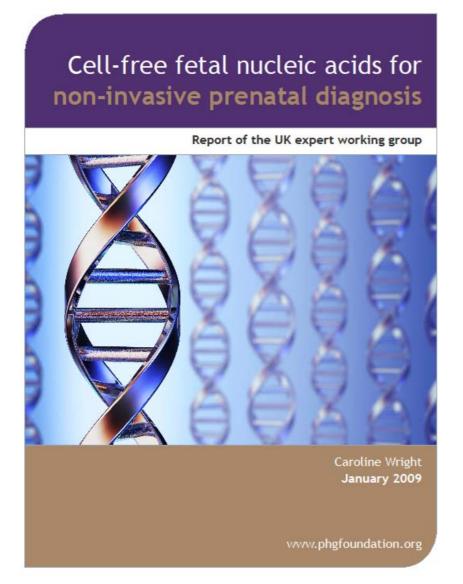
Rapid test



Report & action









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www.rapid.nhs.uk



NIPD at 8 – 10 weeks gestation (All ascertainments or population based) (+ve) (-ve) Abn USS Rapid test? will account for ~80% of abnormalities rapid test confirmations centralised Karyotype • role for cytogenetics in rearrangements & abnormal scans fast turnaround and Report & action rationalisation Report & action

array-cgh/shotgun

sequencing





The End.....

■ Thank you for your attention

