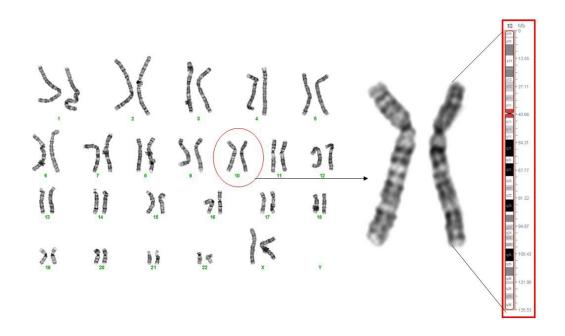


PathWest Education

Conventional Karyotype Vs Chromosomal Microarray



What is a Conventional Karyotype and What Does it Detect

- Karyotyping refers to visualising chromosomes under a microscope
- Special stains are used to help distinguish chromosome regions, most commonly **Giemsa staining** i.e. a **G-Banded Karyotype**
- Assesses ALL chromosomes and identifies the sex chromosome constitution
- Assesses for abnormalities in:
 - Total number of chromosomes (aneuploidy)
 - Size and structure of each chromosome
 - Detects large chromosomal imbalances (e.g. subchromosomal deletion)
 - Detects balanced translocations: where there is no loss of total amount of genetic material
 - Detects unbalanced translocations: where there is a change in the total amount of genetic material
- Multiple cells are assessed (in case an abnormality is not present in all cells known as mosaicism)
- Deletions, duplications or rearrangements must be large enough (approximately >5 Mb) to be visualised under the microscope to be detected by conventional karyotype. Small changes will not be detected.

What is Chromosomal Microarray (CMA)

- There are two types of Chromosomal microarray:
 - Comparative Genomic Hybridization (CGH) array
 - Single Nucleotide Polymorphism (SNP) Array
- At PathWest, we do SNP array only: provides information on copy number variants and genotype
- CMA looks at ALL chromosomes: allows many regions across the genome to be analysed simultaneously
- Allows the identification of smaller copy number changes (microdeletions and microduplications) than conventional karyotype analysis, but analysis can be more complex and usually has a longer turn-around time

What Chromosomal Microarray (CMA) <u>DOES</u> Detect

Chromosomal Microarray is a test that searches for additional (gains) or missing (losses) segments of chromosomes. These include:

- Most changes in total chromosome number: i.e. trisomy (an extra chromosome) or monosomy (a missing chromosome)
- Sub-chromosomal content changes: micro-duplications or micro-deletions of segments of chromosomes: these changes may be too small to see under a microscope (i.e. by karyotype)
- Structural chromosomal changes: unbalanced rearrangements of chromosome structure: e.g. translocations with some loss or gain of genetic material

Other abnormalities that CMA may detect include:

- Mosaicism (where some cells are normal and some cells are abnormal), may be detected if the proportion of abnormal cells is approximately >5-10%
- Duplications of the entire set of chromosomes: e.g. triploidy (3 copies of each chromosome, resulting in 69 chromosomes per cell)
- Regions of homozygosity (may suggest common ancestry, an imprinting disorder, or a recessive disease)

What Chromosomal Microarray (CMA) Does <u>NOT</u> Detect

Chromosomal Microarray does NOT detect:

- Point mutations: a change in a single nucleotide base of a gene i.e. the majority of changes within individual genes
- Tiny deletions or duplications of DNA segments within a single gene
- Balanced rearrangements of chromosomes: e.g. inversions, balanced translocations, insertions
- Low level mosaicism i.e. when the proportion of abnormal cells is below the detection threshold of the technology

CMA is generally not the most appropriate test if there is high suspicion of a single gene disorder.

Risks Associated with Chromosomal Microarray Testing

Variants of Uncertain Significance (VOUS)

- As CMA allows wider analysis of the genome than other more specific genetic testing methods, there is a risk of finding variants of uncertain significance:
 - Testing of family members may help clarify the result
 - Future knowledge may also improve understanding of the result
 - Genetic counselling is advised in this scenario

Unsolicited Findings

- With CMA, there is also a risk of identifying variants that may be unrelated to the reason CMA was originally requested, and may have relevance for other family members
- Genetic counselling is helpful in this scenario

Parental Relatedness

CMA may reveal parental relatedness

Families / patients should therefore be <u>counselled</u> appropriately prior to requesting chromosomal microarray

Should I request Conventional Karyotype or Microarray?

Common Conventional Karyotype Indications:

- Recurrent miscarriage or Infertility
- Suspected common aneuploidy (Trisomy 13, Trisomy 18, Trisomy 21, Turner syndrome, Klinefelter syndrome)
- Family history of a known chromosomal rearrangement detectable by karyotype analysis

Common Microarray indications:

- Postnatal*: Microarray is useful when you have a patient who has features that do not fit a specific known syndrome or when the concerns are more general and include:
 - Developmental delay
 - > Intellectual impairment
 - Autism spectrum disorder
 - ➤ 2 or more congenital abnormalities
- Prenatal^: Microarray is useful when you have a fetus with:
 - > 1 or more congenital abnormalities detected on ultrasound
 - Nuchal Translucency greater than 3.5mm
- Testing of fresh products of conception/fetal tissue in pregnancy loss cases, particularly where there is a history of recurrent miscarriage or fetal anomalies are present

Please see Medicare Benefits Schedule (MBS) online:

*Postnatal: <u>Item 73292 | Medicare Benefits Schedule (health.gov.au)</u>

^Prenatal: Item 73388 | Medicare Benefits Schedule (health.gov.au)

Karyotype vs Microarray Summary

| | Conventional Karyotype | Chromosomal Microarray (SNP) |
|-----------------------------|--|--|
| Variation type detected | Numerical and structural chromosomal anomalies >5Mb | Copy number variants (CNVs) with mean test resolution >27Kb Regions of homozygosity |
| Level of mosaicism detected | Generally ~20% | 5-10% |
| Specimen Requirement | Sodium-Heparin tube | EDTA tube |
| Turn-around time* | 5-18 days | 2 weeks – 3 months |
| Main indications | Infertility Recurrent miscarriage Suspected common aneuploidy Suspected chromosomal rearrangement | Developmental delay Intellectual impairment Autism spectrum disorder 2 or more congenital malformations Fetal ultrasound abnormalities |

^{*}Please note that prenatal and neonatal samples are prioritised. For non-priority samples, turnaround times may be at the longer end of the range specified

Suspected Cytogenomic Syndrome: Which Genetic Test Should I Request?

- Although conventional karyotype and microarray can detect some genetic syndromes, they cannot detect everything
- For some genetic syndromes there may be a more specific genetic test available with a higher detection rate than microarray
- Depending on the clinical features and your index of suspicion for a particular syndrome, starting with a microarray may be the most appropriate 1st step, moving onto more specific testing if the microarray result is unremarkable or does not account for the clinical features in your patient
- Suggested first line testing when you suspect a specific genetic syndrome is described in the table on the next slide

Cytogenomic Syndromes/Conditions: Which Genetic Test Should I Request First-Line at PathWest?

| Suspected Syndrome or Condition | First Line Test Required | Notes |
|--------------------------------------|--|--|
| Trisomy 13 | Conventional karyotype | Can also be detected on SNP microarray* |
| Trisomy 18 | Conventional karyotype | Can also be detected on SNP microarray* |
| Trisomy 21 | Conventional karyotype | Can also be detected on SNP microarray* |
| Monosomy X (Turner Syndrome) | Conventional Karyotype | Can also be detected on SNP microarray* |
| XXY (Klinefelter Syndrome) | Conventional Karyotype | Can also be detected on SNP microarray* |
| 22q11.2 Deletion (DiGeorge Syndrome) | MLPA for 22q11 deletion | Can also be detected on SNP microarray* |
| Williams Syndrome | Microarray | |
| Prader-Willi Syndrome | MS-MLPA for Prader-Willi Syndrome | SNP microarray will detect approximately 80% of cases* |
| Angelman Syndrome | MS-MLPA for Angelman syndrome | |
| Silver-Russell Syndrome | MS-MLPA for Silver-Russel syndrome | |
| Beckwith-Wiedemann Syndrome | MS-MLPA for Beckwith-Wiedemann syndrome | |
| Fragile X syndrome | Fragile X genetic testing (CGG repeat analysis of FMR1 gene) | |
| Spinal Muscular Atrophy | SMA genetic testing | |

^{*} N.B. if less specific phenotype and more general features present (such as developmental delay, autistic spectrum disorder or multiple congenital abnormalities): a SNP microarray may be the more appropriate first-line genetic test

Tips for Requesting Cytogenomic Testing

- Ensure the patient has been counselled and consented appropriately
- Please clearly state the test being requested on the request form
- Highlight if test is prenatal/diagnostic/predictive/carrier testing
- Please provide as much clinical information as possible on the request form
- Please be mindful of the extensive time that goes into analysing and reporting the data:
 - We suggest requesting the genetic test most likely to provide an answer to the clinical query initially
 - Consider clinical genetics service referral if the phenotype is complex or specialist genetic counselling is required
 - Turn-around times are likely to be longer than pathology tests performed by many other departments due to the complex nature of genetic analyses