

# NIGM

## Non-invasive Prenatal Testing (NIPT) of GeneMind

### Simple

Test from a tube of 10ml maternal blood sample as early as 9 gestational weeks

### Accurate

Proven >99% sensitivity based on a test of more than 70,000 pregnancies.

### Fast & Convenient

- Quick test from sample preparation to issue report within 24 hours
- Simplified operation: Automated and localized data analysis and report

### Flexibility

- Compatible with low, medium and high throughput sequencing platforms
- Offering flexible options for both manual and automated workflows



## Introduction to NIGM

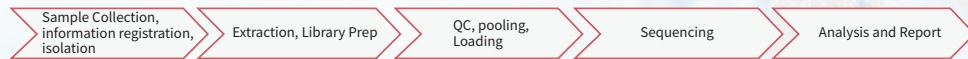
Noninvasive prenatal testing (NIPT) performed with NGS sequencing system GenoLab M\*, FASTASeq 300\* and SURFSeq 5000\* provides reliable screening results for fetal chromosomal aneuploidies as early as 9 gestational weeks—from a single tube of 10 mL maternal blood. Support G tube (Provided by GeneMind) and Streck Tube.

| Parameter                                    |                              | NIPT Basic   | NIPT Standard   | NIPT Plus  | NIPT Pro   |
|--|------------------------------|--|---|--|--|
| Sample Volume                                |                              | 10 mL of a single tube of maternal blood                               |   |  |  |
| Library preparation                          |                              | PCR  |   |  |  |
| Method                                       |                              | Low-pass whole genome sequencing                                       |   |  |  |
| No. of samples per run(1 PC +1 NC included ) | GenoLab M*                   | 1 FCM : 48<br>2 FCM/1 FCH : 96<br>1 FCM+1 FCH : 144<br>2 FCH : 192     | 1 FCM : 32<br>2 FCM/1 FCH : 64<br>1 FCM+1 FCH : 96<br>2 FCH : 128 | 1 FCM : 24<br>2 FCM/1 FCH : 48<br>1 FCM+1 FCH : 72<br>2 FCH : 96 | 1 FCM : 10<br>2 FCM/1 FCH : 20<br>1 FCM+1 FCH : 30<br>2 FCH : 40 |
|  | FASTASeq 300*                | 1 FCH : 48<br>1 FCM : 24   | 1 FCH : 32<br>1 FCM : 16  | 1 FCH : 24<br>1 FCM : 12   | 1 FCH : 10<br>1 FCM : 5  |
|  | SURFSeq 5000*                | 1 FCH : 250  | 1 FCH : 160   | 1 FCH : 96   | 1 FCH : 64   |
| Read length                                  |                              | SE 75 on GenoLab M* and FASTASeq 300*<br>SE 50 on SURFSeq 5000*        |   |  |  |
| Average Unique Reads / sample                | GenoLab M* and FASTASeq 300* | ≥3.5 M   | ≥5 M  | ≥7 M   | ≥16 M  |
|  | SURFSeq 5000*                | ≥5 M   | ≥7 M  | ≥12 M  | ≥20 M  |
| Turnaround time                              |                              | 25 h on GenoLab M*, 21.5 h on FASTASeq 300*, and 26 h on SURFSeq 5000* |   |  |  |
| Report generation                            |                              | Local analysis and report system                                       |   |  |  |

| NIPT Basic  | NIPT Standard  |
|---|--|
| <ul style="list-style-type: none"> <li>▶ Trisomies 21/18/13</li> <li>▶ Sex identification, fetal fraction estimation</li> </ul>   | <ul style="list-style-type: none"> <li>▶ Trisomies 21/18/13</li> <li>▶ 6 Sex Chromosome Aneuploidies (SCAs)<br/>(XO/XXX/XXY/XYY/XO+XY/XXX+XY)</li> <li>▶ Other 19 autosomal Aneuploidies</li> <li>▶ Sex identification, fetal fraction estimation</li> </ul>   |
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\*NIPT Basic and Standard can be performed in twin pregnancies, but Plus or Pro is NOT applicable.

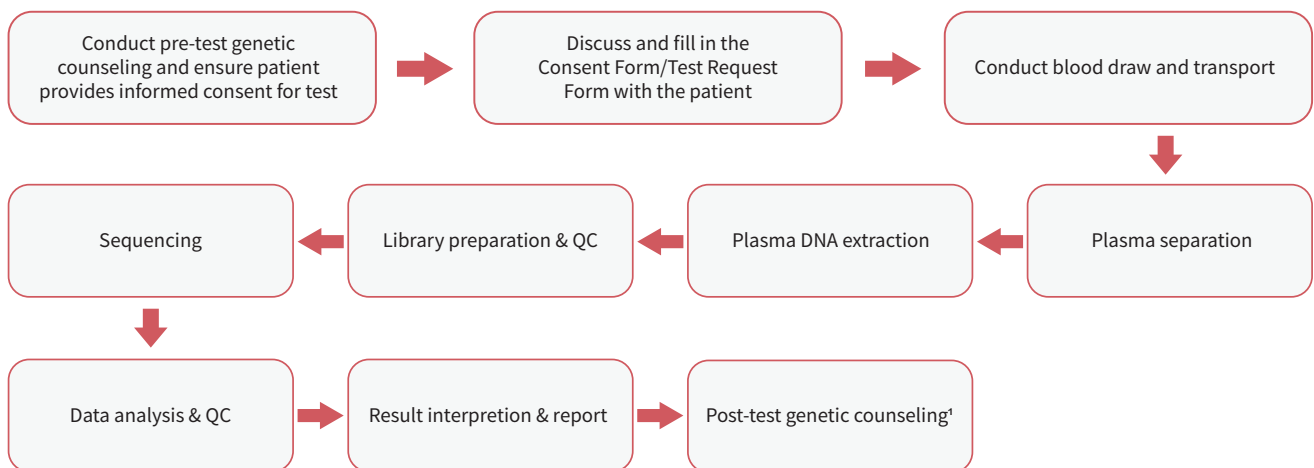
# Support Both Manual and Automated Solutions



|                    | G Tube | Streck Tube | MrLH-96                          |       | GenoLab M*    | AIRS  |                     |
|--------------------|--------|-------------|----------------------------------|-------|---------------|-------|---------------------|
| Automated workflow |        |             |                                  |       |               |       | 25 hours in total   |
|                    | 2 h    |             | 6 h                              | 2.5 h | 13 h          | 1.5 h |                     |
|                    | G Tube | Streck Tube |                                  |       | FASTASeq 300* | AIRS  |                     |
| Manual workflow    |        |             |                                  |       |               |       | 21.5 hours in total |
|                    | 2 h    |             | 8 hours (at least 3 technicians) | 2.5 h | 7.5 h         | 1.5 h |                     |
|                    | G Tube | Streck Tube | MrLH-96                          |       | SURFSeq 5000* | AIRS  |                     |
| Automated workflow |        |             |                                  |       |               |       | 26 hours in total   |
|                    | 2 h    |             | 6 h                              | 2.5 h | 14 h          | 1.5 h |                     |

- ▶ G Tube: provided and validated by GeneMind
- ▶ MrLH-96: Automated Sample Preparation System, consumables (tips and plates) provided and validated by GeneMind
- ▶ GenoLab M\*, FASTASeq 300\* and SURFSeq 5000\* : sequencing device from GeneMind
- ▶ AIRS: Automated Integrated Report Server provided and validated

## The Test Workflow and Advantages

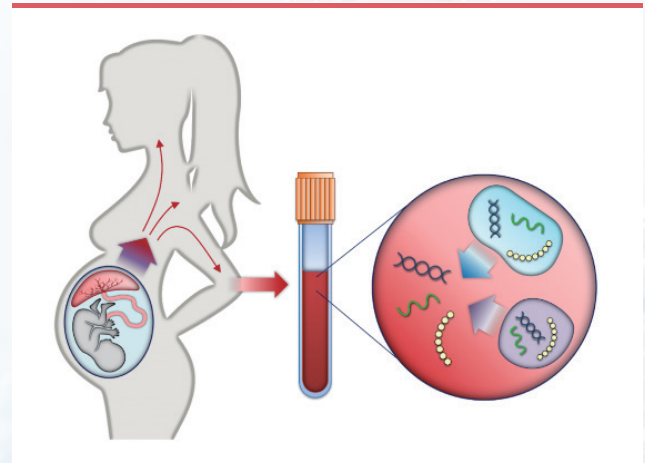


1. non-complimentary

## ● cfDNA Extraction

During pregnancy, cell-free DNA (cfDNA) fragments originate from both the mother and fetus are present in maternal blood circulation. Cell-free fetal DNA (cffDNA) is present only as a minority component of the total cfDNA in maternal plasma, which poses a significant technical challenge for some NIPT detection methods.

Using paramagnetic particle method, our cfDNA extraction process is compatible with a variety of blood collection vessels with the requirement of as little as 200uL plasma. The success rate of cfDNA extraction is above 99.9% if the maternal plasma meets the sampling standard.



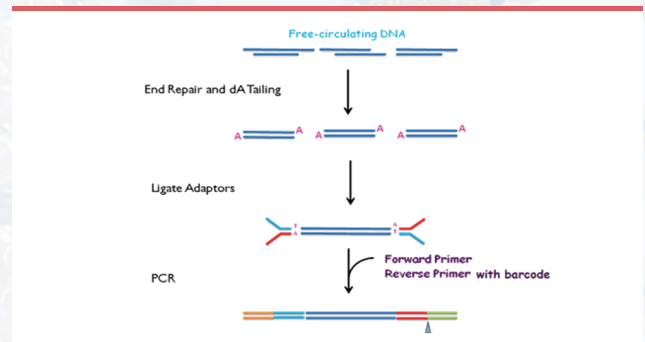
## ● Library preparation

### ● Low-input

The success rate of cfDNA library construction is above 99.9% even if the amount of cfDNA is as little as 100pg.

### ● High conversion efficiency

With significant enrichment effect of fetal concentration, the library yield is sufficient for multiple validation and long-term storage.



## ● Sequencing

We conduct sequencing on High-throughput DNA Sequencing Platform GenoLab M\*.

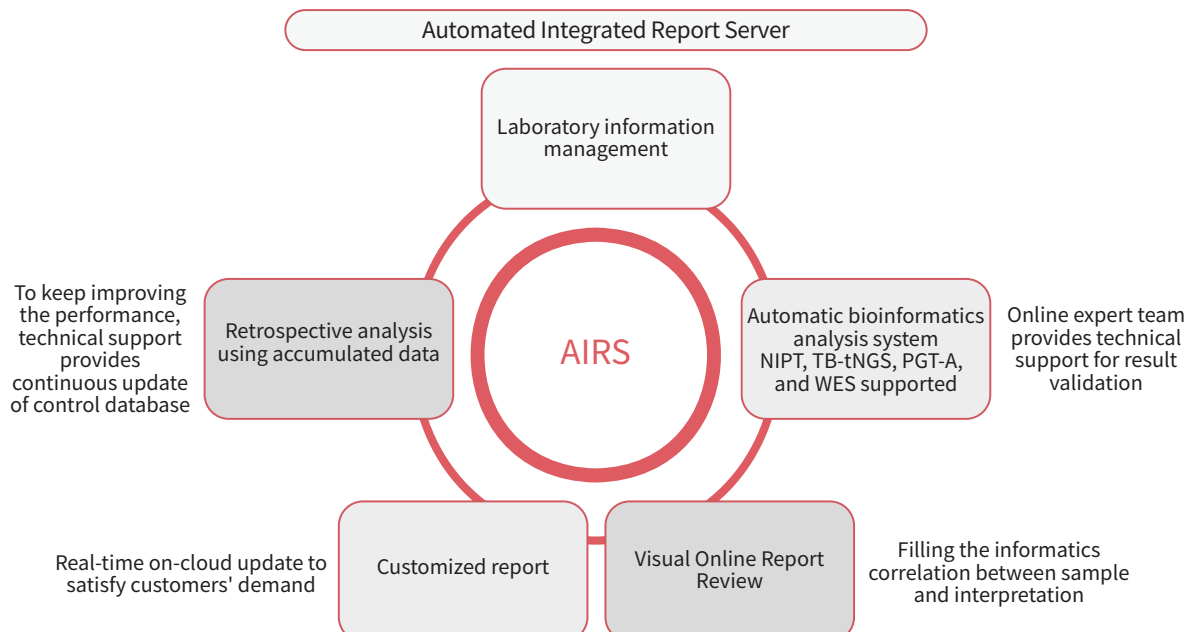
### Accurate

- High-fidelity sequencing chemistry and unique optical system

### Cost Effective

- Competitive instrument and reagent kit prices
- Single-flow cell and dual-flow cell mode give customer more control over per-sample cost

## ● Automated Bioinformatics Analysis and Report



## Sample Requirement

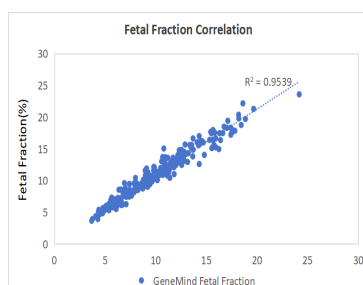
| Sample Type | Quantity               | Requirements   | Shipment   |
|-------------|------------------------|--|--|
| Plasma      | 2ml (4 tubes required) | Stored in 1.5ml Eppendorf tubes, and sealed with 1cm wide parafilm | Stored at -80 °C, shipped with dry ice within one week.                                  |
| Whole Blood | 10ml                   | Gently invert the tube ten times immediately after blood sampling  | Stored and shipped between 6~35 °C within 7 days. Keep the tubes upright during shipping |

## Clinical Validation<sup>2</sup>

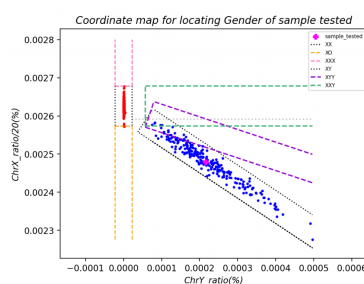
| Trisomy   | Positive | Sensitivity | Specificity | PPV    | NPV     |
|-----------|----------|-------------|-------------|--------|---------|
| T21       | 402      | 100.00%     | 99.94%      | 90.03% | 100.00% |
| T18       | 93       | 100.00%     | 99.98%      | 84.95% | 100.00% |
| T13       | 57       | 100.00%     | 99.95%      | 42.11% | 100.00% |
| Autosomes | 116      | 98.18%      | 99.91%      | 46.55% | 99.99%  |
| SCAs      | 429      | 97.93%      | 99.73%      | 55.24% | 99.99%  |
| Total     | 1097     | 99.21%      | 99.51%      | 69.01% | 99.99%  |

2. validated by 70,075 retrospective clinical samples

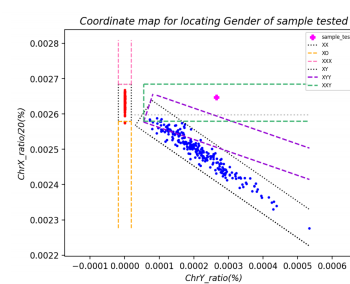
### Performance in Sex Identification, Fetal Fraction Estimation and SCAs Detections



Fetal fraction estimation is highly consistent with existing products



Accuracy of Sex identification is 99.3%



Reliable performance in sex chromosome aneuploidies detection

# About GeneMind

GeneMind Biosciences Co., Ltd. (hereinafter referred to as 'GeneMind') was founded in 2012 and is headquartered in Luohu, Shenzhen. Its facility includes more than 8,000 square meters of research and development laboratories and GMP production lines.

From the company's inception, GeneMind aimed at the research and development of the DNA sequencer: the core instrument of the modern molecular diagnostic platform. GeneMind is committed to building a precision medical ecosystem that serves healthcare through collaborations with genetic testing service providers and medical institutions. We has launched high-throughput sequencing platform GenoLab M\*, FASTASeq 300\*, and SURFSeq 5000\*, which can provide 'instrument-reagent-chip-software' full-platform solutions. With nearly 200 technical patents, covering mainland China, Hong Kong, Europe, and the United States, GeneMind is one of the few companies in the world that possess core DNA sequencing technologies and independent products.

After several years of hard work, GeneMind has mastered the key technologies of the DNA sequencing, and realized production of core materials such as enzymes, nucleic acids, dyes, and biochips in China, providing research and medical institutions with competent domestically produced sequencing solutions.



\*Unless otherwise informed, GeneMind sequencing platform and related sequencing reagents are not available in the USA, Canada, Australia, Japan, Singapore, Western Europe, Southern Europe and Nordic countries yet.

