

Genie-A Solution

Preimplantation Genetic Testing for Aneuploidy



Comprehensive detection

A single solution to detect CNV, Ploidy and ROH (regions of homozygosity)

Default SNP analysis

Genie-A uses Genome-wide SNP information to detect Ploidy and ROH

High Resolution

>4Mb CNVs (unknown), and >1 Mb inherited CNVs (known)

High throughput s Scalability

500 M reads/run, 5 M reads/samples, 1 x 100 bp reads, Up to $48,\,96$ samples/run



Genie-A Fast: Fast workflow, combining WGA and library preparation to reduce cost and time.

Genie-A Advanced: Separate WGA (MDA) followed by a patented library preparation method to support wider genome coverage. Two version of Genie-A kits with different sample preparation methods for greater flexibility, but with the same outcome.

Amplification Genie-A Kits Sequencers Reporting content method Aneuploidy, Triploidy, Haploidy PicoPlex Genie-A Fast • z4Mb CNVs in unknown regions Genie Sequencer ≥1Mb known inherited CNVS DNBSEQ-G400* Mosaicism ≥30% C ≥10Mb Genie-A Advanced MDA Whole chromosome level ROH

Competitive Pricing

Offering significant cost savings without compromising on quality

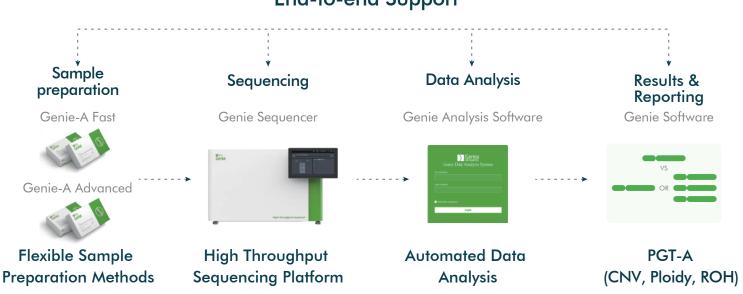
Superior Data Volume and Detection Capabilities

Data volume starts at 5M, allowing for the detection of a wide range of diseases

Supports Localised Deployment

Compatible with various devices and capable of quick report generation

End-to-end Support





Genie-Plus Solution

All-in-one test kit combining PGT-A, PGT-M & PGT-SR



A single solution for combined detection of PGT-M, PGT-SR and PGT-A, including Triploidy, UPD and ROH

Genome-wide coverage

WGA by MDA and a Patented library preparation method for better whole-genome coverage

High Resolution

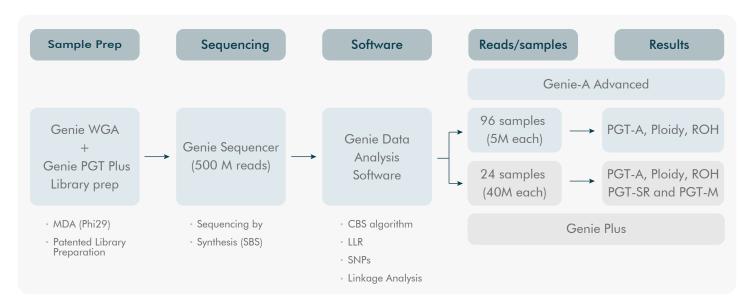
>4Mb CNVs (unknown), and >1 Mb inherited CNVs (known)

High throughput s Scalability

500M reads/run, 40M reads/samples, 2×100 bp reads, up to 24, 48 samples/run



Genie Plus: A Single workflow for PGT



Competitive Pricing

Offering significant cost savings without compromising on quality

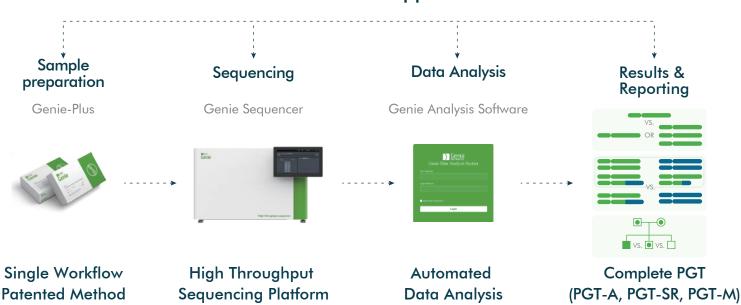
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^[1] Xie P, Hu X, Kong L, et al. Hum Reprod. 2022

^[2] Note: risk warning available when using Genie PGT-Plus Analysis Software QRTM485