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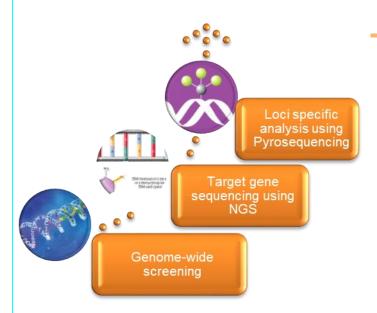


NGS Targeted Sequencing

Targeted Gene Sequencing

EpigenDx is focused on targeted disease gene panel sequencing and targeted gene bisulfite sequencing using Next-Gen Sequencing (NGS) technology.

Coupled with genome-wide DNA methylation screening and loci specific methylation analysis using Pyrosequencing, EpigenDx provides a full solution for genetic and epigenetic biomarker discovery and validation.



The services include:

- Gene Panel Assay Design
- Library Construction
- Template Preparation and QC
- NGS sequencing using Ion S5TM system
- Sequencing analysis
 - o DNA methylation analysis
 - o Disease panel sequencing and SNP genotyping
 - Allele specific expression analysis and allele specific methylation analysis using extensive bioinformatics analysis







Amplicon Library Prep /Templating







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tNGBS vs. Pyrosequencing

Methylation Analysis - Single Base Pair Resolution

	NEXT-GEN SEQUENCING (ION TORRENT)	Pyrosequencing
Technology	Sequencing by Synthesis – Hydrogen Ion Detection	Sequencing by Synthesis – Fluorescence Detection
Methylation Calculation	# of methylated reads/ # total reads (at each CpG)	Average in each sample (Calculated by RLU at each CpG)
Advantages	 Can run many assays on many samples simultaneously Extremely sensitive in detecting small changes in methylation due to ability to look at single reads 	 Rapid Results Reproducible 92 samples per run Can successfully detect ~5-10% differences in methylation
Disadvantages	Longer turnaround time	• Run only 1 assay at a time
When to use/Cost Effectiveness	Best for <u>more</u> than 20 assays with smaller sample sets	Best for <u>fewer</u> than 20 assays with larger sample sets
PCR Bias and Validation	All assays are validated by EpigenDx Methylation Controls	

