Analysis of Reference and Target Gene Data

The EpiQ assay provides a genomic DNA bisulfite conversion format that can be used to compare DNA methylation levels. This allows for the quantification of DNA methylation at specific sequence loci. The EpiQ assay is highly sensitive and can detect small changes in DNA methylation. In addition, the EpiQ assay can be used to identify regions of the genome that are hyper- or hypomethylated. This information can be used to identify potential targets for further investigation.

Conclusion

The EpiQ assay provides a powerful tool for the analysis of DNA methylation at specific sequence loci. This information can be used to identify potential targets for further investigation. The EpiQ assay is highly sensitive and can detect small changes in DNA methylation. This information can be used to identify potential targets for further investigation.