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Memorial Sloan Kettering
Cancer Center

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sample preparation to data analysis and training.

The [Bioinformatics Core Facility](#), led by [Nicholas Soccia](#), assists investigators in analyzing large-scale genomics data. In addition, the facility offers consultation, training, and custom programming, and administers Memorial Sloan Kettering's high-performance research computing resources.

A comprehensive resource for research involving human tissue, the [Pathology Core Facility](#) provides instrumental support to Center investigators conducting genomic investigation of clinical samples. The core staff and pathologists provide a wide range of services — from sample acquisition and banking to tissue-based experimentation, analysis, and data interpretation. The facility is led by [Marija Drobniak](#) and [Victor E. Reuter](#).

The [New York Genome Center \(NYGC\)](#), which launched last year, is a collaboration of 11 New York-based biomedical institutions. Memorial Sloan Kettering is playing a key role in the development of the NYGC, which will be located in lower Manhattan. The NYGC will give scientists at Memorial Sloan Kettering and other institutions enhanced access to high-throughput genome sequencing services as well as computational and bioinformatics support.

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