Illumina NextSeq500 high-throughput sequencing available. Samples now accepted!

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Note that we will potentially adjust pricing and services offered as we gain more experience with this instrument.
High numbers of multiplexed samples should be in multiples of eight (8) or twelve (12) to maximize our robotic library preparation capabilities.

The purchase of an Illumina NextSeq500 genome sequencer was made possible by an NSF-Major Research Instrumentation grant to Dr. Hoyt. The new sequencing instrument is located in the Genomics and Proteomics center of the Henry Bellmon Research Center (Arrival pics here). The award included power backup and tape storage for the large data archiving. For more information on the new instrument go HERE

Pricing varies greatly depending on experimental design. Final pricing will require contact with the Genomics center personnel.

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