

Pippin Prep™

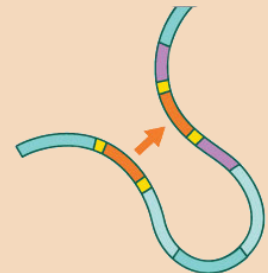
DNA Size Selection



Researchers Citing Pippin Prep in New Methods

TIP-seq: Transposon Insertion Profiling

Vectorette PCR and paired-end Illumina® sequencing are used to map retrotransposon insertions in the human genome.



Steranka, et al. [Transposon insertion profiling by sequencing \(TIPseq\) for mapping LINE-1 insertions in the human genome.](#) *Mobile DNA*. 2019. (DOI: [10.1186/s13100-019-0148-5](#))

Epigenetic ddRAD-seq on Museum Specimens

Successful ddRAD genotyping with bisulfite sequencing on archived mouse skulls up to 76 years old.



Rubi, et al. [Museum epigenomics: characterizing cytosine methylation in historic museum specimens.](#) *BioRxv preprint*. 2019. (DOI: [10.1101/620583](#))

Terminome-Seq: RNA Termini Sequencing

Characterization of primary 5' ends, processed 5' ends, and 3' ends in plant chloroplast transcripts.



Castandet, et al. [Systematic sequencing of chloroplast transcript termini from Arabidopsis thaliana reveals >200 transcription initiation sites and the extensive imprints of RNA-binding proteins and secondary structures.](#) *BioRxiv preprint*. 2019. (DOI: [10.1101/621938](#))

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