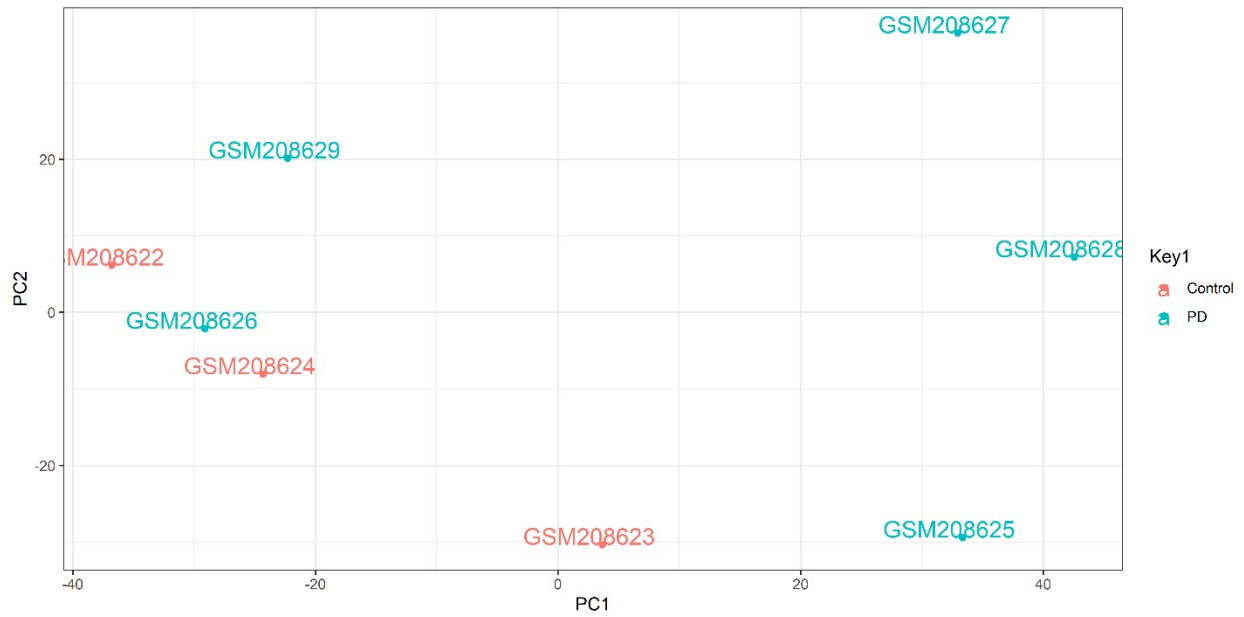
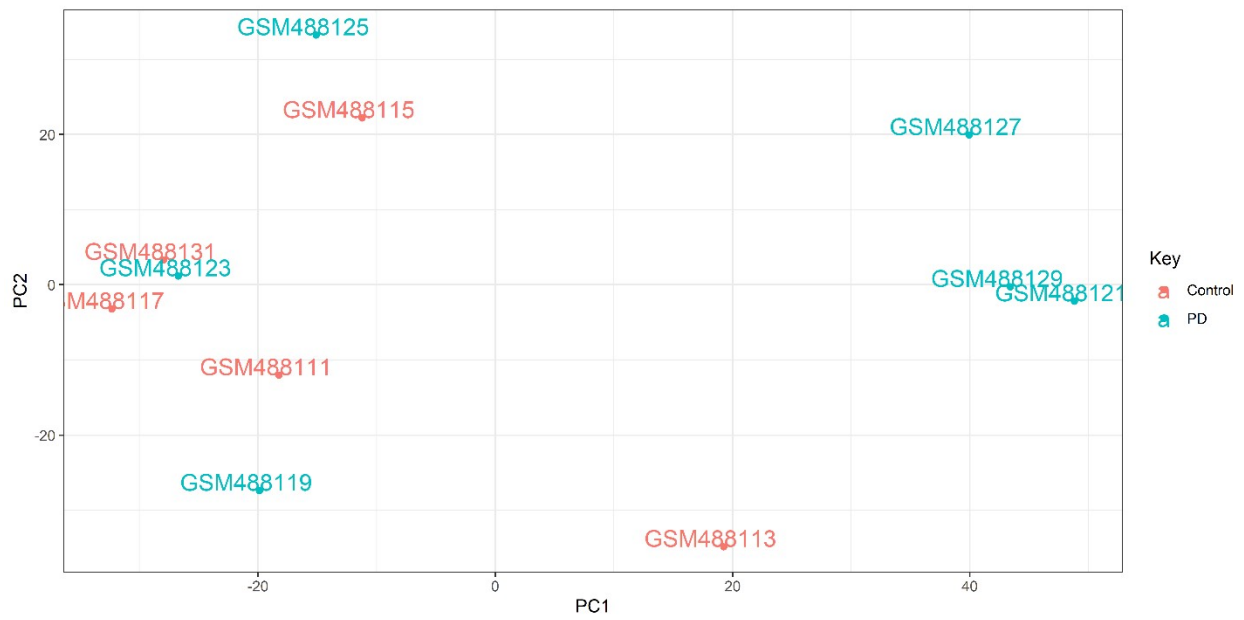


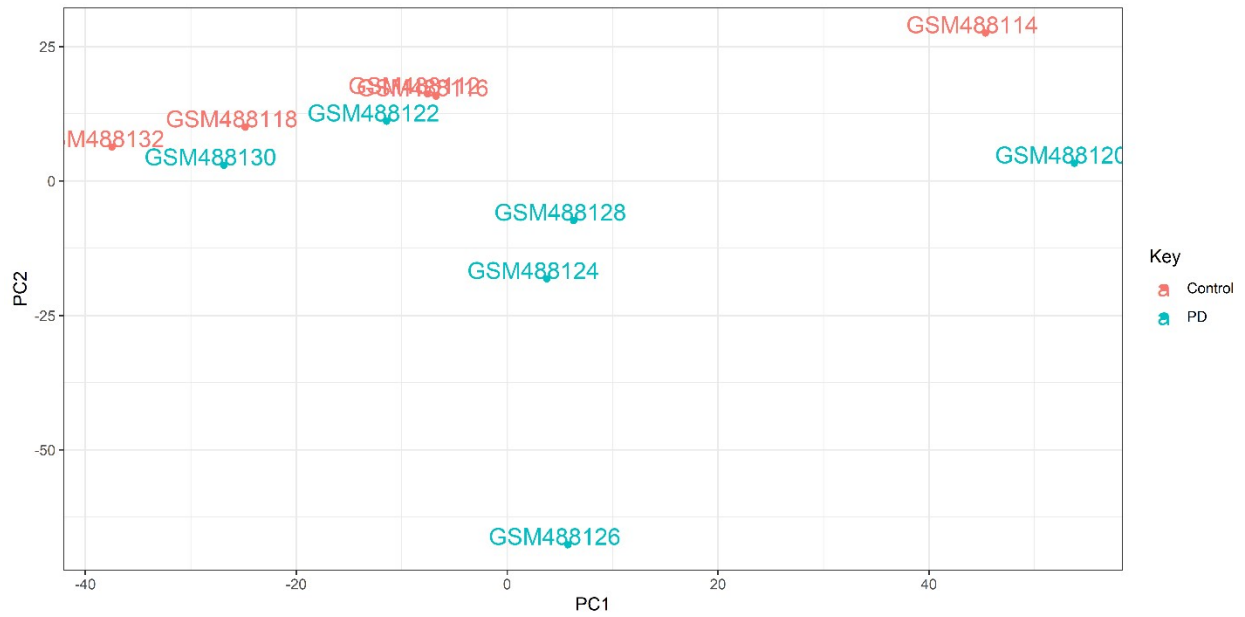
GSE8397: SN



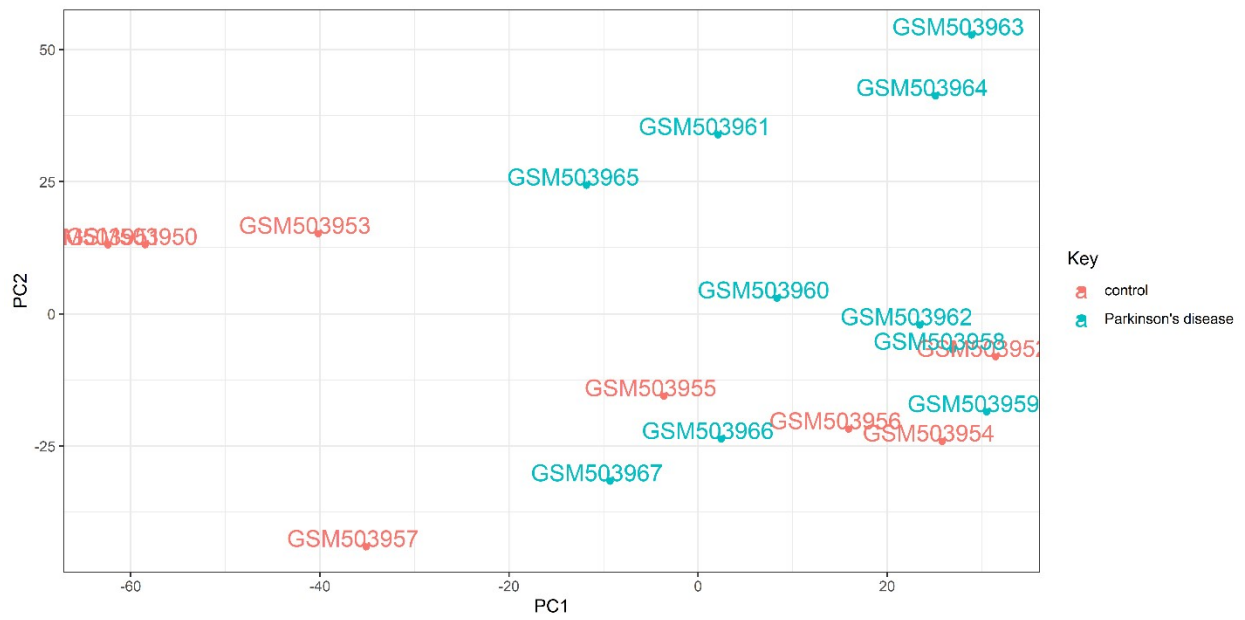
GSE19587: DON



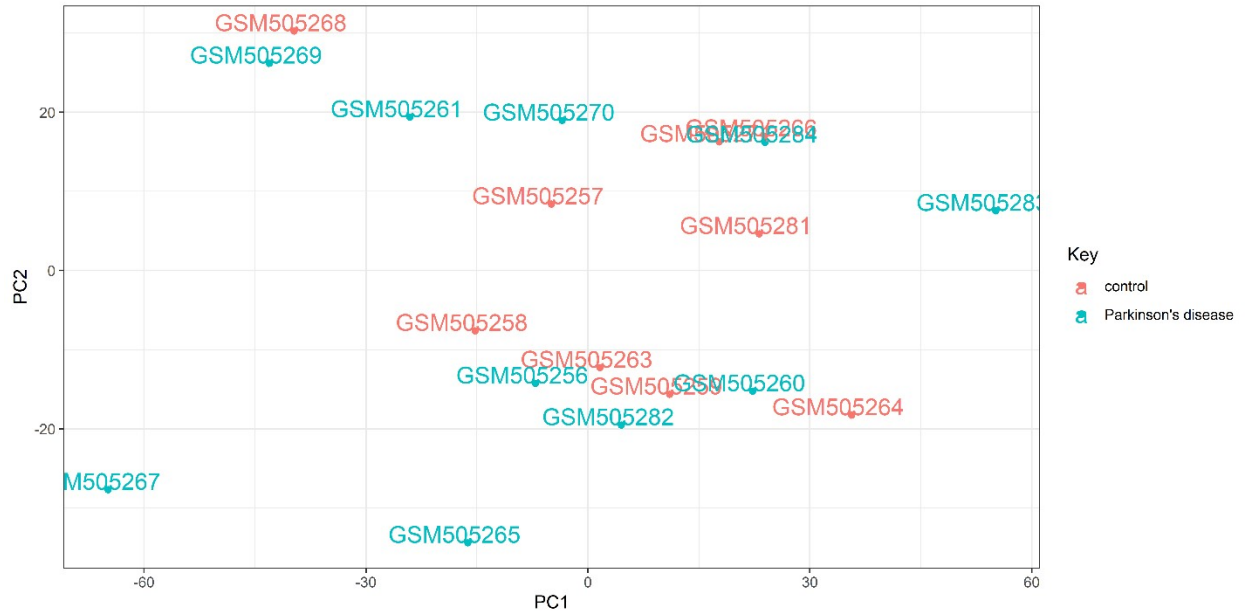
GSE19587: ION



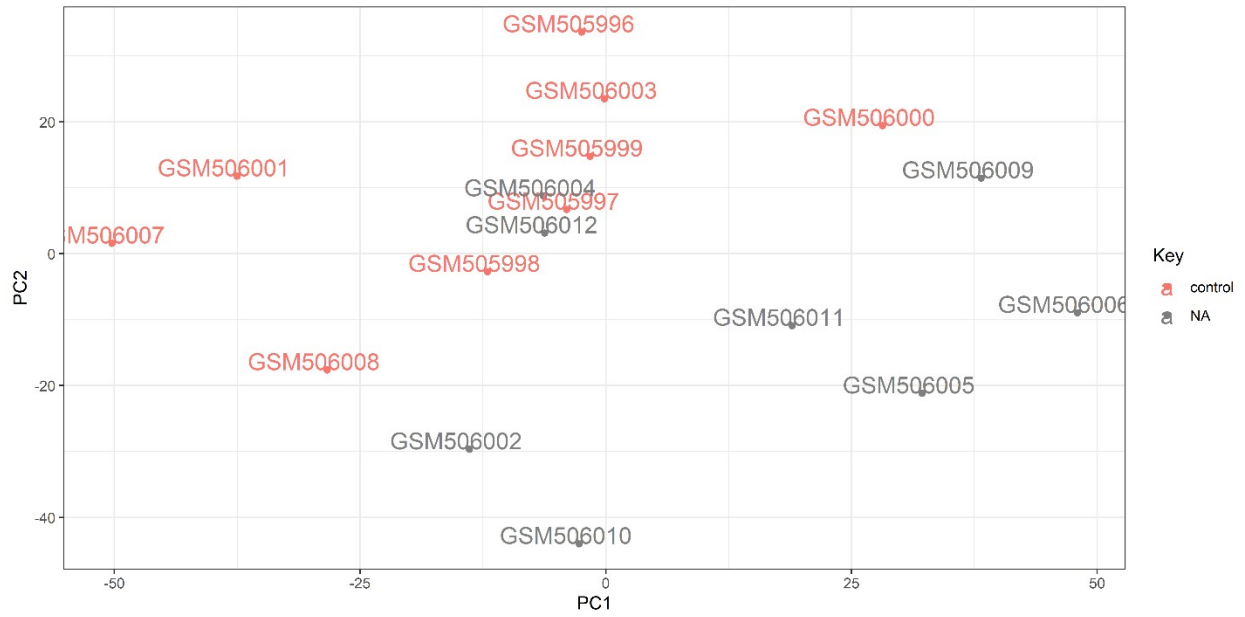
GSE20141: SN

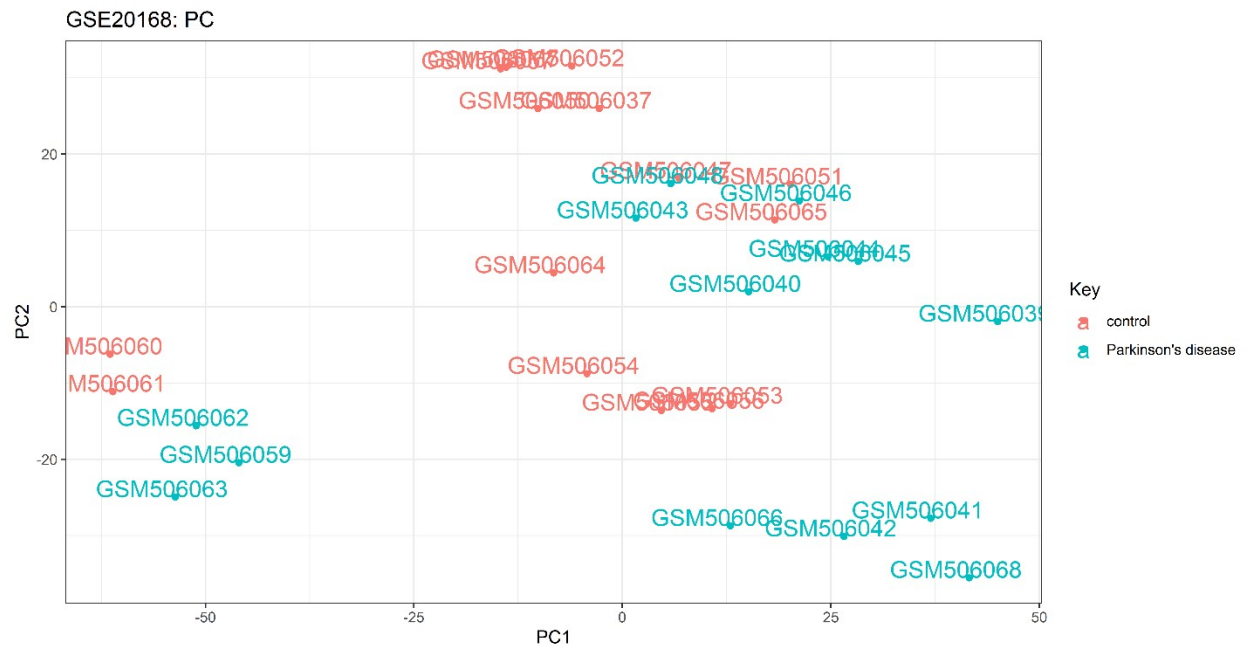
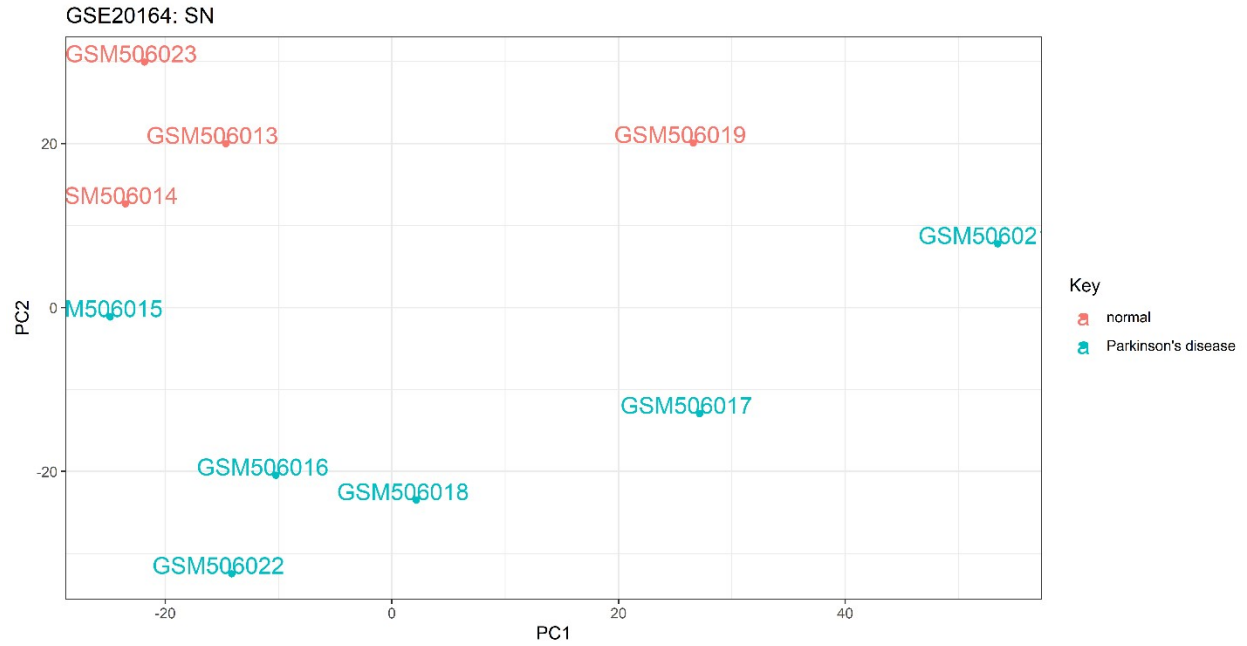


GSE20146: GPi

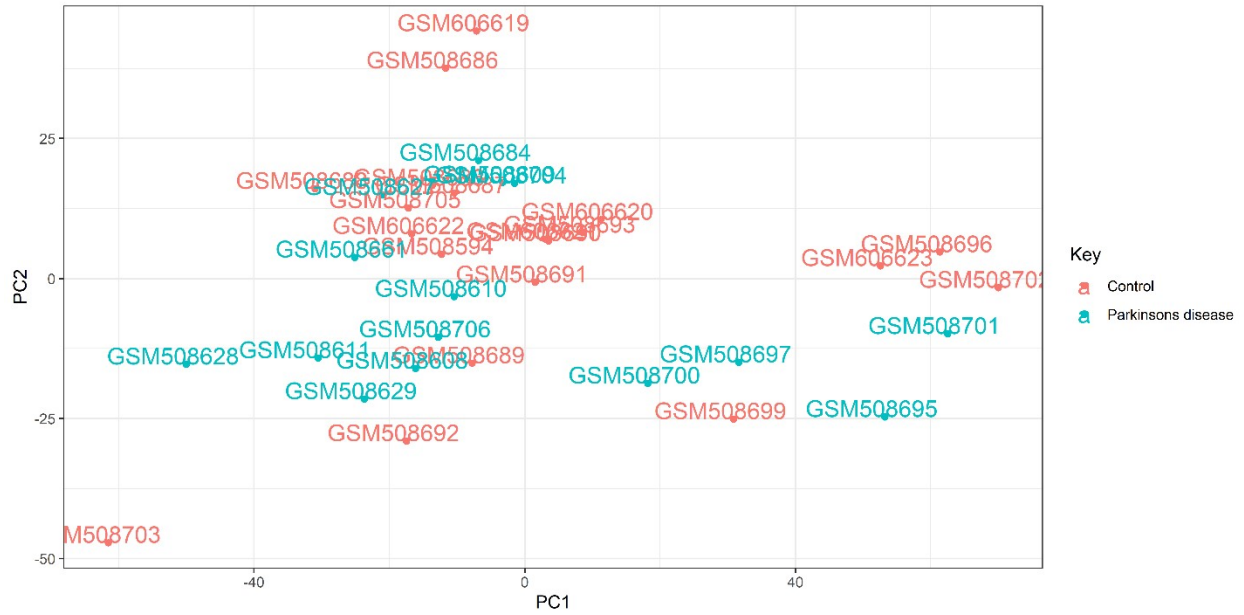


GSE20163: SN

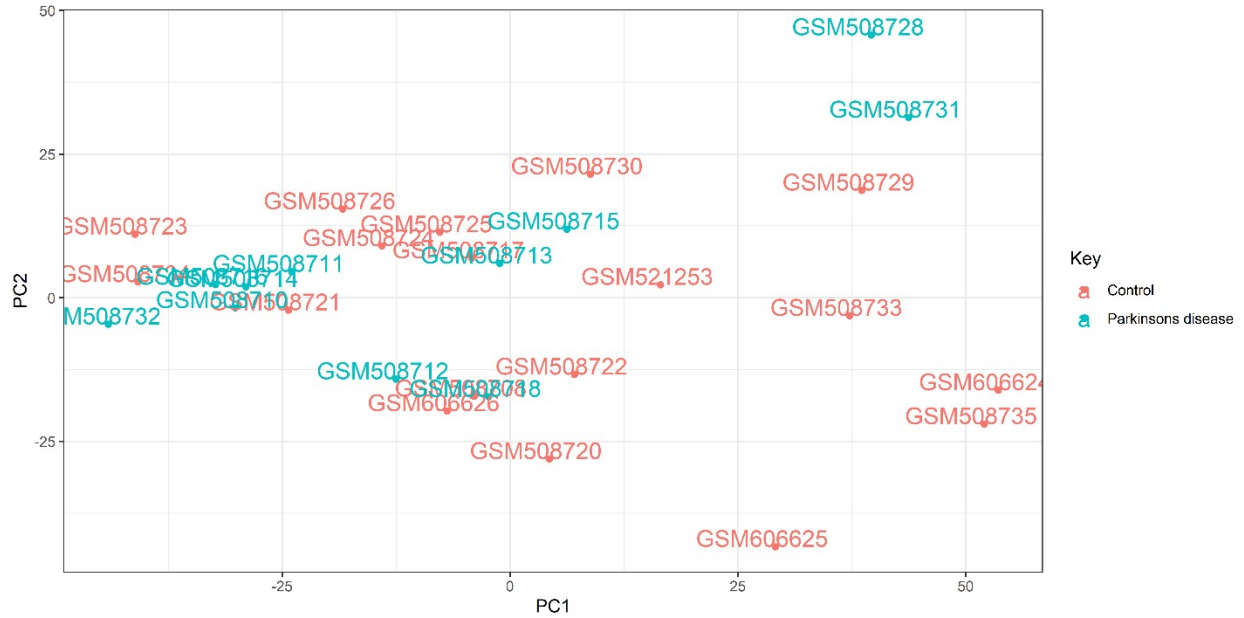


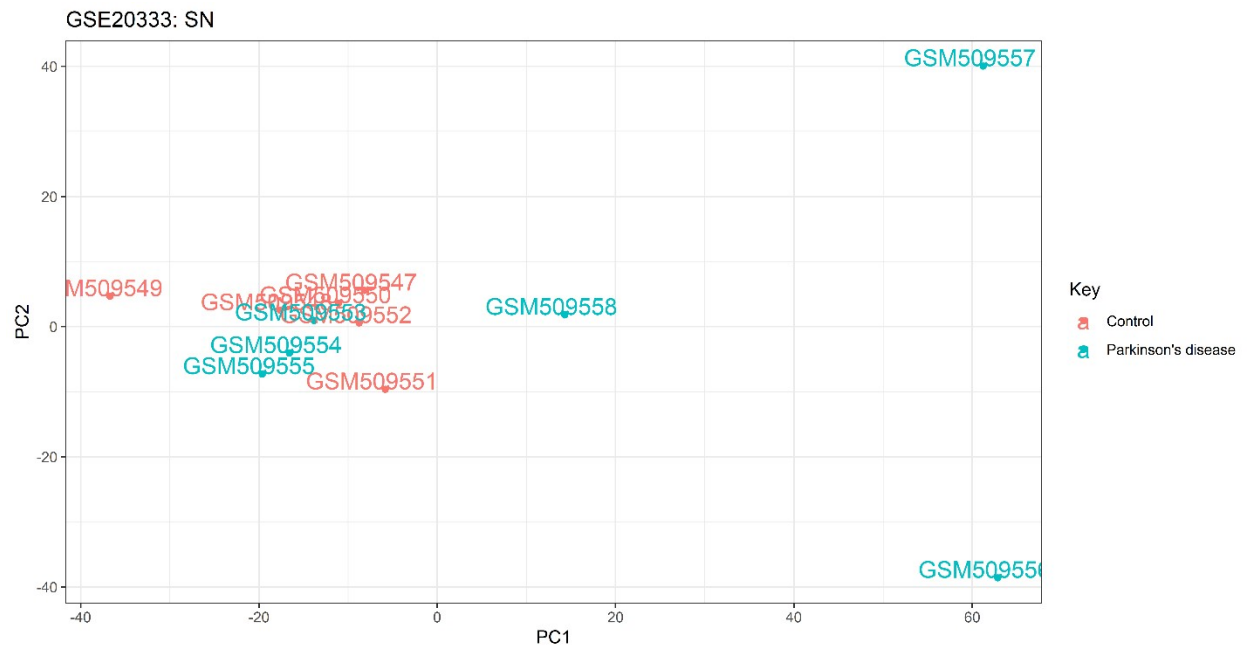
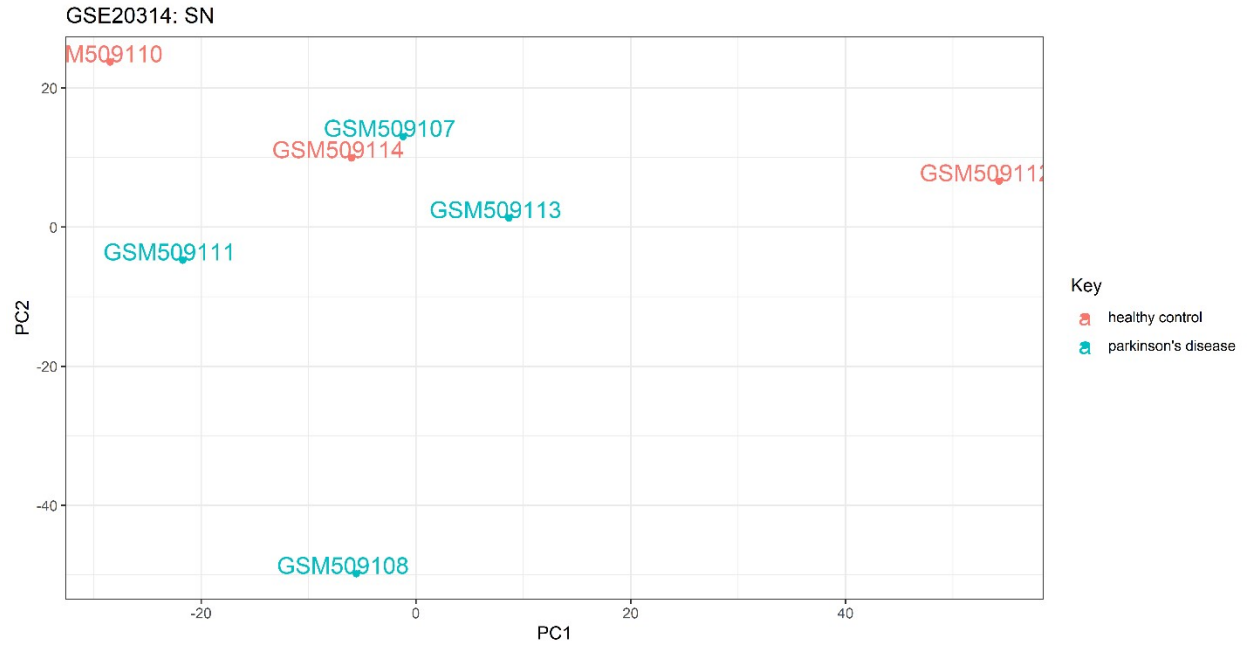


GSE20291: PUT

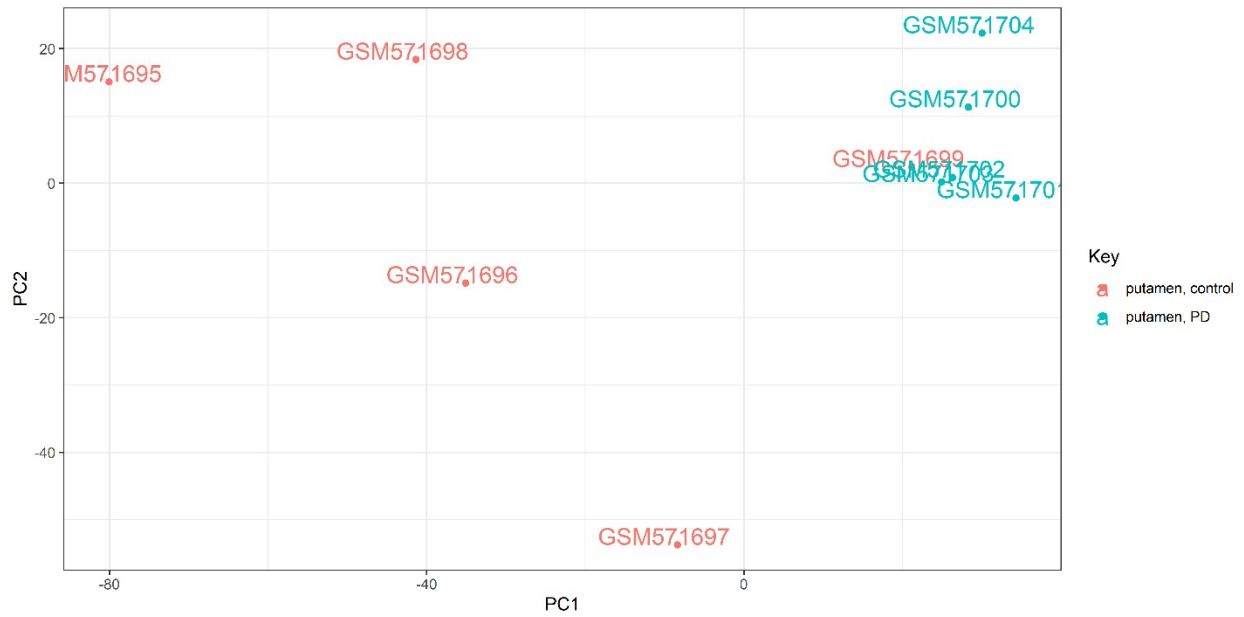


GSE20292: SN

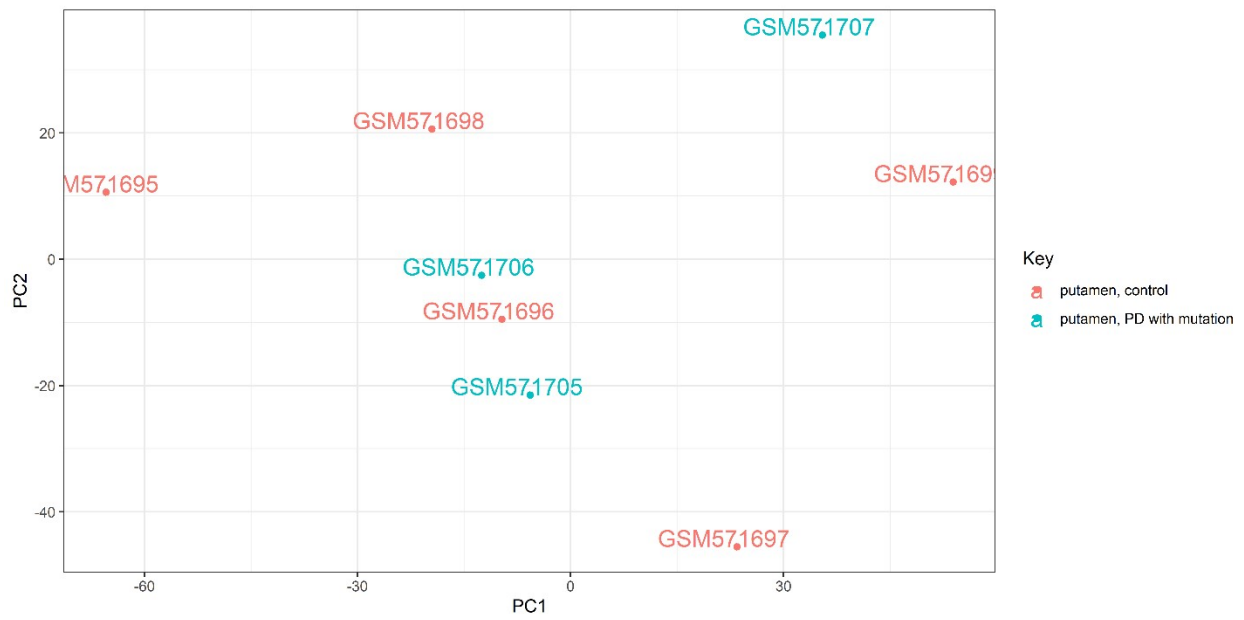




GSE23290: iPUT

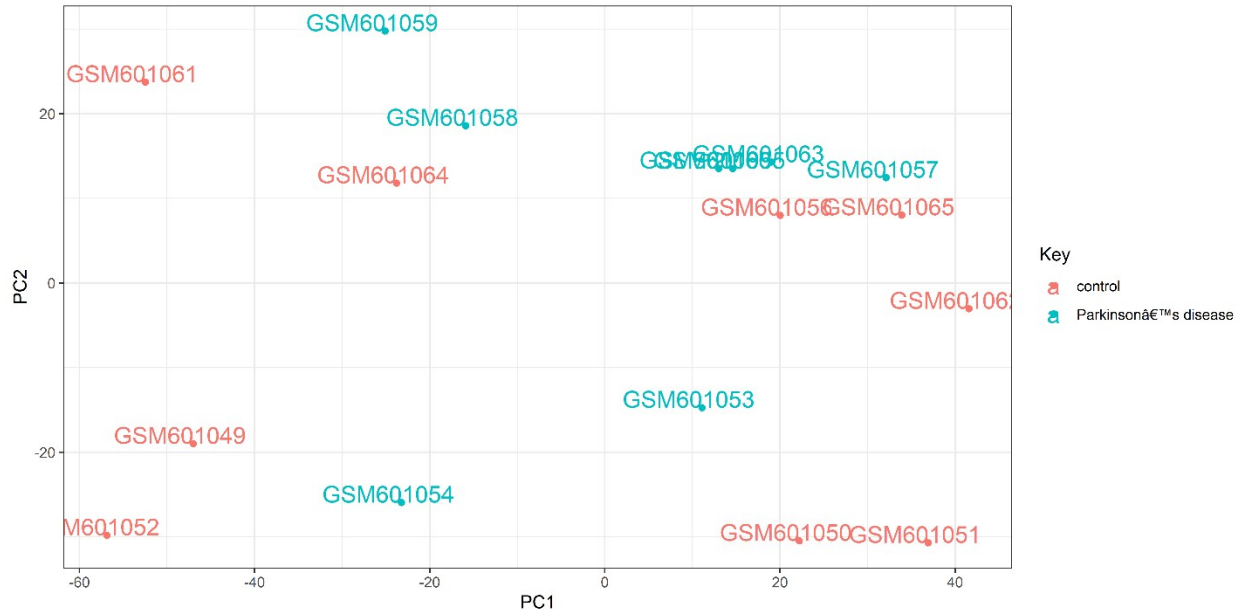


GSE23290: mPUT

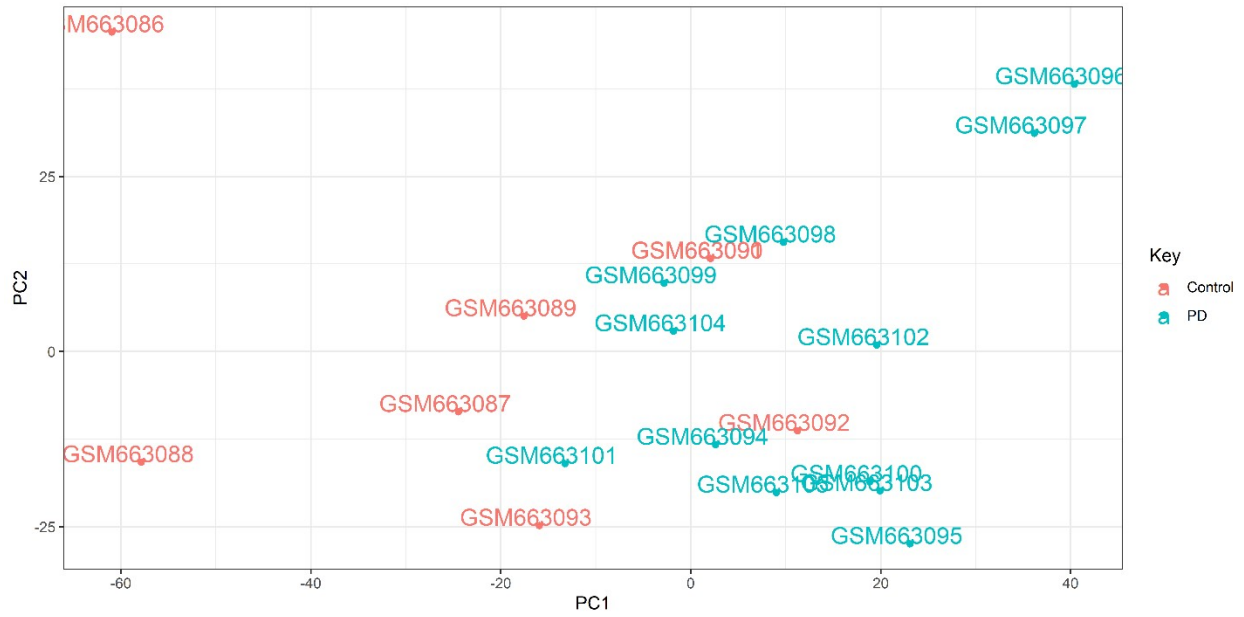




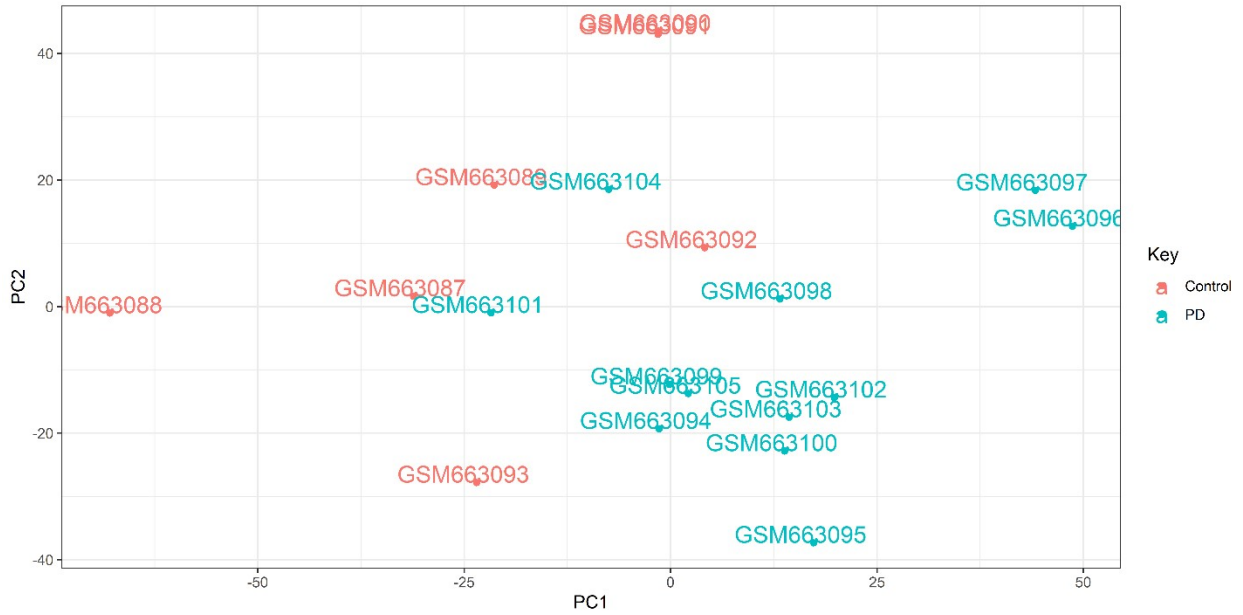
GSE24378: SN



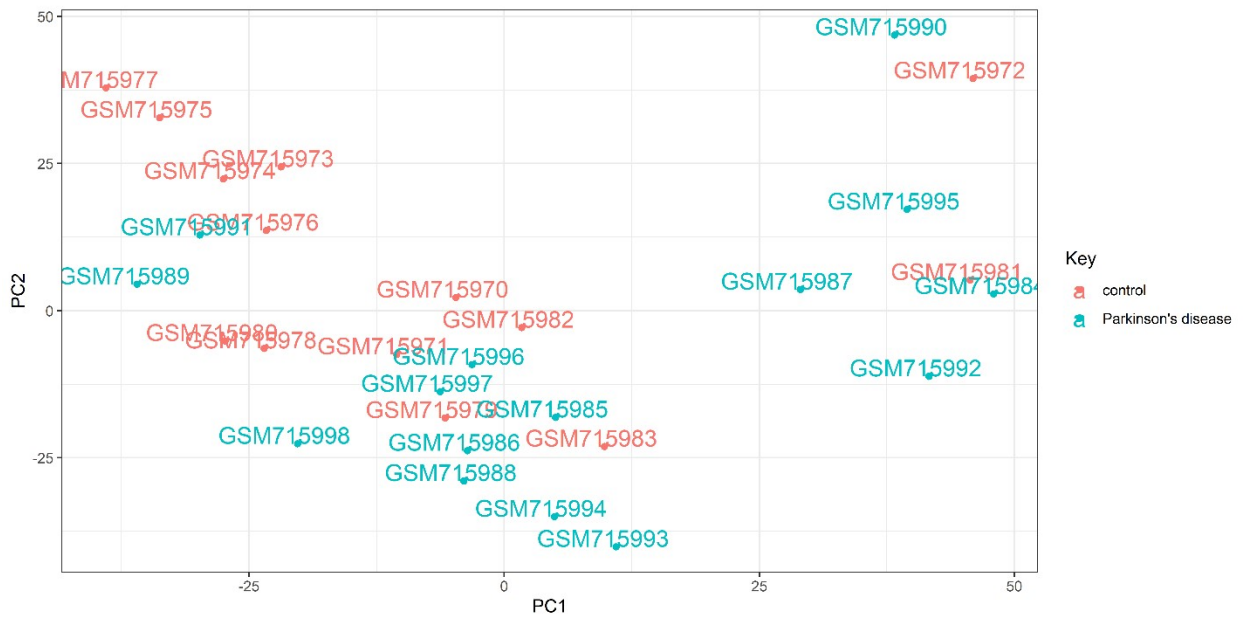
GSE26927: PC



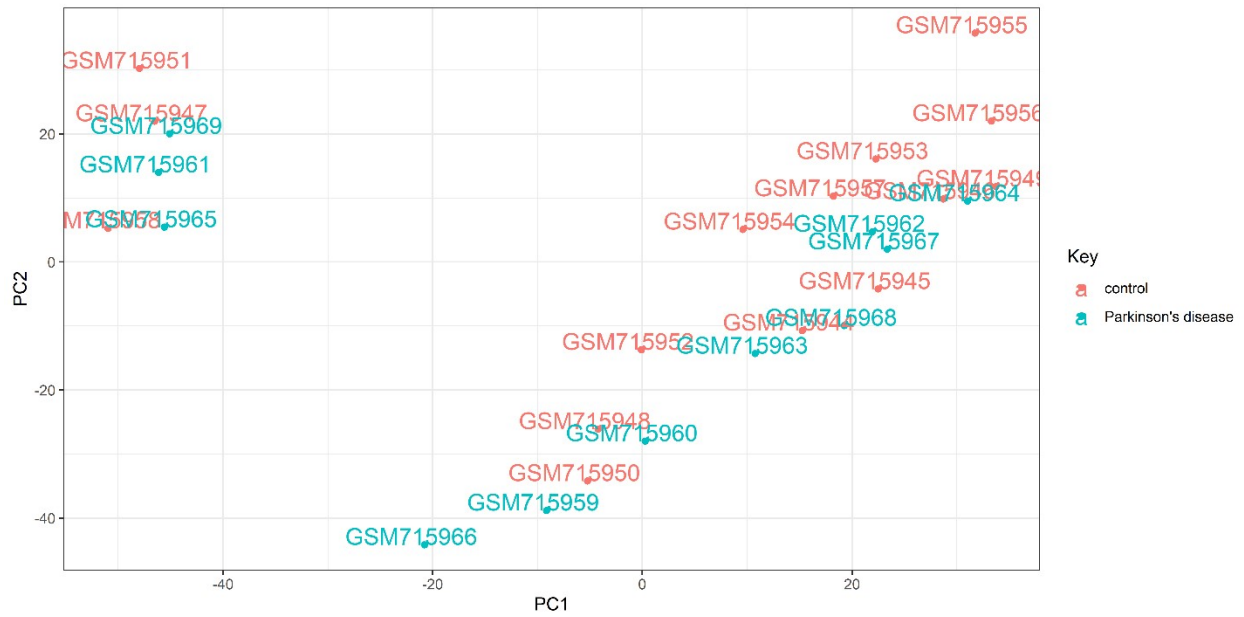
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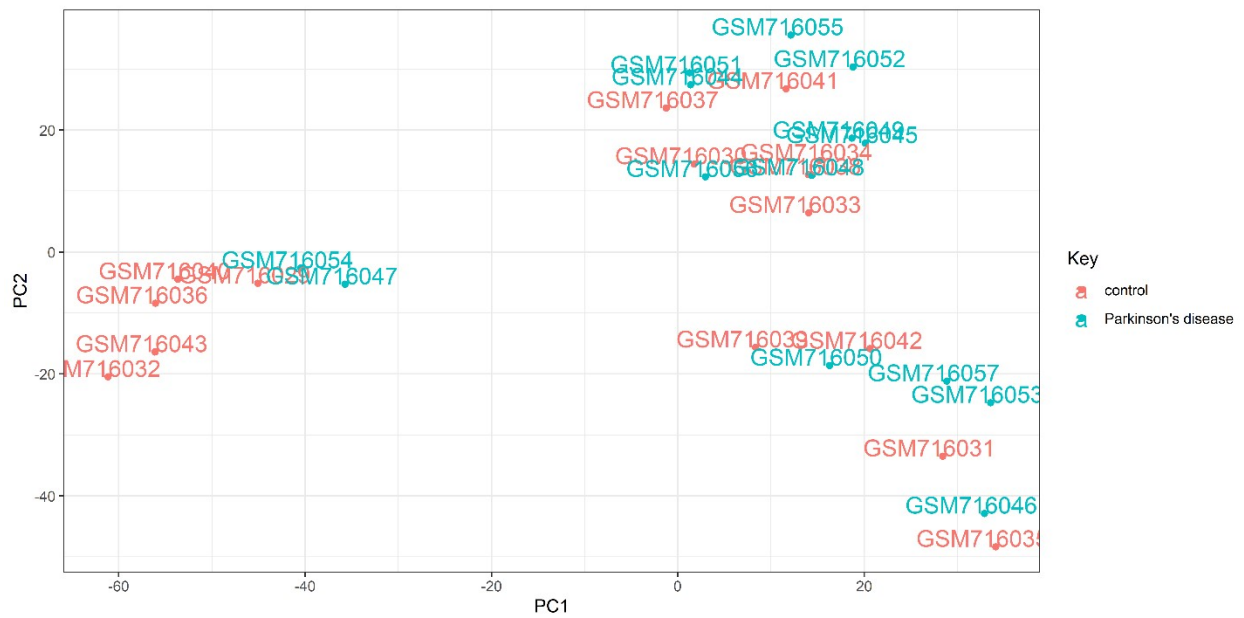
GSE28894: CB



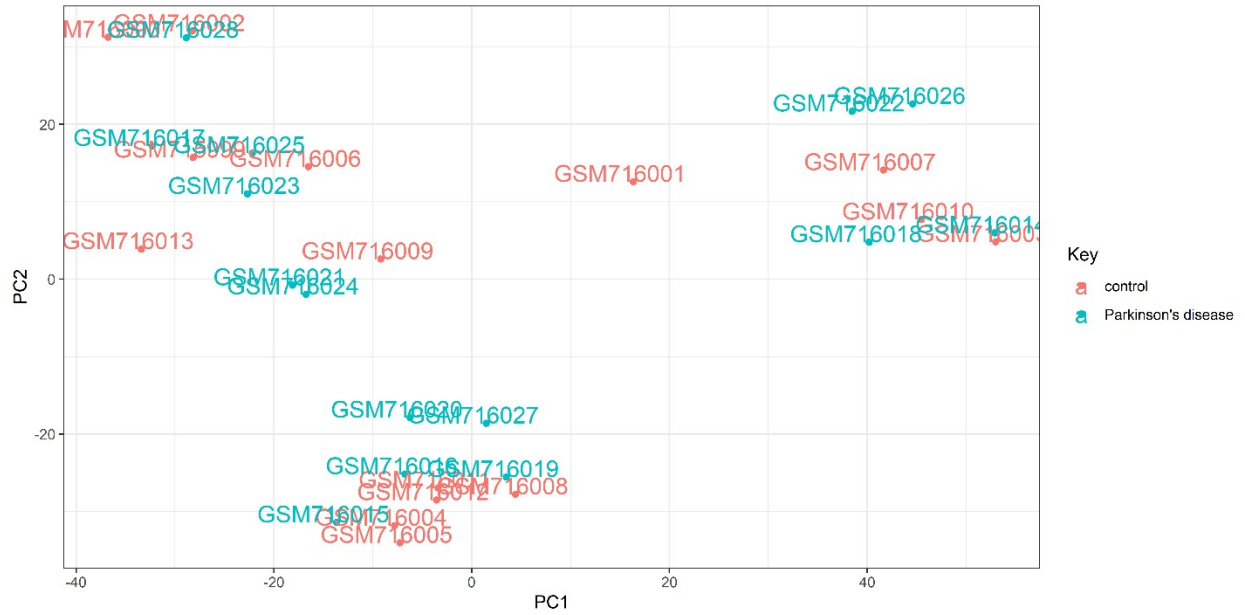
GSE28894: FCX



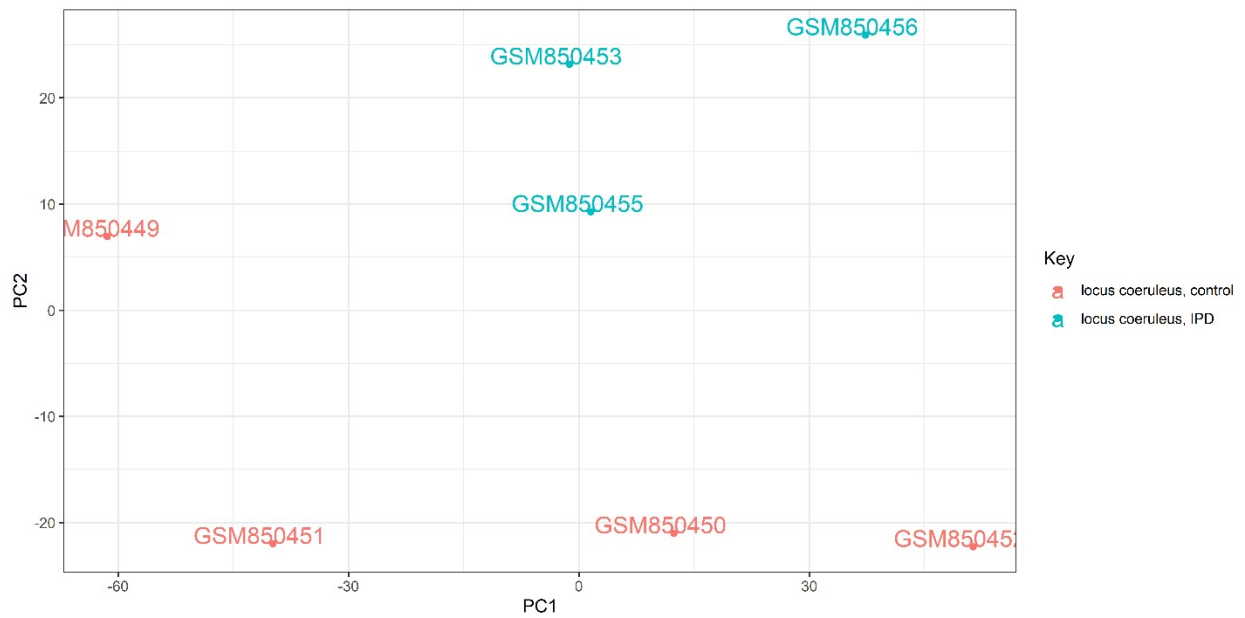
GSE28894: MED



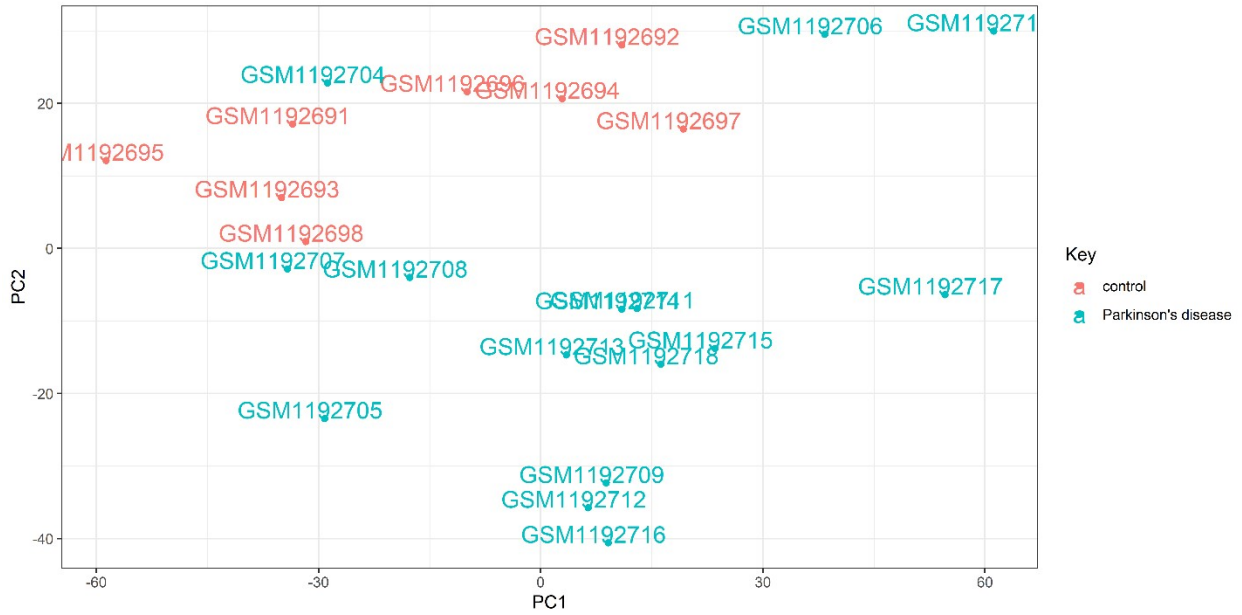
GSE28894: STR



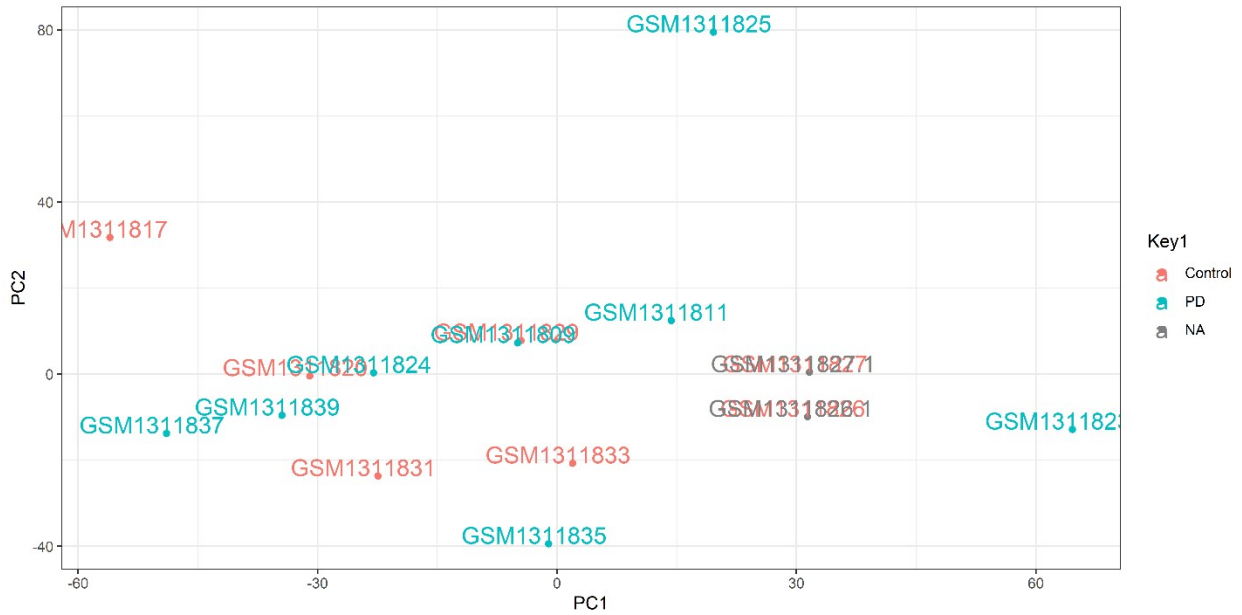
GSE34516: LOC

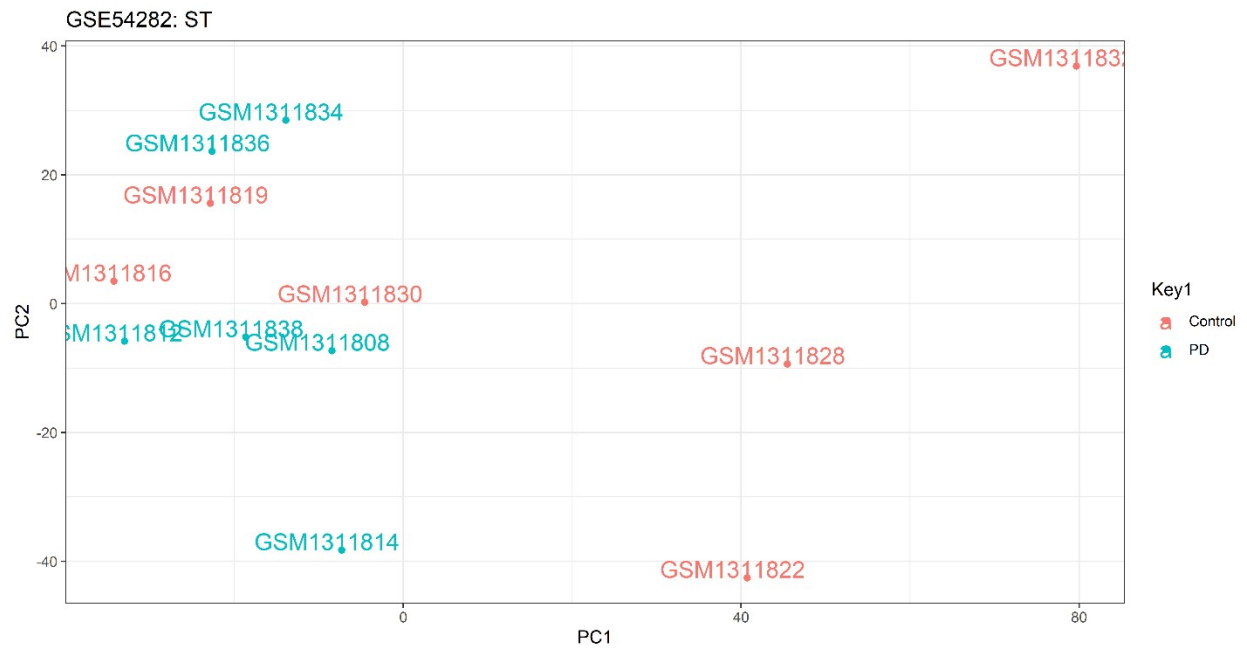
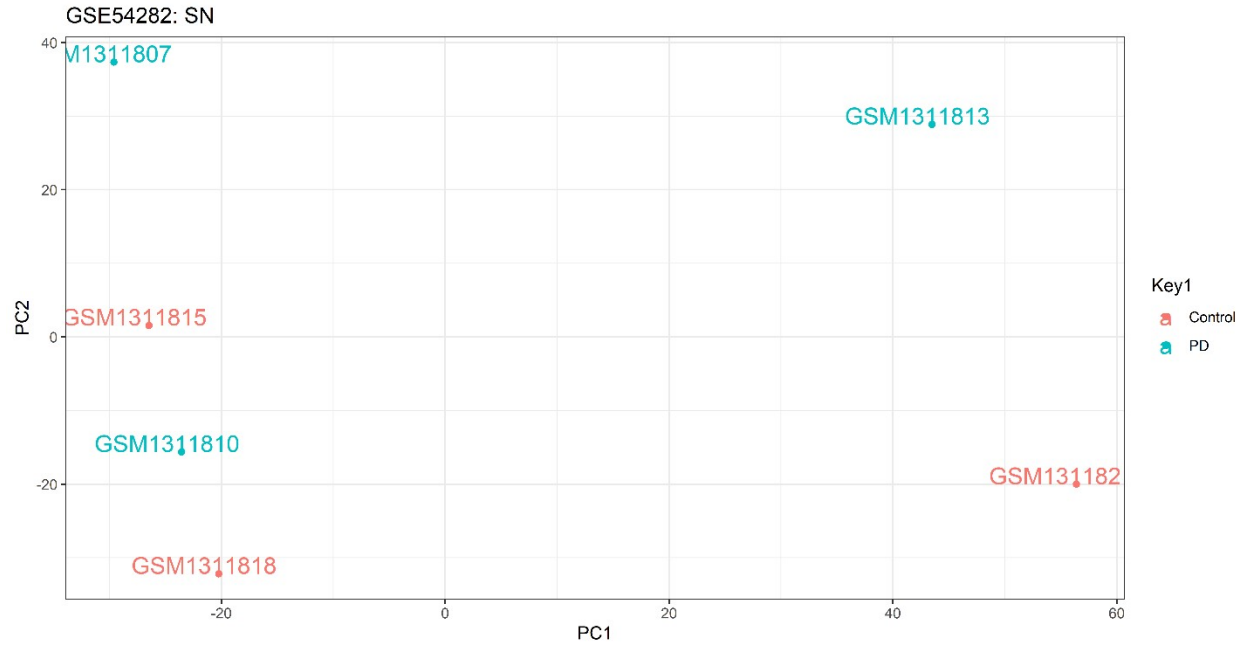


GSE49036: SN

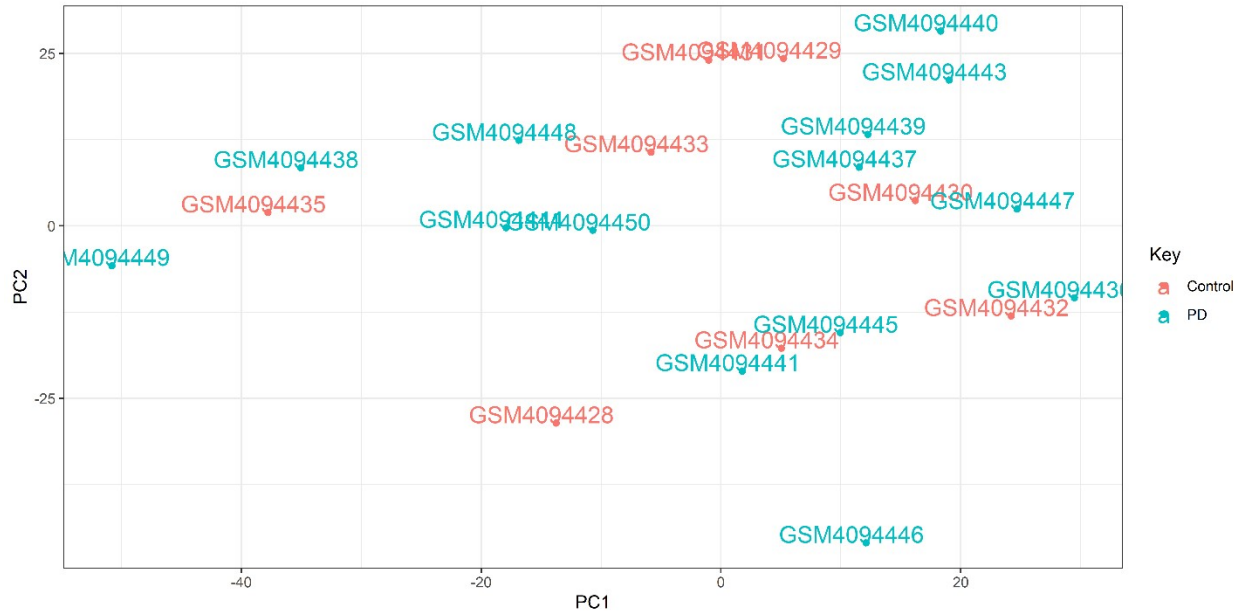


GSE54282: CO

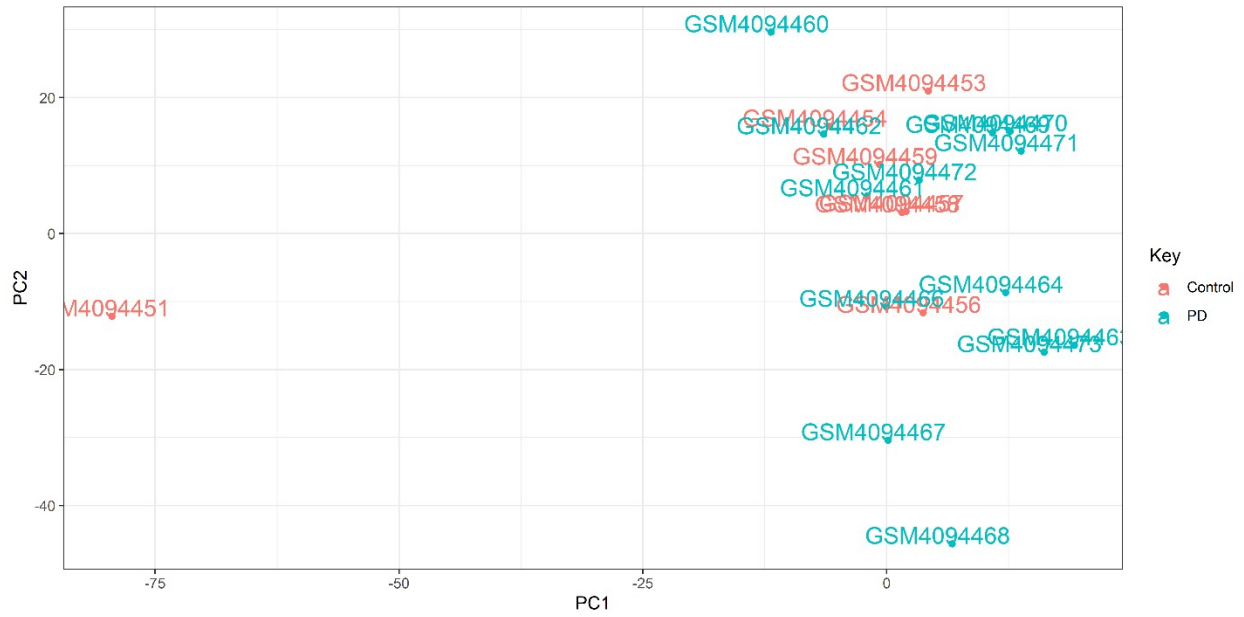




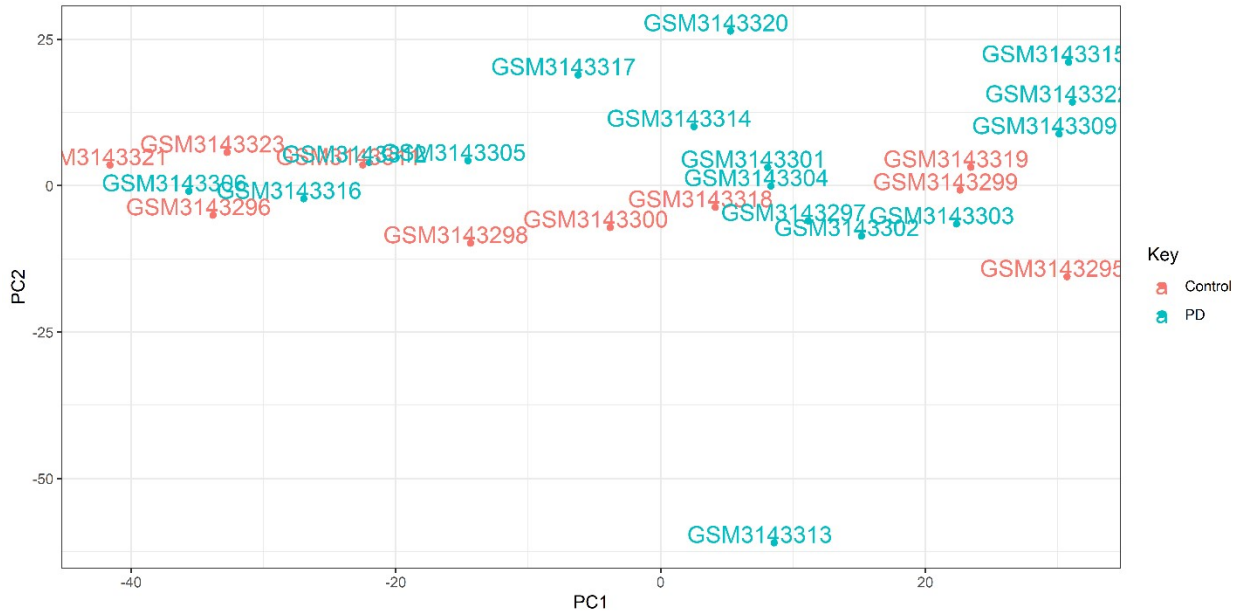
GSE114517: AM



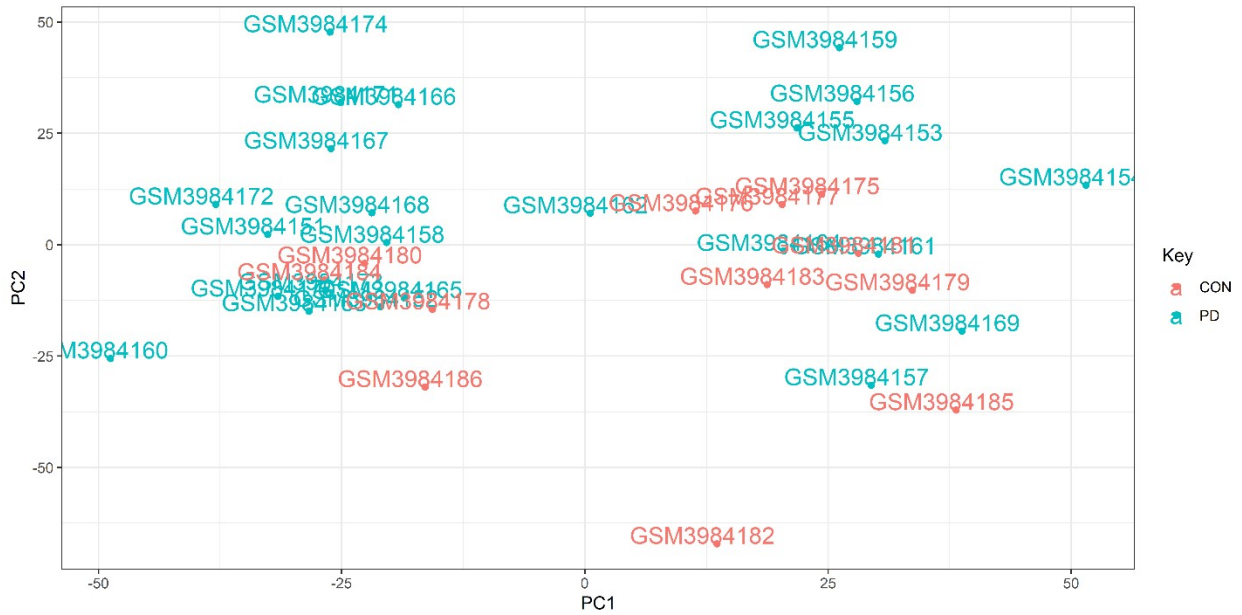
GSE114517: MTG



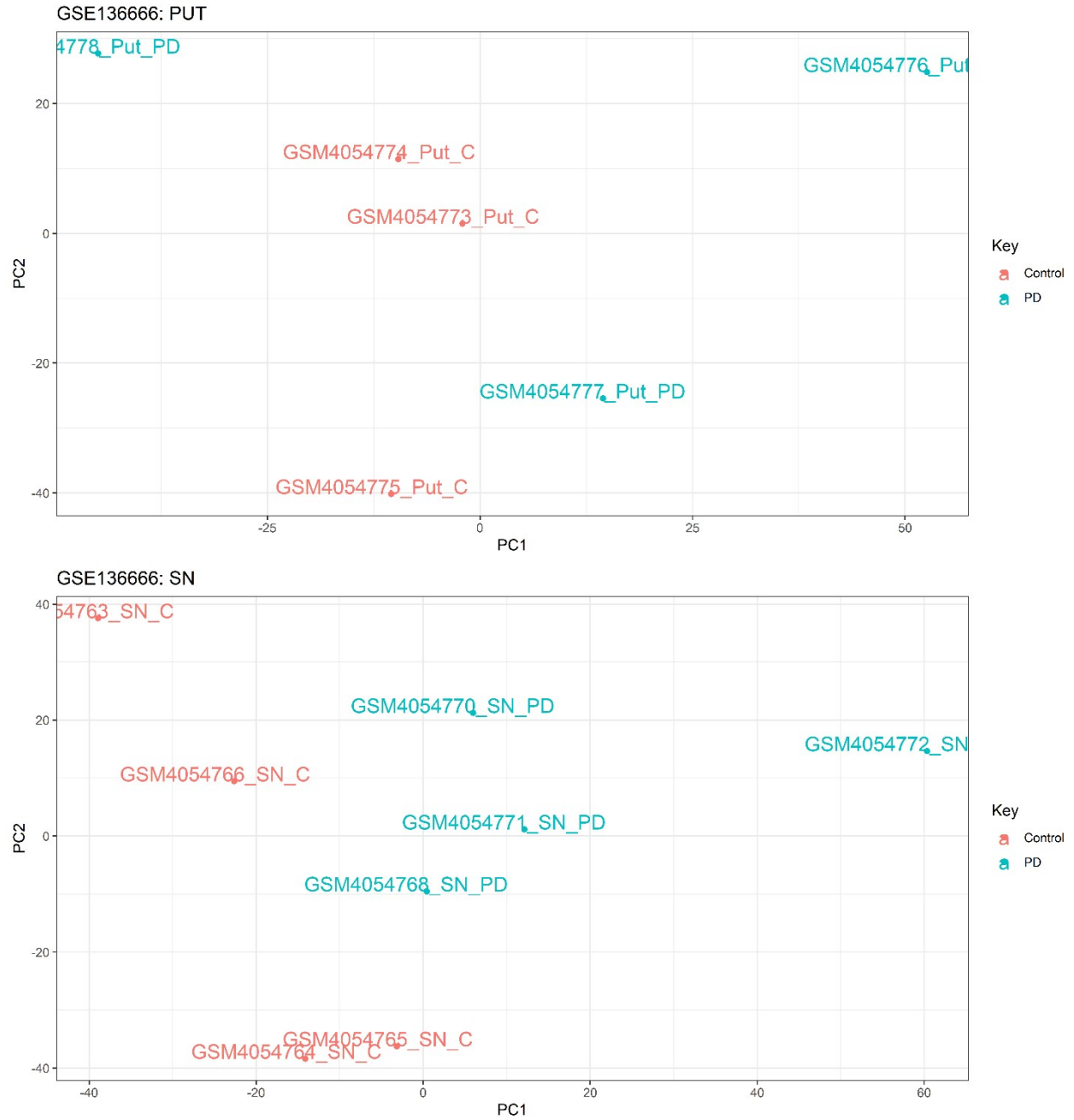
GSE114517: SN



GSE135036: PCX







**Supplementary Figure 1:** PCA plots of the first two principal components performed on the human-derived transcriptome data.