| File ID | Feature type | Data type | Tissue | Age (years) | Sex | Source |
|-------------|--------------------|---------------------------------------|--------|----------------|--------|--------|
| ENCFF516PTT | EZH2 | fold change over control | B cell | 27 | Female | ENCODE |
| ENCFF071CIY | RNF2 | fold change over control | K562 | NA | NA | ENCODE |
| ENCFF857HEZ | RNF2 | fold change over control | K562 | NA | NA | ENCODE |
| ENCFF320VKN | RNF2 | fold change over control | K562 | NA | NA | ENCODE |
| ENCFF847TGB | RNF2 | fold change over control | K562 | NA | NA | ENCODE |
| ENCFF737OJY | H3K27ac | fold change over control | PBMC | 32 | Male | ENCODE |
| ENCFF303YKC | H3K4me3 | fold change over control | PBMC | 32 | Male | ENCODE |
| ENCFF643USH | H3K36me3 | fold change over control | PBMC | 32 | Male | ENCODE |
| ENCFF249WVX | H3K36me3 | fold change over control | PBMC | 28 | Male | ENCODE |
| ENCFF759GIZ | H3K27ac | fold change over control | PBMC | 28 | Female | ENCODE |
| ENCFF412KUE | H3K27me3 | fold change over control | PBMC | 32 | Male | ENCODE |
| ENCFF455IGC | H3K9ac | fold change over control | PBMC | 28 | Male | ENCODE |
| ENCFF457WMB | H3K4me1 | fold change over control | PBMC | 32 | Male | ENCODE |
| ENCFF211ORP | H3K9ac | fold change over control | PBMC | 27 | Male | ENCODE |
| ENCFF171WZC | H3K9me3 | fold change over control | PBMC | 27 | Male | ENCODE |
| ENCFF573QMJ | H3K4me3 | fold change over control | PBMC | 27 | Male | ENCODE |
| ENCFF150RIG | H3K27me3 | fold change over control | PBMC | 28 | Female | ENCODE |
| ENCFF033IPJ | H3K9me3 | fold change over control | PBMC | 28 | Female | ENCODE |
| ENCFF796FFT | H3K4me3 | fold change over control | PBMC | 28 | Female | ENCODE |
| ENCFF100NYH | H3K4me1 | fold change over control | PBMC | 27 | Male | ENCODE |
| ENCFF713QZB | H3K9me3 | fold change over control | PBMC | 32 | Male | ENCODE |
| ENCFF265VZG | H3K27me3 | fold change over control | PBMC | 28 | Male | ENCODE |
| ENCFF319EBK | H3K9me3 | fold change over control | PBMC | 28 | Male | ENCODE |
| ENCFF754LBN | RNA-seq | minus strand signal of unique reads | PBMC | 52 | Female | ENCODE |
| ENCFF398HDS | RNA-seq | plus strand signal of unique reads | РВМС | 52 | Female | ENCODE |
| GSM923447 | Replication timing | Wavelet-transformed signals | IMR90 | NA | Female | GEO |
| GSM1289416 | LaminB1 | Normalised read counts | IMR90 | NA | NA | GEO |