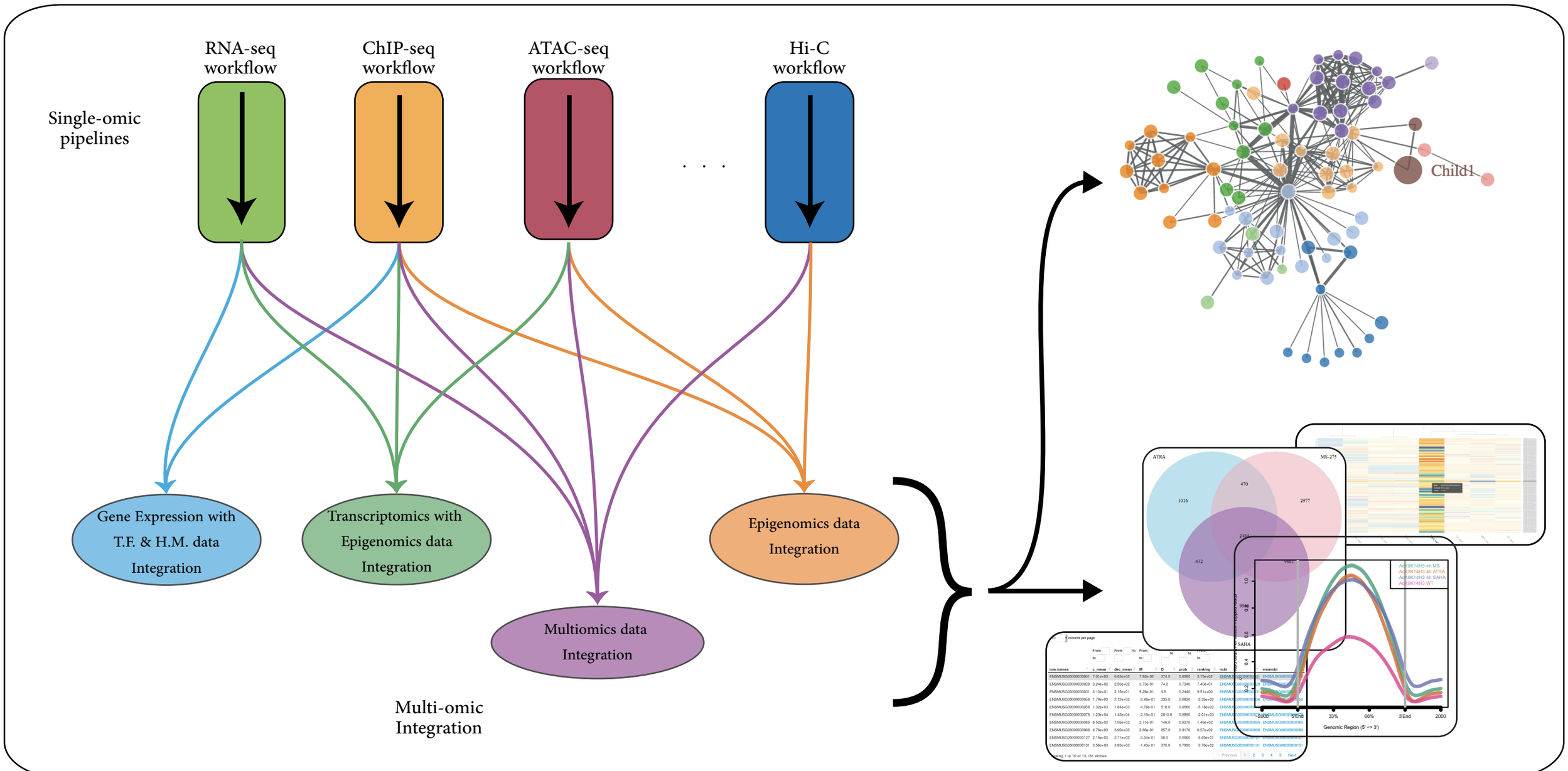
# IntegrHO


Single-omic pipelines
RNA-seq workflow
ChIP-seq workflow
ATAC-seq workflow
Hi-C workflow
. . .
Gene Expression with T.F. & H.M. data Integration
Transcriptomics with Epigenomics data Integration
Multiomics data Integration
Epigenomics data Integration
Multi-omic Integration
Child1
ATRA
MS-275
SAHA
470
1016
2977
2454
432
9881
9986
Ac3K14H3 4h MS
Ac3K14H3 4h ATRA
Ac3K14H3 4h SAHA
Ac3K14H3 WT
5'End
33%
66%
3'End
2000
Genomic Region (5' -> 3')