

Supplementary Figure

Multi-omics approaches for revealing the epigenetic regulation of histone H3.1 during spermatogonial stem cell differentiation in vitro

Figure S1: Extracted ion chromatograms of histone post-translational modification peptides by PRM

(A) Representative extracted ion chromatograms of histone post-translational modification peptides from 0h, 24h, and 48h differentiated SSCs. Heavy peptide with BSA is used as a negative control.

Figure S1



Figure S2: Differentiation trajectory of SSCs compared with another two published datasets.

Figure S2

