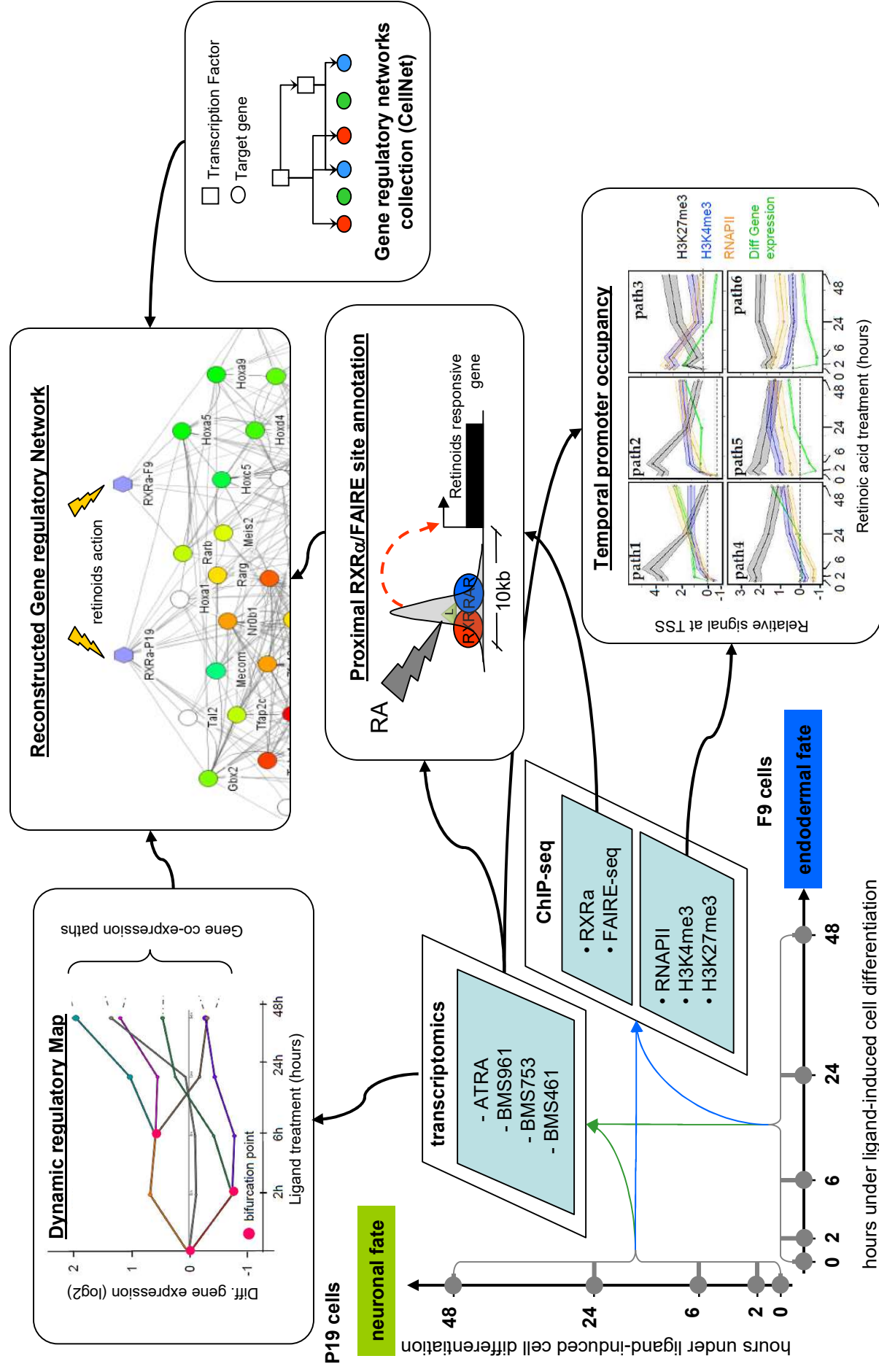




Supplemental Figure S2: Sketch of the data types generated and overall analysis. Schematic representation of the different readouts assessed during RA-induced cell differentiation in P19 and F9 cells, as well as their integration to reconstruct a global gene regulatory view of the neuronal and endodermal cell fate transitions.