



Supplementary Figure 2

(a) Ensembl genome browser screenshot showing normalized RNA-seq read coverage (mean of the two biological replicates) at the *Nr4a2* (*Nurr1*) locus. Plus (green) and minus (blue) strand reads are displayed in separate tracks.

(b) Gene ontology terms associated with 7SK-regulated genes. Enrichment *P*-values were adjusted using the Benjamini and Hochberg multiple testing correction method.