

## Supplementary Material

<b>File</b>	<b>Content</b>	<b>Format</b>	<b>Size</b>
SM1	BED files for SRC-3 and Pol II ChIP-Seq experiments	txt	1.8 MB
SM2	SRC-3 /Pol II ChIP-Seq Binding Sites Analysis: counts and genomic mappings (includes ER and FoxA1 comparative analyses), peak distance distributions (comparison between SRC-3 and ER binding sites), TSS distance density plots, and comparison to known/suspected ER target genes.	xls	144 KB
SM3	Validation Experiments: accompanying ChIP-chip experiments, IP validation by qPCR	doc	312 KB
SM4	MCF-7 E2 Gene Expression Consensus Array and integrated ChIP binding site analysis: database exports for extended promoter (EPR) and gene transcribed (G10) regions, 2-set and 3-set intersections of potential target genes, and gene lists for select expression classes used for the functional classification of gene regulation.	xls	3.2 MB
SM5	Functional Annotation of Gene Regulation: Summary and enriched DAVID annotations for select expression classes	xls	2.1 MB
SM6	STAMP output of enriched motifs for SRC-3 ChIP-Seq binding sites	pdf	3.7 MB
SM7	Potential SRC-3/RFX1 target genes that have a disease annotation	xls	40 KB
SM8	ER/SRC3 IP/MS	pdf	764 KB
SM9	Primer sequences used for quantitative real-time CR	xls	20 KB