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BCell CD133HSC	Human_BCell_Meth Human_CD133HSC_Meth	
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H1BMP4 H1-mesendoderm	Human_H1BMP4_Meth Human_H1-mesendoderm_Meth	1 11
H1-NPC Mesenchymal	Human_H1-NPC_Meth Human_Mesenchymal_Meth	<u> </u>
IMR90 BS 1a IMR90 Cell Line DN	Human_IMR90_Meth NA Methylation by Bisulfite-seq Signal from REMC/UCSD (Library:methylC-seq_imr90_r1a)	
BloodALLL2	Human_BloodALLL1_Meth	n n n
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Layered H3K4Me1	e1 Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE 3K4Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE	
DNase Clusters	DNasel Hypersensitivity Clusters in 125 cell types from ENCODE (V3) tion Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs	
Txn Factor ChIP LNG.IMR90 LNG.IMR90	chromHMM tracks from Roadmap	
Restr Enzymes	Restriction Enzymes from REBASE Non-Human RefSeq Genes	
Salmo hoxo Salmo hoxo Salmo hoxo Danio ho:	c12ba	
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-4 _ Rhesus Mouse	Salmo hoxd12ab >>>>>> Caenorh nob-1 >>>>> UCSC annotations of RefSeq RNAs (NM_* and NR_*)	
Dog Elephant Opossum Chicken	Salmo hoxd12ab >>>>>> Caenorh nob-1 >>>>> UCSC annotations of RefSeq RNAs (NM_* and NR_*) XC12L	
X_tropicalis Zebrafish RepeatMasker	Salmo hoxd12ab >>>>>> Caenorh nob-1 \$>>>> UCSC annotations of RefSeq RNAs (NM_* and NR_*) XC12 CpG Islands (Islands < 300 Bases are Light Green) Placental Mammal Basewise Conservation by PhyloP	