1147 1149 1150 1151 1152 1153 1154 884 885 886 97 98 1101 1103	.		NANOGP1 lincRNA and TUCP transcripts C/D and H/ACA Box snoRNAs, scaRNAs, and microRNAs from snoRNABase and miRBase CTR147 CpG merge methylation level	
53 54 4 5 6 7 8 8 01	11 11		CTR151 CpG merge memylation level	
5 7 3 3 01		111	CTR152 CpG merge methylation level	
7 3 01 03			CTR84 CpG merge methylation level CTR85 CpG merge methylation level CTR85 CpG merge methylation level CTR85 CpG merge methylation level CTR86 CpG merge methylation level	
03		1 1		
			CTR101 CpG merge methylation level	
06 07 08		111 1	CTR106 CpG merge methylation level	
10 32 34		1 111 1	CTR110 CpG merge methylation level CTR132 CpG merge methylation level CTR134 CpG merge methylation level CTR134 CpG merge methylation level	
48 11	 	1111 1	CTR114 CpG merge methylation level	
17			CTR114 CpG merge methylation level CTR17 CpG merge methylation level CTR17 CpG merge methylation level CTR18 CpG merge methylation level	
18 26 27	11		CTR126 CpG merge methylation level	
28 29 31		111 1	CTR128 CpG merge methylation level	
6 03 3 3 11 BS 03			UCSD Adipose Tissue Bisulfite-Seq Donor STL003 EA Release 9	
BS 03 nagus BS 03 3S 96 66			UCSD Esophagus Bisulfite-Seq Donor STL003 EA Release 9	
c BS 03	11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11. 11		BI Fetal Thymus Bisulfite-Seq Donor UW H24943 Library WGBS_Lib 65 EA Release 9 UCSD Gastric Bisulfite-Seq Donor STL003 EA Release 9 UCSD Left Ventricle Bisulfite-Seq Donor STL001 EA Release 9	
3 01 3 03 BS 02			UCSD Left Ventricle Bisulfite-Seq Donor STL003 EA Release 9 UCSD Lung Bisulfite-Seq Donor STL002 EA Release 9	
BS 02 eas BS 03				
S 03 S 03			UCSD Right Atrium Bisulfite-Seq Donor STL003 EA Release 9	
6 03 6 01 6 03			UCSD Sigmoid Colon Bisulfite-Seq Donor STL001 EA Release 9 UCSD Sigmoid Colon Bisulfite-Seq Donor STL001 EA Release 9 UCSD Sigmoid Colon Bisulfite-Seq Donor STL003 EA Release 9	
01 n BS 03			UCSD Small Intestine Bisulfite-Seq Donor STL001 EA Release 9 UCSD Spleen Bisulfite-Seq Donor STL003 EA Release 9 UCSD Spleen Bisulfite-Seq Donor STL003 EA Release 9 UCSD Thymus Bisulfite-Seq Donor STL001 EA Release 9	
Methyl 2 y Methyl 2			DNA methylation in kidney tissue (bigWig)	
nta1 Methyl 2			DNA methylation in placenta (biological replicate 1) (bigWig) DNA methylation in placenta (biological replicate 2) (bigWig) DNA methylation in placenta (biological replicate 2) (bigWig)	
ellum			DNA methylation in placenta (biological replicate 3) (bigWig)	
n alPancreas1	1		Human_NKcells_Meth Human_Sperm_Meth Human_NormalPancreas1_Meth	
alPancreas2			Human_NormalPancreas2_Meth	
rmis-old-sun-ex rmis-old-sun-pro			Human_Epidermis-old-sun-exposed_Meth Human_Epidermis-old-sun-protected_Meth Human_Epidermis-old-sun-protected_Meth Human_Epidermis-young-sun-exposed_Meth	
rmis-young-sun-			Human_Epidermis-young-sun-protected_Meth Human_Buccals_Meth Human_Sperm_Meth	11
Healthy -100yr -Newborn			Human_BloodHealthy_Meth	
BHSC phage			Distinct Human DNA Methylomes from Different Ages, Heyn 2012: Human_PBMC_Meth Changes in Human Hematopoietic Stem Cells, Hodges 2011: Human_CD133HSC_Meth Roadmap 2015: Human_Macrophage_Meth	
BHSC			Roadmap 2015 : Human_NK_Meth Human_BCell_Meth Human_CD133HSC_Meth	
SHSC			Human_HSPC_Meth Human_Neut_Meth Human_Neut_Meth	
P4			Human_H1mesendoderm_Meth Human_H1-mesendoderm_Meth Human_H1-mesendoderm_Meth	
oc achymal			Human Mesenchymal Meth Human JMR90 Meth	
D BS 1a ALLL2 ALLL1		IMR9	D Cell Line DNA Methylation by Bisulfite-seq Signal from REMC/UCSD (Library:methylC-seq_imr90_r1a)	<u> </u>
Cancer		focal DNA hy	Human_IMR90_Meth Human_MCF7_Meth Human_MCF3_Meth	
Cancer 954		Inc	Human Breast Cancer, Hon 2012: Human HCC1954_Meth Human HepG2_Meth	
paticCancer1 paticCancer2 paticCancer3			Human_PancreaticCancer1_Meth Human_PancreaticCancer2_Meth Human_PancreaticCancer3_Meth Human_PancreaticCancer3_Meth	
eaticCancer4			Human_PancreaticCancer4_Meth Human_PancreaticCancer5_Meth Human_PancreaticCancer6_Meth Human_PancreaticCancer6_Meth	
paticCancer6 paticCancer7 paticCancer8			Human PancreaticCancer8 Meth Human PancreaticCancer8 Meth Human PancreaticCancer8 Meth Human PancreaticCancer9 Meth	
paticCancer9 paticCancer10 paticCancer11			Human PancreaticCancer10_Meth Human PancreaticCancer11_Meth	
d H3K4Me1			H3K27Ac Mark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE H3K4Me1 Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE H3K4Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE	
DNase Clusters xn Factor ChIP LNG.IMR90			DNasel Hypersensitivity Clusters in 125 cell types from ENCODE (V3) Transcription Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs chromHMM tracks from Roadmap	
LNG.IMR90 Restr Enzymes Felis NANOG Ovis SLC2A3	····		Restriction Enzymes from REBASE]]
Bos SLC2A3 Mus Slc2a3 SLC2A14 SLC2A14 SLC2A14 SLC2A14			UCSC annotations of RefSeq RNAs (NM_* and NR_*)	· · · · ·
SLC2A14 SLC2A14 SLC2A14 4 al Cons		· · · · · · · · · · · · · · · · · · ·	CpG Islands (Islands < 300 Bases are Light Green) CpG: 52 Placental Mammal Basewise Conservation by PhyloP	
-4 _ Rhesus Mouse Dog			Multiz Alignments of 46 Vertebrates	