Scale chr15:		83,952,50¢	83,95: UCS	1 kb hg19 3,000 83,953,500 83,954,000 83,954,500 83,955,000 83 SC Genes (RefSeq, GenBank, CCDS, Rfam, tRNAs & Comparative Genomics)	s,955,50d
BNC1 CTR147			C/D and	lincRNA and TUCP transcripts  H/ACA Box snoRNAs, scaRNAs, and microRNAs from snoRNABase and miRBase  CTR147 CpG merge methylation level	1:1
CTR149 CTR150				CTR149 CpG merge methylation level  CTR150 CpG merge methylation level	
CTR151 CTR152	. 1	. Non-1. L.		CTR151 CpG merge methylation level	1 1
CTR153				CTR153 CpG merge methylation level	
CTR84			. a.	CTR84 CpG merge methylation level  CTR85 CpG merge methylation level	111
CTR86				CTR86 CpG merge methylation level  CTR97 CpG merge methylation level	1 111
CTR101			<b></b>	CTR98 CpG merge methylation level	
CTR103			. la	CTR103 CpG merge methylation level	İİ
CTR106 CTR107	. d		 àl	CTR106 CpG merge methylation level  CTR107 CpG merge methylation level	1 1
CTR108				CTR108 CpG merge methylation level	1 11
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CTR126 CTR127	-			CTR126 CpG merge methylation level  CTR127 CpG merge methylation level	. 1
CTR128	. 1			CTR128 CpG merge methylation level  CTR129 CpG merge methylation level	
CTR131 AT BS 03				CTR131 CpG merge methylation level UCSD Adipose Tissue Bisulfite-Seq Donor STL003 EA Release 9	1 11
AL BS 3 11 Aorta BS 03	T to Face of the Art	ntina ni ikina a a a a a a a a a a a a a a a a a a	adin .	BI Adult Liver Bisulfite-Seq Donor 3 Library WGBS_Lib 11 EA Release 8  UCSD Aorta Bisulfite-Seq Donor STL003 EA Release 9	
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FML BS 96 66 FT BS 43 65			BI Fet	tal Thymus Bisulfite-Seq Donor UW H24943 Library WGBS_Lib 65 EA Release 9	
Gastric BS 03	r II tarrilla illinia	udia. Bakasa i usa usi usa ina maka u	. دهامت	UCSD Gastric Bisulfite-Seq Donor STL003 EA Release 9  UCSD Left Ventricle Bisulfite-Seq Donor STL001 EA Release 9	
LV BS 03	t blat to	.6		UCSD Left Ventricle Bisulfite-Seq Donor STL003 EA Release 9  UCSD Lung Bisulfite-Seq Donor STL002 EA Release 9	
Lung BS 02 Ovary BS 02			aud Berrein	UCSD Ovary Bisulfite-Seq Donor STL002 EA Release 9	
Pancreas BS 03 PM BS 03	I what he are	11		UCSD Pancreas Bisulfite-Seq Donor STL003 EA Release 9  UCSD Psoas Muscle Bisulfite-Seq Donor STL003 EA Release 9	
RA BS 03				UCSD Right Ventricle Bisulfite-Seq Donor STL003 EA Release 9	
	I trad in the			UCSD Sigmoid Colon Bisulfite-Seq Donor STL001 EA Release 9	
SC BS 03 SI BS 01				UCSD Sigmoid Colon Bisulfite-Seq Donor STL003 EA Release 9  UCSD Small Intestine Bisulfite-Seq Donor STL001 EA Release 9	
Spleen BS 03	total action			UCSD Spleen Bisulfite-Seq Donor STL003 EA Release 9  UCSD Thymus Bisulfite-Seq Donor STL001 EA Release 9	
Thymus BS 01 Brain Methyl 2	. h lan a na c			DNA methylation in brain tissue (bigWig)	
Kidney Methyl 2 Placenta1 Methyl 2				DNA methylation in kidney tissue (bigWig)	
Placenta2 Methyl 2				DNA methylation in placenta (biological replicate 2) (bigWig)	
Cerebellum  Kidney				Human_Cerebellum_Meth Human_Kidney_Meth	
NKcells Sperm				Human_NKcells_Meth Human_Sperm_Meth	
NormalPancreas1 NormalPancreas2			<u></u>	Human_NormalPancreas1_Meth Human_NormalPancreas2_Meth	
93A 93N				Human_93N_Meth  Human_93N_Meth	
Epidermis-old-sun-ex	1 11 1 1			Human_Epidermis-old-sun-exposed_Meth  Human_Epidermis-old-sun-protected_Meth	
Epidermis-young-sun				Human_Epidermis-young-sun-exposed_Meth  Human_Epidermis-young-sun-protected_Meth	
Buccals Sperm		<u> </u>		Human_Buccals_Meth  Human_Sperm_Meth	
BloodHealthy  CD4T-100yr			Distinct Hu	Human_BloodHealthy_Meth	
CD4T-Newborn PBMC		Dit		nan DNA Methylomes from Different Ages, Heyn 2012 : Human_CD4T-Newborn_Meth	
CD133HSC  Macrophage		1 31 21	Change	as in Human Hematopoietic Stem Cells, Hodges 2011 : Human_CD133HSC_Meth  Roadmap 2015 : Human_Macrophage_Meth	
NK BCell				Roadmap 2015 : Human_NK_Meth  Human_BCell_Meth	
CD133HSC		111111111111111111111111111111111111111	1	Human_CD133HSC_Meth  Human_HSPC_Meth	
Neut H1				Human_Neut_Meth  Human_H1_Meth	
H1BMP4 H1-mesendoderm				Human_H1-mesendoderm_Meth	
H1-NPC Mesenchymal				Human_H1-NPC_Meth  Human_Mesenchymal_Meth	
IMR90 BS 1a			Cell Line D	Human_IMR90_Meth  Human_IMR90_Meth  DNA Methylation by Bisulfite-seq Signal from REMC/UCSD (Library:methylC-seq_imr90_r1a)	
BloodALLL2	I litter forein	hilu Tuase, als as as how rather	and the control of	Human_BloodALLL2_Meth  Human_BloodALLL1_Meth	11111111111111
IMR90 MCF7	<u> </u>	11 <mark>8</mark> hij l x 186		Human_IMR90_Meth Human_MCF7_Meth	
MCF7 ColonCancer ColonCancer		<u> Instanta di Gold I ana ilia ilibi II II a lumin</u>	na la i	on and long-range hypomethylation in colorectal cancer, Berman 2012 : Human_ColonCancer_Meth  the till the first transfer of the color	
HCC1954 HepG2	<u> </u>			Human Breast Cancer, Hon 2012 : Human HCC1954_Meth Human HepG2_Meth	
PancreaticCancer1 PancreaticCancer2		allalid double and a lift the beat of the last of the		Human_PancreaticCancer1_Meth  Human_PancreaticCancer2_Meth	
PancreaticCancer2 PancreaticCancer3 PancreaticCancer4		1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1		Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth	
PancreaticCancer4  PancreaticCancer5  PancreaticCancer6		I have all the distribution of the		Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth	
PancreaticCancer7				Human_PancreaticCancer8_Meth Human_PancreaticCancer8_Meth	
PancreaticCancer8  PancreaticCancer9  PancreaticCancer10		Table of the statement of the		Human_PancreaticCancer9_Meth Human_PancreaticCancer10_Meth Human_PancreaticCancer10_Meth	
PancreaticCancer11			H3K27A^	Human_PancreaticCancer11_Meth Human_PancreaticCancer11_Meth Human_PancreaticCancer11_Meth Human_PancreaticCancer11_Meth	
Layered H3K27Ac			H3K4N	Me1 Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE	
Layered H3K4Me3  DNase Clusters				13K4Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  DNasel Hypersensitivity Clusters in 125 cell types from ENCODE (V3)	
Txn Factor ChIP  LNG.IMR90  LNG.IMR90			Transcri	ption Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs  chromHMM tracks from Roadmap	
Restr Enzymes Rattus Bnc1				Restriction Enzymes from REBASE  Restriction Enzymes from REBASE  Non-Human RefSeq Genes	11 11 11 11 11 11 11 11 11 11 11 11 11
Mus Bnc1  BNC1  BNC1		*****************	******	UCSC annotations of RefSeq RNAs (NM_* and NR_*)  CpG Islands (Islands < 300 Bases are Light Green)	
4 _ Mammal Cons -4	CpG: 188	A throw we that I was sufficient of the last last last last last last last last	uhandan open pirapo	CpG Islands (Islands < 300 Bases are Light Green) Placental Mammal Basewise Conservation by PhyloP	معدد والمراجعة المعاونة والمعاونة وا
-4 _ Rhesus Mouse Dog				Multiz Alignments of 46 Vertebrates	
Elephant Opossum Chicken X_tropicalis					
Zebrafish RepeatMasker				Repeating Elements by RepeatMasker	