Scale chr10: PRLHR		PRLH	1 kb   hg19 120,354,000	57,000 120,357	7,500
TR147 TR149		11 :	C/D and H/ACA Box snoRNAs, scaRNAs, and microRNAs from snoRNABase and miRBase  CTR147 CpG merge methylation level  ا المادة الم		
TR150 TR151	I II	11 .	CTR150 CpG merge methylation level	11	
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TR154 TR84 TR85			CTR84 CpG merge methylation level		I
TR86	 		CTR97 CpG merge methylation level  CTR96 CpG merge methylation level  CTR97 CpG merge methylation level		
TR98 TR101	ı 	.1		,      <sub>11</sub>	
TR103	di   	[ ]	CTR103 CpG merge methylation level	      	
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L BS 3 11 orta BS 03			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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V BS 03 ung BS 02			UCSD Lung Bisulfite-Seq Donor STL002 EA Release 9		
vary BS 02 ancreas BS 03	111		UCSD Pancreas Bisulfite-Seq Donor STL003 EA Release 9		
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C BS 03		11 1	UCSD Sigmoid Colon Bisulfite-Seq Donor STL003 EA Release 9  UCSD Sigmoid Colon Bisulfite-Seq Donor STL003 EA Release 9  UCSD Sigmoid Colon Bisulfite-Seq Donor STL001 EA Release 9  UCSD Small Intestine Bisulfite-Seq Donor STL001 EA Release 9		
pleen BS 03			UCSD Spleen Bisulfite-Seq Donor STL003 EA Release 9		
hymus BS 01 rain Methyl 2					
idney Methyl 2 lacenta1 Methyl 2	Hi i		DNA methylation in kidney tissue (bigWig)		Í
lacenta2 Methyl 2	111 1		DNA methylation in placenta (biological replicate 2) (bigWig)		
erebellum	att I	1 3 1	Human_Cerebellum_Meth		
Kcells perm			Human_NKcells_Meth  Human_Sperm_Meth  Human_NormalPancreas1_Meth		
ormalPancreas1 ormalPancreas2			Human_NormalPancreas1_Meth  Human_NormalPancreas2_Meth  Human_NormalPancreas2_Meth  Human_93A_Meth		
3N pidermis-old-sun-ex			Human_93N_Meth  Human_Epidermis-old-sun-exposed_Meth		
pidermis-old-sun-pro pidermis-young-sun			Human_Epidermis-young-sun-exposed_Meth    Human_Epidermis-young-sun-exposed_Meth		
oidermis-young-sun- uccals			Human_Epidermis-young-sun-protected_Meth		
perm podHealthy D4T-100yr	111		Human_Sperm_Meth  Human_BloodHealthy_Meth		
D4T-Newborn BMC			Distinct Human DNA Methylomes from Different Ages, Heyn 2012: Human_PBMC_Meth		
D133HSC acrophage			Changes in Human Hematopoietic Stem Cells, Hodges 2011 : Human_CD133HSC_Meth		
Cell			Roadmap 2015: Human_NK_Meth  [IIII   I   I   I   I   I   I   I   I		
D133HSC SPC			Human_CD133HSC_Meth  Human_HSPC_Meth  Human_HSPC_Meth  Human_Neut_Meth		
eut I IBMP4			Human_Neut_Meth		<u> </u>
-mesendoderm			Human_H1-mesendoderm_Meth  Human_H1-mesendoderm_Meth  Human_H1-NPC_Meth		
esenchymal			Human_Mesenchymal_Meth  Human_MR90_Meth		
R90 BS 1a			IMR90 Cell Line DNA Methylation by Bisulfite-seq Signal from REMC/UCSD (Library:methylC-seq_imr90_r1a)	. 11 in 1 #1 1	
podALLL1			Human_BloodALLL1_Meth		
lonCancer			Human_MCF7_Meth  focal DNA hypermethylation and long-range hypomethylation in colorectal cancer, Berman 2012 : Human_ColonCancer_Meth  Increased methylation variation in epigenetic domains across cancer types. : Human_ColonCancer_Meth		
C1954			Human Breast Cancer, Hon 2012 : Human HCC1954_Meth  Human HepG2_Meth	1 11 11 1 11 1	
ncreaticCancer1	1		Human_PancreaticCancer1_Meth  Human_PancreaticCancer2_Meth		
ncreaticCancer3			Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth		
ncreaticCancer5			Human_PancreaticCancer5_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth	11	
creaticCancer8			Human_PancreaticCancer7_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer9_Meth		
creaticCancer9	<u> </u>		Human_PancreaticCancer9_Meth		
ered H3K27Ac ered 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		
ered H3K4Me3			H3K4Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  DNasel Hypersensitivity Clusters in 125 cell types from ENCODE (V3)		
DNase Clusters Txn Factor ChIP LNG.IMR90			Transcription Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs  chromHMM tracks from Roadmap		
LNG.IMR90 Restr Enzymes Bos PRLHR	TTI		Restriction Enzymes from REBASE		. 1 ■
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o LOC100136462 o si:dkey-202l22.3	G	allus PRI			
PRLHR 4 _ mmal Cons		CpG:	CpG Islands (Islands < 300 Bases are Light Green)  172  Placental Mammal Basewise Conservation by PhyloP		ada J
-4 _ Rhesus Mouse			Multiz Alignments of 46 Vertebrates		
Dog Elephant Opossum Chicken X_tropicalis Zebrafish					
Zebrafish RepeatMasker			Repeating Elements by RepeatMasker		