SSTR1 →→I	38,678,50d		CSC Genes (RefSeq, GenBank, CCDS, Rfam, tRNAs & Comparative Genomics)	38,682,000
		C/Da	lincRNA and TUCP transcripts Ind H/ACA Box snoRNAs, scaRNAs, and microRNAs from snoRNABase and miRBase CTR147 CpG merge methylation level CTR149 CpG merge methylation level CTR149 CpG merge methylation level	
)			CTR150 CpG merge methylation level	
!		ka ka naha	CTR151 CpG merge methylation level CTR152 CpG merge methylation level	
	- va 1		CTR153 CpG merge methylation level	
		المال معالم المالية ال	CTR84 CpG merge methylation level	
			CTR86 CpG merge methylation level	
		التي المنظمة المنظمة التي التي التي التي التي التي التي التي	CTR97 CpG merge methylation level	
			CTR101 CpG merge methylation level	
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,		a	CTR107 CpG merge methylation level	
· ·			CTR110 CpG merge methylation level	
!		4 a l 1 1.	CTR132 CpG merge methylation level L fl CTR134 CpG merge methylation level	
•		In	CTR148 CpG merge methylation level	
.			CTR113 CpG merge methylation level CTR114 CpG merge methylation level	
			CTR117 CpG merge methylation level	
;	- i	. 1 L	CTR126 CpG merge methylation level	
			CTR127 CpG merge methylation level CTR128 CpG merge methylation level	
,		.1.3	CTR129 CpG merge methylation level	
13	L:	Line of the selection o	UCSD Adipose Tissue Bisulfite-Seq Donor STL003 EA Release 9 BI Adult Liver Bisulfite-Seq Donor 3 Library WGBS_Lib 11 EA Release 8	
S 03			BI Adult Liver Bisulfite-Seq Donor 3 Library WGBS_Lib 11 EA Release 8	
gus BS 03		and the second second	UCSD Esophagus Bisulfite-Seq Donor STL003 EA Release 9	
96 66 3 65			tal Muscle Leg Bisulfite-Seq Donor UW H24996 Library WGBS_Lib 66 EA Release 9	
BS 03			UCSD Gastric Bisulfite-Seq Donor STL003 EA Release 9	
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3 02		AND COLUMN TO STATE OF THE PARTY OF THE PART	UCSD Lung Bisulfite-Seq Donor STL002 EA Release 9	
S 02			UCSD Ovary Bisulfite-Seq Donor STL002 EA Release 9 UCSD Pancreas Bisulfite-Seq Donor STL003 EA Release 9	
03			UCSD Psoas Muscle Bisulfite-Seq Donor STL003 EA Release 9	
03		Jan ka .toa	UCSD Right Atrium Bisulfite-Seq Donor STL003 EA Release 9 UCSD Right Ventricle Bisulfite-Seq Donor STL003 EA Release 9	
01		and the state of t	UCSD Sigmoid Colon Bisulfite-Seq Donor STL001 EA Release 9	
03	itti I		UCSD Sigmoid Colon Bisulfite-Seq Donor STL003 EA Release 9 UCSD Small Intestine Bisulfite-Seq Donor STL001 EA Release 9	
SS 03		المارية المستخدمة المستخدم المستخدمة المستخدم المستخد	UCSD Spleen Bisulfite-Seq Donor STL003 EA Release 9	
BS 01		Annual of the same	UCSD Thymus Bisulfite-Seq Donor STL001 EA Release 9 DNA methylation in brain tissue (bigWig)	
ethyl 2 Methyl 2			DNA methylation in kidney tissue (bigWig)	
a1 Methyl 2	.		DNA methylation in placenta (biological replicate 1) (bigWig)	
a2 Methyl 2 a3 Methyl 2	h .m. .		DNA methylation in placenta (biological replicate 3) (bigWig)	
lum	- 1 mL11 . h		Human_Kidney_Meth	
			Human_NKcells_Meth	
Pancreas1	1 1 1 1		Human_NormalPancreas1_Meth	
Pancreas2	des. of kn 1		Human_NormalPancreas2_Meth Human_93A_Meth	
iis-old-sun-ex			Human_93N_Meth Human_Epidermis-old-sun-exposed_Meth	
is-old-sun-pro			Human_Epidermis-old-sun-protected_Meth Human_Epidermis-young-sun-exposed_Meth	
iis-young-sun			Human_Epidermis-young-sun-protected_Meth Human_Buccals_Meth	
agithus			Human_Sperm_Meth Human_BloodHealthy_Meth	
00yr		The first of cooling terms of the cooling	Human DNA Methylomes from Different Ages, Heyn 2012 : Human_CD4T-100yr_Meth	
ewborn			luman DNA Methylomes from Different Ages, Heyn 2012 : Human_CD4T-Newborn_Meth	
HSC anage		Chai	nges in Human Hematopoietic Stem Cells, Hodges 2011 : Human_CD133HSC_Meth Roadmap 2015 : Human_Macrophage_Meth	
			Roadmap 2015 : Human_NK_Meth Human_BCell_Meth	
HSC			Human_CD133HSC_Meth	
			Human_HSPC_Meth Human_Neut_Meth	
4			Human_H1_Meth Human_H1BMP4_Meth	
endoderm			Human_H1-mesendoderm_Meth Human_H1-mesendoderm_Meth	
hymal			Human_Mesenchymal_Meth	
3S 1a			Human_IMR90_Meth DNA Methylation by Bisulfite-seq Signal from REMC/UCSD (Library:methylC-seq_imr90_r1a)	
LL2		.a.a	Human_BloodALLL2_Meth Human_BloodALLL1_Meth	
			Human_IMR90_Meth	
ıncer			Human_MCF7_Meth	eth
ncer 4	n yi ha asilimilik ili	Increased m	ethylation variation in epigenetic domains across cancer types. : Human_ColonCancer_Meth	
ticCancer1			Human_HepG2_Meth Human_PancreaticCancer1_Meth	
icCancer2	1 1 1 1 1 1		Human_PancreaticCancer2_Meth	
icCancer3		. III. dan adal bersala . II. II. and an adal	Human_PancreaticCancer3_Meth Human_PancreaticCancer4_Meth	
icCancer5	11 11 11 11		Human_PancreaticCancer5_Meth Human_PancreaticCancer6_Meth	
ticCancer7		. 1 100k	Human_PancreaticCancer7_Meth Human_PancreaticCancer8_Meth	
icCancer8		. <u>1 kata ata</u> . <u>1 kata bi ira</u>	Human_PancreaticCancer9_Meth	
icCancer10	1 h		Human_PancreaticCancer10_Meth Human_PancreaticCancer11_Meth	
H3K27Ac			Ac Mark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE (4Me1 Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE	
H3K4Me1 H3K4Me3			H3K4Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE	
ase Clusters Factor ChIP		Trans	DNasel Hypersensitivity Clusters in 125 cell types from ENCODE (V3) cription Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs	
LNG.IMR90 LNG.IMR90			chromHMM tracks from Roadmap Restriction Enzymes from REBASE	
			Restriction Enzymes from REBASE	
Mus Sstr1 Rattus Sstr1	Canis SSTR1		**************************************	
Mus Sstr1 Rattus Sstr1	Sus SSTR1 Gallus SS			
Mus Sstr1 Rattus Sstr1		sstr1a →→→	UCSC annotations of RefSeq RNAs (NM_* and NR_*)	
Mus Sstr1 Rattus Sstr1 SSTR1 CpG: 187 4_	Gallus SS Oncorhy s	sstr1a →→→	UCSC annotations of RefSeq RNAs (NM_* and NR_*) CpG Islands (Islands < 300 Bases are Light Green) Placental Mammal Basewise Conservation by PhyloP	
SSTR1	Gallus SS Oncorhy s	sstr1a →→→	CpG Islands (Islands < 300 Bases are Light Green)	سفطند وإضارا فارقاه فللفظاف المخافضة فالعرباء وسطنت فالله استدرب