Scale chr5: GPRIN1 GPRIN1/SQ	LLARFLGPPF	23,900 RKAAGDPP RKAAGDPP	ATRVSGSRGPGA		I A <mark>M</mark> GL V E V E MAAGY V E	:WT <mark>M</mark> GKED\	VSVDRVPE	176,024,150 APEPPAVAAALA APEPPAVAAALA
CTR147	LLAKFLGPPF	RKAAGDPP		lincRNA and TUCP transc. H/ACA Box snoRNAs, scaRNAs, and microRN/ CTR147 CpG merge methylar CTR149 CpG merge methylar	oripts As from snoRNABase and miR tion level			
CTR149 CTR150				CTR150 CpG merge methylar				
CTR151 CTR152				CTR151 CpG merge methylar				
CTR153		=	-	CTR153 CpG merge methylat CTR154 CpG merge methylat				1 11 _
CTR84	— — — — — —			CTR84 CpG merge methylati			-	
CTR85		+		CTR86 CpG merge methylati	ion level			
CTR97				CTR97 CpG merge methylati				
CTR101 CTR103				CTR101 CpG merge methylat	L			
CTR104	-			CTR104 CpG merge methylar CTR106 CpG merge methylar		•	-	- - - - - - - - - -
CTR106 CTR107			- - -	CTR107 CpG merge methylar	tion level			
CTR108 CTR110				CTR108 CpG merge methylat		_		
CTR132 CTR134				CTR132 CpG merge methylar				
CTR148				CTR148 CpG merge methylar		-		
CTR113		 		CTR113 CpG merge methylar				
CTR114 CTR117	 - I			CTR114 CpG merge methylat				
CTR118 CTR126				CTR118 CpG merge methylar CTR126 CpG merge methylar				
CTR127				CTR127 CpG merge methylar				
CTR129				CTR129 CpG merge methylat		■ -		
AT BS 03	<u>.</u>			CTR131 CpG merge methylat UCSD Adipose Tissue Bisulfite-Seq Donor				
AL BS 3 11	ā		┲ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒ ╒	BI Adult Liver Bisulfite-Seq Donor 3 Library WC	■■			
Aorta BS 03 Esophagus BS 03		<u>- </u>		UCSD Esophagus Bisulfite-Seq Donor S				
FML BS 96 66		-	.	Muscle Leg Bisulfite-Seq Donor UW H24996 Li				
FT BS 43 65 Gastric BS 03		<u> </u>		UCSD Gastric Bisulfite-Seq Donor STL	003 EA Release 9			
LV BS 01		+		UCSD Left Ventricle Bisulfite-Seq Donor S				
LV BS 03 Lung BS 02		+		UCSD Lung Bisulfite-Seq Donor STL0	02 EA Release 9		• • -	
Ovary BS 02		+	~ + + * + + - + + + + * +	UCSD Ovary Bisulfite-Seq Donor STL0				
Pancreas BS 03 PM BS 03	al .1	_	- • - -	UCSD Pancreas Bisulfite-Seq Donor ST UCSD Psoas Muscle Bisulfite-Seq Donor S				
RA BS 03		+	<u>+</u>	UCSD Right Atrium Bisulfite-Seq Donor S UCSD Right Ventricle Bisulfite-Seq Donor S				
RV BS 03 SC BS 01		-		UCSD Right Ventricle Bisulfite-Seq Donor S	■■ _			
SC BS 03		-	_	UCSD Sigmoid Colon Bisulfite-Seq Donor S				- - - - - - - -
SI BS 01 Spleen BS 03		_ ■■ .	_	UCSD Spleen Bisulfite-Seq Donor STL	003 EA Release 9	I _		• - • - • -
Thymus BS 01		; ;		UCSD Thymus Bisulfite-Seq Donor STL			- - -	
Brain Methyl 2 Kidney Methyl 2				DNA methylation in kidney tissu				
Placenta1 Methyl 2	III 111		H . I I .	DNA methylation in placenta (biological r	1.1.		. 1 .	
Placenta2 Methyl 2 Placenta3 Methyl 2	1		1. 1	DNA methylation in placenta (biological r	eplicate 3) (bigWig)			
Cerebellum	111 1	- • - •	 <u> </u>	Human_Cerebellum_Mo		1	11.	<u>, , , , , , , , , , , , , , , , , , , </u>
NKcells				Human_NKcells_Meth	1			
Sperm NormalPancreas1				Human_NormalPancreas1				
NormalPancreas2 93A				Human_NormalPancreas2	_Meth			
93N Epidermis-old-sun-ex				Human_93N_Meth Human_Epidermis-old-sun-expo	psed_Meth			
Epidermis-old-sun-pro				Human_Epidermis-old-sun-prote		1 1		
Epidermis-young-sun-	11 11	<u> </u>	11	Human_Epidermis-young-sun-pro	stected_Meth	1 1	111	11111
Buccals Sperm	111 11			Human_Buccals_Meth				
BloodHealthy CD4T-100yr			Distinct Hu	Human_BloodHealthy_N uman DNA Methylomes from Different Ages, He		r_Meth		
CD4T-Newborn			<u> </u>	nan DNA Methylomes from Different Ages, Heyr Human DNA Methylomes from Different Ages,				
PBMC CD133HSC				es in Human Hematopoietic Stem Cells, Hodges	2011 : Human_CD133HSC_N			
Macrophage NK				Roadmap 2015 : Human_Macrop Roadmap 2015 : Human_Ni				
BCell CD133HSC				Human_BCell_Meth Human_CD133HSC_M	eth			
HSPC Neut				Human_HSPC_Meth Human_Neut_Meth				
H1	ш .1		11.1.1	Human_H1_Meth Human_H1BMP4_Met	h	<u> </u>		1
H1-mesendoderm	111 1 1			Human_H1-mesendoderm	_Meth		111	
H1-NPC Mesenchymal	111		11 1 1 1 1 1	Human_H1-NPC_Met	h			
IMR90 IMR90 BS 1a			IMR90 Cell Line D	Human_IMR90_Meth		eq_imr90_r1a)		
BloodALLL2			┡ ┼ ┡ ┼ ┡ ┼	Human_BloodALLL2_M	eth			
BloodALLL1 IMR90				Human_BloodALLL1_M Human_IMR90_Meth				
MCF7 ColonCancer		foca	al DNA hypermethylation	Human_MCF7_Meth on and long-range hypomethylation in colorecta		n_ColonCance	r_Meth	
ColonCancer HCC1954			Increased meth	ylation variation in epigenetic domains across c		ancer_Meth		
HepG2	111 11		11	Human_HepG2_Meth				
PancreaticCancer1 PancreaticCancer2			<u> </u>	Human_PancreaticCancer1	Meth			
PancreaticCancer3 PancreaticCancer4				Human_PancreaticCancer3 Human_PancreaticCancer4				
PancreaticCancer6				Human_PancreaticCancer5				
PancreaticCancer7				Human_PancreaticCancer?	<u> </u>	•		
PancreaticCancer8 PancreaticCancer9				Human_PancreaticCancer\$				
PancreaticCancer10 PancreaticCancer11				Human_PancreaticCancer1				
Layered H3K27Ac				Mark (Often Found Near Active Regulatory Element Mark (Often Found Near Regulatory Element)				
Layered H3K4Me1				Me1 Mark (Often Found Near Regulatory Elemental Mark (Often Found Near Promoters) o				
DNase Clusters			Transcri	DNasel Hypersensitivity Clusters in 125 cell ty ption Factor ChIP-seq Clusters (161 factors) from		Motifs		
Txn Factor ChIP LNG.IMR90 LNG.IMR90				chromHMM tracks from Roa	admap			
Restr Enzymes Mus Gprin1				Restriction Enzymes from R Non-Human RefSeq Gei				
Mus Gprin1 GPRIN1			***************************************	UCSC annotations of RefSeq RNAs () CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC				
CpG: 91 4 _ Mammal Cons		B.Ba, e, b _a ll _a e, b		Placental Mammal Basewise Conserv				_{/*} -4.884-,48 ₈ 4-,1
				Multiz Alignments of 46 Vert	ebrates			
-4 _ Rhesus Mouse								
Rhesus					رو روسي			