	72,756,75d	72,756,80d	100 bases	q, GenBank, CCDS, Rfam, tRNAs & Comp	hg19 72,756,95d arative Genomics)	72,757,00d
LOC100132891 -	<del>&gt;&gt;&gt;&gt;&gt;&gt;&gt;</del>	<del>&gt;&gt;&gt;&gt;&gt;&gt;&gt;</del>	**************************************	lincRNA and TUCP transcripts  NAs, scaRNAs, and microRNAs from snoR	<del>&gt;</del>	<del>&gt;</del>
CTR147 CTR149		_ • •	•	CTR147 CpG merge methylation level  CTR149 CpG merge methylation level		al _
CTR150 CTR151	_			CTR150 CpG merge methylation level CTR151 CpG merge methylation level		7.5
CTR152 CTR153		<b>-</b> -		CTR152 CpG merge methylation level CTR153 CpG merge methylation level		_0 _
CTR154		-		CTR154 CpG merge methylation level CTR84 CpG merge methylation level		- i
CTR85	_			CTR85 CpG merge methylation level CTR86 CpG merge methylation level		
CTR97	_	 I		CTR97 CpG merge methylation level		-• • • -
CTR101	-		-	CTR101 CpG merge methylation level		 
CTR104	-	- 1 -		CTR104 CpG merge methylation level		- <b>1 -</b>
CTR107				CTR107 CpG merge methylation level  CTR108 CpG merge methylation level	11	
CTR110	-			CTR110 CpG merge methylation level		<b>8</b> - <b>8</b>
CTR148	_			CTR134 CpG merge methylation level CTR148 CpG merge methylation level		
CTR111			• •	CTR111 CpG merge methylation level	<b>I</b>	•
CTR113				CTR114 CpG merge methylation level	 I	.1 -
CTR117 CTR118				CTR118 CpG merge methylation level CTR126 CpG merge methylation level		_8 _
CTR126 CTR127	_	+		CTR127 CpG merge methylation level		_8
CTR128 CTR129			+	CTR128 CpG merge methylation level  CTR129 CpG merge methylation level  CTR131 CpG merge methylation level		<b>.1</b> .
CTR131 AT BS 03				CTR131 CpG merge methylation level se Tissue Bisulfite-Seq Donor STL003 EA F		
AL BS 3 11 Aorta BS 03				sulfite-Seq Donor 3 Library WGBS_Lib 11 E		
Esophagus BS 03				phagus Bisulfite-Seq Donor STL003 EA Re		
FML BS 96 66 FT BS 43 65			BI Fetal Thymus Bisulfi	te-Seq Donor UW H24943 Library WGBS_L	Lib 65 EA Release 9	
Gastric BS 03		1 .		astric Bisulfite-Seq Donor STL003 EA Rele Ventricle Bisulfite-Seq Donor STL001 EA R		. i
LV BS 03				Ventricle Bisulfite-Seq Donor STL003 EA R		
Lung BS 02 Ovary BS 02				Lung Bisulfite-Seq Donor STL002 EA Relea		-• •
Pancreas BS 03				ncreas Bisulfite-Seq Donor STL003 EA Rel s Muscle Bisulfite-Seq Donor STL003 EA R		
RA BS 03		• <u>-</u>		t Atrium Bisulfite-Seq Donor STL003 EA Re		3.1
RV BS 03 SC BS 01	_	- • I		oid Colon Bisulfite-Seq Donor STL001 EA F		11
SC BS 03 SI BS 01	-	iii		oid Colon Bisulfite-Seq Donor STL003 EA F		93
Spleen BS 03	- -			pleen Bisulfite-Seq Donor STL003 EA Rele		3.1
Thymus BS 01 Brain Methyl 2	_			DNA methylation in brain tissue (bigWig)	<b></b>	-1 1
Kidney Methyl 2 Placenta1 Methyl 2				NA methylation in kidney tissue (bigWig)	pigWig) <sup>-</sup> -	•
Placenta2 Methyl 2	+			ylation in placenta (biological replicate 2) (but in placenta (biological replicate 3) (but in placenta (biological replicate 2) (but in placenta (biological replicate 3) (but in placenta 4) (but		·
Placenta3 Methyl 2  Cerebellum	-			Human_Cerebellum_Meth Human_Kidney_Meth		1. 1
Kidney NKcells Sperm				Human_NKcells_Meth Human_Sperm_Meth		
NormalPancreas1  NormalPancreas2				Human_NormalPancreas1_Meth  Human_NormalPancreas2_Meth		
93A  - 				Human_93A_Meth Human_93N_Meth		
Epidermis-old-sun-ex - Epidermis-old-sun-pro				luman_Epidermis-old-sun-exposed_Meth uman_Epidermis-old-sun-protected_Meth		-
Epidermis-young-sun- Epidermis-young-sun-				man_Epidermis-young-sun-exposed_Meth		
Buccals -				Human_Buccals_Meth Human_Sperm_Meth		
BloodHealthy  CD4T-100yr			Distinct Human DNA Methy	Human_BloodHealthy_Meth	man_CD4T-100yr_Meth	<del></del>
CD4T-Newborn PBMC		-		mes from Different Ages, Heyn 2012: Hum		
CD133HSC Macrophage				atopoietic Stem Cells, Hodges 2011 : Huma admap 2015 : Human_Macrophage_Meth	an_CD133HSC_Meth	
NK - BCell				Roadmap 2015 : Human_NK_Meth  Human_BCell_Meth		
CD133HSC HSPC				Human_CD133HSC_Meth  Human_HSPC_Meth		
Neut _				Human_Neut_Meth Human_H1_Meth		
H1BMP4 -H1-mesendoderm				Human_H1BMP4_Meth  Human_H1-mesendoderm_Meth		
H1-NPC  Mesenchymal				Human_H1-NPC_Meth  Human_Mesenchymal_Meth		
IMR90 - IMR90 BS 1a		IMR90	O Cell Line DNA Methylation t	Human_IMR90_Meth  by Bisulfite-seq Signal from REMC/UCSD (L	_ibrary:methylC-seq_imr90_	r1a)
BloodALLL2  BloodALLL1				Human_BloodALLL2_Meth Human_BloodALLL1_Meth		
IMR90 MCF7			<u> </u>	Human_IMR90_Meth Human_MCF7_Meth		
ColonCancer  ColonCancer	-			hypomethylation in colorectal cancer, Bern epigenetic domains across cancer types.	<u> </u>	
HCC1954 HepG2			Human Br	east Cancer, Hon 2012 : Human_HCC1954 Human_HepG2_Meth	4_Meth	11 1
PancreaticCancer1 PancreaticCancer2			1	Human_PancreaticCancer1_Meth  Human_PancreaticCancer2_Meth	***************************************	•
PancreaticCancer3 PancreaticCancer4				Human_PancreaticCancer3_Meth Human_PancreaticCancer4_Meth		
PancreaticCancer5 PancreaticCancer6				Human_PancreaticCancer5_Meth Human_PancreaticCancer6_Meth		
PancreaticCancer7 PancreaticCancer8				Human_PancreaticCancer7_Meth  Human_PancreaticCancer8_Meth		
PancreaticCancer9 PancreaticCancer10				Human_PancreaticCancer9_Meth  Human_PancreaticCancer10_Meth		
PancreaticCancer11  Layered H3K27Ac			H3K27Ac Mark (Often Fou	Human_PancreaticCancer11_Meth  Ind Near Active Regulatory Elements) on 7 of	cell lines from ENCODE	
Layered H3K4Me1				Found Near Regulatory Elements) on 7 cell		
Layered H3K4Me3  DNase Clusters			DNasel Hypers	on 7 cell lines ensitivity Clusters in 125 cell types from EN- eseq Clusters (161 factors) from ENCODE	ICODE (V3)	
Txn Factor ChIP LNG.IMR90 LNG.IMR90				chromHMM tracks from Roadmap	SONO DOUN IVIUUIIS	
Restr Enzymes  Bos MSC				Restriction Enzymes from REBASE  Non-Human RefSeq Genes  annotations of RefSeq RNAs (NM * and NF	R †)	
MSC-AS1 - MSC-AS1 - 4_	**************************************	<del>&gt;                                    </del>	>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>	annotations of RefSeq RNAs (NM_* and Nf	**************************************	<del>,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,</del>
Mammal Cons -4 _ Rhesus			┇╬┸╉ <sub>┯</sub> ┺╌┙╅ <sub>┱</sub> ┯ <sub>╇</sub> ╬┧ <mark>┼</mark> ╗	Multiz Alignments of 46 Vertebrates		⋗ <sub>⋜</sub> ⋥⋞⋕∊⋰⋫⋇⋇⋎⋰⋰ <mark>⋑</mark> ∊⋏⋖⋞⋷
Mouse Dog Elephant Opossum Chicken				neannach (197		
X_tropicalis= Zebrafish RepeatMasker				Repeating Elements by RepeatMasker		