chr12:		129	9,337,000	129	),337,50	od		1 kb hg19 29,338,00¢ 129,338,50¢ 129,339,00¢ 129,339,50¢ 129,340,00¢ Genes (RefSeq, GenBank, CCDS, Rfam, tRNAs & Comparative Genomics) GLT1D1 GLT1D1	129,340,500
CTR147 CTR149	à.		hr		<b>l</b> i 1 <b>1</b> 1		] <sub>].</sub>	/ACA Box snoRNAs, scaRNAs, and microRNAs from snoRNABase and miRBase CTR147 CpG merge methylation level CTR149 CpG merge methylation level	
CTR150 CTR151	<b>1</b>		111		, II b l II	•       1       1		CTR150 CpG merge methylation level	1, 1, 1, 1, 1
CTR153	h H	1	DI Di		1 <b>1</b> 1 1 <b>1</b> 1	<b>l</b>   1	   <b>  </b>	CTR152 CpG merge methylation level  CTR153 CpG merge methylation level  CTR154 CpG merge methylation level	Uri Hii I
CTR154 CTR84 CTR85	<b>h</b> ı		]     <sub>1</sub>			1     1       1		CTR84 CpG merge methylation level	 
CTR86	1. 11		111 			1     1     1		CTR86 CpG merge methylation level	
CTR101	11 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	, III	ıll	<b> </b>   1	1 <b>l</b> i 1 li	<b>1</b>		CTR98 CpG merge methylation level  Liki  CTR101 CpG merge methylation level  CTR103 CpG merge methylation level	## ## 1 ####
CTR104 CTR106	1	. h	dl li		1 I	         		CTR104 CpG merge methylation level  CTR106 CpG merge methylation level	
CTR107	1 1	n n	.lı H.	] ]		4   1     1		CTR107 CpG merge methylation level  CTR108 CpG merge methylation level	
CTR110 CTR132	Ì	"	hi	   	<b> </b>	 	11	CTR110 CpG merge methylation level  CTR132 CpG merge methylation level  CTR134 CpG merge methylation level	, " "
CTR134  CTR148  CTR111	1	.			II Ii	<b>1</b>	hn.   h n	I I	
CTR113		1 11	lı ul T	]   	<b>l</b> í lí	I		CTR114 CpG merge methylation level	11
CTR117 CTR118	- 1	1	llı	] ]	<b>I</b> ı			CTR117 CpG merge methylation level  CTR118 CpG merge methylation level	"
CTR126 CTR127 CTR128	ŀ	1	n.	Ì			] <b>,</b>	CTR126 CpG merge methylation level  CTR127 CpG merge methylation level  CTR128 CpG merge methylation level	1
CTR129		1	     	1	1 1	]   	. I	CTR129 CpG merge methylation level	11
AT BS 03 AL BS 3 11	III	ı    ı	1) a)	          	 	•         		UCSD Adipose Tissue Bisulfite-Seq Donor STL003 EA Release 9	
Aorta BS 03 Esophagus BS 03			11 				I III i	UCSD Aorta Bisulfite-Seq Donor STL003 EA Release 9  UCSD Esophagus Bisulfite-Seq Donor STL003 EA Release 9	
FML BS 96 66 FT BS 43 65			ur dr dr			BI	<sub> </sub> BI Fetal	Uscle Leg Bisulfite-Seq Donor UW H24996 Library WGBS_Lib 66 EA Release 9	
Gastric BS 03						# (   		UCSD Gastric Bisulfite-Seq Donor STL003 EA Release 9  UCSD Left Ventricle Bisulfite-Seq Donor STL001 EA Release 9	
LV BS 03 Lung BS 02						#                	1	UCSD Left Ventricle Bisulfite-Seq Donor STL003 EA Release 9  UCSD Lung Bisulfite-Seq Donor STL002 EA Release 9	
Ovary BS 02 Pancreas BS 03	11 11 1							UCSD Ovary Bisulfite-Seq Donor STL002 EA Release 9  UCSD Pancreas Bisulfite-Seq Donor STL003 EA Release 9	
PM BS 03							Hili	UCSD Psoas Muscle Bisulfite-Seq Donor STL003 EA Release 9  UCSD Right Atrium Bisulfite-Seq Donor STL003 EA Release 9	
RV BS 03 SC BS 01	li li	 	 					UCSD Right Ventricle Bisulfite-Seq Donor STL003 EA Release 9  UCSD Sigmoid Colon Bisulfite-Seq Donor STL001 EA Release 9	
SC BS 03			d  		 		 	UCSD Sigmoid Colon Bisulfite-Seq Donor STL003 EA Release 9	
SI BS 01 Spleen BS 03			di Ur	 	 	 		UCSD Spleen Bisulfite-Seq Donor STL003 EA Release 9  UCSD Thymus Bisulfite-Seq Donor STL001 EA Release 9	
Thymus BS 01 Brain Methyl 2	[				 	 	1 11/1	DNA methylation in brain tissue (bigWig)	
Kidney Methyl 2 Placenta1 Methyl 2			<u> </u>				1 1114	DNA methylation in kidney tissue (bigWig)  DNA methylation in placenta (biological replicate 1) (bigWig)  DNA methylation in placenta (biological replicate 2) (bigWig)	
Placenta2 Methyl 2 Placenta3 Methyl 2					.		1 11 l . 1 l . l .	DNA methylation in placenta (biological replicate 2) (bigWig)	
Cerebellum Kidney NKcells								Human_Cerebellum_Meth  Human_Kidney_Meth  Human_NKcells_Meth	
Sperm NormalPancreas1								Human_Sperm_Meth  Human_NormalPancreas1_Meth	
NormalPancreas2								Human_NormalPancreas2_Meth  Human_93A_Meth  Human_93N_Meth	
93N Epidermis-old-sun-ex Epidermis-old-sun-pro	1							Human_93N_Meth  Human_Epidermis-old-sun-exposed_Meth  Human_Epidermis-old-sun-protected_Meth	
Epidermis-young-sun	<u> </u>							Human_Epidermis-young-sun-exposed_Meth Human_Epidermis-young-sun-protected_Meth	
Buccals Sperm								Human_Buccals_Meth  Human_Sperm_Meth  Human_BloodHealthy_Meth	
BloodHealthy CD4T-100yr CD4T-Newborn						Disti	nct Hur	Human_BloodHealthy_Meth  an DNA Methylomes from Different Ages, Heyn 2012: Human_CD4T-100yr_Meth  DNA Methylomes from Different Ages, Heyn 2012: Human_CD4T-Newborn_Meth	
PBMC CD133HSC		1					1 1.1.	luman DNA Methylomes from Different Ages, Heyn 2012 : Human_PBMC_Meth	
Macrophage NK	1.							Roadmap 2015 : Human_Macrophage_Meth  Roadmap 2015 : Human_NK_Meth  Human_RCell_Meth	
BCell CD133HSC HSPC	11							Human_BCell_Meth Human_CD133HSC_Meth Human_HSPC_Meth	
Neut								Human_Neut_Meth  Human_H1_Meth	1111111111
H1BMP4 H1-mesendoderm								Human_H1BMP4_Meth  Human_H1-mesendoderm_Meth	
H1-NPC  Mesenchymal  IMR90								Human_H1-NPC_Meth  Human_Mesenchymal_Meth  Human_IMR90_Meth	11 11 11 11
IMR90 BS 1a BloodALLL2	<u>l</u> l 1				IMR90	D Cell	Line DN	A Methylation by Bisulfite-seq Signal from REMC/UCSD (Library:methylC-seq_imr90_r1a)	
BloodALLL1								Human_BloodALLL1_Meth  Human_IMR90_Meth	
MCF7 ColonCancer				focal D	LÚ		ĹШı	Human_MCF7_Meth    And long-range hypomethylation in colorectal cancer, Berman 2012 : Human_ColonCancer_Meth    And long-range hypomethylation in colorectal cancer, Berman 2012 : Human_ColonCancer_Meth    And long-range hypomethylation in colorectal cancer types: Human_ColonCancer_Meth	
ColonCancer HCC1954 HepG2			<u>                                     </u>		nicr	Juseco	uny	ation variation in epigenetic domains across cancer types:: Human_GolonCancer_Meth	1.111.
PancreaticCancer1 PancreaticCancer2								Human_PancreaticCancer2_Meth  Human_PancreaticCancer2_Meth	
PancreaticCancer3 PancreaticCancer4								Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth	
PancreaticCancer5 PancreaticCancer6 PancreaticCancer7								Human_PancreaticCancer5_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer7_Meth	
PancreaticCancer8 PancreaticCancer9					1 (1)			Human_PancreaticCancer8_Meth  Human_PancreaticCancer9_Meth	
PancreaticCancer10 PancreaticCancer11								Human_PancreaticCancer10_Meth  Human_PancreaticCancer11_Meth  Human_PancreaticCancer11_Meth	
Layered H3K27Ac							H3K4M	lark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE  11 Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE	
Layered H3K4Me3  DNase Clusters						Te		K4Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  DNasel Hypersensitivity Clusters in 125 cell types from ENCODE (V3)  on Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs	
Txn Factor ChIP  LNG.IMR90  LNG.IMR90						113	الإدر.	chromHMM tracks from Roadmap  Restriction Enzymes from REBASE	
Restr Enzymes		<b>                                  </b>							**************************************
LOC100128276	<del></del>					<del></del>		UCSC annotations of RefSeq RNAs (NM_* and NR_*) GLT1D1 GLT	**************************************
							Сре	GLT1D1 GL	**************************************
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-4_								Multiz Alignments of 46 Vertebrates	
Ammal Cons  -4 _  Rhesus  Mouse  Dog  Elephant  Opossum  Chicken								Multiz Alignments of 46 Vertebrates	