Scale chr14:		60,978,05d	100 bases 60,978,100 UCSC 6	hg19 60,978,150 60,978,200 60,978,250 enes (RefSeq, GenBank, CCDS, Rfam, tRNAs & Comparative Genomics)	60,978,300
SIX6			C/D and H/A	lincRNA and TUCP transcripts A Box snoRNAs, scaRNAs, and microRNAs from snoRNABase and miRBase CTR147 CpG merge methylation level	
CTR149 CTR150	-			CTR149 CpG merge methylation level CTR150 CpG merge methylation level	
CTR151				CTR151 CpG merge methylation level	. <b>I.</b>
CTR152 CTR153	-			CTR152 CpG merge methylation level  CTR153 CpG merge methylation level	 
CTR154	-			CTR154 CpG merge methylation level  CTR84 CpG merge methylation level	
CTR85	-	-		CTR85 CpG merge methylation level	
CTR86 CTR97	-	-	<b>-</b>	CTR86 CpG merge methylation level	
CTR98	-	<del>-</del>	-	CTR98 CpG merge methylation level	-
CTR103	-	-	<u>-</u> -	CTR103 CpG merge methylation level	l
CTR104 CTR106	-			CTR104 CpG merge methylation level  CTR106 CpG merge methylation level	
CTR107 CTR108		•		CTR107 CpG merge methylation level  CTR108 CpG merge methylation level	
CTR110	-			CTR110 CpG merge methylation level	 
CTR132 CTR134				CTR134 CpG merge methylation level	
CTR148				CTR148 CpG merge methylation level  CTR111 CpG merge methylation level	■
CTR113	-		-	CTR113 CpG merge methylation level	<b>_</b>
CTR114 CTR117	_	_	-	CTR114 CpG merge methylation level  CTR117 CpG merge methylation level	·
CTR118	-	<del>-</del>		CTR118 CpG merge methylation level  CTR126 CpG merge methylation level	
CTR126 CTR127	-			CTR127 CpG merge methylation level	-
CTR128 CTR129		_		CTR128 CpG merge methylation level CTR129 CpG merge methylation level	
CTR131	-			CTR131 CpG merge methylation level	
AT BS 03 AL BS 3 11	-	-		dult Liver Bisulfite-Seq Donor 3 Library WGBS_Lib 11 EA Release 8	
Aorta BS 03		•		UCSD Aorta Bisulfite-Seq Donor STL003 EA Release 9	
Esophagus BS 03			BI Fetal Mus	UCSD Esophagus Bisulfite-Seq Donor STL003 EA Release 9	
FML BS 96 66 FT BS 43 65				ymus Bisulfite-Seq Donor UW H24943 Library WGBS_Lib 65 EA Release 9	
Gastric BS 03				UCSD Gastric Bisulfite-Seq Donor STL003 EA Release 9	
LV BS 01 LV BS 03			■ -	UCSD Left Ventricle Bisulfite-Seq Donor STL001 EA Release 9  UCSD Left Ventricle Bisulfite-Seq Donor STL003 EA Release 9	
LV BS 03 Lung BS 02				UCSD Lung Bisulfite-Seq Donor STL002 EA Release 9	
Ovary BS 02				UCSD Ovary Bisulfite-Seq Donor STL002 EA Release 9	
Pancreas BS 03 PM BS 03	-	-		UCSD Pancreas Bisulfite-Seq Donor STL003 EA Release 9 UCSD Psoas Muscle Bisulfite-Seq Donor STL003 EA Release 9	[                       <u>+</u> -
RA BS 03				UCSD Right Atrium Bisulfite-Seq Donor STL003 EA Release 9	
RV BS 03		- -		UCSD Right Ventricle Bisulfite-Seq Donor STL003 EA Release 9  UCSD Sigmoid Colon Bisulfite-Seq Donor STL001 EA Release 9	••
SC BS 01 SC BS 03	-	-		JCSD Sigmoid Colon Bisulfite-Seq Donor STL003 EA Release 9	-1
SI BS 01		<u>-</u>	=	JCSD Small Intestine Bisulfite-Seq Donor STL001 EA Release 9	
Spleen BS 03	-	-	<b> ■</b> .	UCSD Spleen Bisulfite-Seq Donor STL003 EA Release 9 UCSD Thymus Bisulfite-Seq Donor STL001 EA Release 9	
Thymus BS 01 Brain Methyl 2	-	-	_ •	DNA methylation in brain tissue (bigWig)	••
Kidney Methyl 2	-			DNA methylation in kidney tissue (bigWig)	
Placenta1 Methyl 2 Placenta2 Methyl 2	-	-		DNA methylation in placenta (biological replicate 1) (bigWig)  DNA methylation in placenta (biological replicate 2) (bigWig)	- •
Placenta3 Methyl 2	-	-		DNA methylation in placenta (biological replicate 3) (bigWig)	- •
Cerebellum Kidney				Human_Cerebellum_Meth Human_Kidney_Meth	1
NKcells				Human_NKcells_Meth  Human_Sperm_Meth	
Sperm NormalPancreas1				Human_NormalPancreas1_Meth	<u> </u>
NormalPancreas2				Human_NormalPancreas2_Meth  Human_93A_Meth	
93N				Human_93N_Meth  Human_Epidermis-old-sun-exposed_Meth	
Epidermis-old-sun-ex Epidermis-old-sun-pro	-			Human_Epidermis-old-sun-protected_Meth	
Epidermis-young-sun-				Human_Epidermis-young-sun-exposed_Meth  Human_Epidermis-young-sun-protected_Meth	
Buccals				Human_Buccals_Meth	•
Sperm BloodHealthy				Human_Sperm_Meth  Human_BloodHealthy_Meth	
CD4T-100yr CD4T-Newborn				DNA Methylomes from Different Ages, Heyn 2012 : Human_CD4T-100yr_Meth  NA Methylomes from Different Ages, Heyn 2012 : Human_CD4T-Newborn_Meth	
PBMC				an DNA Methylomes from Different Ages, Heyn 2012 : Human_PBMC_Meth	
CD133HSC Macrophage			Changes in	Human Hematopoietic Stem Cells, Hodges 2011 : Human_CD133HSC_Meth  Roadmap 2015 : Human_Macrophage_Meth	
NK				Roadmap 2015 : Human_NK_Meth  Human_BCell_Meth	-
BCell CD133HSC				Human_CD133HSC_Meth	
HSPC Neut				Human_HSPC_Meth  Human_Neut_Meth	
H1 H1BMP4				Human_H1_Meth  Human_H1BMP4_Meth	
H1BMP4 H1-mesendoderm				Human_H1-mesendoderm_Meth	
H1-NPC Mesenchymal				Human_H1-NPC_Meth  Human_Mesenchymal_Meth	
IMR90		I I I	1R90 Cell Line DNA	Human_IMR90_Meth  Methylation by Bisulfite-seq Signal from REMC/UCSD (Library:methylC-seq_imr90_r1a)	<del></del>
IMR90 BS 1a BloodALLL2				Human_BloodALLL2_Meth	
BloodALLL1				Human_BloodALLL1_Meth  Human_IMR90_Meth	
MCF7				Human_MCF7_Meth	
ColonCancer	<u> </u>	focal DNA		d long-range hypomethylation in colorectal cancer, Berman 2012 : Human_ColonCancer_Meth	
HCC1954 HepG2		1		Human Breast Cancer, Hon 2012 : Human_HCC1954_Meth  Human_HepG2_Meth	
				Human_PancreaticCancer1_Meth	
PancreaticCancer1					
				Human_PancreaticCancer2_Meth  Human_PancreaticCancer3_Meth	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4					
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3				Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth  Human_PancreaticCancer5_Meth  Human_PancreaticCancer6_Meth	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer5				Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth  Human_PancreaticCancer5_Meth	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer5 PancreaticCancer6 PancreaticCancer7 PancreaticCancer7 PancreaticCancer8 PancreaticCancer9				Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth  Human_PancreaticCancer5_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer7_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer8_Meth	1
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer5 PancreaticCancer6 PancreaticCancer7 PancreaticCancer8				Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth  Human_PancreaticCancer5_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer7_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer10_Meth  Human_PancreaticCancer11_Meth	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer5 PancreaticCancer6 PancreaticCancer7 PancreaticCancer8 PancreaticCancer9 PancreaticCancer10 PancreaticCancer11 Layered H3K27Ac				Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer7_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer9_Meth	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer5 PancreaticCancer6 PancreaticCancer7 PancreaticCancer8 PancreaticCancer9 PancreaticCancer10 PancreaticCancer11 Layered H3K27Ac	1		H3K4Me1	Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth  Human_PancreaticCancer5_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer7_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer10_Meth  Human_PancreaticCancer11_Meth  Coften Found Near Active Regulatory Elements) on 7 cell lines from ENCODE	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer5 PancreaticCancer6 PancreaticCancer6 PancreaticCancer7 PancreaticCancer8 PancreaticCancer9 PancreaticCancer10 PancreaticCancer11 Layered H3K27Ac Layered H3K4Me1 Layered H3K4Me3 DNase Clusters			H3K4Me1 H3K4 DI	Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer7_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer10_Meth  Human_PancreaticCancer11_Meth  Goften Found Near Active Regulatory Elements) on 7 cell lines from ENCODE	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer5 PancreaticCancer6 PancreaticCancer6 PancreaticCancer7 PancreaticCancer8 PancreaticCancer9 PancreaticCancer10 PancreaticCancer11 Layered H3K27Ac Layered H3K4Me1 Layered H3K4Me3			H3K4Me1 H3K4 DI	Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth  Human_PancreaticCancer5_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer10_Meth  Human_PancreaticCancer11_Meth  (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE  Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Mes Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  asel Hypersensitivity Clusters in 125 cell types from ENCODE (V3)  Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs  chromHMM tracks from Roadmap	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer5 PancreaticCancer6 PancreaticCancer6 PancreaticCancer7 PancreaticCancer8 PancreaticCancer9 PancreaticCancer10 PancreaticCancer11 Layered H3K27Ac Layered H3K4Me1 Layered H3K4Me3 DNase Clusters Txn Factor ChIP LNG.IMR90			H3K4Me1 H3K4 DI	Human_PancreaticCancer3_Meth  Human_PancreaticCancer4_Meth  Human_PancreaticCancer5_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer10_Meth  Human_PancreaticCancer11_Meth  (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE  Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Mes Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  asel Hypersensitivity Clusters in 125 cell types from ENCODE (V3)  Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer6 PancreaticCancer6 PancreaticCancer7 PancreaticCancer8 PancreaticCancer9 PancreaticCancer10 PancreaticCancer11 Layered H3K27Ac Layered H3K4Me1 Layered H3K4Me3 DNase Clusters Txn Factor ChIP LNG.IMR90 LNG.IMR90 Restr Enzymes Bos SIX6 Rattus Six61 Mus Six6	1		H3K4Me1 H3K4 DI	Human_PancreaticCancer3_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer7_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer10_Meth  Human_PancreaticCancer11_Meth  (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE  Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  asel Hypersensitivity Clusters in 125 cell types from ENCODE (V3)  Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs  chromHMM tracks from Roadmap  Restriction Enzymes from REBASE	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer4 PancreaticCancer5 PancreaticCancer6 PancreaticCancer6 PancreaticCancer7 PancreaticCancer8 PancreaticCancer9 PancreaticCancer10 PancreaticCancer11 Layered H3K27Ac Layered H3K4Me1 Layered H3K4Me3 DNase Clusters Txn Factor ChIP LNG.IMR90 LNG.IMR90 LNG.IMR90 Restr Enzymes Bos SIX6 Rattus Six6 Mus Six6 SIX6I CpG: 178			H3K4Me1 H3K4 DI	Human_PancreaticCancer3_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer10_Meth  Human_PancreaticCancer11_Meth  (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE  Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Mes Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Seasel Hypersensitivity Clusters in 125 cell types from ENCODE (V3)  Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs  chromHMM tracks from Roadmap  Restriction Enzymes from REBASE  Non-Human RefSeq Genes	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer6 PancreaticCancer6 PancreaticCancer7 PancreaticCancer8 PancreaticCancer9 PancreaticCancer10 PancreaticCancer11 Layered H3K27Ac Layered H3K4Me1 Layered H3K4Me3 DNase Clusters Txn Factor ChIP LNG.IMR90 LNG.IMR90 Restr Enzymes Bos SIX6 Rattus Six6 Mus Six6 SIX6 CpG: 178 4 Mammal Cons			H3K4Me1 H3K4 DI	Human_PancreaticCancer3_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer8_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer10_Meth  Human_PancreaticCancer11_Meth  It (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE  Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me4 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me5 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me6 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me7 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  Me8 Mark (Often Found Nea	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer5 PancreaticCancer6 PancreaticCancer7 PancreaticCancer8 PancreaticCancer9 PancreaticCancer10 PancreaticCancer11 Layered H3K27Ac Layered H3K4Me1 Layered H3K4Me3 DNase Clusters Txn Factor ChIP LNG.IMR90 LNG.IMR90 LNG.IMR90 Restr Enzymes Bos SIX6 Rattus Six6 Mus Six6 SIX6I CpG: 178 Mammal Cons -4 Rhesus Mouse Dog Elephant			H3K4Me1 H3K4 DI	Human_PancreaticCancer3_Meth  Human_PancreaticCancer5_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer7_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer10_Meth  Human_PancreaticCancer11_Meth  (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE  Med Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  assel Hypersensitivity Clusters in 125 cell types from ENCODE (V3)  Factor ChiP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs  chromHMM tracks from Roadmap  Restriction Enzymes from REBASE  Non-Human RefSeq Genes  UCSC annotations of RefSeq RNAs (NM_* and NR_*)  CpG Islands (Islands < 300 Bases are Light Green)  Placental Mammal Basewise Conservation by Phylop	
PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer6 PancreaticCancer6 PancreaticCancer7 PancreaticCancer7 PancreaticCancer9 PancreaticCancer10 PancreaticCancer11 Layered H3K27Ac Layered H3K4Me1 Layered H3K4Me3 DNase Clusters Txn Factor ChIP LNG.IMR90 LNG.IMR90 Restr Enzymes Bos SIX6 Rattus Six61 Mus Six6 SIX61 CpG: 178 4 Mammal Cons -4 Rhesus Mouse Dog			H3K4Me1 H3K4 DI	Human_PancreaticCancer3_Meth  Human_PancreaticCancer5_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer6_Meth  Human_PancreaticCancer7_Meth  Human_PancreaticCancer9_Meth  Human_PancreaticCancer10_Meth  Human_PancreaticCancer11_Meth  (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE  Med Mark (Often Found Near Promoters) on 7 cell lines from ENCODE  assel Hypersensitivity Clusters in 125 cell types from ENCODE (V3)  Factor ChiP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs  chromHMM tracks from Roadmap  Restriction Enzymes from REBASE  Non-Human RefSeq Genes  UCSC annotations of RefSeq RNAs (NM_* and NR_*)  CpG Islands (Islands < 300 Bases are Light Green)  Placental Mammal Basewise Conservation by Phylop	