Table	74,729
	1 1
Second	
	.
	.11 1 1 11
May	
18	.
	l dia Lati
	1
1	
1900	
	,
The content will be content with the content will be content wil	.
Company	
## 12	
Design Hydroc DNA Monthbooks to the Design Hydroc DNA Monthbooks to the Design Hydroc DNA Monthbooks to the DN	
Company in Fundament Membrace (2017) 1	
Received 2015 Filteral INC Media Receiv	
Name of Section 2012	
A HARDER HT 1990 THE	
Harran, Fit means of control (Amb) Harran, Fit means of control (
NROP Cell Line DNA Methylation by Boutflewood Spoal from REVICUSD Alberty-weety/C-seq_inrig0_flag NROP Cell Line DNA Methylation by Boutflewood Spoal from REVICUSD Alberty-weety/C-seq_inrig0_flag Harman BloodALLI Meth Harman B	
MRSD Cet Live DNA Merry patients by Boudleward Spring from RRMCUCSD (Library particul)—cet, and 0, 1s) Harman, Block ALL J. Merry Harman, Bl	
Harman_NCF_Memin Noted DNA hypermethylation and large-range hyper	1 11
The second secon	
Human Presst Cancer, Hon 2012; Human LPC1954, Meth Human Report Cancer I Meth Human Pencrest Cancer I M	
Human Pancreatic Cancers J. Meth	
Human_PancreatcCancer3 Meth Human_PancreatcCancer4 Meth Human_PancreatcCancer5_Meth Human_PancreatcCancer6_Meth Human_PancreatcC	
is in the process of	
Human_PancreaticCancer9_Meth	
icCancer10 icCancer11 H3K27Ac H3K4Me1 H3K27Ac Mark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE H3K4Me1 H3K4Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE H3K4Me3 DNasel Hypersensitivity Clusters in 125 cell types from ENCODE Transcription Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs Encor ChIP LNG.IMR90 LNG.IMR90 LNG.IMR90 Str Enzymes Restriction Enzymes from REBASE str Enzymes Restriction Enzymes from REBASE Non-Human RefSeq Genes Rattus Lbx2 Mus Lbx2 Bos LBX2 UCSC annotations of RefSeq RNAs (NM_* and NR_*)	
H3K27Ac Mark (Often Found Near Active Regulatory Elements) on 7 cell lines from ENCODE H3K4Me1 H3K4Me1 Mark (Often Found Near Regulatory Elements) on 7 cell lines from ENCODE H3K4Me3 H3K4Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE DNasel Hypersensitivity Clusters in 125 cell types from ENCODE (V3) ase Clusters Factor ChIP ChromHMM tracks from Roadmap LNG.IMR90 LNG.IMR90 LNG.IMR90 Restriction Enzymes from REBASE Str Enzymes Restriction Enzymes from REBASE Non-Human RefSeq Genes Mus Lbx2 Mus Lbx2 Bos LBx2 UCSC annotations of RefSeq RNAs (NM_* and NR_*)	
H3K4Me3 Mark (Often Found Near Promoters) on 7 cell lines from ENCODE DNasel Hypersensitivity Clusters in 125 cell types from ENCODE (V3) Transcription Factor ChIP-seq Clusters (161 factors) from ENCODE with Factorbook Motifs ChromHMM tracks from Roadmap LNG.IMR90 LNG.IMR90 Str Enzymes TRESTRICTION FOR ENCODE WITH Factorbook Motifs Restriction Enzymes from REBASE Non-Human RefSeq Genes Mus Lbx2 Bos LBX2 UCSC annotations of RefSeq RNAs (NM_* and NR_*) LBX2 LBX2 LBX2	
Factor ChIP LNG.IMR90 LNG.IMR90 LNG.IMR90 Restriction Enzymes from REBASE str Enzymes Rattus Lbx2 Mus Lbx2 Bos LBX2 UCSC annotations of RefSeq RNAs (NM_* and NR_*) LBX2 LBX2 LBX2	
Restriction Enzymes from REBASE Restriction Enzymes from REBASE Non-Human RefSeq Genes Mus Lbx2 Bos LBX2 LBX2 LBX2 LBX2 LBX2 LBX2	
Mus Lbx2 Bos LBX2 Control Cont	
	< < < < < < < <
Cons CpG: 161 Placental Mammal Basewise Conservation by PhyloP Cons	والمراجعة
-4	