| Scale chr11: PAX6 | 31,824,000 31,82 | | | hg19 ,826,500 31,827,000 trative Genomics) | 31,827,50¢ |
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| AT BS 03 AL BS 3 11 Aorta BS 03 Esophagus BS 03 FML BS 96 66 FT BS 43 65 Gastric BS 03 LV BS 01 LV BS 01 LV BS 03 Lung BS 02 Ovary BS 02 Pancreas BS 03 PM BS 03 | The first decision of the control of | BI Adult Liver B BI Fetal Thymus Bisulf BI Fetal Thymus Bisulf UCSD Left UCSD Psoc UCSD Pso | ose Tissue Bisulfite-Seq Donor STL003 EA Reliabilitie-Seq Donor 3 Library WGBS_Lib 11 E Reliabilitie-Seq Donor STL003 EA Reliabiliti | A Release 8 If any of the second of the sec | All the first of the continue |
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| Sperm NormalPancreas1 NormalPancreas2 93A 93N Epidermis-old-sun-ex Epidermis-young-sun Epidermis-young-sun Buccals Sperm BloodHealthy CD4T-100yr CD4T-Newborn PBMC CD133HSC Macrophage | | Distinct Human DNA Methylo Distinct Human DNA Methylo Changes in Human Hen | Human_NormalPancreas1_Meth Human_NormalPancreas1_Meth Human_NormalPancreas2_Meth Human_93A_Meth Human_93A_Meth Human_93N_Meth Human_Epidermis-old-sun-exposed_Meth Iman_Epidermis-old-sun-protected_Meth Human_Epidermis-young-sun-exposed_Meth Human_Epidermis-young-sun-protected_Meth Human_Buccals_Meth Human_BloodHealthy_Meth Human_BloodHealthy_Meth Plomes from Different Ages, Heyn 2012 : Human_Buccals_Heyn 2013 : Human_Buccals_Heyn 2015 : Human_Macrophage_Meth | man_CD4T-100yr_Meth an_CD4T-Newborn_Meth Human_PBMC_Meth | |
| NK BCell CD133HSC HSPC Neut H1 H1BMP4 H1-mesendoderm H1-NPC Mesenchymal IMR90 IMR90 BS 1a BloodALLL2 BloodALLL1 IMR90 MCF7 | | IMR90 Cell Line DNA Methylation | Roadmap 2015: Human_NK_Meth Human_BCell_Meth Human_CD133HSC_Meth Human_HSPC_Meth Human_Neut_Meth Human_H1_Meth Human_H1-Meth Human_H1-NPC_Meth Human_Mesenchymal_Meth Human_MR90_Meth by Bisulfite-seq Signal from REMC/UCSD (L | | |
| ColonCancer ColonCancer HCC1954 HepG2 PancreaticCancer1 PancreaticCancer2 PancreaticCancer3 PancreaticCancer4 PancreaticCancer5 PancreaticCancer6 PancreaticCancer7 PancreaticCancer7 PancreaticCancer8 PancreaticCancer9 PancreaticCancer10 PancreaticCancer11 Layered H3K27Ac | | focal DNA hypermethylation and long-rang increased methylation variation Human B | Human_PancreaticCancer2_Meth Human_PancreaticCancer3_Meth Human_PancreaticCancer4_Meth Human_PancreaticCancer6_Meth Human_PancreaticCancer6_Meth Human_PancreaticCancer7_Meth Human_PancreaticCancer8_Meth | Human_ColonCancer_Meth Human_ColonCancer_Meth Meth Meth Mill | |
| Macaca PAX6 Mus Pax6 Sus PAX6 Sus PAX6 Sus PAX6 Rattus Pax6 Canpus Pax6 Xenopus | ************************************** | H3K4Me3 Mark (0 DNasel Hyper Transcription Factor Chlf | Pound Near Regulatory Elements) on 7 cell lines Sensitivity Clusters in 125 cell types from EN P-seq Clusters (161 factors) from ENCODE v chromHMM tracks from Roadmap Restriction Enzymes from REBASE Non-Human RefSeq Genes | from ENCODE CODE (V3) | |
| Astyana pax6 Saccogl pax6 Drosoph toy Caenorh vab-3 | ************************************** | | annotations of RefSeq RNAs (NM_* and NR | | |
| PAX6 | | | Islands (Islands < 300 Bases are Light Gree CpG: 81 tall Mammal Basewise Conservation by Phy Multiz Alignments of 46 Vertebrates | loP | CpG: 28 |