Scale chr6: POU3F2		PHHADHH	99,283,		QP P P		100 bases 99,283,500 UCSC G	Senes (RefSeq, G	99,283,550 enBank, CCDS, Rf DEDTPTSDDL	fam, tRNAs & Co	99,283,600 omparative Geno	g19 omics) FQADVC	99,283,65 GLALGTLYC	ľ	99,283,70
CTR147		PUBLISHED			-			CA Box snoRNAs CTF	lincRNA and TUCF , scaRNAs, and mi R147 CpG merge n	P transcripts icroRNAs from s nethylation level	snoRNABase and I 				
CTR149 CTR150						_ I	- =	CTE	R149 CpG merge n	methylation level	,	= -			
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CTR127									R127 CpG merge n R <u>1</u> 28 CpG merge n						
CTR128 CTR129								СТ	R129 CpG merge n	nethylation level	•				
CTR131 AT BS 03	_ ■_								issue Bisulfite-Seq			I _			
AL BS 3 11 Aorta BS 03				-			- BI	UCSD Aorta	e-Seq Donor 3 Libi	or STL003 EA R	telease 9	B .			
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Pancreas BS 03	 			_		 		UCSD Pancre	as Bisulfite-Seq Do	onor STL003 EA	Release 9				
RA BS 03			- -	_			- -	UCSD Psoas Mu UCSD Right Atr	uscle Bisulfite-Seq					+	
RV BS 03	 		- - -	†	_	++ ++ +	- -	UCSD Right Ven	tricle Bisulfite-Seq			I _		+ +	
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Cerebellum			- +	-		- -	1		Human_Cerebel		- - -	• -		- -	
NKcells Sperm									Human_NKcel						
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93A									Human_93A	_Meth	-				
93N Epidermis-old-sun-ex									n_Epidermis-old-s	un-exposed_Me					
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Sperm BloodHealthy									Human_Sperr	althy_Meth		100	40		
CD4T-100yr CD4T-Newborn							tinct Human I	DNA Methylomes	from Different Age	es, Heyn 2012 : H	Human_CD4T-N	ewborn_N			
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Macrophage NK									padmap 2015 : Human_		leth				
BCell CD133HSC							-		Human_BCel						
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H1	-				1				Human_H1_	_Meth					
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H1-NPC Mesenchymal									Human_Mesench	nymal_Meth					
IMR90 BS 1a					IMI	R90 C€	ell Line DNA	Methylation by Bi	Human_IMR9 sulfite-seq Signal fi	rom REMC/UCS	SD (Library:methy	ylC-seq_in	nr90_r1a)		
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ColonCancer	11	•	1 .	foca		Ω.			pomethylation in co					th	
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Bos POU3F2	>>>>		>>>>>	>>>>>	>>>>> >>>>>	>>>>>	>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>	>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>	Non-Human Refs	Seq Genes	>>>>>>	>>>>>>	>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>	>>>>>>	***************************************
Rattus Pou3f2 Xenopus pou3f2.L Xenopus pou3f2.S Coturni POU3F2 Pan POU3F2	>>>	>>>>>>> >>>>>>>>>>>>>>>>>>>>>>>>>>>>>>	>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>		>>>>> >>>>>> >>>>>>>>>>>>>>>>>>>>>>>>>	······································	·>>>>>> ·>>>>>>>>>>>>>>>>>>>>>>>>>>>>>	>>>>>> >>>>>>>>>>>>>>>>>>>>>>>>>>>>>>				>>>>>)))))	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,
Pan POU3F2 Xenopus pou3f2 POU3F2	H	}}}}	>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>	******** *******	>>>>> >>>>>>	>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>>	****	UCSC anno	otations of RefSeq	RNAs (NM_* an	d NR_*)				
CpG: 323 4 _ Mammal Cons		4 ,5 5 1 5 5 4 4							ds (Islands < 300 E						
-4 _ Rhesus Mouse Dog								Mu	ultiz Alignments of	46 Vertebrates					
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X_tropicalis Zebrafish RepeatMasker		1 ST S. S.			ani l			Rep	eating Elements b	y RepeatMasker					التكليب بد