Scale chr10:		81,022,000	81,022,500	81,023,000	1 kb	81,024,000	mm10 81,024,500	81,025,000	81,025,500	81,026,000
ATACseq,brain,ATACemb		01,022,000	01,022,000			seq,brain,ATACembryoBrain1		0.,025,000	01,020,000	01,020,000
ATACseq,brain,ATACemb	Щ				ATACs	seq,brain,ATACembryoBrain2				
ATACseq,liver,ATACembra	Ų,			الألا	ATAC	seq,liver,ATACembryoLiver1				
ATACseq,liver,ATACembr					ATAC:	seq,liver,ATACembryoLiver2				
ChIPseq,brain,CHIPembry					ChIPs	eq,brain,CHIPembryoBrain1				
ChIPseq,brain,CHIPembry					ChIPs	eq,brain,CHIPembryoBrain2			Married Co.	
ChIPseq,liver,CHIPembry					ChlPs	seq,liver,CHIPembryoLiver1				
ChlPseq,liver,CHIPembry					ChiPs	seq,liver,CHIPembryoLiver2				
RNAseq,brain,PRNAembr			J/"-[		RNAse	eq,brain,PRNAembryoBrain1				
RNAseq,brain,PRNAembr					RNAse	eq,brain,PRNAembryoBrain2		<del></del>		
RNAseq,liver,PRNAembry					RNAs	seq,liver,PRNAembryoLiver1				
RNAseq,liver,PRNAembry					RNAs	seq,liver,PRNAembryoLiver2				
Diras1 ■				GEI	NCODE VM23 Compreher	nsive Transcript Set (only Bas	sic displayed by default	<del>(                                    </del>	<del>&lt;</del>	
Diras1/NM 145217.2■		NC NC	BI RefSeq genes, curated	d subset (NM_*, N	R_*, NP_* or YP_*) - Anno	otation Release NCBI Mus mu	usculus Updated Anno	tation Release 108.2020	00622 (2020-07-09)	