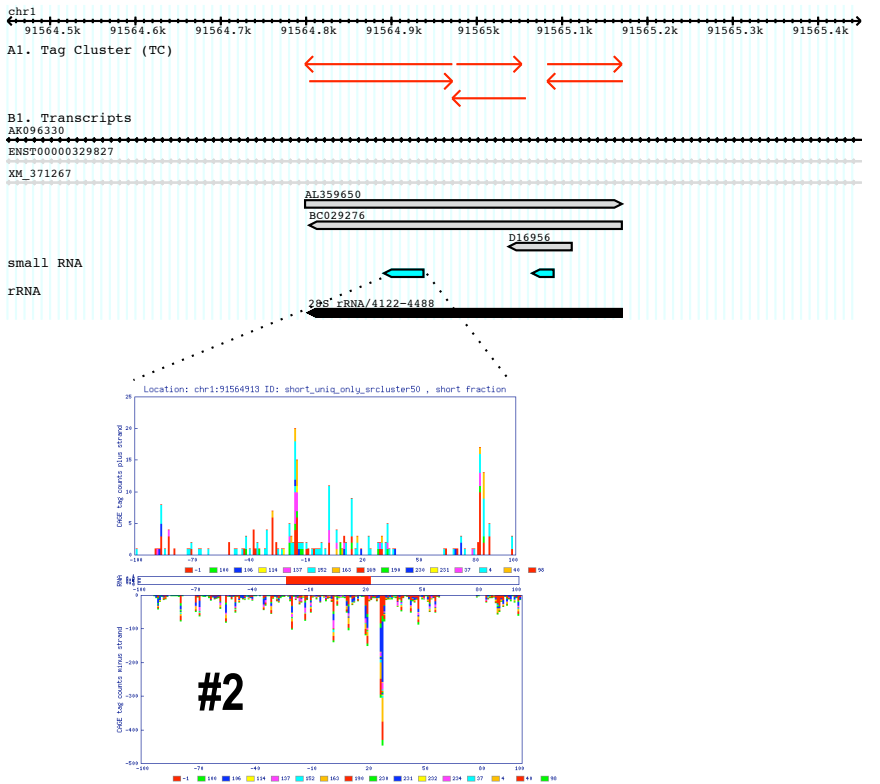


Different colors in the histograms represent CAGE tags deriving from different tissues. Tissue name for each ID are described here:

ID	Tissue name
-1	UNDEFINED_TISSUE_TYPE
100	large intestine
106	liver
114	mammary gland
137	pancreas
152	rectum
163	small intestine
169	stomach
190	urinary bladder
230	ureter
231	renal artery
232	prostate gland
234	epididymis
37	cecum
4	adipose
40	cerebrum
98	kidney



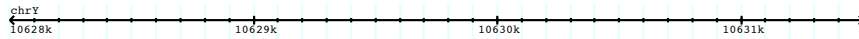
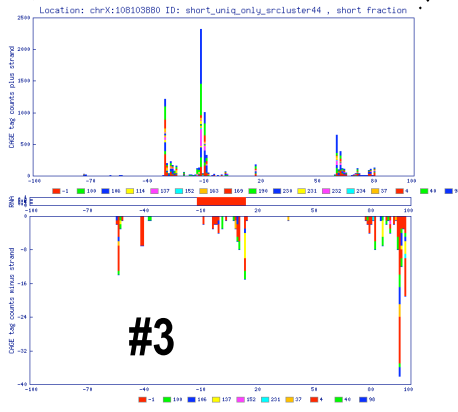
B1. Transcripts

small RNA

rRNA

D17136

28S rRNA/3625-4102



A1. Tag Cluster (TC)

B1. Transcripts

small RNA

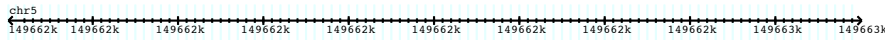
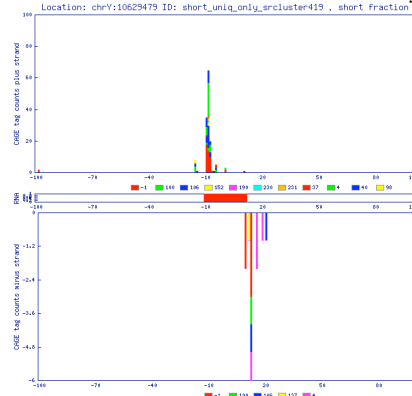
rRNA

18S rRNA/0-1869

CR749662

5\_S rRNA/1-152

**#4**



A1. Tag Cluster (TC)

B1. Transcripts

AK122641

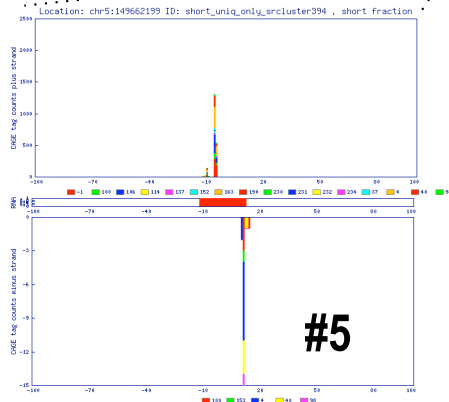
ENST00000328668

ENST00000341819

XM\_291139

small RNA

rRNA



zsa22518a10t3.scf	223	243	n=1	CGCCAAGGGCCACCACCGGC
zsa22501c05t3.scf	184	205	n=1	CGCCGAGGGCGCACCACCGGC
zsa22505g19t3.scf	316	336	n=2	CGCCGAGGGCGCACCACCGGC
zsa22506g19t3.scf	328	348	n=1	CGCCGAGTGCACACCACCGGC
zsa22507i24t3.scf	455	477	n=1	CGCCGAGGGCGCACCACCGGC <sup>TC</sup>
zsa22508a24t3.scf	227	251	n=1	CGCCGAGGGCGCACCACCGGC <sup>CCG</sup>
zsa22514a20t3.scf	550	569	n=1	CGCCGAGGGCGCACCA <sup>TCGG</sup>
rDNA(U13369):9368-9391				CGCCGAGGGCGCACCACCGGC <sup>CCG</sup>
chr5:149662187-149662211				CGCCGAGGGCGCACCACCGGC <sup>CCG</sup>

