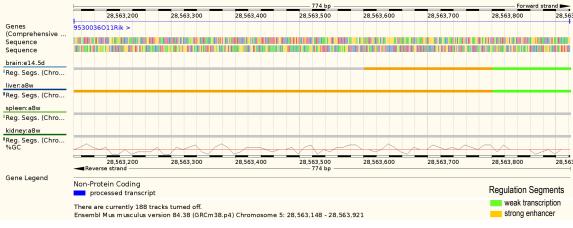
SBE6.1



В

SBE6.2

				1.07	KD			— Forward strand 🗩
	28,567,400	28,567,600		28,567,800	28,568,000		28,568,200	28,568,4
Genes Comprehensive	9530036011Rik >							
rain:e14.5d								
leg. Segs. (Chro								
ver:a8w								
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pleen:a8w								
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idney:a8w								
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	28,567,400	28,567,600	~	28,567,800	28,568,000		28,568,200	28,568,
	Reverse strand —	20,507,000		1.07			20,500,200	20,500,
iene Legend	Non-Protein Coding	cript					Regu	lation Segments
								weak transcription
C	There are currently 18 Ensembl Mus musculu		m38.p4) Chromos	ome 5: 28,567,395	- 28,568,460		3	strong enhancer
C #2			m38.p4) Chromos	ome 5: 28,567,395	28,568,460		5	strong enhancer
C #2			m38.p4) Chromos	ome 5: 28,567,395			2	trong enhancer
C #2			m38.p4) Chromos			29,034,250	29,034,500	
	Ensembl Mus musculu	is version 84.38 (GRC		2.01	kb	29,034,250		— Forward strand ∎
rain:e14.5d	Ensembl Mus musculu	is version 84.38 (GRC		2.01	kb	29,034,250		— Forward strand ∎
rain:e14.5d .eg. Segs. (Chro ver:a8w	Ensembl Mus musculu	is version 84.38 (GRC		2.01	kb	29,034,250		— Forward strand ∎
rain:e14.5d .eg. Segs. (Chro ver:a8w	Ensembl Mus musculu	is version 84.38 (GRC		2.01	kb	29,034,250		— Forward strand ∎
rain:e14.5d eg. Segs. (Chro ver:a8w eg. Segs. (Chro pleen:a8w	Ensembl Mus musculu	is version 84.38 (GRC		2.01	kb	29,034,250		— Forward strand ∎
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rain:e14.5d eg. Segs. (Chro ver:a8w eg. Segs. (Chro pleen:a8w eg. Segs. (Chro idney:a8w	Ensembl Mus musculu	is version 84.38 (GRC		2.01	kb	29,034,250		— Forward strand ∎
C #2 rain:e14.5d eg. Segs. (Chro ver:a8w eg. Segs. (Chro idney:a8w eg. Segs. (Chro idney:a8w eg. Segs. (Chro sGC	Ensembl Mus musculu	is version 84.38 (GRC		2.01	kb	29,034,250		— Forward strand ■
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rain:e14.5d .eg. Segs. (Chro ver:a8w .eg. Segs. (Chro pleen:a8w .eg. Segs. (Chro idney:a8w .eg. Segs. (Chro	29,033,000	29,033,250	29,033,500 29,033,500	29,033,750 29,033,750 29,033,750 29,033,750 2.01	29,034,000 29,034,000 29,034,000 kb	29,034,250	29,034,500	— Forward strand ∎

Supplementary Figure 3. Chromatin state discovery and characterization (ChromHMM) in SBE6.1, SBE6.2 and peak #2.

Ensembl Mus musculus genome version 84.38 (GRCm38.p4) view of regulatory feature tracks from ChromHMM for the regions corresponding to (A) SBE6.1, (B) SBE6.2 and (C) putative peak #2 from Figure 3)