Scale chr19:	55,770,800 I 55,7	0,900 l 200 bases	ove expectation e Sequence Alignments ons
RFPL4AL1		19q13.42 GENCODE V47	
		RefSeq genes from I	NCBI
RefSeq Curated RFPL4AL1	Basic Gene Annotation Set from GENCODE Version 28 (Ensembl 92)		
KI F L4ALT		Pseudogene Annotation Set from GENCOL Geneid Gene Predic	DE Version 28 (Ensembl 92)
chr19_1362.1	***************************************	Genscan Gene Predi	<del>, , , , , , , , , , , , , , , , , , , </del>
chr19.1341	Alternative Solicina. Alternative Promoter and Similar Events in UCSC Genes		
OMIM Alleles	OMIM Allelic Variant Phenotypes		
RFPL4AL1	Gene Expression in 33 TCGA Cancer Tissues (GENCODE v23)		
RFPL4AL1	Transcript-level Expression in 33 TCGA Cancer Tissues (GENCODE v23)		
100 _		ClinGen CNVs: Benign Gai	n Coverage
Benign Gain			
100 _		ClinGen CNVs: Benign Los	s Coverage
Benign Loss			
0		ClinGen CNVs: Patho	ogénic
nssv13649600_unk nssv13643100_unk			
nssv13639328_unk nssv13646143_unk			
nssv576289_unk nssv576381_unk			
nssv578812_unk nssv578809 unk	<del></del>		
nssv578810_unk nssv578811_unk			
nssv578813_unk nssv582233_unk			
nssv3394913_unk		ClinGen CNVs: Pathogenic G	Cala Calvaraga
100 _		Clingen Chvs. Pathogenic G	rain Coverage
Path Gain			
0			
100 _		ClinGen CNVs: Pathogenic Lo	oss Coverage
Path Loss			
_			
0		Catalogue of Somatic Mutations	s in Cancer V82
Catalogue of Somatic Mutations in Cancer V82 Protein Interactions from Curated Databases and Text-Mining RFPL4AL1: UBC NHGRI-FRI Catalog of Published Genome-Wide Association Studies			pases and Text-Mining
DEDI 441.4		NHGRI-EBI Catalog of Published Genome Gene Expression in 53 tissues from GTEx RNA-s	eq of 8555 samples (570 donors)
RFPL4AL1		Transcript Expression in 53 tissues from GTEx RN	A-seq of 8555 samples/570 donors
L4AL1/ENST00000341750 100 _		H3K27Ac Mark (Often Found Near Regulatory Elem	nents) on 7 cell lines from ENCODE
Layered H3K27Ac			
0		DNase I Hypersensitivity Peak Clusters fro	m ENCODE (95 cell types)
DNase Clusters		GeneHancer Regulatory Elements and Gene Interactions	
GH Reg Elems (DE)	GeneCards genes TSS (Double Elite) Interactions between GeneHancer regulatory elements and genes (Double Elite)		
	Interactions between GeneHancer regulatory elements and genes (Double Elite)		
		Clustered interactions of GeneHancer regulatory e Regulatory elements from NCBI RefSeg Functional	elements and genes (Double Elite) ORegAnno Elements