

Scale  
chr7:

5 kb | hg38  
155,805,000 | 155,810,000

Transcript structures for LncExpDB and LncBook databases

HSALNT0344402

HSALNT0131333

HSALNT0131331

HSALNT0131332

Perfect Matches to Short Sequence ()  
GENCODE V47 (4 items filtered out)

ENSG00000296779

SHH

GTEX V8 RNA-Seq Read Coverage by Tissue

- Whole Blood
- Vagina
- Uterus
- Thyroid
- Testis
- Stomach
- Spleen
- Small Intestine
- Skin sun exp
- Skin not sun exp
- Prostate
- Pituitary
- Pancreas
- Ovary
- Nerve Tibial
- Muscle Skeletal
- Minor Saliv Gland
- Lung
- Liver
- Kidney Medulla
- Kidney Cortex
- Heart Left Ventr
- Heart Atr Append
- Fallopian Tube
- Esoph Muscularis
- Esoph Mucosa
- Esoph Gastroes Junc
- Colon Transverse
- Colon Sigmoid
- Cervix Endocerv
- Cervix Ectocerv
- Cells fibrobl cult
- Cells EBV lymphoc
- Breast Mammary
- Brain Subst nigr
- Brain Spinal cord cerv
- Brain Put bas gang
- Brain Nucl acc bas gang
- Brain Hypothal
- Brain Hippocamp
- Brain Front Cortex
- Brain Cortex
- Brain Cereb Hemisph
- Brain Cereb
- Brain Caud bas gangl
- Brain Ant cin cort
- Brain Amygd
- Bladder
- Artery Tibia
- Artery Coron
- Artery Aorta
- Adren Gland
- Adip Visc Om
- Adip Subcut



TSS peaks

FANTOM5: DPI peak, robust set

UCSC 100 Vertebrates - 100 vertebrate genomes aligned with MultiZ by the UCSC Browser Group

Cons 100 Verts

Multiz Align

Repeating Elements by RepeatMasker

RepeatMasker