

Product datasheet for RC204733L4V

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Thyroxine Binding Globulin (SERPINA7) (NM_000354) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type: Lentiviral Particles

Product Name: Thyroxine Binding Globulin (SERPINA7) (NM_000354) Human Tagged ORF Clone Lentiviral

Particle

Symbol: Thyroxine Binding Globulin

Synonyms: TBG; TBGQTL

Mammalian Cell Puromycin

Selection:

Vector:

pLenti-C-mGFP-P2A-Puro (PS100093)

Tag: mGFP

ACCN: NM_000354 **ORF Size:** 1245 bp

ORF Nucleotide

The ORF insert of this clone is exactly the same as(RC204733).

OTI Disclaimer:

Sequence:

The molecular sequence of this clone aligns with the gene accession number as a point of reference only. However, individual transcript sequences of the same gene can differ through naturally occurring variations (e.g. polymorphisms), each with its own valid existence. This clone is substantially in agreement with the reference, but a complete review of all prevailing

variants is recommended prior to use. More info

OTI Annotation: This clone was engineered to express the complete ORF with an expression tag. Expression

varies depending on the nature of the gene.

RefSeq: <u>NM 000354.3</u>

RefSeq Size: 1600 bp
RefSeq ORF: 1248 bp
Locus ID: 6906
UniProt ID: P05543
Cytogenetics: Xq22.3
Domains: SERPIN





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Protein Families: Druggable Genome, Secreted Protein

MW: 46.3 kDa

Gene Summary: There are three proteins including thyroxine-binding globulin (TBG), transthyretin and

albumin responsible for carrying the thyroid hormones thyroxine (T4) and 3,5,3'-

triiodothyronine (T3) in the bloodstream. This gene encodes the major thyroid hormone transport protein, TBG, in serum. It belongs to the serpin family in genomics, but the protein has no inhibitory function like many other members of the serpin family. Mutations in this gene result in TGB deficiency, which has been classified as partial deficiency, complete deficiency, and excess, based on the level of serum TBG. Alternatively spliced transcript variants encoding different isoforms have been found, but the full-length nature of these

variants has not been determined.[provided by RefSeq, Jun 2012]