

OriGene Technologies, Inc.

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Product datasheet for RC226384L3V

FANCI (NM_001113378) Human Tagged ORF Clone Lentiviral Particle

Product data:

Product Type:	Lentiviral Particles
Product Name:	FANCI (NM_001113378) Human Tagged ORF Clone Lentiviral Particle
Symbol:	FANCI
Synonyms:	KIAA1794
Mammalian Cell Selection:	Puromycin
Vector:	pLenti-C-Myc-DDK-P2A-Puro (PS100092)
Tag:	Myc-DDK
ACCN:	NM_001113378
ORF Size:	3984 bp
ORF Nucleotide Sequence:	The ORF insert of this clone is exactly the same as(RC226384).
OTI Disclaimer:	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. <u>More info</u>
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 001113378.1</u>
RefSeq ORF:	3987 bp
Locus ID:	55215
UniProt ID:	<u>Q9NVI1</u>
Cytogenetics:	15q26.1
MW:	149.1 kDa



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GRIGENE FANCI (NM_001113378) Human Tagged ORF Clone Lentiviral Particle – RC226384L3V

Gene Summary:The Fanconi anemia complementation group (FANC) currently includes FANCA, FANCB,
FANCC, FANCD1 (also called BRCA2), FANCD2, FANCE, FANCF, FANCG, FANCI, FANCJ (also
called BRIP1), FANCL, FANCM and FANCN (also called PALB2). The previously defined group
FANCH is the same as FANCA. Fanconi anemia is a genetically heterogeneous recessive
disorder characterized by cytogenetic instability, hypersensitivity to DNA crosslinking agents,
increased chromosomal breakage, and defective DNA repair. The members of the Fanconi
anemia complementation group do not share sequence similarity; they are related by their
assembly into a common nuclear protein complex. This gene encodes the protein for
complementation group I. Alternative splicing results in two transcript variants encoding
different isoforms. [provided by RefSeq, Jul 2008]

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