

## OriGene Technologies, Inc.

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## Product datasheet for SR302456

## Integrin alpha 1 (ITGA1) Human siRNA Oligo Duplex (Locus ID 3672)

## **Product data:**

Product Type:	siRNA Oligo Duplexes
Purity:	HPLC purified
Quality Control:	Tested by ESI-MS
Sequences:	Available with shipment
Stability:	One year from date of shipment when stored at -20°C.
# of transfections:	Approximately 330 transfections/2nmol in 24-well plate under optimized conditions (final conc. 10 nM).
Note:	Single siRNA duplex (10nmol) can be ordered.
RefSeq:	<u>NM 181501</u>
UniProt ID:	<u>P56199</u>
Synonyms:	CD49a; VLA1
Components:	ITGA1 (Human) - 3 unique 27mer siRNA duplexes - 2 nmol each (Locus ID 3672) Included - SR30004, Trilencer-27 Universal Scrambled Negative Control siRNA Duplex - 2 nmol Included - SR30005, RNAse free siRNA Duplex Resuspension Buffer - 2 ml
Summary:	This gene encodes the alpha 1 subunit of integrin receptors. This protein heterodimerizes with the beta 1 subunit to form a cell-surface receptor for collagen and laminin. The heterodimeric receptor is involved in cell-cell adhesion and may play a role in inflammation and fibrosis. The alpha 1 subunit contains an inserted (I) von Willebrand factor type I domain which is thought to be involved in collagen binding. [provided by RefSeq, Jul 2008]



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Performance Guaranteed:	OriGene guarantees that at least two of the three Dicer-Substrate duplexes in the kit will provide at least 70% or more knockdown of the target mRNA when used at 10 nM
	concentration by quantitative RT-PCR when the TYE-563 fluorescent transfection control
	duplex (cat# SR30002) indicates that >90% of the cells have been transfected and the HPRT
	positive control (cat# SR30003) provides 90% knockdown efficiency.
	For non-conforming siRNA, requests for replacement product must be made within ninety
	(90) days from the date of delivery of the siRNA kit. To arrange for a free replacement with
	newly designed duplexes, please contact Technical Services at techsupport@origene.com.
	Please provide your data indicating the transfection efficiency and measurement of gene expression knockdown compared to the scrambled siRNA control (quantitative RT-PCR data

required).

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