

Product datasheet for LY300178

OriGene Technologies, Inc.

9620 Medical Center Drive, Ste 200 Rockville, MD 20850, US Phone: +1-888-267-4436 https://www.origene.com techsupport@origene.com EU: info-de@origene.com CN: techsupport@origene.cn

GNAQ Human Knockdown Lysate

Product data:

Product Type: Knockdown Lysates

Description: WB-validated GNAQ Knockdown HeLa Cell Lysate

Species: Human Expression Host: HeLa

Tag: Tag Free

Synonyms: GNAQ; G Protein Subunit Alpha Q; GAQ; G-ALPHA-Q; Guanine Nucleotide Binding Protein (G

Protein), Q Polypeptide; Guanine Nucleotide-Binding Protein G(Q) Subunit Alpha; Guanine Nucleotide-Binding Protein Alpha-Q; Epididymis Secretory Sperm Binding Protein; CMAL;

CMC1; SWS

Predicted MW: 42 kDa

Components: 1 vial of 100 ug WT HeLa cell lysate

1 vial of 100 ug GNAQ KD HeLa cell lysate

Storage: Store at -20 °C for two years.

Concentration: Lot-specific

Buffer: IntactProtein Cell-Tissue Lysis buffer

Locus ID: 2776 **UniProt ID:** P50148

Protein Families: Druggable Genome

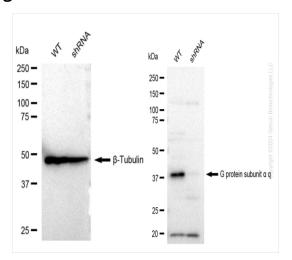
Protein Pathways: Alzheimer's disease, Calcium signaling pathway, Gap junction, GnRH signaling pathway,

Huntington's disease, Long-term depression, Long-term potentiation, Melanogenesis,

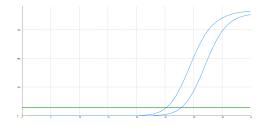
Vascular smooth muscle contraction



Product images:



Western blotting analysis. GNAQ protein expression in wild-type (WT) and shRNA knockdown (KD) HeLa cells was detected using Western blotting. β -Tubulin served as a loading control. The blots were incubated with primary antibodies (Cat#63616, 1:5,000) against GNAQ and β -Tubulin , respectively, followed by incubating with HRP-conjugated goat anti-rabbit secondary antibody (Cat#201, 1:20,000). Images were developed using FeQTM ECL Substrate Kit (Cat#226).



| Genotype | Ct Value |
|---|----------|
| Wild-Type | 25.10 |
| Knock-Down | 27.66 |
| Δ Ct (Ct _{KD} -Ct _{WT}) | 2.56 |
| % mRNA Reduction | ₽ 83% |

RT-qPCR analysis. HeLa cells were infected with GNAQ-specific shRNA lentiviral particles, total RNA was extracted from wild-type and knockdown cells, RT-qPCR was performed using gene-specific primers. Δ Ct (CtKD-CtWT) was used to calculate mRNA reduction (%) between wild-type and knockdown cells using the following formula: $(1-1/2\Delta$ Ct) x 100%.