

## Product datasheet for **MG227194**

### Pik3cg (NM\_001146200) Mouse Tagged ORF Clone

#### Product data:

Product Type:	Expression Plasmids
Product Name:	Pik3cg (NM_001146200) Mouse Tagged ORF Clone
Tag:	TurboGFP
Symbol:	Pik3cg
Synonyms:	5830428L06Rik; p110gamma; PI3Kgamma
Mammalian Cell Selection:	Neomycin
Vector:	pCMV6-AC-GFP (PS100010)
E. coli Selection:	Ampicillin (100 ug/mL)
ORF Nucleotide Sequence:	>MG227194 representing NM_001146200 Red=Cloning site Blue=ORF Green=Tags(s)

TTTTGTAATACGACTCACTATAGGGCGCCGGGAATTCGTCGACTGGATCCGGTACCGAGGAGATCTGCC  
GCC**CGATCGCC**

ATGGAGCTGGAGAACTATGAACAACCGGTGGTTCTAAGAGAGGACAACCTCCGCCGGCGCCGGAGGATGA  
AGCCACGCAGCGCAGCAGGCAGCCTGTCTCCATGGAGCTCATCCCCATTGAGTTCGTAAGTCCACCAG  
CCAGCGCATCAGCAAGACTCCAGAAACAGCGCTGCTGCATGTGGCTGGCCATGGCAATGTGGAACAGATG  
AAAGCTCAGGTGTGGCTGCGCGCACTGGAGACCAGTGTGGCTGCGGAGTCTACCACCGATTGGGCCGG  
ACCAATTCCTCCTGCTCTACCAGAAGAAAGGACAATGGTATGAGATCTATGACAGGTACCAAGTGGTGCA  
GACCCTAGACTGCCTGCATTACTGGAAGTTGATGCACAAGAGCCCTGGCCAGATCCACGTGTACAGCGA  
CACGTACCTTCTGAGGAGACCTTGGCTTTCCAGAAGCAGCTCACCTCCCTGATTGGCTATGACGTCACTG  
ACATCAGCAATGTGCACGATGATGAGCTAGAGTCACTCGCCCGCTGTGGTTACGCCCCGCATGGCTGA  
AGTGGCTGGCCGGGATGCCAAACTCTATGCTATGCACCCTTGGGTAACGTCCAAACCTCTCCAGACTAC  
CTGTCAAAAAGATTGCCAAACTGCATCTTCATCGTCATCCACCGCGTACCACCAGCCAAACCATCA  
AGGTCTCCGCAGATGATACTCCTGGTACCATCTCCAGAGCTTCTCACCAAGATGGCCAAGAAGAAGTC  
CCTAATGAATATCTCAGAAAGTCAAAGTGAGCAGGATTTTGTATTGCGGGTTTGTGGCCCGATGAGTAC  
CTGGTGGGTGAAACACCCCTCAAAAATTTCCAGTGGGTGAGGCAGTGCCTCAAGAACGGAGATGAAATAC  
ACCTGGTGTCTCGACACCGCTCCAGACCCAGCCCTTGTGAGGTGAGGAAGGAAGAATGGCCGCTGGTGGA  
TGAAGTCACTGGAGTCAACCGCTACCACGAGCAGCTGACCATCCATGGCAAGACCACGAGAGTGTGTTT  
ACAGTGTCTTTGTGGACTGCGACCGAAAGTTCAGGGTCAAGATCAGAGGCATTGATATCCCTGTCTGCTGC  
CTCGGAACACCGACCTCACTGTGTTTGTGGAAGCGAACATCCAGCACGGGCAACAAGTCTCTGCCAAAG  
GAGAACCAGCCCTAAGCCCTTCGAGAAGAGGTAAGTCTGGAATGTGTGGCTGGAGTTTGGCATCAAATC  
AAAGACTGCCAAAGGGGCTCTATTGAACCTACAGATCTACTGCTGCAAAACCCATCACTGTCCAGCA  
AGGCTTCTGCAGAGACTCCAGGCTCCGAGTCCAAGGGCAAGCCAGCTTCTCTATTACGTGAACCTTGT  
GTTAATAGACCACCGTTTCTCCTCCGCCAGGGGACTATGTGCTCCACATGTGGCAGATATCTGGCAAG



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GCAGAGGAGCAGGGCAGCTTCAATGCTGACAAGCTCACATCCGCAACCAATCCTGACAAGGAGAACTCAA  
 TGTCCATTTCCATCCTGCTGGACAATTACTGTACCCCATAGCTTTGCCTAAGCACCGGCCACCCCTGA  
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 AGCATCCGAAGGCTTACCCTAAGCTATTCAGCTCAGTAAATGGGGCAGCAAGAAATGTTGCCAAAAC  
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 GATCGCTCAAAATTCAGCAAAGCACAGTGGGTAACACGGGGCATTCAAAGATGAAGTCTGAATCACTGG  
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 GAGAGAGTGCCCTTCGTCTAACCCAGACTTCTGTTTGTGATGGGATCTTCTGGAAAAAGACAAGTC  
 CACACTCCAGAAATCCAGGATGTCTGTGTTAGAGCTTACCTAGCTCTTCCGCATCACACAAACCTGTT  
 GATCATCTTGTCTCCATGATGCTGATGACAGGAATGCCCCAGCTGACAAGCAAAGAGGACATTGAATAT  
 ATCCGGGATGCCCTACCGTGGGAAAAAGCGAGGAGGACGCTAAGAAATATTTCTTGTGATCAGATCGAAG  
 TCTGCAGAGACAAAGGATGGACTGTGCAGTTTAACTGGTCTACATCTTGTCTTGGCATCAAACAAGG  
 AGAAAAGCACTCCGCT

ACGCGTACGCGGGCCGCTCGAG - GFP Tag - GTTTAA

**Protein Sequence:**

>MG227194 representing NM\_001146200  
 Red=Cloning site Green=Tags(s)

MELENYEQPVVLRDNLRRRRRMRKPRSAAGSLSSMELIPIEFVLPSTQRISKTPETALLHVAGHGNVEQM  
 KAQVWLRALETSVAAEFYHRLGPDQFLLLYQKKGQWYEIYDRYQVVQTLDCLEHYWKLMMKSPGQIHVVQR  
 HVPSEETLAFQKQLTSLIGYDVTDISNVHDELEFTRRRLVTPRMAEVAGRDAKLAMHPVWTSKPLPDY  
 LSKKIANNCFIVIHRTTSQTIKVSADDTPTGILQSFFTKMAKKSLMNISESQSEQDFVLRVCGRDEY  
 LVGETPLKNFQWVRQCLKNGDEIHLVLDTPDPALDEVRKEEWPLVDDCTGVTGYHEQLTIHGKDHEVSVF  
 TVSLWDCDRKFRVKIRGIDIPVLPNTDLTVFVEANIQHQQVLCQRRTPSPKPF AEEVLWNVWLEFGIKI  
 KDLPKGALLNLQIYCKTPSLSSKASAETPGSESKGAQLLYVNLILLIDHRFLLRHGDYVLLHMWQISGK  
 AEEQGSFNADKLT SATNPDKENSMSISILLDNYCHPIALPKHRPTDPEGDRVRAEMPQLRKQLEAIIA  
 TDPLNPLTAEDKELLWHFRYESLKHPKAYPKLFSSVKWGQQEIVAKTYQLLARREIWDQSALDVGLTMQL  
 LDCNFSDENVRAIAVQKLESLEDDDLHYLLQLVQAVKFEPYHDSALARFLLKRGLRNKRIGHFLFWFLR  
 SEIAQSRHYQQRFAVILEAYLRGCGTAMLQDFTQQVHVIEMLQKVTIDIKSLSAEKYDVSSQVISQLKQK  
 LESLQNSNLPESFRVPYDPGLKAGTLVIEKCKVMASKKKPLWLEFKCADPTVLSNETIGIIFKHGDDLRLQ  
 DMLILQILRIMESIWETESLDLCLLPYGCISTGDKIGMIEIVKDATTIAQIQSTVGTGAFKDEVLNHW  
 LKEKCPIEEKFAAVERFVYSCAGYCVATFVLGIGDRHNDNIMISSETGNLFHIDFGHILGNYKSFLLGINK  
 ERVPFVLPDFLVMGSSGKTSPHFQKQFQDVCVFRAYLALRHHTNLLIILFSMMLMTGMPQLTSKEDIEY  
 IRDALTVGKSEEDAKKYFLDQIEVCRDKGWTQVFNWFLHLVGLGIKQGEKHS

TRTRPLE - GFP Tag - V

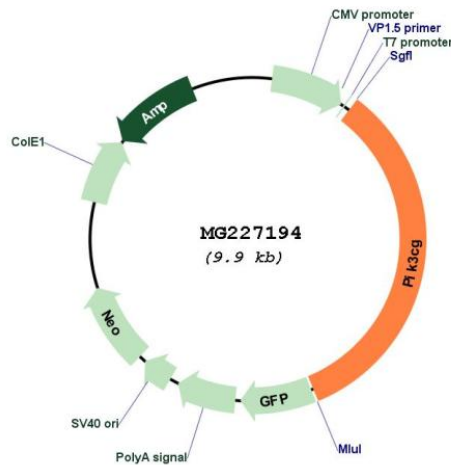
**Restriction Sites:**

Sgfl-MluI

Cloning Scheme:



Plasmid Map:



ACCN: NM\_001146200

ORF Size: 3306 bp

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. [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.

<b>Components:</b>	The ORF clone is ion-exchange column purified and shipped in a 2D barcoded Matrix tube containing 10ug of transfection-ready, dried plasmid DNA (reconstitute with 100 ul of water).
<b>Reconstitution Method:</b>	<ol style="list-style-type: none"><li>1. Centrifuge at 5,000xg for 5min.</li><li>2. Carefully open the tube and add 100ul of sterile water to dissolve the DNA.</li><li>3. Close the tube and incubate for 10 minutes at room temperature.</li><li>4. Briefly vortex the tube and then do a quick spin (less than 5000xg) to concentrate the liquid at the bottom.</li><li>5. Store the suspended plasmid at -20°C. The DNA is stable for at least one year from date of shipping when stored at -20°C.</li></ol>
<b>RefSeq:</b>	<u>NM_001146200.2</u>
<b>RefSeq Size:</b>	6634 bp
<b>RefSeq ORF:</b>	3309 bp
<b>Locus ID:</b>	30955
<b>UniProt ID:</b>	<u>Q9JHG7</u>
<b>Cytogenetics:</b>	12 A3

**Gene Summary:**

Phosphoinositide-3-kinase (PI3K) that phosphorylates PtdIns(4,5)P<sub>2</sub> (Phosphatidylinositol 4,5-bisphosphate) to generate phosphatidylinositol 3,4,5-trisphosphate (PIP<sub>3</sub>). PIP<sub>3</sub> plays a key role by recruiting PH domain-containing proteins to the membrane, including AKT1 and PDK1, activating signaling cascades involved in cell growth, survival, proliferation, motility and morphology. Links G-protein coupled receptor activation to PIP<sub>3</sub> production. Involved in immune, inflammatory and allergic responses. Modulates leukocyte chemotaxis to inflammatory sites and in response to chemoattractant agents. May control leukocyte polarization and migration by regulating the spatial accumulation of PIP<sub>3</sub> and by regulating the organization of F-actin formation and integrin-based adhesion at the leading edge. Controls motility of dendritic cells. Together with PIK3CD is involved in natural killer (NK) cell development and migration towards the sites of inflammation. Participates in T-lymphocyte migration. Regulates T-lymphocyte proliferation and cytokine production. Together with PIK3CD participates in T-lymphocyte development. Required for B-lymphocyte development and signaling. Together with PIK3CD participates in neutrophil respiratory burst. Together with PIK3CD is involved in neutrophil chemotaxis and extravasation. Together with PIK3CB promotes platelet aggregation and thrombosis. Regulates alpha-IIb/beta-3 integrins (ITGA2B/ITGB3) adhesive function in platelets downstream of P2Y<sub>12</sub> through a lipid kinase activity-independent mechanism. May have also a lipid kinase activity-dependent function in platelet aggregation. Involved in endothelial progenitor cell migration. Negative regulator of cardiac contractility. Modulates cardiac contractility by anchoring protein kinase A (PKA) and PDE3B activation, reducing cAMP levels. Regulates cardiac contractility also by promoting beta-adrenergic receptor internalization by binding to GRK2 and by non-muscle tropomyosin phosphorylation. Also has serine/threonine protein kinase activity: both lipid and protein kinase activities are required for beta-adrenergic receptor endocytosis. May also have a scaffolding role in modulating cardiac contractility. Contribute to cardiac hypertrophy under pathological stress. Through simultaneous binding of PDE3B to RAPGEF3 and PIK3R6 is assembled in a signaling complex in which the PI3K gamma complex is activated by RAPGEF3 and which is involved in angiogenesis (By similarity).[UniProtKB/Swiss-Prot Function]