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**gk<sup>K414E</sup> - Mouse Strain RES192****Mouse Information**

<b>Common Name:</b>	gk <sup>K414E</sup>
<b>MGI Official Name:</b>	Gck <sup>tm2Mgn</sup>
<b>Description:</b>	This line of mice contains a gk <sup>K414E</sup> point mutation that was introduced by gene knock-in. This point mutation was identified in a MODY-GK pedigree. These mice will be useful for studies of sustained hyperglycemia since they contain only a single mutation and are congenic with the C57BL/6J strain.
<b>Categories:</b>	None specified.

**Genetic Alterations****1) Targeted Mutagenesis**

<b>Type of Allele</b>	Global Mutation
<b>Targeted Gene</b>	Glucokinase (Gck - <a href="#">NCBI GeneID:103988</a> )
<b>Targeted Allele</b>	targeted mutation 2 (Gck <sup>tm2Mgn</sup> - <a href="#">MGI:3701697</a> )
<b>Description of Targeting Vector</b>	A single base pair mutation was introduced into exon 9 via site specific mutagenesis to change amino acid 414 from lysine to glutamic acid. Genotype by DNA PCR using primers 5'-TGT CTC AAT TTG CTG TGT CCT CCA-3' and 5'-ATG TGT GAG TGT GCC AAT ATG AGT-3'. These primers will amplify a 636 bp fragment from the wild type allele and a 741 bp fragment from the targeted (mutant) allele. These animals are homozygous lethal. Heterozygous mice are viable. They are hyperglycemic and hypoinsulinemic when compared to wild type.
<b>Targeting Vector Genbank File</b>	<a href="#">pBOB2_K414E.gb</a>

Citations	PubMedID	Citation
	<a href="#">17353190</a>	<a href="#">Glucokinase thermolability and hepatic regulatory protein binding are essential factors for predicting the blood glucose phenotype of missense mutations.</a> (2007) <i>J Biol Chem</i> <b>282</b> : 13906-16 (Added 2008-03-29 16:59:08)


**Strain Information**

<b>Strain Type:</b>	Congenic Strain
<b>Chimera/Founder Genetic Background:</b>	129S6/SvEvTac
<b>Current Genetic Background:</b>	C57BL/6J (date recorded: Not provided)
<b>Strain Description:</b>	After achieving germline transmission mice carrying the gk <sup>K414E</sup> allele were backcrossed for eleven generations into a C57BL/6J strain.


**Associated Images**

Image 1	Description:
	Gene targeting strategy used to generate point mutations in the gk gene. Uppermost map is

**Access Status**

 This resource is publicly viewable.

**Request this Resource**

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Primary contributor: [Magnuson Lab](#)

**Resource Tags**

Gck, gk<sup>K414E</sup>, mouse, mouse strain

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**Resource History & Actions**

Approved on Feb 02, 2007  
Last modified on Sep 20, 2007

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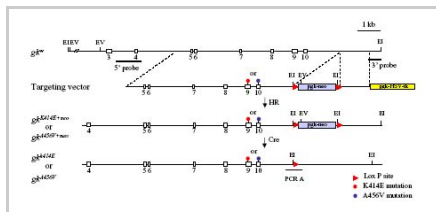
**Related resources****BCBC**

No matching resources

**Other Consortia**

No matching resources

Data courtesy of [dkCOIN](#). Only public resources are displayed.



a diagram of the mouse *gk* gene showing locations of exons 3 to 10 (indicated by open boxes). The locations of the DNA fragments used as 5' and 3' hybridization probes are shown. E1, EcoRI; EV, EcoRV. Second map from top is of the gene targeting vector carrying either the K414E mutation in exon 9 depicted by the red circle or the A456V mutation in exon 10 as indicated by the blue circle. A neomycin resistance cassette (pgk-neoR), which is flanked with two loxP sites depicted by red triangles, and a HSV-thymidine kinase cassette (pgk-HSV-TK), were used for positive and negative selection, respectively. Third map from top is of the recombinant *gk* allele after homologous recombination (HR) carrying a floxed pgk-neoR cassette and the respective point mutation in exon 9 or 10. Fourth map from top is of the mutant *gk* allele after Cre recombination.

**Reference:**  
17353190

## Repositories

MMRRC

Request via [www.mmrrc.org](http://www.mmrrc.org) website

Stock #: 015201-UCD

Availability Notes: *Not provided*

BCBC members may [Login](#) to request this resource.

## Contact Information

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## Associated Publications

*No publications associated*

## Comments

*There are no comments for this entry.*

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