

**Gene:** Il6ra

**Colony prefix:** DASU

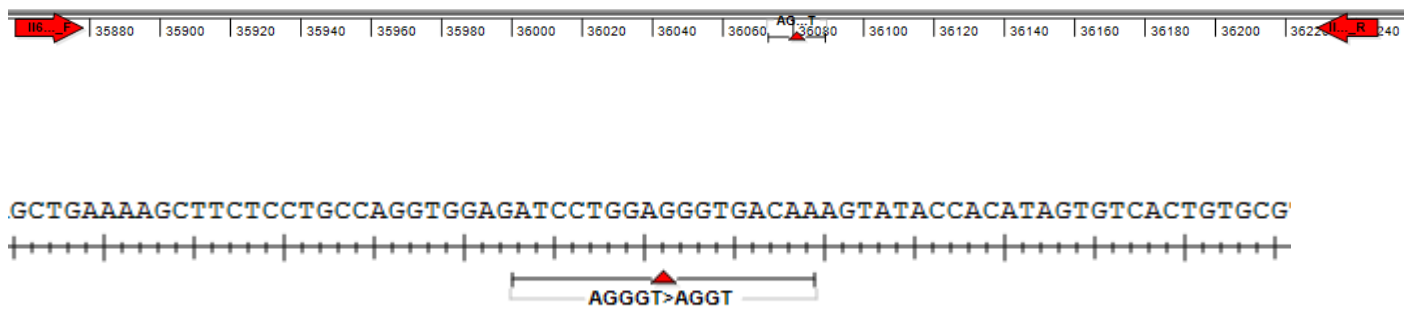
**Allele:** *Il6ra*<sup>em3(IMPC)Wtsi</sup>

**Allele type:** Crispr/Cas9 mediated Point mutation

**Allele information:**

Single base deletion. Expect mixed sequence after the deletion in heterozygous mice. Use <https://synthego.com/products/bioinformatics/crispr-analysis> to aid in analysis.

Further information about the allele can be found on the 'International Mouse Phenotyping Consortium' (IMPC) web site at <http://www.mousephenotype.org/>



**Mouse QC information**

|                              |    |                              |      |
|------------------------------|----|------------------------------|------|
| Loss of WT Allele (LoA) qPCR | na | Mutation Sequence confirmed  | pass |
| Mutant Specific SR-PCR       | na | Off-target analysis complete | na   |

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## Guide RNAs and mutant oligos used in initial experiment

| Sequence  | Chr | Chr Start | Chr End  |
|---|-----|-----------|----------|
| GGTATACTTTGTCACCCTCCAGG   | 3   | 89794405  | 89794430 |
| CTGGACTTGCTTCCCACACTGTTTGCAACGCACAGTGACACTAT<br>GTGGTATACTTTGTCACCCTCCAGGATCTCCACCTGGCAGGAGA<br>AGCTTTTCAGCTGCTGAGAATACTGACAGGGCA | 3   | 89794355  | 89794487 |

### Mutant allele sequence:

TGTCCCACAGCAACACCACCAACGGGAAGAGTGACTTCCAGGTGCCCTGTCAGTATTCTCAGCAGCT  
GAAAAGCTTCTCCTGCCAGGTGGAGATCCTGGAGGTGACAAAGTATACCACATAGTGTCACTGTGC  
GTTGCAAACAGTGTGGGAAGCAAGTCCAGCCACAACGAAGCGTTTCACAGCTTAAAAATGGGTAA  
GG

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## Genotyping by end-point PCR

### PCRs primer pairs and expected size bands

| Assay Type   | Assay      | Forward Primer | Reverse Primer | Expected Size Band (bp) |
|--------------|------------|----------------|----------------|-------------------------|
| Standard PCR | Screening* | Il6ra_PM_EM3_F | Il6ra_PM_EM3_R | 388                     |

\*The screening PCR flanks the SNP region and can be used for sequence verification of the allele. The PCR will not distinguish wild type from mutant mice, however, as a product will be amplified in all cases.

**We recommend that mice are sequence-verified with the screening primers to confirm the genotype.**

### Primer sequences

| Primer Name    | Primer Sequence (5' > 3') |
|----------------|---------------------------|
| Il6ra_PM_EM3_F | GCAGACACTGGCTGAGTTGA      |
| Il6ra_PM_EM3_R | CACATCCTCCCTGCCTTG        |

### Reaction setup

| Reagent                   | µl   |
|---------------------------|------|
| DNA (~50-100 ng)          | 1    |
| 10x Buffer                | 2    |
| MgCl <sub>2</sub> (50mM)  | 0.6  |
| Platinum Taq (Invitrogen) | 0.2  |
| dNTP's (100mM)            | 0.2  |
| Primer 1 (10 µM)          | 0.4  |
| Primer 2 (10 µM)          | 0.4  |
| ddH <sub>2</sub> O        | 15.2 |
| Total                     | 20   |

### Amplification conditions

| Step | Conditions            | Time     |
|------|-----------------------|----------|
| 1    | 94°C                  | 5 min    |
| 2    | 94°C                  | 30 sec   |
| 3    | 58°C                  | 30 sec   |
| 4    | 72°C                  | 1:30 sec |
| 5    | Go to '2' + 34 cycles | -        |
| 6    | 72°C                  | 5 min    |
| 7    | 12°C                  | Forever  |

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## Links to information and frequently asked questions

MGP mouse phenotype data:

<http://www.mousephenotype.org>

How the "critical" exon is decided:

<http://www.i-dcc.org/kb/entry/102/>

## Relevant publications

White, J.K., Gerdin, A.-K., Karp, N.A., Ryder, E., Buljan, M., Bussell, J.N., Salisbury, J., Clare, S., Ingham, N.J., Podrini, C., et al. (2013). Genome-wide Generation and Systematic Phenotyping of Knockout Mice Reveals New Roles for Many Genes. *Cell* 154, 452–464.

Mali P, Yang L, Esvelt KM, et al (2013) RNA-guided human genome engineering via Cas9. *Science* 339:823–6. doi: 10.1126/science.1232033

Jinek M, Chylinski K, Fonfara I, et al (2012) A programmable dual-RNA-guided DNA endonuclease in adaptive bacterial immunity. *Science* 337:816–21. doi: 10.1126/science.1225829

Cong L, Ran FA, Cox D, et al (2013) Multiplex genome engineering using CRISPR/Cas systems. *Science* 339:819–23. doi: 10.1126/science.1231143

Singh P, Schimenti JC, Bolcun-Filas E (2014) A Mouse Geneticist's Practical Guide to CRISPR Applications. *Genetics* genetics.114.169771–. doi: 10.1534/genetics.114.169771

Brandl C, Ortiz O, Röttig B, et al (2015) Creation of targeted genomic deletions using TALEN or CRISPR/Cas nuclease pairs in one-cell mouse embryos. *FEBS Open Bio* 5:26–35. doi: 10.1016/j.fob.2014.11.009

Zhou J, Wang J, Shen B, et al (2014) Dual sgRNAs facilitate CRISPR/Cas9 mediated mouse genome targeting. *FEBS J*. doi: 10.1111/febs.12735

Kraft K, Geuer S, Will AJ, et al (2015) Deletions, Inversions, Duplications: Engineering of Structural Variants using CRISPR/Cas in Mice. *Cell Rep*. doi: 10.1016/j.celrep.2015.01.016

Shen B, Zhang J, Wu H, et al (2013) Generation of gene-modified mice via Cas9/RNA-mediated gene targeting. *Cell Res* 23:720–3. doi: 10.1038/cr.2013.46

Wang H, Yang H, Shivalila CS, et al (2013) One-step generation of mice carrying mutations in multiple genes by CRISPR/Cas-mediated genome engineering. *Cell* 153:910–8. doi: 10.1016/j.cell.2013.04.025

Yang H, Wang H, Shivalila CS, et al (2013) One-Step Generation of Mice Carrying Reporter and Conditional Alleles by CRISPR/Cas-Mediated Genome Engineering. *Cell* 154:1370–1379. doi: 10.1016/j.cell.2013.08.022

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