

Gpihbp1 Cas9-KO Strategy

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Project Overview

Project Name

Gpihbp1

Project type

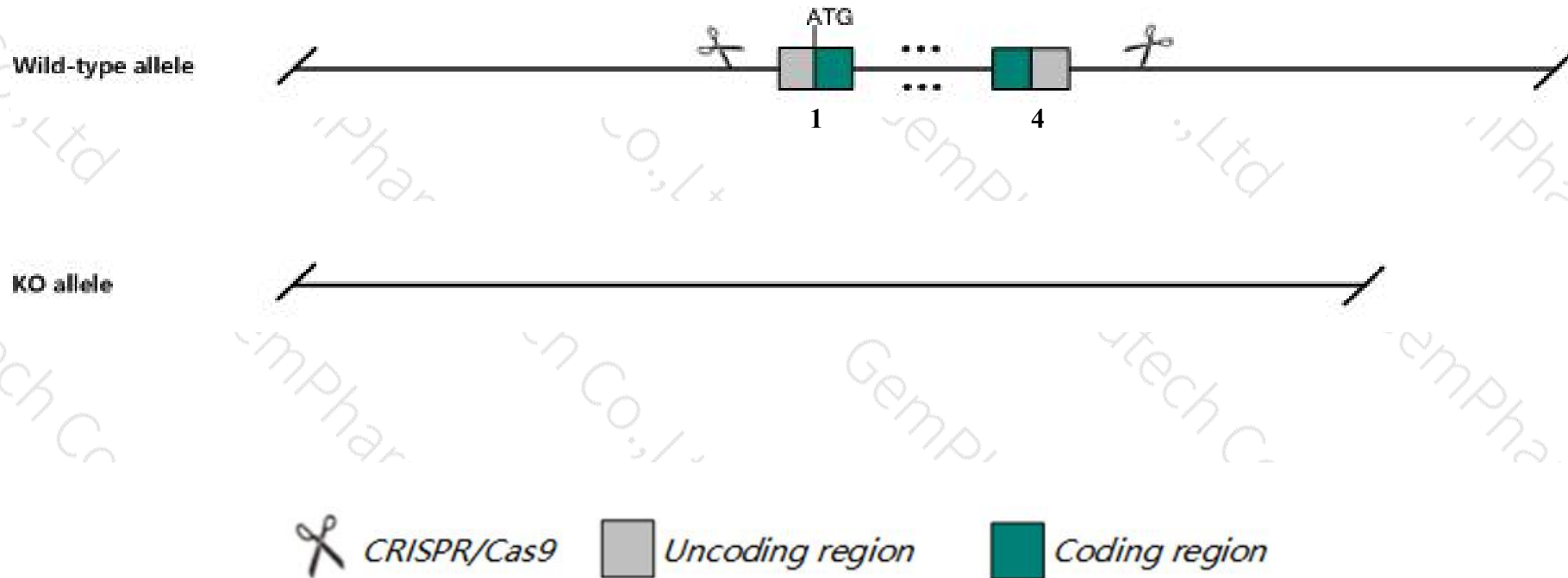
Cas9-KO

Strain background

C57BL/6JGpt

Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Gpihbp1* gene. The schematic diagram is as follows:



- The *Gpihbp1* gene has 5 transcripts. According to the structure of *Gpihbp1* gene, exon1-exon4 of *Gpihbp1-201* (ENSMUST00000023243.10) transcript is recommended as the knockout region. The region contains all of the coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Gpihbp1* gene. The brief process is as follows: CRISPR/Cas9 system

- According to the existing MGI data, Mice homozygous for a null allele exhibit milky plasma due to increased triglyceride and cholesterol levels.
- The *Gpihbp1* gene is located on the Chr15. If the knockout mice are crossed with other mice strains to obtain double gene positive homozygous mouse offspring, please avoid the two genes on the same chromosome.
- This Strategy is designed based on genetic information in existing databases. Due to the complexity of biological processes, all risk of the gene knockout on gene transcription, RNA splicing and protein translation cannot be predicted at the existing technology level.

Gene information (NCBI)

Gpihbp1 GPI-anchored HDL-binding protein 1 [*Mus musculus* (house mouse)]

Gene ID: 68453, updated on 24-Aug-2019

Summary

Official Symbol Gpihbp1 provided by [MGI](#)
Official Full Name GPI-anchored HDL-binding protein 1 provided by [MGI](#)
Primary source [MGI:MGI:1915703](#)
See related [Ensembl:ENSMUSG00000022579](#)
Gene type protein coding
RefSeq status VALIDATED
Organism [Mus musculus](#)
Lineage Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as GPI-HBP1; 1110002J19Rik
Expression Biased expression in lung adult (RPKM 157.2), subcutaneous fat pad adult (RPKM 146.6) and 10 other tissues [See more](#)
Orthologs [human](#) [all](#)

Genomic context

Location: 15; 15 D3

See Gpihbp1 in [Genome Data Viewer](#)

Exon count: 4

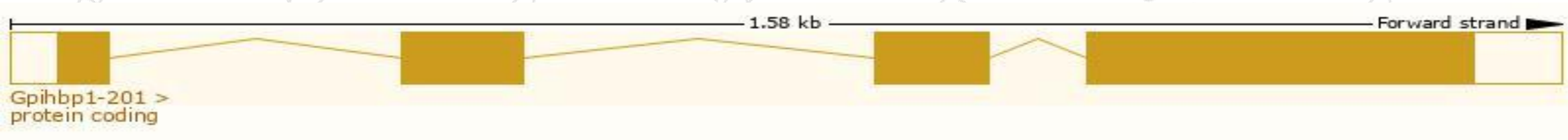
Annotation release	Status	Assembly	Chr	Location
108	current	GRCm38.p6 (GCF_000001635.26)	15	NC_000081.6 (75596630..75598213)
Build 37.2	previous assembly	MGSCv37 (GCF_000001635.18)	15	NC_000081.5 (75427088..75428643)

Transcript information (Ensembl)

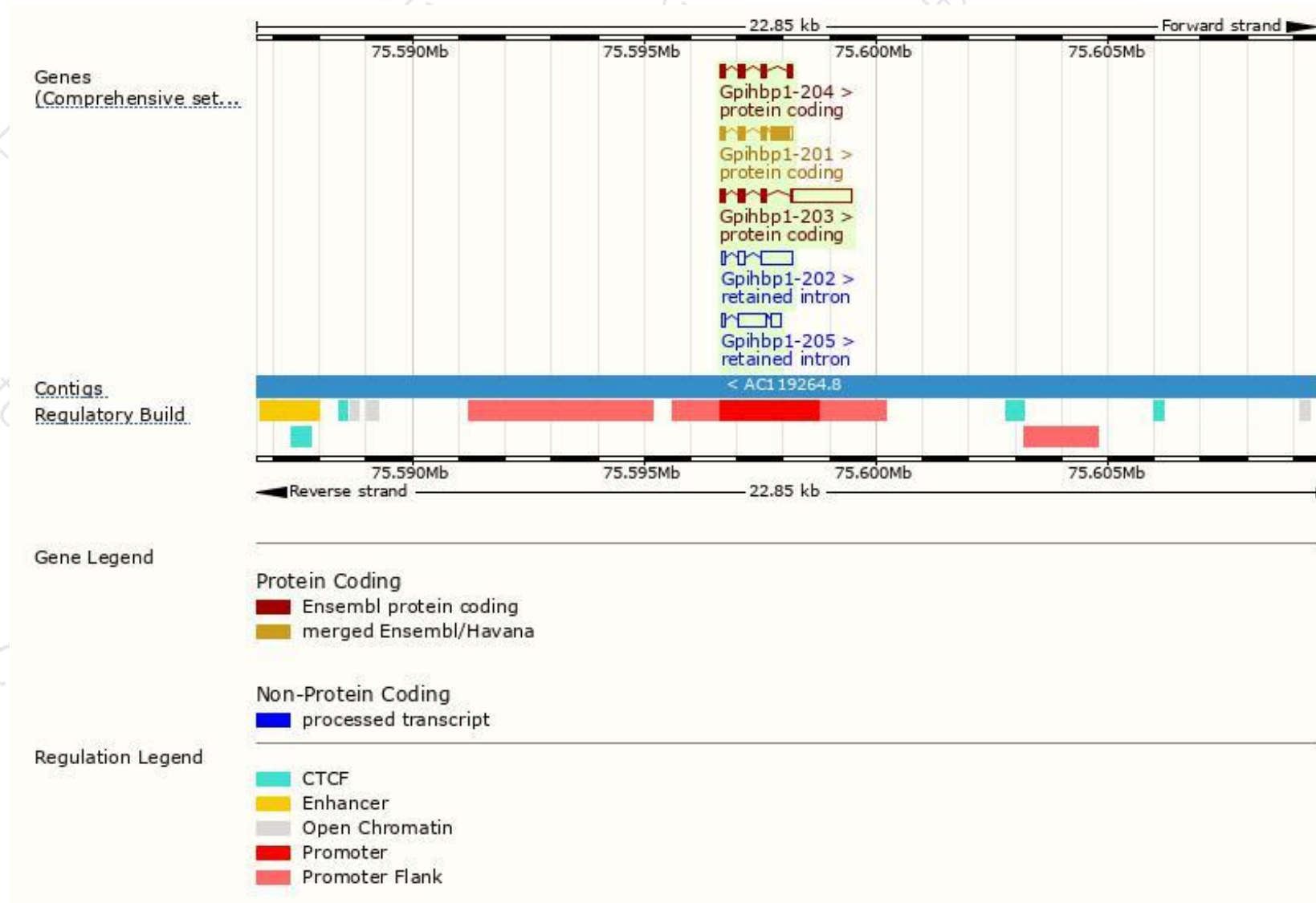
The gene has 5 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Gpihbp1-201	ENSMUST00000023243.10	825	228aa	Protein coding	CCDS27545	Q9D1N2	TSL:1 GENCODE basic APPRIS P1
Gpihbp1-204	ENSMUST00000189944.6	466	123aa	Protein coding	CCDS84173	A0A087WNV7	TSL:2 GENCODE basic
Gpihbp1-203	ENSMUST00000189874.1	1652	111aa	Protein coding	-	A0A087WPI6	TSL:1 GENCODE basic
Gpihbp1-205	ENSMUST00000190119.6	877	No protein	Retained intron	-	-	TSL:5
Gpihbp1-202	ENSMUST00000186934.1	862	No protein	Retained intron	-	-	TSL:2

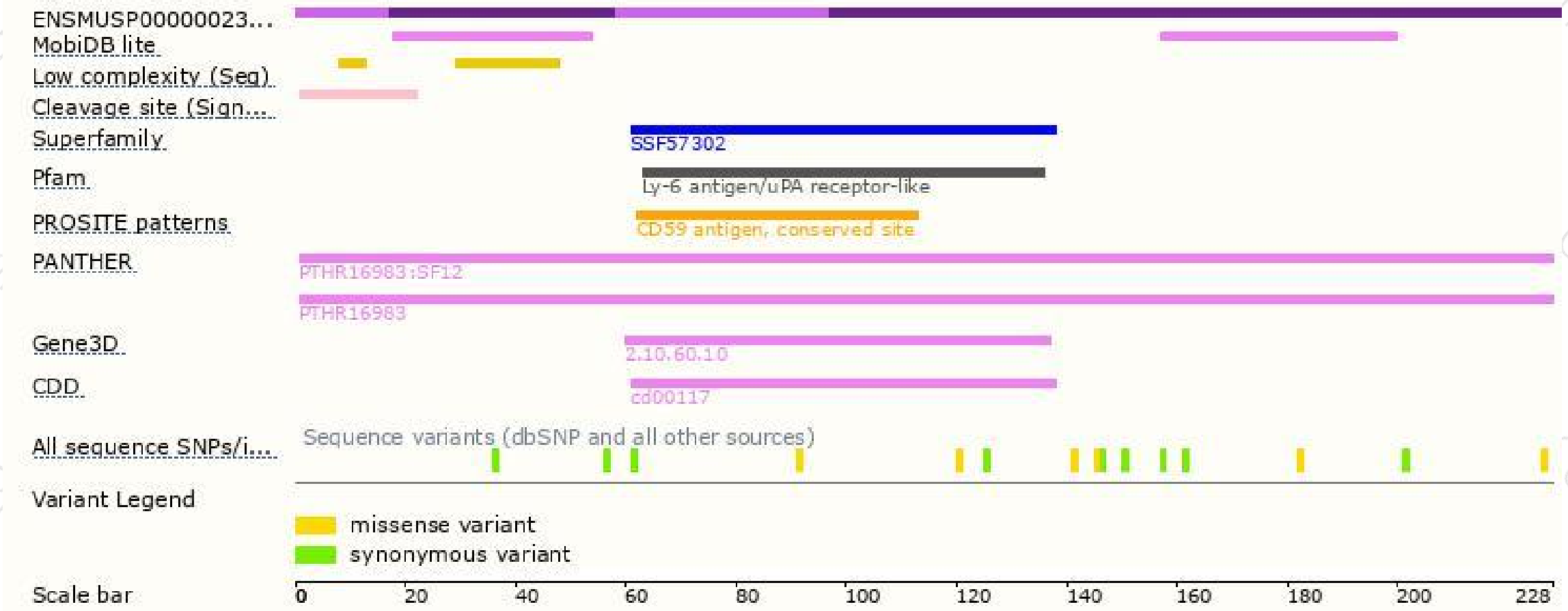
The strategy is based on the design of *Gpihbp1-201* transcript,The transcription is shown below



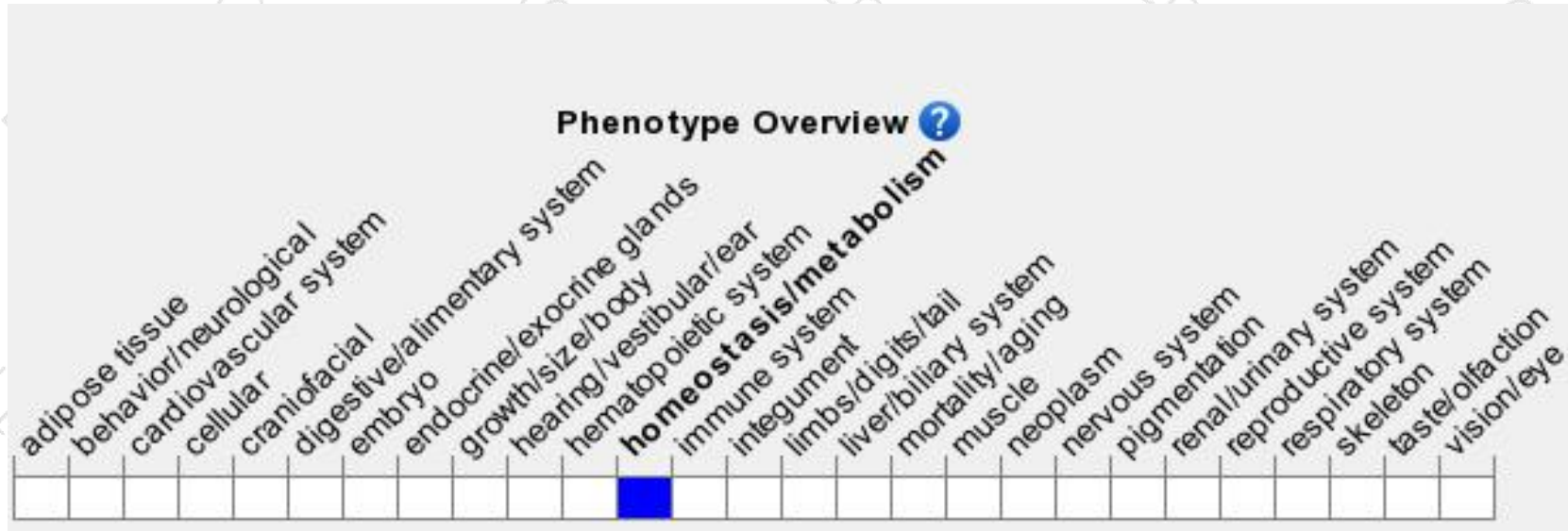
Genomic location distribution



Protein domain



Mouse phenotype description(MGI)



Phenotypes affected by the gene are marked in blue. Data quoted from MGI database(<http://www.informatics.jax.org/>).

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If you have any questions, you are welcome to inquire.

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