

# *Alox5* Cas9-KO Strategy

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**Reviewer:**

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# Project Overview

**Project Name**

*Alox5*

**Project type**

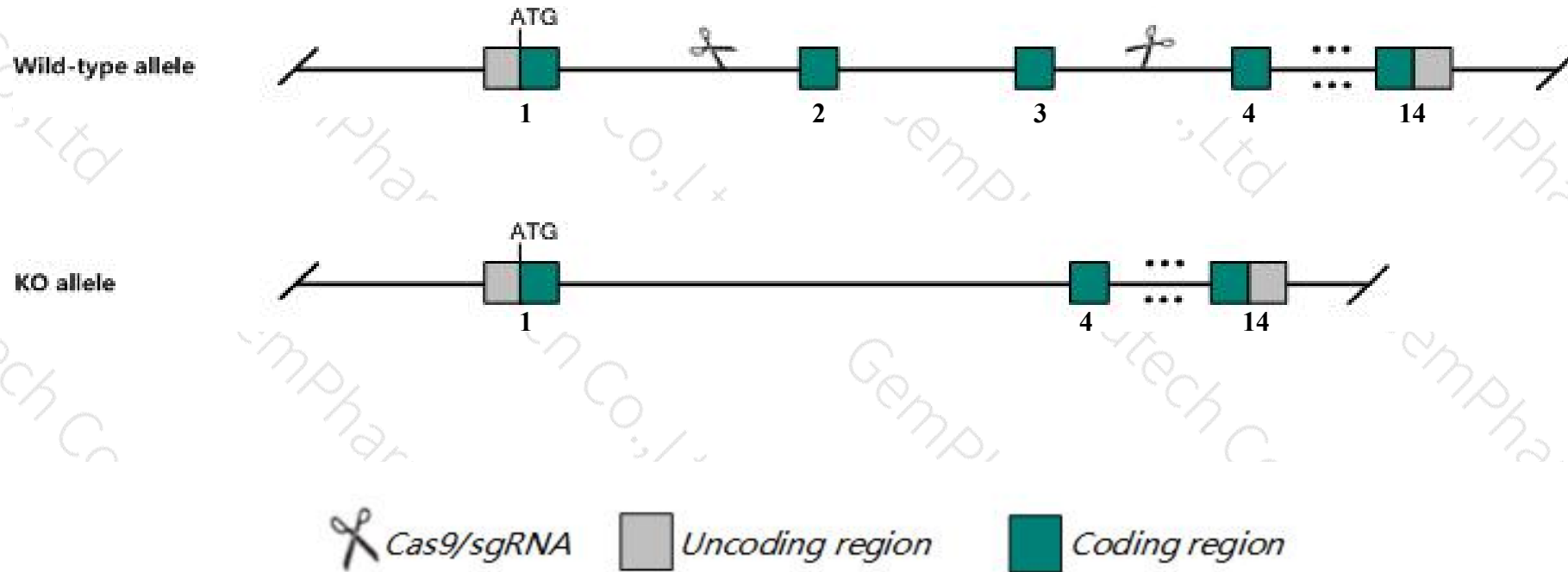
**Cas9-KO**

**Strain background**

**C57BL/6J**

# Knockout strategy

This model will use CRISPR/Cas9 technology to edit the *Alox5* gene. The schematic diagram is as follows:



- The *Alox5* gene has 8 transcripts. According to the structure of *Alox5* gene, exon2-exon3 of *Alox5-201* (ENSMUST00000026795.12) transcript is recommended as the knockout region. The region contains 281bp coding sequence. Knock out the region will result in disruption of protein function.
- In this project we use CRISPR/Cas9 technology to modify *Alox5* gene. The brief process is as follows: sgRNA was transcribed in vitro. Cas9 and sgRNA were microinjected into the fertilized eggs of C57BL/6J mice. Fertilized eggs were transplanted to obtain positive F0 mice which were confirmed by PCR and sequencing. A stable F1 generation mouse model was obtained by mating positive F0 generation mice with C57BL/6J mice.

- According to the existing MGI data, nullizygous mice show altered inflammatory responses. One null mutation causes resistance to lethal anaphylaxis, abnormal eicosanoid production and neutrophil recruitment while another leads to increased body fat, bone density, leptin and VLDL cholesterol levels and resistance to autoimmune uveitis.
- The *Alox5* gene is located on the Chr6. 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)

## Alox5 arachidonate 5-lipoxygenase [ *Mus musculus* (house mouse) ]

Gene ID: 11689, updated on 4-Dec-2019

### Summary

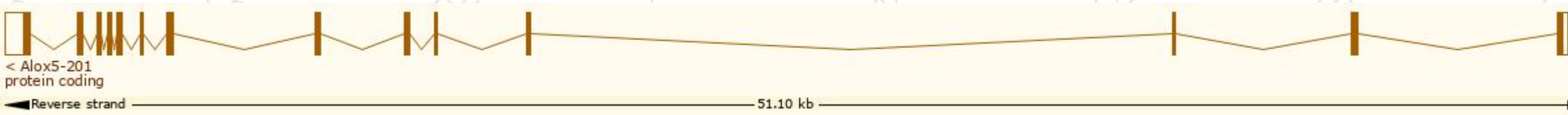
Official Symbol	Alox5 provided by <a href="#">MGI</a>
Official Full Name	arachidonate 5-lipoxygenase provided by <a href="#">MGI</a>
Primary source	<a href="#">MGI:MGI:87999</a>
See related	<a href="#">Ensembl:ENSMUSG000000025701</a>
Gene type	protein coding
RefSeq status	VALIDATED
Organism	<a href="#">Mus musculus</a>
Lineage	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Euarchontoglires; Glires; Rodentia; Myomorpha; Muroidea; Muridae; Murinae; Mus; Mus
Also known as	5LO; 5LX; 5-LO; 5-LOX; AI850497; F730011J02
Expression	Broad expression in lung adult (RPKM 3.4), heart adult (RPKM 2.7) and 22 other tissues <a href="#">See more</a>
Orthologs	<a href="#">human</a> <a href="#">all</a>

# Transcript information (Ensembl)

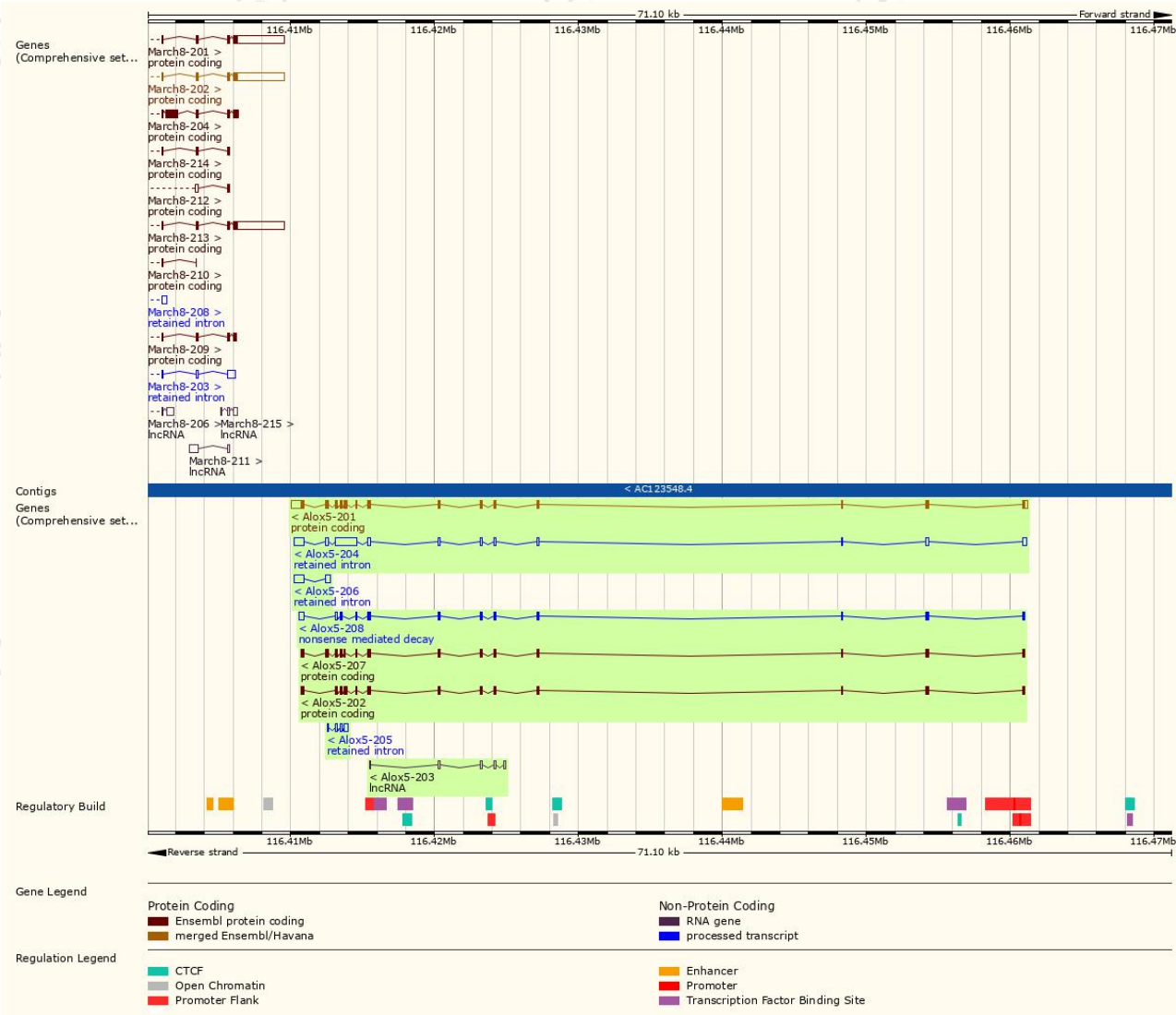
The gene has 8 transcripts,all transcripts are shown below:

Name	Transcript ID	bp	Protein	Biotype	CCDS	UniProt	Flags
Alox5-201	<a href="#">ENSMUST00000026795.12</a>	2821	<a href="#">674aa</a>	Protein coding	<a href="#">CCDS20452</a>	<a href="#">P48999</a>	TSL:1 GENCODE basic APPRIS P1
Alox5-207	<a href="#">ENSMUST00000170186.1</a>	1929	<a href="#">642aa</a>	Protein coding	-	<a href="#">E9Q6H6</a>	TSL:5 GENCODE basic
Alox5-202	<a href="#">ENSMUST00000164547.7</a>	1854	<a href="#">617aa</a>	Protein coding	-	<a href="#">E9QA93</a>	TSL:5 GENCODE basic
Alox5-208	<a href="#">ENSMUST00000203722.2</a>	1787	<a href="#">434aa</a>	Nonsense mediated decay	-	<a href="#">A0A0N4SW45</a>	TSL:5
Alox5-204	<a href="#">ENSMUST00000167447.7</a>	3576	No protein	Retained intron	-	-	TSL:2
Alox5-206	<a href="#">ENSMUST00000169625.1</a>	956	No protein	Retained intron	-	-	TSL:1
Alox5-205	<a href="#">ENSMUST00000167585.1</a>	546	No protein	Retained intron	-	-	TSL:2
Alox5-203	<a href="#">ENSMUST00000167174.1</a>	659	No protein	lncRNA	-	-	TSL:3

The strategy is based on the design of *Alox5-201* transcript, the transcription is shown below:

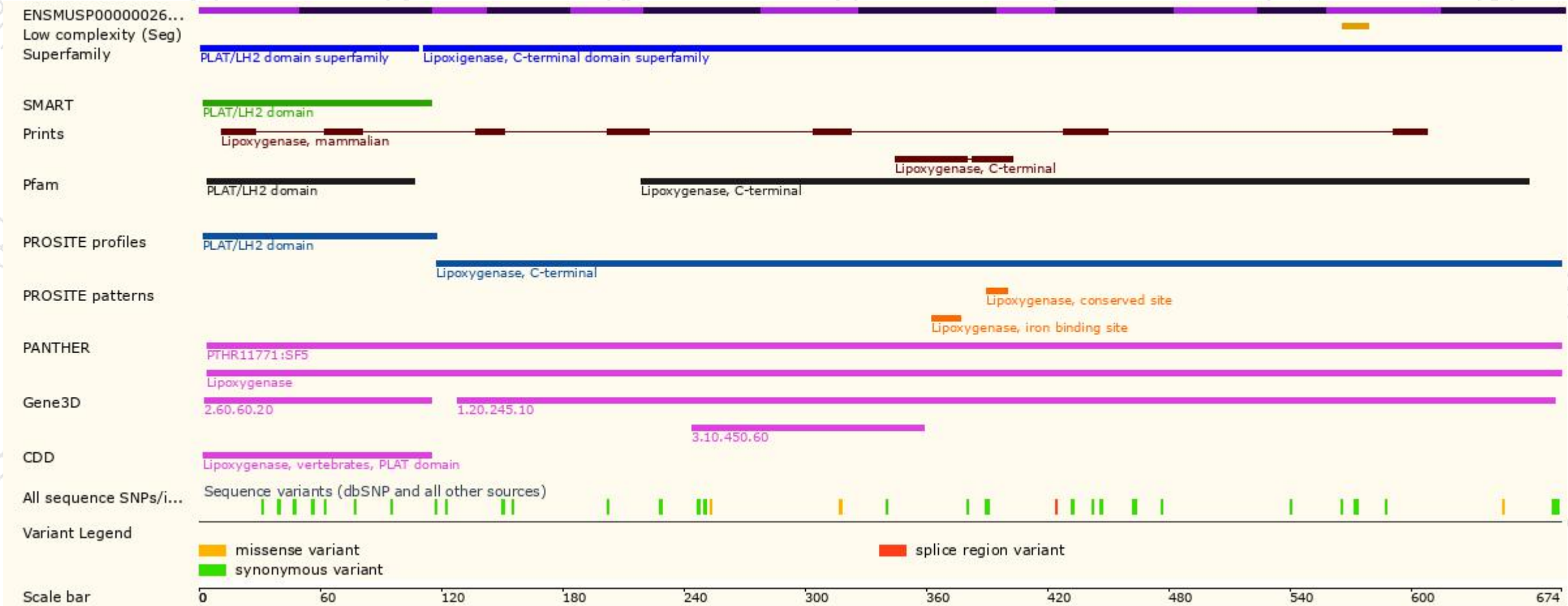


# Genomic location distribution

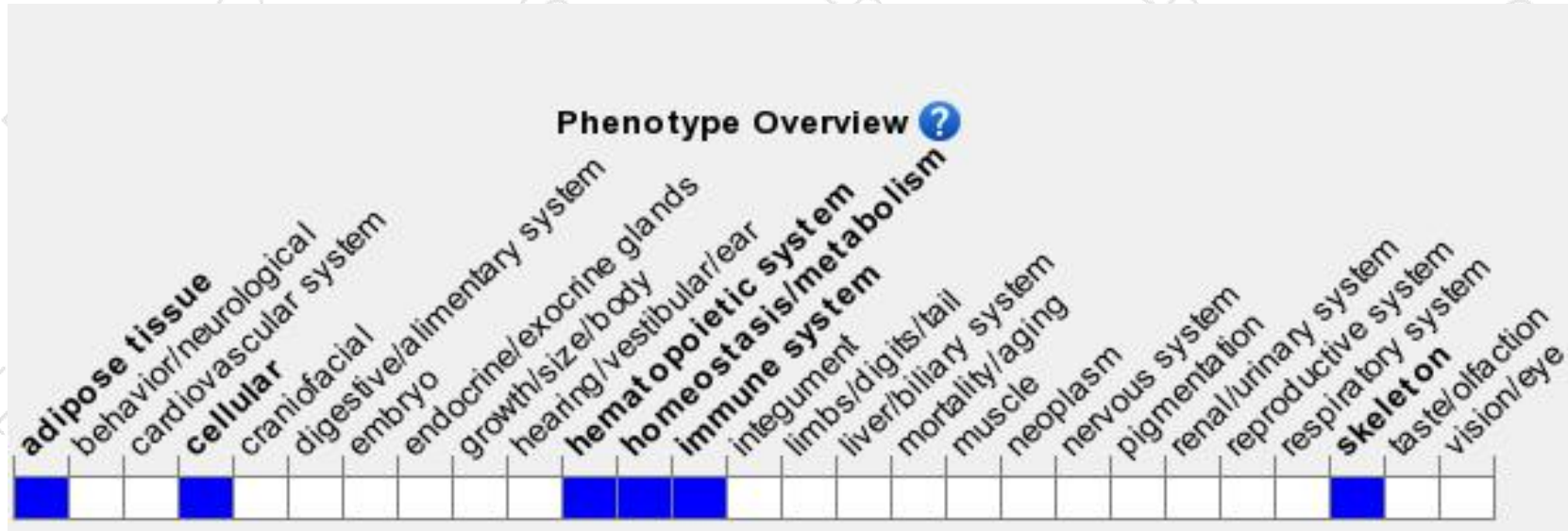




# Protein domain



# Mouse phenotype description(MGI )



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

According to the existing MGI data, nullizygous mice show altered inflammatory responses. One null mutation causes resistance to lethal anaphylaxis, abnormal eicosanoid production and neutrophil recruitment while another leads to increased body fat, bone density, leptin and VLDL cholesterol levels and resistance to autoimmune uveitis.

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

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