Byeonghwi Lim¹, Chiwoong Lim¹, Seung-Hoon Lee¹, Keon Bong Oh^{2*} and Jun-Mo Kim^{1*}

Functional Genomics & Bioinformatics Lab. Department of Animal Science and Technology Chung-Ang University

JUN-MO KIM



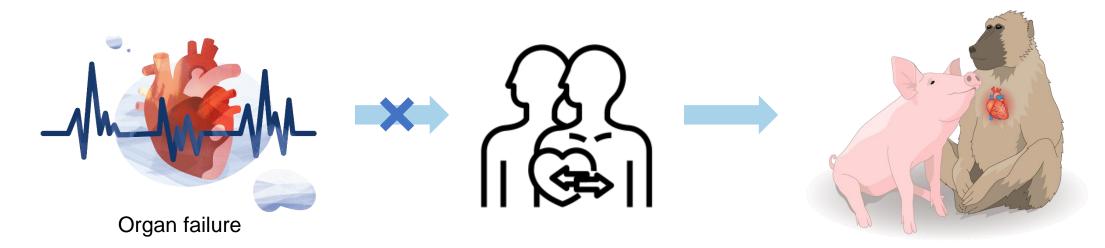


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Xenotransplantation

- Procedure that involves the transplantation, implantation or infusion into a recipient of either live cells, tissues, or organs from other animal source
- Genetically engineered pig hearts are a potential alternative for heart failure due to limited human donors



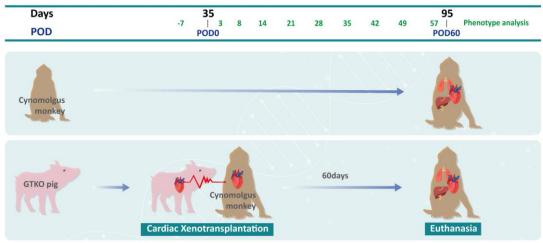
To elucidate the transcriptomic mechanisms occurring in other tissues subsequent to pig-tomonkey xenotransplantation.



Byeonghwi Lim¹, Chiwoong Lim¹, Seung-Hoon Lee¹, Keon Bong Oh^{2*} and Jun-Mo Kim^{1*}

¹Department of Animal Science and Technology, Chung-Ang University, Anseong, Gyeonggi-do 17546, Republic of Korea ²Animal Biotechnology Division, National Institute of Animal Science, RDA, Wanju, Jeollabuk-do 55365, Republic of Korea

1. Animal and experimental design



(Hetero heart xenotransplantation)

2. RNA-seq

- Library & sequencing

- Wile heart (Not xenotransplanted heart)
- RNA extraction from 24 samples (WH [3], WA [3], WLi [3], WLu [3], XTH [3], XTA [3], XTLi [3], XTLu [3])
- cDNA libraries were constructed with the TruSeq Stranded Total RNA LT Sample Prep Kit
- 150 bp paired-end sequencing by Illumina HiSeq X Ten

- Raw data processing
 - Quality check (FastQC-0.11.8)
 - Trimming (Trimmomatic-0.39)
 - Mapping / Sorting (HISAT2-2.1.0 / SAMtools-1.9)
 - Counting (featureCounts (Subread-1.6.3))
- Annotation
 - Macaca fascicularis 5.0.101

3. Differentially expressed gene (DEG) analysis

- Normalization: The gene expression level relative abundances were obtained as the trimming means of M values (TMM) (edgeR-3.22.5)
- Cutoff: absolute log_2 fold change \geq 1 and FDR < 0.05

4. Functional annotation

- Gene Ontology (GO) database
- Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway database



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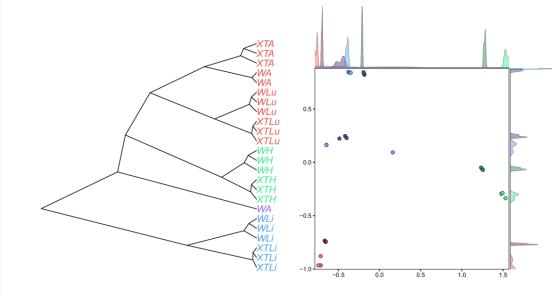


Fig 1. Cluster and multi-dimensional scaling (MDS) plot.

Liver and lung had relatively higher changes after heart xenotransplantation

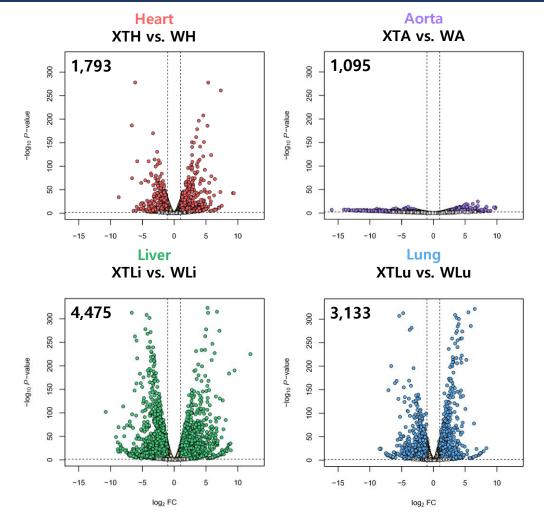


Fig 2. Volcano plots. The number of DEGs are indicated top left corner.

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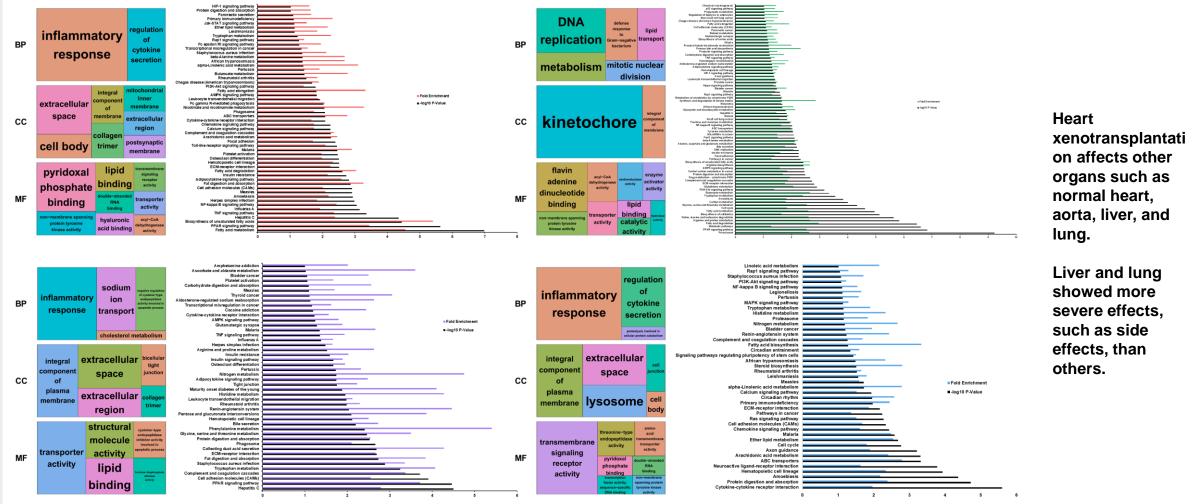


Fig 3. Functional annotation in heart (red), aorta (purple), liver (green), and lung (blue) samples

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