

# Analysis of mRNA data of patients with atherosclerosis<sup>1</sup>

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## Abstract

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### 2 Weighted Gene Co-Expression Analysis

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#### 2.3 Interconnectivity analysis

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#### 3.2 Logistic regression

### 4 Experiments

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#### 4.2 Results

### 5 Conclusion

### 6 Future Work

## References

- [1] Lingxue Zhang, Seyoung Kim, "Learning Gene Networks under SNP Perturbations Using eQTL Datasets"  
<http://dx.doi.org/10.1371%2Fjournal.pcbi.1003420>

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