Machine Learning-based Genome Wide

Association Studies of Rheumatoid Arthritis

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Introduction

Rheumatoid arthritis (RA) is an autoimmune disease affecting the membrane between the

joints causing pain, joint damage, and eventually severe disability. Although RA has been

identified as a multifactorial disease (meaning that there is more than one factor causing

it)[1], this paper will specifically focus on the genetic components.

To look for these genetic components, researchers perform what is known as a genome-

wide association study, or GWAS, which the National Human Genome Research Institute

defines as "an approach that involves rapidly scanning markers across the complete sets of

DNA, or genomes, of many people to find genetic variations associated with a particular

disease." The results of these studies, however, have been generally inconclusive collec-

tively. For example, Takahashi et. al.'s genome-wide association study (GWAS) studying

approximately 1,400 Japanese females shared data on 87 of the top 100 single nucleotide

polymorphisms (SNPs) found in Sharma et. al.'s [2] racially diverse GWAS of 419 fami-

lies, however, only one of those SNPs showed significant association in Takahashi et. al.'s

GWAS [3]. The goal of my project is to study the usage and efficacy of machine learning

for GWAS of scoliosis.

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Background

Related Work

Although there have not been any studies done to date using machine learning for GWAS of idiopathic scoliosis, there have been many studies using machine learning for other phenotypes including IgM and rheumatoid arthritis, as mentioned above, in addition to myocardial infarction, coronary artery calcification, and anti-cyclic citrullinated peptide[4]. These studies will provide the basis for my methodology, specifically D'Angelo et. al.'s [5] and Tang et. al.'s [6] GWASs of rheumatoid arthritis and Stassen et. al.'s GWASs of IgM. Since there are no prior machine learning-based GWASs of AIS, I plan replicate their respective methodologies as best as possible, adapting where necessary for the specifics of scoliosis and the data sets I use.

References

[1] Y. Alamanos and A. A. Drosos, "Epidemiology of adult rheumatoid arthritis," *Autoimmunity Reviews*, vol. 4, no. 3, pp. 130–136, 2005.