PREDICTION OF BREAST CANCER RELAPSE TIME IN CONTINUOUS SCALE BASED ON TYPE-2 TSK FUZZY MODEL

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By

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Chairman : Associate Professor Mohammad Hamiruce Marhaban, PhD
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Recently, microarray analysis and gene expression profiles have been widely applied in diagnosis and classification of different types of cancer such as liver, colon or breast cancer. As the number of breast cancer cases increased dramatically in many countries including Malaysia in recent decades, different types of studies have been done to control the disease or reduce the cost of their treatments. Gene expression profiles, which can screen the behavior of a large number of genes simultaneously, have been used in some studies to extract the significant genes related to breast cancer. Tumor classification, Estrogen Receptor status recognition or survival analysis has been usually considered as important objectives in these studies. Due to the fact that studies in survival analysis of breast cancer can reduce the cost of treatments and side effects of the adjuvant therapy,
different methods for predicting the outcome of the disease have been proposed by previous researcher.

The two major objectives of this research are to propose a fuzzy classifier to discriminate breast cancer tumors into two classes, which are high risk and low risk by some interpretable rules similar to linguistic words, and to predict the relapse time of breast cancer by TSK fuzzy models in continuous scale. For this reason, breast cancer dataset has been applied for training the models and two other independent samples have been used for validating the results. In addition, K-fold Cross Validation, B632 and B632+ methods have been used for error estimation.

In the first objective of the thesis, a lemma has been proven and a new hybrid algorithm based on Fuzzy Association Rule Mining has been proposed to gather some selected genes and generate fuzzy rules for classification.

In the second one, a method for generating the fuzzy rules to discriminate the samples of breast cancer into the different groups have been proposed and applied to predict the relapse time of samples in continuous scale while handling the uncertainties in linguistic terms of the rules.
The relapse time of two available independent samples of breast cancer have been predicted by the model and the results show the superiority of the proposed model with respect to the previous study. Finally 46 significant genes and 16 fuzzy rules have been introduced which can be used in a Type-2 TSK fuzzy model as a predictor.
ABSTRAK

Abstrak tesis dikemukakan kepada Senat Universiti Putra Malaysia sebagai memenuhi keperluan untuk ijazah Doktor Falsafah

RAMALAN MASA RELAPS DALAM SKALA SELANJAR BAGI KANSER PAYUDARA BERDASARKAN MODEL KABUR TSK JENIS-2

Oleh

SAYED HAMID MAHMOUDIAN

November 2010

Pengerusi: Profesor Madya Mohammad Hamiruce Marhaban, PhD
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Kebelakangan ini, analisa jujukan mikro dan profil ekspresi gen telah digunakan dengan meluas dalam diagnosis dan pengelasan beberapa jenis kanser seperti kanser hati, kolon atau payudara. Disebabkan kes kanser payudara telah meningkat bilangannya dalam dekad ini di kebanyakan negara termasuk Malaysia, pelbagai kajian telah dijalankan untuk mengawal penyakit tersebut atau mengurangkan kos rawatannya. Profil ekspresi gen berupaya untuk mencerminkan tingkah laku gen berjumlah besar dengan serentak, telah digunakan dalam beberapa kajian mengekstrak gen-gen penting yang berkaitan dengan kanser payudara. Pengelasan tumor, pengecaman status penerima estrogen atau analisa peluang hidup biasanya dianggap sebagai objektif utama untuk kajian-kajian ini. Memandangkan kajian tentang peluang hidup daripada kanser payudara boleh
mengurangkan kos rawatan dan kesan sampingan daripada terapi adjuvan, pelbagai kaedah untuk meramal hasil penyakit tersebut telah dicadangkan oleh para penyelidik.

Dua objektif utama kajian ini ialah untuk mencadangkan satu pengelas kabur dalam membezakan tumor kanser payudara kepada dua kelas iaitu kelas berisiko tinggi dan kelas berisiko rendah, mengikut peraturan yang boleh diinterpretasi, menggunakan kata-kata linguistik serta meramalkan masa relaps bagi kanser payudara menggunakan model kabur TSK dalam skala selanjur.

Untuk itu, set data kanser payudara van’t Veer telah digunakan dalam melatih model-model yang dicadangkan serta 2 lagi sampel bebas yang diterbitkan oleh van’t Veer dan van de Vijver telah digunakan untuk mengesahkan keputusan yang diperolehi. Kaedah keesahan silang, B632 dan B632+ telah digunakan untuk menganggarkan ralat.

Bagi objektif pertama, satu lemma telah dibuktikan dan satu algoritma hibrid berdasarkan Perlombongan Peraturan Berkaitan Kabur telah dicadangkan untuk mengumpul beberapaa gen yang terpilih dan menjana peraturan kabur untuk pengelasan.
Dalam objektif kedua, satu kaedah untuk menjana peraturan kabur dalam membezakan sampel-sampel kanser payudara kepada kumpulan-kumpulan yang berbeza telah dicadangkan dan diaplikasikan untuk meramal masa relaps sampel-sampel tersebut dalam skala selanjar serta mengendali ketidakpastian dalam terma linguistik peraturan-peraturan.

Masa relaps bagi 2 sampel kanser payudara yang bebas telah diramalkan oleh model tersebut dan hasil menunjukkan model yang dicadangkan jauh lebih baik daripada model-model dari kajian sebelum ini. Subset gen-gen yang penting dan peraturan kabur yang dijana untuk ramalan masa relaps juga dikemukakan. Akhirnya, 46 gen penting dan 16 aturan kabur telah dicadangkan yang mana ia boleh digunakan oleh model kabur TSK Jenis-2 sebagai peramal.
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I certify that a Thesis Examination Committee has met on (July 2010) to conduct the final examination of Hamid Mahmoodian on his thesis entitled “Prediction of Breast Cancer Relapse Time in Continuous Scale Based on Type-2 Fuzzy Model” in accordance with the Universities and University Colleges Act 1971 and the Constitution of the Universiti Putra Malaysia [P.U.(A) 106] 15 March 1998. The Committee recommends that the student be awarded the Doctor of Philosophy.

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DECLARATION

I declare that the thesis is my original work except for quotations and citations which have been duly acknowledged. I also declare that it has not been previously, and is not concurrently, submitted for any other degree at Universiti Putra Malaysia or at any other institutions.

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Date: 29 November 2010
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