

Forensic Age Estimation Model Using Left Hand Bones Based On GA-ANN Hybrid Approach For Hispanic Population

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ABSTRAK

Model anggaran umur ialah bantuan berfungsi untuk membezakan identiti si mati atau individu hidup dari segi antropologi forensik. Model anggaran umur konservatif hanya terhad untuk penduduk spesifik serta variabiliti tinggi. Kajian ini memberi tumpuan dalam membina model anggaran umur menggunakan pendekatan hybrid GA-ANN (Genetic Algorithm-Artificial Neural Network) bergantung pada 19 panjang tulang tangan kiri individu yang berumur dari 0 hingga 18 tahun. Gambar X-ray tangan kiri dikumpul dari dataset penduduk Hispanik. 19 panjang tulang diukur menggunakan Photo Pos Pro dan pendekatan hybrid dilaksanakan mengguna Python skrip Python dan Matlab. GA adalah untuk memilih tulang paling sesuai manakala ANN menganggarkan usia. Kajian ini dikira berjaya sekira keputusan pendekatan hybrid dicadang dapat mengatasi penanda aras.

ABSTRACT

Age estimation model serves as an aid to distinguish the identities of deceased or alive individuals in term of forensic anthropology. The existing age estimation models are population specific and high variability. This research focuses on using Genetic Algorithm-Artificial Neural Network (GA-ANN) hybrid approach on age estimation models depend on the lengths of 19 left hand bones of alive individuals who ages from 0 to 18 years old. X-ray images of left hands are collected from Image Processing and Informatics Lab of University of Southern California. The lengths of the 19 bones are assessed using Photo Pos Pro while the hybrid approach is implemented using Python scripts and Matlab. GA approach is applied to select the appropriate bones while ANN approach is used to estimate age. It is believed that the research is successful as the results of GA-ANN hybrid approach outperform the ANN approach.

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CHAPTER 1

INTRODUCTION

1.1 INTRODUCTION

There are many unidentified decedent cases in world and we may need some models or approaches to predict features or physical characteristics of unknown corpses. Lynnerup & Villa (2014) stated that evaluation of age at death is involved in the identification of human remains.

Age estimation is essential since it is possible that the examiner or investigator is requested to make an educated guess and hypothesize the age of an individual as a part of the investigation. Stavrianos, Mastagas, Stavrianou, & Karaiskou (2008) stated that age is the significant component in clinical practice, research, and court of law.

Age estimation serves in minimizing the scope of the probe feasibility for unknown deceased or living individuals for constitutional reasons. For people who thought to be guilty of crime with unrevealed chronological age, who are committed to a grave crime, the precisions of age estimation models are required for absolute justice (Duangto, Janhom, Prasitwattanaseree, Mahakkanukrauh, & Iamaroon, 2017). An entry of large number of asylum seekers, unlawful immigrants, and also more young perpetrators being subjected to trials in court have caused the grow desire for age estimation (Sykes, Bhayat, & Bernitz, 2017).

Age estimation in forensic medicine has progressively obtained significance since there are inevitable situations like individuals do not have or having falsified identification documents (Auf der Mauer et al., 2019). There is some current published literature with a particular attention allocate to age estimation to be used for refugees

and individuals involved in human trafficking (Franklin, Flavel, Noble, Swift, & Karkhanis, 2015).

There is a lack of studies on age estimation model for Hispanic population. It is believed the field of forensic science prefers more age estimation models for different ethnic groups. Therefore, this research focuses on Hispanic population.

1.2 PROBLEM STATEMENT

The problems that this paper aims to tackle are as follows:

- i. There are only few age estimation models for Hispanic population.
- ii. There are difficulties in getting research studies that applying GA-ANN hybrid approach on age estimation model.
- iii. The field of forensic science demands more age estimation models for different ethnic groups.

1.3 OBJECTIVES

The goal of this project is to apply GA-ANN hybrid approach on age estimation model using left hand bones for Hispanic population with the following objectives:

- i. To investigate the adoption of GA approach for bone selection
- ii. To hybridize the GA approach with ANN approach for age estimation
- iii. To evaluate the GA-ANN hybrid approach with ANN approach

1.4 SCOPE

This research focuses on using GA-ANN hybrid approach on age estimation models depend on the lengths of 19 left hand bones of Hispanic population who ages from 0 to 18 years old. The lengths of the 19 bones are assessed using Photo Pos Pro while the hybrid approach is implemented using Python scripts and Matlab.

1.5 REPORT ORGANISATION

This thesis consists of five chapters. Chapter 1 will discuss on introduction of this thesis while chapter 2 will discuss on the literature review for this research paper. Chapter 3 is about methodology while chapter 4 is the implementation and discussion. Chapter 5 is the conclusion of this research.

CHAPTER 2

LITERATURE REVIEW

2.1 INTRODUCTION

Age estimation is pivotal in law enforcement and legal justice systems all over the world (Asif et al., 2019). Other than that, to be able to precisely determine the age of children and young adults has always been a problem in forensics since there are cases where youths without proper identification documents are requested to be assessed for their criminal responsibility, commonly regarding the range of their possible age, and also to determine if a residency permit can be given, and refugees that may look too old for their claimed age. (Bartolini et al., 2018).

Stern & Ursekter (2016) stated that it is critical to the fields of clinics and forensics that the evaluation of the age of individuals is accurate and precise, as medicine released from clinics places emphasis on the predicting the age to determine endocrinological diseases in children or in pediatric surgery while medicine in forensics emphasizes on age estimation under the condition where no distinct clues can be found to help in estimating the age, such as the age of minor refugees that has lost their identification documents.

In forensics, the search for victims can be simplified by creating a range for age through age prediction. The search for possible contributors of crime scene traces can also be made simpler by predicting the age range for possible suspects. Prediction intervals are always preferred when approximating age in a police investigation. (Smeers, Decorte, Van de Voorde, & Bekaert, 2018).

Other than that, it is essential to have age estimation of unspecified human bodies at the crime scene since the mortality age, birth date and other details allow investigators to get the true identity of a human body by providing potential counterparts. The determination of human bodies is always a difficulty to investigators when there is no sign about the identity from particular data (Alkass et al., 2009). The age of an individual is always an elemental piece of data relating to forensic identification of unknown human bodies (Lynnerup & Villa, 2014).

2.2 THE IMPORTANCE OF AGE ESTIMATION

There are several factors that make age estimation an important technique.

2.2.1 CRIMINAL RESPONSIBILITY

In much of the first-world nations, in example New Zealand and Australia, 10 years old is the standard age of criminal responsibility. Meanwhile it is 7 to 8 years old in India, Indonesia and Jordan. A formal statement made by the United Nations (UN) that children under the legal age of criminal responsibility must not be subjected to criminal accusations and nations should start to revise themselves if they have set criminal responsibilities for children under the age of 12 years and below. *Doli incapax* assumes that a child could not hope to commit a crime under common law. This is a presumption that is employed to children between the age of 13 or 14 years, depending on jurisdiction, and is a more controversial concept (Franklin et al., 2015).

17 to 21 years would be the standard range of age for young people which age between *doli incapax* and adulthood, depending on jurisdiction are ideally subjected to rehabilitation than punishment, as the full criminal liability of an adult is not carried by them. However, they must still face the consequence of their actions, albeit the consequences are lighter. (Franklin et al., 2015).

2.2.2 CHILD PORNOGRAPHY

Any activities relating to child pornographic files, especially having ownership to such files is a crime in several countries. Therefore, there are frequent requests for forensics analysis in computer storage devices to verify the presence of child pornography (Polastro & Eleuterio, 2012).

Macedo, Costa, & A. dos Santos (2019) proposed a combined method to detect child pornography using a child face detection module and a pornography detector. The results showed the viability of the strategy of combining different detectors and classifiers to perform detection of child pornography, but it requires much more improvement, especially in age estimation and pornography classification.

2.2.3 FALSIFICATION OF AGE

Falsification of age or also known as age cheating is a common practice in domain of sports and public service, and some members of society may have a different moral perspective and find this completely acceptable as they view it a vital underhandedness, or a means of survival in a situation of destitution (Jerome Tosam, 2015). It is important to have age estimation in the sports academics and military services at the entry and also retirement (Bakthula & Agarwal, 2014).

It is possible for some children in certain countries, while being born to this world, do not have their births registered and therefore a birth record is not made. There happened the alterations of births on official documents of players in hopes to obtain the right to compete with other countries in sports industry. International Olympic Committee has come out with a consensus statement since the matter of age verification in youth sports is a great concern (Malina, 2011).

To prevent issues of falsification of age and to protect children from recruitment in hostilities, General Assembly launched the Optional Protocol to the Convention on the Right of the Child on the involvement of children in armed conflict (OPAC). Office of the Special Representative of the Secretary-General for Children and Armed Conflict adopted the campaign Zero under 18 in order to support OPAC (UN, n.d.).

2.2.4 HUMAN TRAFFICKING

Kiss, Yun, Pocock, & Zimmerman (2015) stated that every year, many children fall victim to the grave crime of human trafficking. 5.7 million boys and girls have been estimated to work without their, 1.2 million are sold illegally, and about 1.8 million have been forced to sexual slavery.

Child labour robs children away from their childhood, potential and dignity, and can also cause harm to mental and physical development. Worst forms of child labour are trafficking, forced labour, sexual exploitation and use of underage children for security and military purposes (UNICEF, The Global Compact, & Save the Children, 2012).

Children with low socioeconomic contexts are unprotected to human trafficking and once trafficked, these children are frequently involved in criminal acts and therefore being inspected by local legal system with no or falsified identity documents (Franklin et al., 2015).

2.2.5 REFUGEE AND ASYLUM SEEKERS

According to United Nations High Commissioner for Refugees (UNHCR), almost half of victims of natural disasters, internal conflict, and war are vulnerable children and are now seeking international protection. The majority of UN members

supports and signed The Convention on the Rights of the Child, which declares without discrimination, the rights of every child, defined as individuals under 18 years old, within any States Parties' jurisdiction will be respected and ensured (OHCHR, n.d.).

Sykes, Bhayat, & Bernitz (2017) states the asylum seeker crisis has contributed to the necessity of having age prediction, causing an increase in such cases, which is most likely caused by a sudden increase in trials involving refugees, illegal immigrants and juvenile offenders. This makes the legal authorities the main users who need age estimation tests in support of their investigation.

2.2.6 UNKNOWN DECEASED CHILDREN AND MISSING CHILDREN

Age estimation is the most precise biological identifier that a forensic anthropologist can give in the case of an unknown deceased child (Lewis & Flavel, n.d.). Lacking of documents, clothing or other personal belongings are often main issues of identification in historical grave findings. Thus, the focus will be on age estimation, so as sex and stature determination. Generally, the principal criteria to estimate age of deceased children and teenagers are the appearance of ossification centre, epiphyseal union, the development of teeth, and also bones lengths. Age estimation is more reliable to children and teenagers than adults as children and teenagers are based on skeletal and dental maturation while adults focus on degenerative changes (Huomonen et al., 2016).

Nowadays, missing children becomes a common phenomenon in world. Society today has becoming progressively worse for citizens especially children. Missing children is complicated occurrence and having many dimensions for it to deal with. There is two possible ways that a child can go missing, which are either the child has been kidnapped by someone against the assent of the child and also law, or, the child leaves the domination and protection of guardians without acknowledgment (Vij, 2016).

A crucial process that can bring closure in a tragic case is the cross-matching reports of missing people with unknown bodies. A successful matching helps criminal investigations and completes the civil procedures, and thus those who left behind can mourn their beloved one (People & Paper, 2011). Alkass et al. (2009) stated that age estimation is offering significant potentiality to police authorities and forensics pathologist in determining the time of death as well as the identity of unknown bodies or individuals.

2.3 THE HUMAN PARTS USED FOR AGE ESTIMATION

Hand bone, dental, knee, facial, and pelvis are several parts of human body that have mostly been used for age estimation. Details of each part will be explained in following sections.

2.3.1 HAND BONES

Radiography analysis of the staging of skeletal development has been a common method to estimate the timing of pubertal growth, so as to know the growth velocity and the proportion of growth remaining. In most cases, ossification of bones of the hand and wrist has been used in determining skeletal development since each type of bones can have a varying availability in the areas. The pattern of skeletal growth is almost the same in every individual, just the initiation, duration, and also amount of growth varies greatly due to the sudden, mandatory growth in puberty, and therefore, it becomes a significant consideration. Genetic and environment factors has impacted the skeletal development in every individual while the timing and ossification sequence of skeletal maturity within the hand-wrist area have shown polymorphism and sexual dimorphism that restricts clinical predictive use. Generally, there are two common methods to assess hand-wrist radiograph. The first method are the comparison approaches of Greulich and Pyle and Tanner et al. Geulich and Pyle adopt an atlas as the standard of comparison while Tanner et al. approach compares individual to the radiographic standards of skeletal development of common children in the same group

of gender and age. The second approach to access hand-wrist radiograph is using particular indicators to associate skeletal development to growth curve of puberty. This method is focused on the skeletal development evaluation of an individual than the mean values (Flores-mir, Orth, Nebbe, & Ortho, n.d.).

Many researchers have proposed divergent approaches to estimate age using various parts of bones like knees, spinal cord, femur, foot bones, skull, rib, pelvis, Carpals and Epiphyses of the Ulna and Radius, and left hand bones. Normally, measurements of bone lengths, angles and shape variations are the main focuses of many forensic studies. All measurements are vary from observation to observation and scientist to scientist. Forensic research has driven into various domain of interests such as studies of dead skeleton bones, half decomposed corpses in crime investigation or skeleton development and medical diagnosis (Bakthula & Agarwal, 2014).

An accepted sequence of ossification for the carpal, metacarpal, and phalangeal bones that is extraordinarily constant and similar for both sexes of healthy children. Capitate is the first ossification center that comes in hand and wrist radiographs while sesamoid of adductor pollicis of thumb is usually the last one. The epiphyseal center of the distal radius is the first comes into existence, then following by those of the proximal phalanges, the metacarpals, the middle phalanges, the distal phalanges, and the ulna. Despite that, two main bones are to be exempted from the sequence, which are the epiphysis of the distal phalanx of thumb and of the metacarpals usually come into existence at the same time, and the epiphysis of middle phalanx of fifth finger is often the last to ossify (Gilsanz & Raitb, 2005).

2.3.2 DENTAL

The tooth, another part which can be used for age estimation is also the most rigid tissue that can be found in the physiology of humans. Physical and chemical stresses along with nutritional deficiencies can be resisted by teeth for a long time. Other than DNA profiling and fingerprint comparison, dental evaluation is vital in

forensic identification of an unknown deceased individual. Therefore, it is important to have a valid and reliable approach of dental age estimation (Asif et al., 2019).

Asif et al. (2019) investigated the relationship between chronological age and pulp/tooth volume ratio in maxillary right central incisors and maxillary canines in order to derive a regression equation for dental age estimation for Malaysian population. The study sample of 300 cone-beam computed tomography (CBCT) scanned data belonging to 179 Malays and 121 Chinese, which stored in the Oral and Maxillofacial Imaging Division, Faculty of Dentistry, which were selected based on the image acquiring parameters, age of patients and the quality. The subjects were divided into 5 age groups, ranging from 16 to 65 years old, where each age group of 10 years range in order to ensure balanced sample distribution across the 5 groups. 300 intact teeth, which are 100 maxillary left canines, 100 maxillary right canines and 100 maxillary right central incisors, all with no caries and fully developed roots were eventually identified from database. Only one tooth per subject was selected in order to avoid bias in data acquired from the same patient.

By using i-CAT Cone Beam 3D Dental Imaging System, CBCT data was acquired and were then imported to the Mimics software for the analysis of pulp-tooth volume ratio. Images were first oriented in axial, coronal and sagittal planes. After setting different grayscale threshold values for each of the investigated teeth, new masks were created for the pulp cavity and tooth. Masks were cropped in 3 planes in order to separate the tooth from surrounding structures. There were slice by slice of manual checking of masks for segmentation and separation from surrounding structures. The pulp cavity and tooth were grown in the 'Region growing phase' of the software after editing mask in multiple slice editing phase. Three dimensional models of pulp tissue and tooth were created and the software calculated the volume of the pulp cavity and tooth. A strong inverse relationship between chronological age and pulp/tooth volume ratio has been indicated by the results and showed that this method of dental age estimation is indeed gender independent (Asif et al., 2019).

2.3.3 FACIAL

Important information related to identity, gender, age and ethnicity can be conveyed by human face. These attributes are used in facial image analysis applications and age estimation from facial image has become the active research area in computer vision community. It is because of its real world applications in multimedia communication, Human Computer Interaction (HCI), security and law enforcement (Sawant & Bhurchandi, 2019). Existing facial age estimation systems are roughly divided into two key components, which are face representation and age estimator learning (Liu, Lu, Feng, & Zhou, 2017).

Ouloul, Moutakki, Afdel, & Amghar (2019) proposed a new descriptor called Local Matched Filter Binary Pattern (LMFBP) which is designed specifically for the detection and extraction of skin wrinkles and is based on exploiting both the Matched Filter and the texture operator Local Binary Pattern (LBP). The Matched Filter will handle the detection of wrinkles using template matching between approximate shape of wrinkles using template matching between the approximate shape of wrinkles and the face image patches while LBP will encode the response of the Matched Filter into pattern codes to build the histogram of skin aging feature. Hierarchical approach has adopted in the learning phase in order to consider varying aging process from one age stage to another. It has been tested on both FGnetAD, HQfaces and PAL datasets, and the results proved the efficiency of the proposed approach when compared to the state-of-the-art age estimation methods.

2.3.4 KNEE

The distal end of femur, the proximal ends of tibia and fibula, and the patella are the four bony structures of human knee. All of those bones are present from birth, excluded patella. Patella is develop with the tendons and ligaments of knee from a cartilage precursor that only begins to ossify into bone around 3 years old while other bones are developed by epiphyses where the ends of bones from cartilage growth plates

during skeletal development. Many studies have examined the age estimation of knee and or elements that make up the knee joint such as radiography (X-ray) studies and magnetic resonance imaging (MRI) studies (Maggio, 2017).

Maggio (2017) had conducted studies that showed the varying age estimation potential of the knee joint, and the elements that contained therein. Since there is no overlapping bony structures, it is easier to image the knee. The knee can be imaged at low doses of radiation for MRI, X-ray and ultrasonography. The potential of MRI for determining of legal majority has been demonstrated by many studies while there is still lack of information about the potential of ultrasonography of the knee for age determination.

Age and sex of a person might be estimated by the shape of knee in forensic medicine. Huang et al. (2018) had using geometric morphometric analysis of ten osteometric landmarks on three-dimensional reconstructions of 259 knees in Chinese population to study the differences of distal femur in term of sexual dimorphic and age. In order to identify the differences, General Procrustes analysis, principal component analysis (PCA), and other discriminant analysis had been conducted. PCA distinguished a significant difference of the distal femur between male and female while osteometric analysis showed the differences between the three age-related subgroup which are below 40 years, in between 40 to 60 years, and above 60 years.

2.3.5 CHARACTERISTICS OF SECONDARY SEXUAL

There are issues of secondary sexual characteristics associated with assessment such as privacy and cultural matters even though self-assessments have been implemented. The secondary sexual characteristics are pubic hair and genitalia in males while pubic hair and menarche in females (Malina, 2011).

Normal (2007) studied the yearly assessments of breast, genital, and pubic hair growth and those assessments had aided in analyzing the ages of individuals in each sexual maturity stage and the time it needed to go from one stage to the next. The findings from this study are essential in helping researchers to know the normative variation in the timing and also change in secondary sexual characteristics during puberty.

The timing of sexual maturity of a large group of healthy Danish children had been studied by TEILMANN et al. (2005) in order to assess the dissimilarities between USA and Denmark, and also to study the possible secular trends in sexual maturity development. The study showed that prepubescent child is easily affected by surrounding elements that may affect the endogenous hormonal milieu, and eventually impact the pubertal maturation.

2.3.6 THE SELECTED HUMAN PART TO DETERMINE AGE IN THIS RESEARCH

The most frequently traits that forensic anthropologists are going to assess in the identification of deceased bodies will be the age and the gender. This study focuses on the hand bones since other parts of the body have larger possibility to be missing before the criminal investigation is conducted. There are 19 bones in left hand and therefore it is harder to have no any hand bone remains at all. The widely used models to estimate age is based on the conservative method of observation of hand bones morphology which are Greulich and Pyle (GP) model and Tanner and Whitehouse (TW) model. The restrictions of these models are the estimated age generated is rely on the specialists or anthropologists and also the adopted subjects from certain population cannot ensure for other populations to implement these models in estimating age. (Cantekin, Celikoglu, Miloglu, Dane, & Erdem, 2012; Gungor, Sari, Gungor, Kale, & Celikoglu, 2015; Koc, Karaoglanoglu, Erdogan, Kosecik, & Cesur, 2001) have conducted studies of age estimation using these models for other populations and deduced that some factors like different surroundings circumstances and various ethnic

background have made these models not sufficiently great. Furthermore, genetic, growth, health and lifestyle, nutrient, and so on are factors that may impact the performance of the models to other populations (Malina, 1994).

There are few studies (Cameriere & Ferrante, 2008; Cameriere, Ferrante, Mirtella, & Cingolani, 2006) conducted the multiple linear regressions model for age estimation using the bone measurement. A new age indicator which is based on quantitative dataset of bone measurement that is going to be a substitution input in order to reduce the dependency of the specialists in estimating age.

2.4 GENERAL APPROACHES IN AGE ESTIMATION USING HAND BONES

There are several general approaches in age estimation using hand bones discussed in the following sections.

2.4.1 GREULICH AND PYLE (GP)

GP method, also known as the atlas method, is an improvement of the protocol chronicled by Todd in 1930s. GP method had thrived on the American White Children from the area of Cleveland, OH, USA, who were born between 1917 and 1942. This method used plates as standards, which represents the birth to maturity of 31 boys and 29 girls. It allocated the skeletal age (SA) to each bone of the hand-wrist, which in total is 29 bones. Pragmatically, GP SAs are in general, while unseemly, it excluding variation among bones since it constructed from the SA of the standard plate to the youth closest matches. There is no allocation of SA to each of the 29 bones once a person has attained skeletal maturity. This method has been adopted in the survey of players in Asian Youth under 16 championships (Malina, 2011).

2.4.2 TANNER AND WHITEHOUSE (TW)

The output shown by earlier researches about the adoption of TW in forensic field and Pinchi et al. (2014) analysed and demonstrated that the TW2 method is the worst among the methods of GP, TW2, and TW3 as it has a tendency to be biased, causing it to evaluate the age with a dissatisfactory precision. The study stated the TW2 method is not fitting to forensic age estimation since there is very high risk caused by large overestimation. The higher overestimations and lower accuracy value of TW2 implicit higher risks than GP and TW3 methods of false positives.

Pinchi et al. (2014) took 266 out of the sample of 307 X-rays of left wrist-hand bones in order to analyse the bone maturation of children for orthodontic initiatives while 41 taken for auxological initiatives, which collected from private dental practices and paediatric hospital respectively. There were 145 females and 162 males in the sample. The birthdates and radiological examinations were all accessible. The chronological age of subjects fluctuated from 6 to 20 years. There are three methods using in the study to estimate bone age and found out TW2 is not valid for age estimation as it given the highest overestimation in both genders while GP and TW3 are fit to estimate age with similar values of accuracy. GP and TW3 are more reliable for males than females and TW3 is a way more dependable than GP particularly in criminal cases because of the extraordinary overestimation trend of GP.

2.4.3 MULTIPLE LINEAR REGRESSIONS

Darmawan et al.(2015) had carried out a study that involved 333 X-ray images of left hand bones from 166 males and 167 females of Asian population. These X-ray images were gathered from Children's Hospital Los Angeles where been grouped in nineteen age groups for female and male.

The left hand bones are split into four groups which are distal phalax, middle phalax, proximal phalax and metacarpal. The middle phalanx group consists of four bones while another three groups consist of five bones respectively. The total left hand bones is 19. The data set ranged from newborn to 18 years old with no records of any bone illnesses.

Analysis of variance of intra-observer and inter-observer for repeated measures had performed to prove the measured data are reproducible and no significant statistical difference between those measurements. Each measurement in each image was repeated thrice and regression models are designed using the mean of that three measurements. There were two observers for inter-observer trial and 50 X-ray images were randomly chosen and measured by these observers ten times. It shows that no significant difference between two observers by generating another analysis of variance for measures.

SPSS statistical tool version 16.0 had been utilized for age estimation based on the analysis of bone lengths. The single bone method is age estimation using 11 regression models on each left hand bone, which are Linear, Logarithmic, Inverse, Quadratic, Cubic, Compound, Power, S-curve, Growth, Exponential, and Logistic. Multiple linear regressions had been applied on the 19 left hand bones for all bone method to develop the regression model. MSE value, R-square value and parameter will be generated by each regression model. MSE value is the average of the squares of the difference between the actual age and estimated age. The correlation between age and all bone lengths are represented by R-square value which will generate an equation for age estimation. There were 11 equations for each bone since the total number of bones for both genders is 19. For all bone method, there are only two generated equations while 209 equations for single bone method. Regression model with highest R-square value of female and male was selected as the best correlation with age while the equation with lowest MSE value was chosen for age estimation.

The best regression model based on highest R-square value of 0.960 for male and 0.900 for female is the S-curve regression while the best regression model based on MSE value is multiple linear regressions applied to all bones, which its MSE values are 1.654 years for male and 3.006 years for female. The best method shown in the study is the equation from multiple linear regression, where the left hand bones are complete. Single bone method will be implemented using the equation from regression models with the lowest MSE value based on the availability of left hand bones.

2.4.4 GENETIC ALGORITHM (GA)

GA is a tool for optimizing purposes that developed by Holland and was influenced significantly by the mechanism of natural selection and also the continual growth of all life forms across eons. GA method is not using any specific data to do searching since it is a probabilistic approach while objective function evaluation is needed in every decision in order to do proceeding. In accordance with what is generally done in GA, individual in populations, namely candidate solutions, steadily meet the most favorable solutions over time. 0 and 1 s are chromosomes which forms one linear string that proposes solution of each candidate. Every iteration has its optimizing procedure that initiated generations, which are population sizes. In GA, reproduction, crossover, and mutation are three basic genetic operators that generate following generation. A reproduction operator is the method or selecting the ideal chromosomes following their scaled values taking into account of the given standard of fitness. The selected chromosomes will be sent immediately to the next generation. Crossover operator is particular parts of parents or individuals merging each other and produce new individuals while mutation operator is having an uncertain substitution in elements of chromosomes (Khandelwal et al., 2018).

There are mainly six phases in genetic algorithm, which are initial population, fitness function, selection, crossover, mutation, and termination (Jain, 2019). The process of initial population phase starts with all the dataset as population, where each

record is characterized by a set of features known as genes. Genes will join to form chromosomes.

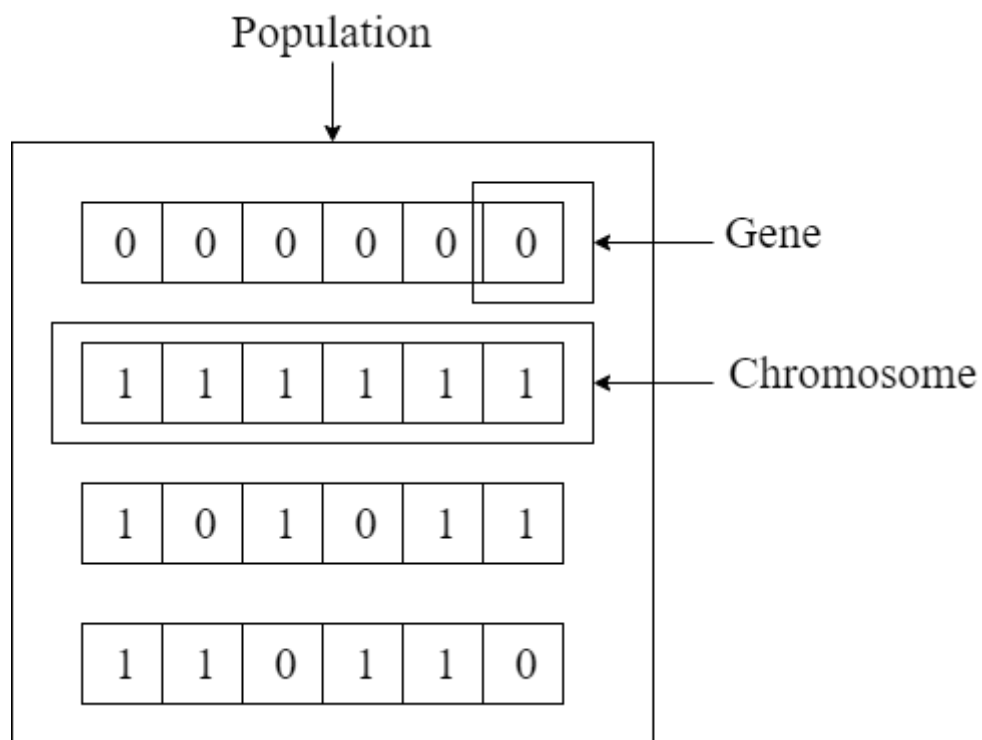
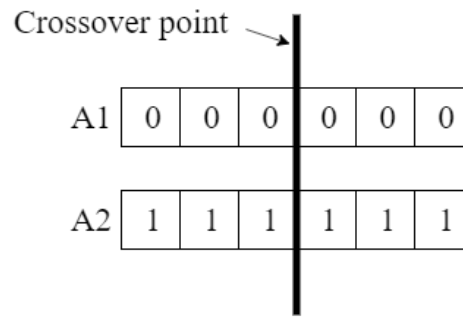


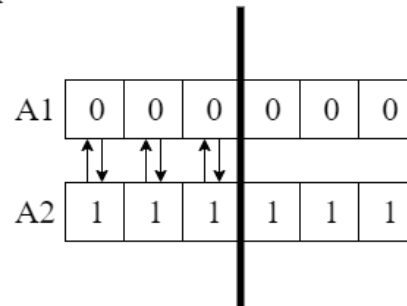
Figure 2.1: Gene, chromosome, and population of initial population phase

The fitness function phase is to measure the fitness of an individual where each of the individual is made up of some genes but not all of them. Features which having a major impact will be determined by finding the fitness of individual. In the selection phase, the two fittest individuals will be chosen for reproduction. The individuals chosen in selection phase will be used to form new individuals by changing some genes in individuals in crossover phase.

Step 1:



Step 2:



Step 3:

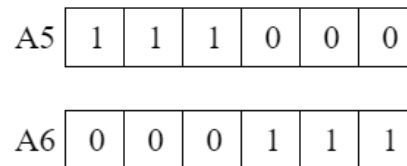


Figure 2.2: Three steps in crossover phase

In mutation phase, some genes are changed by adding some randomness. In termination phase, the genetic algorithm will be terminated when the population no longer generates any new individuals.

2.4.5 ARTIFICIAL NEURAL NETWORKS (ANN)

ANN is a function of estimation that is simulated from the way of sending information of the human brain. ANN is capable of being adopted even in extremely complex and non-linear contact circumstances (Khandelwal et al., 2018). Feed-

Forward Neural Network (FF-NN) is the extensively applied network to date. It is because of its simplicity and intelligibility compared to other networks and so as its potentiality to learn the implied governing relationship among the inputs and outputs when there is sufficient training data. A network structure like FF-NN will only propagate the signal or information in single direction (Sciences, 2015).

Typically, input, hidden and output layers are the three layers in FF-NN. It can possess hidden layers that make up more than one, yet a single layer is plenty to approach any function to a desired degree of accuracy. The number of neurons of input and output layers depend on the specific problem. In most cases, systemic trial and error is the finest procedure to find out the optimal number of neurons in hidden layer. Inputs will be fed through the input layer, then being multiplied by synaptic weights and are sent to the hidden layer. Normally, the selected logistic or the hyperbolic tangent is the nonlinear activation function that transforms the weighted sum of inputs in the hidden neurons. The same way goes through each hidden layers until the result attain the output neuron. On the other hand, the linear activation function is regularly implemented to the output layer. The most well received training algorithms that are extensively implemented for training FF-NN are back-propagation (BP) algorithms since it is easy to understand and applicable (Sciences, 2015).

The ANN compose a group of processing elements known as perceptrons which can be used to build a network structure via distinct arrangements. The network's layers are the location where these perceptrons reside. The releasing output, the managing input, and the input itself are obtained by every of the perceptron. They assimilate the releasing output, the managing input and the input itself, which makes these perceptrons the core components. There are two forms of input obtained by a perceptron, either an original input data is fed into the network, or the output data computed by another perceptron. Similarly, the output of a perceptron can be the final output of the network layer or to serve as an input into another perceptron. A single layer network must consists of at least one input layer and one output layer to establish a working network structure.

To construct the ANN, weights are utilized to link visible and hidden units with at least one other perceptron to the perceptrons in other layers. A hidden layer in a neural network is visualized with Figure 2.3 below.

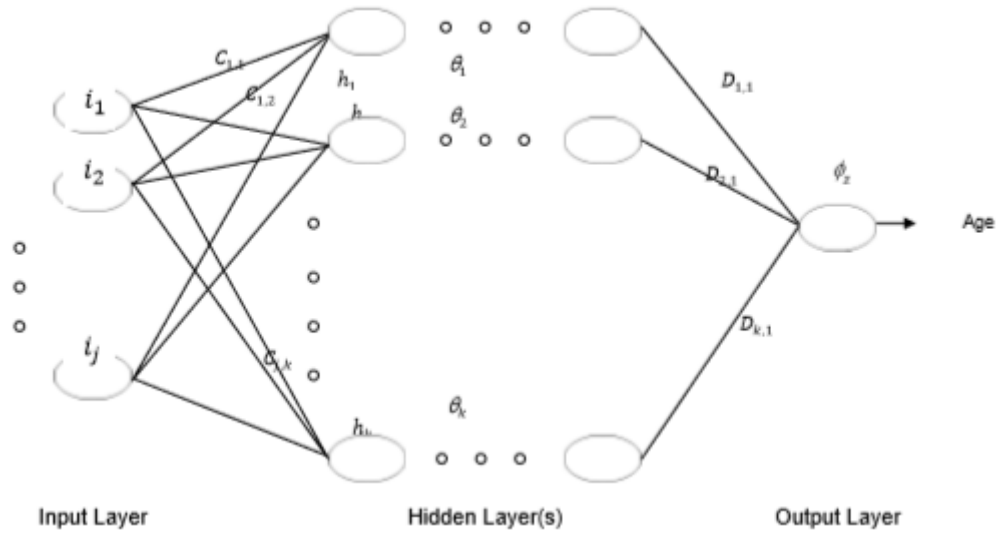


Figure 2.3: A conceptual design of a multi-layer neural network. Each layer are interlinked and the weights (c) indicates the links between the perceptron nodes.

The net input to unit k in the hidden layer is computed by implementing Equation 2.1 under the condition that a hidden layer network is utilized as in Figure 2.3.

$$net_hidden = \sum_{j=1}^J C_{j,k} i_j + \theta_k$$

Where J is nodes number of the input layer, θ_k is the biases of the hidden nodes, i_j is the input variables (the nineteen bones) and $C_{j,k}$ is the weights between the input nodes and hidden nodes.

Equation 2.1 Formula to compute the net input in hidden layer

Therefore, the net output from unit z in the output layer can be computed with Equation 2.2.

$$net_output = \sum_{k=1}^K D_{k,z} h_k + \phi_z$$

Where K is number of node in hidden layer, ϕ_z is the biases on the outputs nodes, h_k is the value of the output for hidden nodes and $D_{k,z}$ is the weight between hidden and output nodes.

Equation 2.2 Formula to compute the net output in output layer

Consequently, Equation 2.3 represents the result of hidden nodes and Equation 2.4 represents the result of output nodes which can be done with Equations 2.1 and 2.2. f is the transfer function to estimate the age.

$$h_k = f(net_hidden)$$

Equation 2.3 Formula to compute result of all hidden nodes

$$o_z = f(net_output) = age$$

Equation 2.4 Formula to compute result of all output nodes

To successfully construct an ANN, some variables are taken into consideration (Mohd, Haron, & Sharif, 2010) and the process trial and error is undergone. Researchers commonly implement the ANN in diverse, distinct areas, including forensics. However, an absence of detailed, recognized practices and procedures that could act as a standard guideline to establish the ideal approach. As a result, the effectiveness is ultimately reliant upon the operation of attempting different combinations for each variables and obtaining the best result. As a consequence, the procedure of trial and error will determine the effectiveness and capability of ANN. Therefore, the capability of the ANN could be impacted and affected by eight elements. These elements are the structure of the network, ratio of training and testing data, normalization of data, network algorithm, transfer function, performance function,

training function, and learning function. With trial and error, the eight elements can decide the perfect and faultless ANN.

2.4.5.1 STRUCTURE OF THE NETWORK

Nodes (neurons) and layers are the basic components of the ANN. The input layer, hidden layer and the output layer is comprised of the underlying structure of the ANN. On a small note, the hidden layer is an optional layer while the input layer and output layer are mandatory in the ANN structure. Figure 2.4 visualizes an ANN with $(i - j - k \dots - l - 1)$ structure. In the figure, the i nodes represent the nodes in the input layer, the j nodes in the first hidden layer, the k nodes in the second hidden layer, the l nodes in the m^{th} hidden layer and at least one node in the output layer.

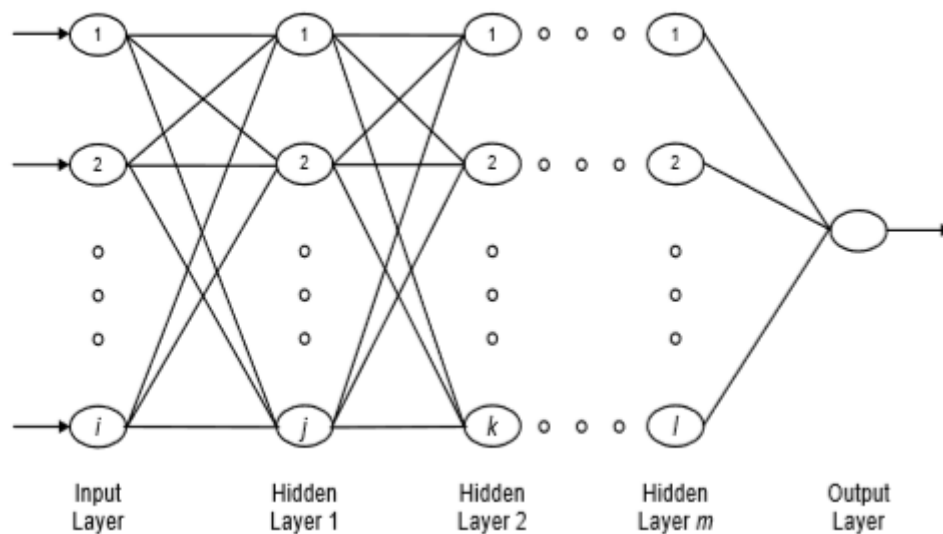


Figure 2.4: The ANN with $(i - j - k \dots - l - 1)$ structure is visualized

Author	(Kohail, 2012)	(Kumar, P, Sriram, & Vijay, 2013)	(Sharma & Venugopalan, 2014)
Network Structure	95-50-50-50-1, 50-25-25-25-1	68-29-4-4	136-68-36-4, 136-68-36-2

Normalization Data	-	-	-
Performance Function	Accuracy, R value	MSE, Accuracy	MSE
Training Function	-	<i>trainscg</i>	Various training function
Network Algorithm	Feed-forward BP	Feed-forward BP	Feed-forward BP
Learning Function	-	-	-
Transfer Function	Sigmoid	-	Sigmoid
Ratio of Training, Validation, and Testing Data	-	60 : 20 : 20	-

Table 2.1: Significant elements of the construction of ANN taken from an earlier study on age estimation

The network structure are dissimilar according to earlier research in Table 2.1. This is a result of the randomness of the trial and error process while constructing the ANN for the prime results where the layer's number and node's number of all hidden layers are modified. With this, it is established that any value to the number of hidden layers and the number of nodes in every hidden layer can be attempted. Be that as it may, the possibility that the number of hidden layer and the number of nodes in every hidden layer is affected by the desired data control effect, the computation time, computer memory and the complexity of the mapping cannot be denied. It is stated by Al-Ahmari (2007) that the expected data control effect may not be obtained with a small number of nodes, while an excess of computation time and computer memory can be a result of excess nodes. The number of nodes in the hidden layer which is " $2n + 1$ ", " $2n$ ", " $1n$ ", and " $n/2$ ", where the n is the number of input nodes can be computed with a formula suggested by (Zhang, Patuwo, & Hu, 1998) and (Mohd et al., 2010).

2.4.5.2 RATIO OF TRAINING AND TESTING DATA

Another factor to be attended to when constructing the ANN is the ratio of training and testing data. A fundamental principle of the ANN is that the accuracy of the ANN is linear to the amount of the training data, where the accuracy of the ANN increases when the amount of the training data increases. A portion from a normalized set of data of the nineteen bones and age of each subject is selected as training data for estimating age.

No rules are needed in selecting the ratio of training and testing data. From Table 2.1, the ratio of training and testing dataset differs from each other while some of the ratios are not mentioned, and thus the researchers have full authority over the selection of ratio for training and testing dataset, under the condition where the training data must be more than the testing data, based on statement by Mohd et al. (2010). On the other hand, a solution to this problem proposed by Zhang, Patuwo, & Hu (1998) suggests that the ratio of training and testing data is replaced with percentages that equals to 100%, such as 80%:20%, 85%:15%, or 90%:10%.

2.4.5.3 DATA NORMALIZATION

Before training and testing ANN, data normalization is generally carried out. A standard range, such as 0 to 1 or -1 to 1, is generally used when normalizing the data. It is a mandatory rule that if the output nodes employs nonlinear transfer functions such as the logistic sigmoid, the format of the expected output values must be converted to the same format of the range of the expected outputs of the network structure. Although a linear output is implemented, normalizing the outputs and the inputs to avoid problems from the computational process can be advantageous. When the issue regarding the data input/output normalization is taken into consideration, another issue that deserves attention is the method of deciding the normalization of the input/output data with the acquired size data. Suggested by (C. Sanjay; & C. Jyothi, 2006) and

(Mohd et al., 2010), a normalization equation for the input and output data is shown in Equation 2.5.

$$x_i = \frac{0.8}{d_{max} - d_{min}} (d_i - d_{min}) + 0.1$$

Where d_i is the input and output data at i th, d_{min} is the minimum value of input and output data, and d_{max} is the maximum value of the input and output data.

Equation 2.5 Normalization equation for the input and output data

2.4.5.4 NETWORK ALGORITHM

Different algorithms aimed at modelling the ANN such as Self-Organizing Map, Radial Basis, Perceptron, Time-delay BP, Elman BP, Cascadeforward BP and so on, have been developed by previous researches. The most common algorithm would be the feed-forward backpropagation algorithm, where it is implemented by earlier studies in age approximation according to Table 2.1. However, the questions remains as to which algorithm is best suited to construct the ANN to produce the finest estimation results. With several choices considered, however, the feed-forward backpropagation is still the most generally used for countless problems compared to other network algorithms, according to a statement made by Mohd et al. (2010).

2.4.5.5 TRANSFER FUNCTION

Different transfer functions for age approximation are implemented, which can be hypothesized by observing Table 2.1. Applicable transfer functions are the linear transfer function (purelin), log-sigmoid transfer function (logsig), hard limit transfer function (hardlim), hyperbolic tangent transfer function (tansig) and so on. Given the many choices of transfer functions, one of them must be selected to produce the finest

approximation result. Several options can be adopted, but no precise statements have been made that any notable effects are observed when different transfer functions are implemented.

According to the User's Guide of Neural Network Toolbox 6 written by (Demuth & Hagan, 2009), there are three transfer functions generally implemented together with the feed-forward backpropagation algorithm which are the tansig, logsig, and linear transfer function. The essence of the issue must align with the aim of the transfer function (Nalbant, Gökkaya, Toktaş, & Sur, 2009). As such, the nonlinear transfer function such as sigmoid and tansig is a fitting option when a non-linear relationship between input and output is applied for training the approximation model. The two key components, the simple derivative and self-limiting are the reason the sigmoid function has gained popularity. The end result is not able to develop infinitely large or small. The logsig and tansig are two transfer functions that have been applied by (Kohli & Dixit, 2005), and it is found that both results produced by the transfer functions are almost the same with each other. The sigmoid transfer function is widely implemented by researchers for age estimation, in which can be deduced from Table 2.1. Equation 2.6 is applied to compute the sigmoid transfer function while Equation 2.7 is employed to compute the tansig transfer function.

$$f = \frac{1}{(1+e^{-net})}$$

Equation 2.6 Formula to compute the sigmoid transfer function

$$f = \frac{2}{(1+e^{-2(net)})} - 1$$

Equation 2.7 Formula to compute the tansig transfer function

Using Equation 2.8, the hidden node's output is computed with the sigmoid function. Additionally the output for hidden nodes with the hyperbolic tangent transfer function is computed by implementing Equation 2.9. Equation 2.10 calculates the

output nodes employing the sigmoid function and the output nodes using the hyperbolic tangent function can be computed as clarified in Equation 2.11.

$$h_k(\text{sigmoid}) = f(\text{net_input}) = \frac{1}{1 + e^{-\sum_{j=1}^J c_{j,k} i_j + \theta_k}}$$

Equation 2.8 The hidden nodes' output with sigmoid transfer function is computed with the formula

$$h_k(\text{hyperbolic tangent}) = f(\text{net_input}) = \frac{2}{1 + e^{-2(\sum_{j=1}^J c_{j,k} i_j + \theta_k)}} - 1$$

Equation 2.9 The hidden nodes' output with hyperbolic tangent transfer function with the formula

$$o_z(\text{sigmoid}) = f(\text{net_output}) = \frac{1}{1 + e^{-\sum_{k=1}^K D_{k,z} h_k + \phi_z}}$$

Equation 2.10 Formula to compute the output nodes' output with sigmoid transfer function

$$o_z(\text{hyperbolic tangent}) = f(\text{net_output}) = \frac{2}{1 + e^{-2 \sum_{k=1}^K D_{k,z} h_k + \phi_z}} - 1$$

Equation 2.11 Formula to compute the output nodes' output with hyperbolic tangent transfer function

2.4.5.6 PERFORMANCE FUNCTION

According to Table 2.1, there is a widespread use of MSE value performance function among researchers for age estimation. To determine the MSE value, the basic equation is shown in Equation 2.12.

$$MSE = \frac{1}{n} \sum_{n=1}^n (actual\ output_n - predicted\ output_n)^2$$

Equation 2.12 Formula to compute the MSE value

A suitable performance function can be the performance measurement for the ANN, in which several options can be considered such as MSE value for age estimation in Table 2.1.

2.4.5.7 TRAINING FUNCTION

The error value must be minimized in the backpropagation approach. To do so, two main components should be assessed, the learning function and training function. To construct the greatest model, the mathematical specification software known as Matlab contains a toolbox of the ANN model that can facilitate the computation of the momentum (α) and learning rate (η) value, where the range of 0 to 1 will be used as a benchmark for the values of these parameters. Some training functions such as `trainscg` (Scaled Conjugate Gradient algorithm), `traingd` (gradient descent backpropagation), `trainlm` (LevenbergMarquardt backpropagation), `trainbr` (Bayesian regularization backpropagation) and others have been applied by earlier researchers, and among these training functions, one must be selected as the top. Unfortunately, the absence of a precise statement where different training functions could bring forth different performances of the model implies that no guidelines are available to select the best training function, based on the earlier case study.

2.4.5.8 LEARNING FUNCTIONS

Once again, same as the selection of training functions, a definite statement is not given if different network performance are yielded from different learning functions, and various learning functions that are commonly seen would be `learngd` (gradient

descent weight/bias), and learngdm (gradient descent with momentum weight/bias). Researchers have not stated the learning function they have chosen in Table 2.1.

2.4.6 HYBRID GENETIC ALGORITHM-ARTIFICIAL NEURAL NETWORKS (GA-ANN)

Although ANN has a lot of favorable features, there are problems like slow convergence and fail to progress at local minimum. In order to solve these issues, researchers proposed a hybrid approach of GA-ANN to improve the performance of ANN to attain the global minimum. GA-ANN contemplating a suitable fitness function and therefore, significant parameters of ANN will be deduced and modified. The intention of adopting hybrid GA-ANN model is to alter a set of weights and biases to lessen objective function (Armaghani, Mohamad, Monjezi, Faradonbeh, & Majid, 2016).

GA is implemented to enhance the accomplishment of artificial intelligence (AI) techniques. GA is extensively applied to choose neural network topology like improving relevant feature subset, deducing the most favorable number of hidden layers and elements in process. Architectural factors of ANN such as feature subset, number of hidden layers, number of elements in process of hidden layers, activation functions, and so as the connection weights among layers are going to be deduced in advance. Nearly all of existing studies were emphasized on the enhancement of the learning algorithms itself (Kim & Han, 2000).

2.5 COMPARISON AND JUSTIFICATION

Source	Approach	Advantages	Disadvantages
(Pinchi et al., 2014)	GP	Simple and effective	Tend to overestimate
(Cantekin et al., 2012)		Contains reliable references for different ages	Require professional anthropologists to execute

(Koc et al., 2001)			Accuracy is dependent on proficiency of the anthropologist
(Spampinato, Palazzo, Giordano, Aldinucci, & Leonardi, 2017) (Benjavongkulchai & Pittayapat, 2018) (Pinchi et al., 2014)	TW	More effective than GP for some ages Contains reliable references for different ages	Require professional anthropologists to execute More complex and time-consuming Accuracy is dependent on proficiency of the anthropologist
(Susan L. King, 1999) (Darmawan et al., 2015)	MLR	Understands the relationship between the variables	Dependent on the dataset as some data can be irrelevant to the prediction Require a predefined relationship between the variables Works only if the relationship between variables are linear
(Susan L. King, 1999) (Rucci, Coppini, Nicoletti, Cheli, & Valli, 1995) (Bocchi, Ferrara, Nicoletti, & Valli, 2003)	ANN	Require no relationships between the data to predict Find both linear and non-linear relationships between variables	Highly dependent on the dataset to make an estimation Require the data to be completely numerical Does not filter the input features as some input features may be irrelevant and negligible to the estimation

Table 2.2: Sources, advantages and disadvantages of approaches in age estimation using hand bones

The source column in Table 2.2 records the documents used as references in reaching these comparisons. The approach column states the names of the approaches. The advantage and disadvantage columns states the strong and weak points for each approach. As a conclusion, the ANN approach is the best approach in terms of estimating the age of the Hispanic population. One of the disadvantage of ANN approach is that input features are not filtered, which may cause waste of data and inaccurate estimations. The problem can be overcome by introducing a GA approach to select the best features among the input features to reduce the load and increase the accuracy of the estimation of the ANN approach. As such, a GA-ANN hybrid approach has been proposed, where the GA approach will be implemented first to select the best features and the ANN approach will estimate the age of the Hispanic population based on the best features.

2.6 SUMMARY

The conservative model for age estimation is reliant upon the examination results of bone morphology from the X-ray images of the left hand bones by forensic anthropologists. The main problem is that this model requires the professionalism of forensic anthropologists in estimating the age, and the variability may vary from one to another. In other words, different levels of experiences of forensic anthropologists in age estimation will result the high variable between the actual age and the estimated age. The next problem is the models of this study are population specific. This means that the models used for age estimation is not applicable for various populations. The third issue is only a few studies using soft computing model for age estimation. The target of the current research is to construct an age estimation model for Hispanic population.

There are advantages and disadvantages of different approaches on age estimation using hand bones. Through the comparison and justification, a conclusion is made and ANN approach has been known as the best approach to estimation the age. In the meantime, there is a biggest disadvantage of ANN approach, where there is no

filter applied on input features, which may lead to waste of data and inaccurate estimation. Therefore, a GA-ANN hybrid approach is proposed, where GA approach is implemented to select features before the ANN approach.

CHAPTER 3

METHODOLOGY

3.1 INTRODUCTION

This chapter reviews the methodology that will be applied during data collection, data normalization, age estimation model using ANN approach, age estimation using GA-ANN approach, data denormalization, and validation and evaluation of results.

3.2 RESEARCH FLOW

The research flow of this study involves seven phases which are the problem definition, data collection, data normalization, age estimation model using ANN approach, age estimation using GA-ANN approach, data denormalization, and validation and evaluation of results.

3.3 PROBLEM DEFINITION

Problem definition is the first phase in this study as stated in Section 3.2. Problem definition requires a direct, specific, and incisive statement of issues to be explored and probed with the objective of getting a solution. In this study, problem definition has been stated in Section 1.2.

3.4 DATA COLLECTION

The original dataset is constructed from an online dataset taken from <http://ipilabmysql.usc.edu/newindex.php>. There are subjects of Caucasian, African American, Hispanic, and Asian in the online dataset. Only Hispanic subjects are adopted in this study as described in the Section 1.2.

There are 365 X-ray images of left hand bones of Hispanics, in which out of total, 182 are males while 183 are females. This dataset varied from age of 0 to 18 years without any problem or disease of bone like fractures, genetic bone problems and so on. Bones with those issues will be excluded since they are regarded as likely to be weak and easy to break which may influence the assessments or measurements. All the X-ray images were collected from Children's Hospital Los Angeles together with demographic data of patients as well as the reading of radiologists, dispersed into nineteen groups which are 0 to 18 years old, for both genders. The details of each subjects are the respective image name, race, gender, chronological age, date of birth, exam date, tanner, height (cm), weight (kg), trunk HT (cm), reading 1, and reading 2. Reading 1 and reading 2 are the estimated age evaluated by experienced radiologist using GP model (Gertych, Zhang, Sayre, Pospiech-Kurkowska, & Huang, 2007). In the end of this study, these estimated age generated using GP model will be adopted as the first benchmark for this study in order to validate and evaluate the proposed model.

The National Institute of Health has funded this X-ray images collection that gathered from the Image Processing and Informatics Lab of University of Southern California and is confined to only purposes of open research and education. Institutional review board has agreed with inspecting candidates for clinical investigations to be inspected for studies. The radiographs were digitalized to 2k x 2k images by applying a laser film scanner (Array, Tokyo, Japan) while patients' demographic data had manually inserted using the scanner GUI (graphical user interface) and saved as DICOM file (Gertych et al., 2007).

The 19 left hand bones are grouped into four main groups which are distal phalanx, middle phalanx, proximal phalanx, and metacarpal. There are four bones in the middle phalanx while five bones respectively in the other three groups. There are 19 bones in hand and Figure 3.1 shows the label of each bone in left hand. Epiphyses and diaphysis are two major parts of long bone during childhood. The former is at the tail of a long bone that has started to disengage from the main bone by a layer of cartilage and is joint to main bone through ossification during adulthood while the central shaft that having mostly of compact bones neighbouring a cavity. Table 3.1 shows the six key phases during childhood and adolescence (Gilsanz & Raitb, 2005).

Phase	Male (age)	Female (age)
Infancy	Newborn to 14 months	Newborn to 10 months
Toddlers	14 months to 3 years	10 months to 2 years
Pre-puberty	3 years to 9 years	2 years to 7 years
Early and Mid-puberty	9 years to 14 years	7 years to 13 years
Late Puberty	14 years to 16 years	13 years to 15 years
Post-puberty	16 years to 19 years	15 years to 17 years

Table 3.1: The six key phases during childhood and adolescence for both gender

Left hand bones lengths in normalization form is the input that adopted in this study. A free photo editor, Photo Pos Pro, Power of Software Company Ltd. is utilized in order to assess the nineteen bones lengths in each X-ray image by constructing a line on each bone, starting from base-middle point to end-middle point of the bone on each X-ray image. These lines are constructed by disregarding the epiphyseal in bone even if it exists for infant phage while lines will be constructed by involving the epiphyseal for other phases even if it is only a small epiphyseal occurred in the X-ray image.

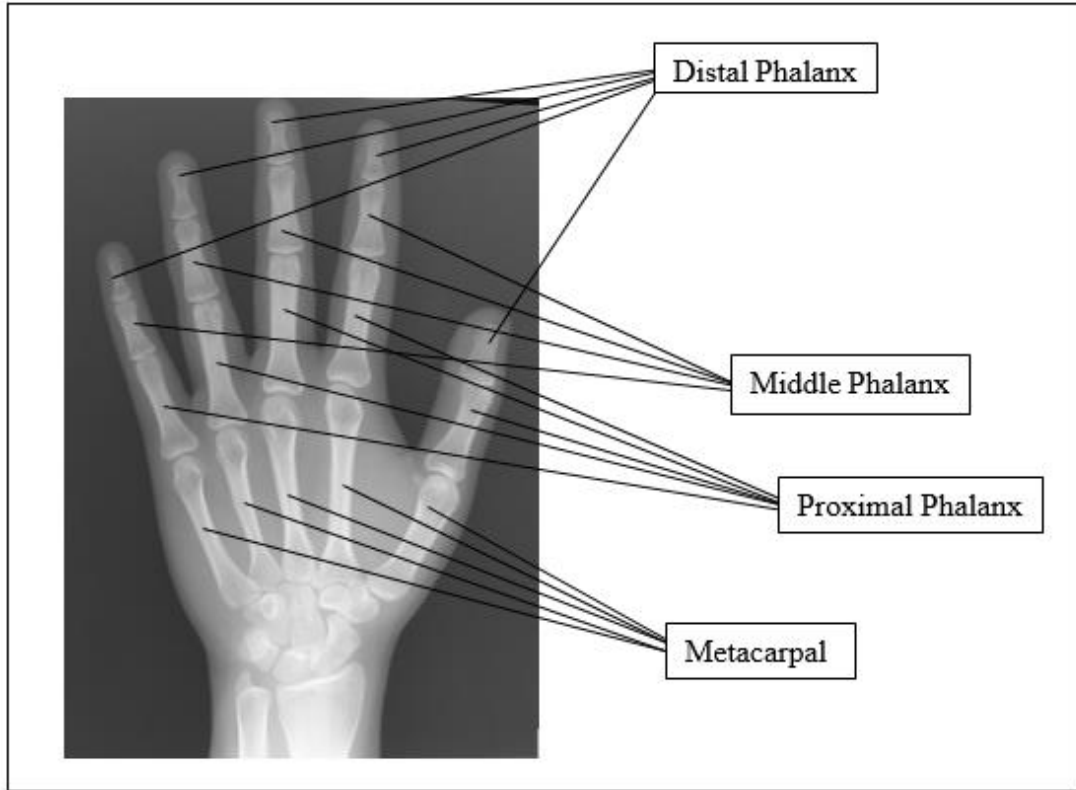


Figure 3.1: Left hand bone with labels

3.5 DATA NORMALIZATION

The proposed approach will adopt the normalized dataset in order to estimate the age. The dataset will be normalized based on Equation 2.5, which contains the dataset in the range of 0.1 to 0.9. The normalization aims to remove any computing errors during the training of the ANN, since the ANN uses mathematical equations to estimate the age. Datasets which are not normalized may produce errors in estimating.

3.6 AGE ESTIMATION MODELS USING ANN APPROACH

The dataset will be distributed into two groups: training dataset and testing dataset. In order to train the age estimation models using ANN approach, we need to have a training dataset while in order to validate and evaluate the models, we require

testing dataset. Samui & Dixon (2012) have adopted different data division between training dataset and testing dataset and stated that there is no a strictly accurate principle. In this study, 70% of total dataset will be used as training dataset while another 30% will be adopted as testing dataset. The selected subjects for the training dataset and testing dataset are going to be determined chaotically by implementing the function *dividerand* in Matlab.

An estimated age in the conservative way of age estimation is determined based on qualitative data such as observation of bone morphology which requiring the inspection of forensic anthropologists. Bone length will be used as the quantitative age indicator in this study in order to reduce the dependency towards forensic anthropologists in estimating age while ANN approach has been proven in many research studies since it showing strong pattern classification, pattern recognition and estimation capabilities when quantitative data being used. Therefore, the adoption of bone length as quantitative age indicator to ANN approach is able to be trusted.

There are eight factors that discussed in Section 2.6.2 in creating age estimation models using ANN approach in order to let the models produce the finest result on the estimation. Test regression value and test set MSE value will be compared in order to choose the finest network structure for age estimation in ANN model for both genders.

3.7 AGE ESTIMATION MODELS USING GA-ANN APPROACH

GA approach is an algorithm to solve restricted and unconstrained optimization problems that rely on natural selection or biological evolution. A population of individual solutions will be altered time after time in this algorithm. The GA approach will generate a population at the beginning, which is completely randomized and then the next generation will be produced. The approach will use the individuals in the current generation in order to generate next population at each step. In order to generate the new population, GA approach will achieve each member of the current population by calculating its fitness value and these values are known as the raw fitness scores. A

more utilizable range of values called expectation values is transformed by scaling the raw fitness scores and parents or the members are selected based on the expectation values. Individuals that having lower fitness in current population will be selected as elite and will be inherited to next generation. Two methods are used to generate children from parents: mutation which randomly modifies one parent and crossover which combines vector entries between two parents. The next generation is then comprised of all children of the parents in the earlier generation. The model will terminate when one of the stopping criteria is fulfilled. The approach will ultimately generate the best input features to be inserted into an ANN for training.

The age estimation model using GA-ANN hybrid approach is the combination of GA approach and ANN approach, where the purpose of GA approach is to select the most appropriate bones of left hand to be used as input in ANN approach while ANN approach is to estimate the age. This is because ANN approach is a black box learning approach that cannot perform the correlation between input and output. Figure 3.2 illustrates the flow of the implementation of the GA-ANN approach.

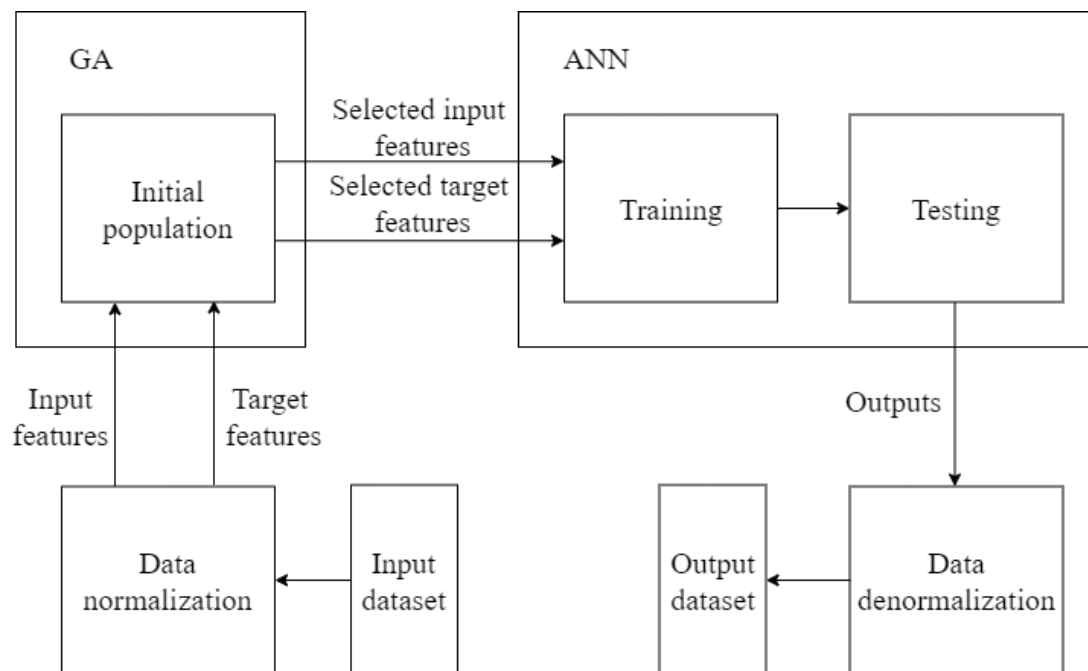


Figure 3.2: The flow of the implementation of the GA-ANN approach

The dataset is first normalized to avoid computation errors and fed into the GA approach. Through crossover and mutation, a new population containing the best input features will be generated. The best input features and target features will then be fed into the ANN for training and testing. The outputs will be denormalized to obtain the real estimated age and the errors will then be calculated. The process of denormalization and evaluation of results will be discussed later.

3.8 DATA DENORMALIZATION

The test set data consists of the input data, targeted output data, the estimated output data and the errors. The input data will not be taken as it is part of the normalized dataset and does not contribute in evaluating the results. The errors are generated from the ANN, in which the errors are generated based on the normalized inputs. These errors will not be used as a metric to evaluate the results, but instead it is used to validate the targeted output data and the estimated output data. This is done by subtracting the targeted output data with the estimated output data and the difference is matched with the errors.

The targeted output data and the estimated output data will be denormalized to obtain the actual age. The denormalization formula will be the inverse of the formula mentioned in the previous section, Section 3.5. The denormalization will convert the data into the range of 0 to 18. Once the data is denormalized, the targeted output data and the estimated output data will be used for the evaluation of results.

3.9 VALIDATION AND EVALUATION OF RESULTS

The test regression value, r and the test set MSE value will be used as the evaluation metric for the results for both approaches. These two evaluation metric are used for the majority of neural networks. The test regression value is a coefficient of correlation, where the closer the value of r is to 1, the better the correlation (Creative

Research Systems, 2016). The higher the test regression value, the better the performance for that approach. The test regression value is the accuracy of the approach, which reflects how well the test data was able to map to the ANN. On the other hand, the lower the test set MSE value, the better the performance for that approach. The test set MSE value is the average of all the test errors by subtracting the denormalized targeted output data with the denormalized estimated output data. The less errors predicted by the ANN, the lesser the test set MSE value, which increases the accuracy of the ANN. Below is a figure of the research flow.

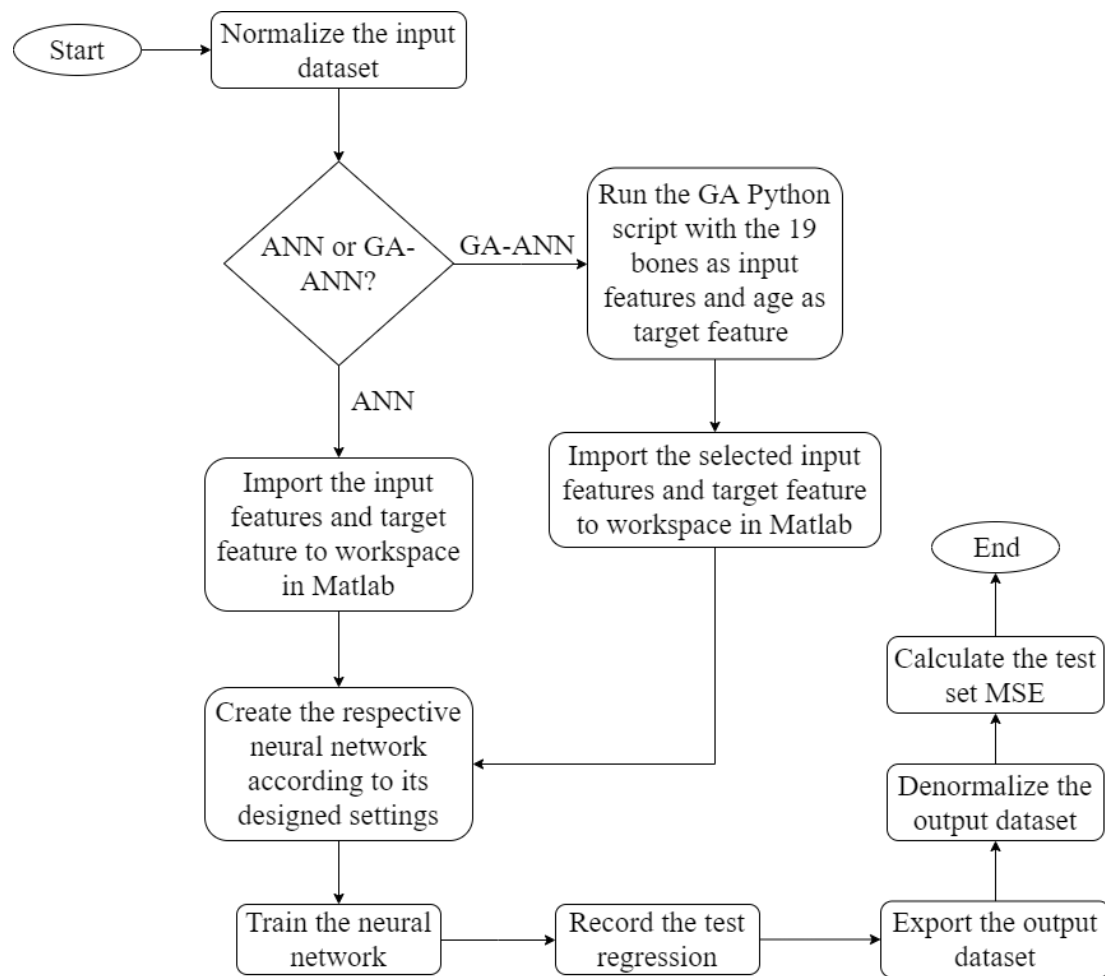


Figure 3.3: Flowchart of this research flow

In the flowchart above, the input is normalized first. For the ANN approach, the ANN will directly receive the normalized input, create and train the ANN based on the normalized input. The test regression and the test set MSE value is calculated to

validate the performance of the ANN approach. For the GA-ANN hybrid approach, the normalized inputs will be reduced to the best input features through feature selection in the GA approach. The same steps with the ANN approach are repeated, where the difference is that the input will be the best inputs. The test regression value and the test set MSE value will be calculated for the GA-ANN hybrid approach. The test regression values and the test set MSE value will be compared from both approaches.

3.10 HARDWARE AND SOFTWARE

Tools in the configuration of hardware and software are compulsory to have in order to aid this research carry out in a smooth way. The hardware tool that adopted to perform the implementation is particularly efficacious and well-organized. The computer that operated for implementation of this research performs on Microsoft Windows 10 Pro with the specifications of Intel Core i5-6200U, 2.6GHz, 64-bit operating system, x64-based processor, and 8GB of random-access memory (RAM).

The software tools that are going to be utilized are a free photo editor, Photo Pos Pro, Power of Software Company Ltd. in order to assess the bones lengths, Microsoft Excel 2013 for data collection, Microsoft Word 2013 for documentation, and Matlab R2012a for constructing the approaches. Python scripts and the Miniconda tool are used to develop the GA approach.

3.11 GANTT CHART

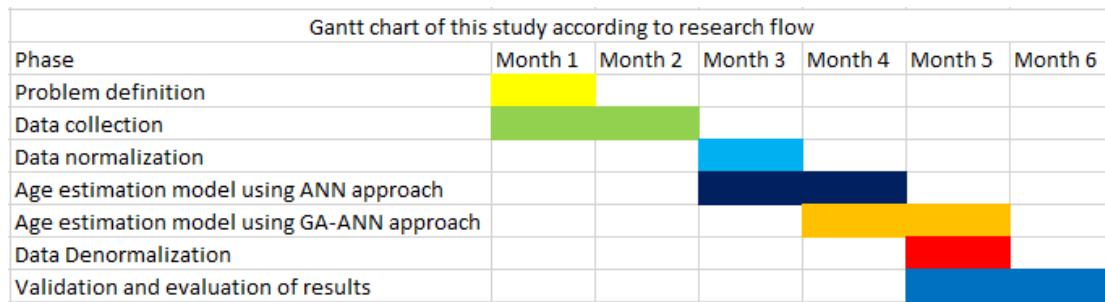


Figure 3.4 Gantt chart of project

3.12 SUMMARY

This chapter explored about the methodology used in age estimation models using GA approach, ANN approach, and hybrid GA-ANN approach. The flows and steps to implement approaches are discussed and the methods to validate and evaluate results have also been indicated.

CHAPTER 4

ARTIFICIAL NEURAL NETWORK MODELS AND HYBRID GENETIC ALGORITHM-ARTIFICIAL NEURAL NETWORK MODELS

4.1 INTRODUCTION

This chapter discusses the implementation of applying ANN approach and GA-ANN hybrid approach on age estimation models for both female and male population of Hispanic population, the results of the models, the discussion, and the justification of the mentioned results.

4.2 IMPLEMENTATION

The original data is constructed from an online dataset taken from <http://ipilabmysql.usc.edu/newindex.php> and only Hispanic subjects are adopted in this study. Photo Pos Pro, a free photo editor, Power of Software Company Ltd. is used in order to assess the nineteen bones lengths in each radiograph, by constructing a line on each bone, starting from base-middle point to end-middle point of the bone of each radiograph. The lines are constructed by disregarding the epiphyseal in bone even if it exists for infant phase while for other phases, lines are constructed by involving the epiphyseal even if it is only a small epiphyseal occurred in the radiograph. Below are the nineteen bones and its particular ages used as input features and target feature respectively during the implementation of ANN approach and the part of GA approach of the GA-ANN hybrid approach. For the input features, which are bones, each of them will be represented by an integer number, starting from 1 to 19, or feature name, starting from Bone 1 to Bone 19, while target feature, age, will be labelled as 1 or Age.

	ANN	GA
Features	Integer number	Feature name
First Metacarpal	1	Bone 1
Second Metacarpal	2	Bone 2
Third Metacarpal	3	Bone 3
Fourth Metacarpal	4	Bone 4
Fifth Metacarpal	5	Bone 5
First Proximal Phalanx	6	Bone 6
Second Proximal Phalanx	7	Bone 7
Third Proximal Phalanx	8	Bone 8
Fourth Proximal Phalanx	9	Bone 9
Fifth Proximal Phalanx	10	Bone 10
Second Middle Phalanx	11	Bone 11
Third Middle Phalanx	12	Bone 12
Fourth Middle Phalanx	13	Bone 13
Fifth Middle Phalanx	14	Bone 14
First Distal Phalanx	15	Bone 15
Second Distal Phalanx	16	Bone 16
Third Distal Phalanx	17	Bone 17
Fourth Distal Phalanx	18	Bone 18
Fifth Distal Phalanx	19	Bone 19
Age	1	Age

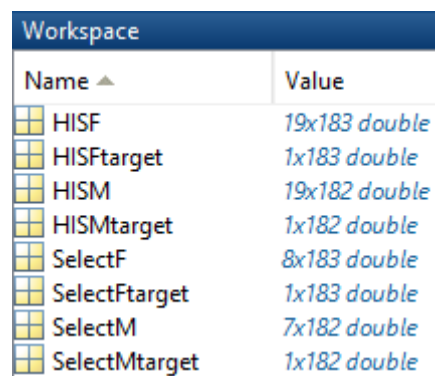
Table 4.1: The features and its represented integer number and feature name in ANN approach and part of the GA approach of GA-ANN hybrid approach respectively

In the implementation of ANN approach, the nineteen bones are used as input features and each of the bones is represented by an integer number, starting from 1 to 19. During the implementation of the part of GA approach of GA-ANN hybrid approach, the nineteen bones are used as input features and each of them is represented by its respective feature name. For example, first metacarpal is represented by integer number 1 in the implementation of ANN approach, and Bone 1 in the implementation of the part of GA approach of GA-ANN hybrid approach. The feature, age has been represented by integer number 1 in the implementation of ANN approach while it remains the same in the implementation of the part of GA approach of GA-ANN hybrid approach. First metacarpal and age, both features can be represented by the same integer number 1 since the feature of age is imported as another variable in the workspace in Matlab.

Before all the models are trained, the data is normalised to prevent any technical errors during the training of the models. The normalization formula is Equation 2.9 in Chapter 2.6.2. The normalised data can be found in the appendix.

4.2.1 ARTIFICIAL NEURAL NETWORK

The input data and target data for all models are imported to the workspace in Matlab.



Workspace	
Name ▲	Value
HISF	19x183 double
HISFtarget	1x183 double
HISM	19x182 double
HISMtarget	1x182 double
SelectF	8x183 double
SelectFtarget	1x183 double
SelectM	7x182 double
SelectMtarget	1x182 double

Figure 4.1: The imported input data and target data for all models in the workspace of Matlab

HISF and HISFtarget are the input features and target feature of age estimation model using ANN approach for female Hispanic population while HISM and HISMtarget are for the male Hispanic population. SelectF and SelectFtarget are the input features and target feature of age estimation model using the GA-ANN approach for female Hispanic population while SelectM and SelectMtarget are for the male Hispanic population. All of these input features and target features are imported as variables in the workspace in Matlab. The data or the input features, and the target feature of SelectF, SelectM, SelectFtarget and SelectMtarget have gone through the process of feature selection using GA approach.

Below is the command in Matlab to open the Network/Data Manager window which can import, create, use, and export neural networks and data.

```
>> nntool
```

Figure 4.2: The command to open Network/Data Manager window in Matlab

The command in Figure 4.2 is entered in the command window in Matlab in order to open the Network/Data Manager window which has been shown in Figure 4.5.

A network can be created by giving a name to the network, select the network type, input data, target data, training function, adaption learning function and performance function, customize the number of layers and also the number of neurons for Layer 1, and transfer function for Layer 1 and Layer 2. Below is the table of the settings for each neural network to be created.

		Female		Male	
Name		HISFnet	SelectFnet	HISMnet	SelectMnet
Network type		Feed-forward backprop	Feed-forward backprop	Feed-forward backprop	Feed-forward backprop
Input data		HISF	SelectF	HISM	SelectM
Target data		HISFtarget	SelectFtarget	HISMtarget	SelectMtarget
Training function		TRAINLM	TRAINLM	TRAINLM	TRAINLM
Adaption learning function		LEARNGDM	LEARNGDM	LEARNGDM	LEARNGDM
Performance function		MSE	MSE	MSE	MSE
Number of layers		2	2	2	2
Properties for Layer 1	Number of neurons	13	6	13	6
	Transfer function	PURELIN	PURELIN	PURELIN	PURELIN
Properties for Layer 2	Transfer function	TANSIG	TANSIG	TANSIG	TANSIG

Table 4.2: The settings of each neural network in this paper

HISFnet and HISMnet are the neural networks where only ANN approach has been applied while SelectFnet and SelectMnet are the neural networks where their

respective input data and target data have gone through the process of feature selection using GA approach.

HISFnet is the feed-forward backpropagation neural network with the input data, HISF and target data, HISFtarget. The training function, adaption learning function, and performance function of HISFnet are TRAINLM, LEARNGDM and MSE respectively. The number of layers of HISFnet is two while the number of neurons in Layer 1 is 13. The transfer function for Layer 1 and Layer 2 are PURELIN and TANSIG respectively.

SelectFnet is the feed-forward backpropagation neural network with the input data, SelectF and target data, SelectFtarget. The training function, adaption learning function, and performance function of SelectFnet are TRAINLM, LEARNGDM and MSE respectively. The number of layers of SelectFnet is two while the number of neurons in Layer 1 is 6. The transfer function for Layer 1 and Layer 2 are PURELIN and TANSIG respectively.

HISMnet is the feed-forward backpropagation neural network with the input data, HISM and target data, HISMtarget. The training function, adaption learning function, and performance function of HISMnet are TRAINLM, LEARNGDM and MSE respectively. The number of layers of HISMnet is two while the number of neurons in Layer 1 is 13. The transfer function for Layer 1 and Layer 2 are PURELIN and TANSIG respectively.

SelectMnet is the feed-forward backpropagation neural network with the input data, SelectM and target data, SelectMtarget. The training function, adaption learning function, and performance function of SelectMnet are TRAINLM, LEARNGDM and MSE respectively. The number of layers of SelectMnet is two while the number of neurons in Layer 1 is 6. The transfer function for Layer 1 and Layer 2 are PURELIN and TANSIG respectively.

Below are the Network/Data Manager windows to create neural network, and also to import, export, and use the neural networks and data.

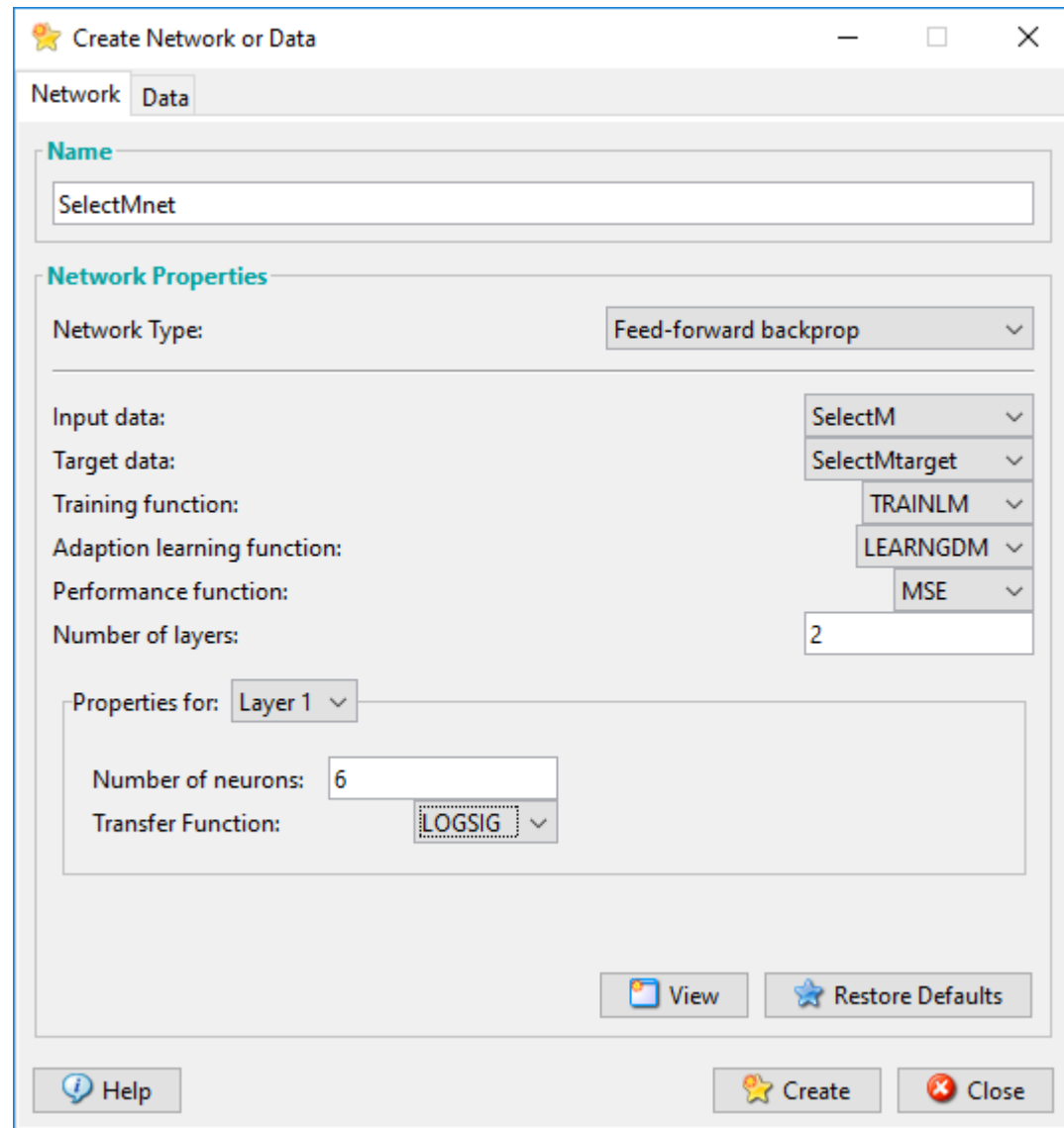


Figure 4.3: Example of the Network/Data Manager window to create one of the neural networks

Figure 4.3 is the example of Network/Data Manager window where a neural network can be created by selecting its network type, input data, target data, training function, adaption learning function and performance function, and customizing the number of layers, the number of neurons of Layer 1 and also the transfer function for Layer 1 and Layer 2. All networks, which are HISFnet, HISMnet, SelectFnet and

SelectMnet will be created with this window. Figure 4.4 has shown the example of customizing the transfer function for Layer 2 of one of the neural networks.

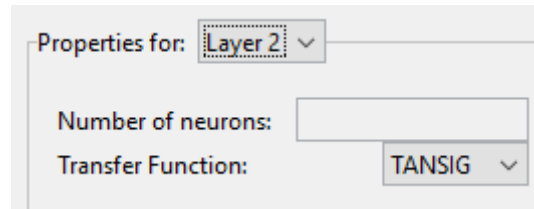


Figure 4.4: Example of customizing the transfer function for Layer 2 of one of the neural networks

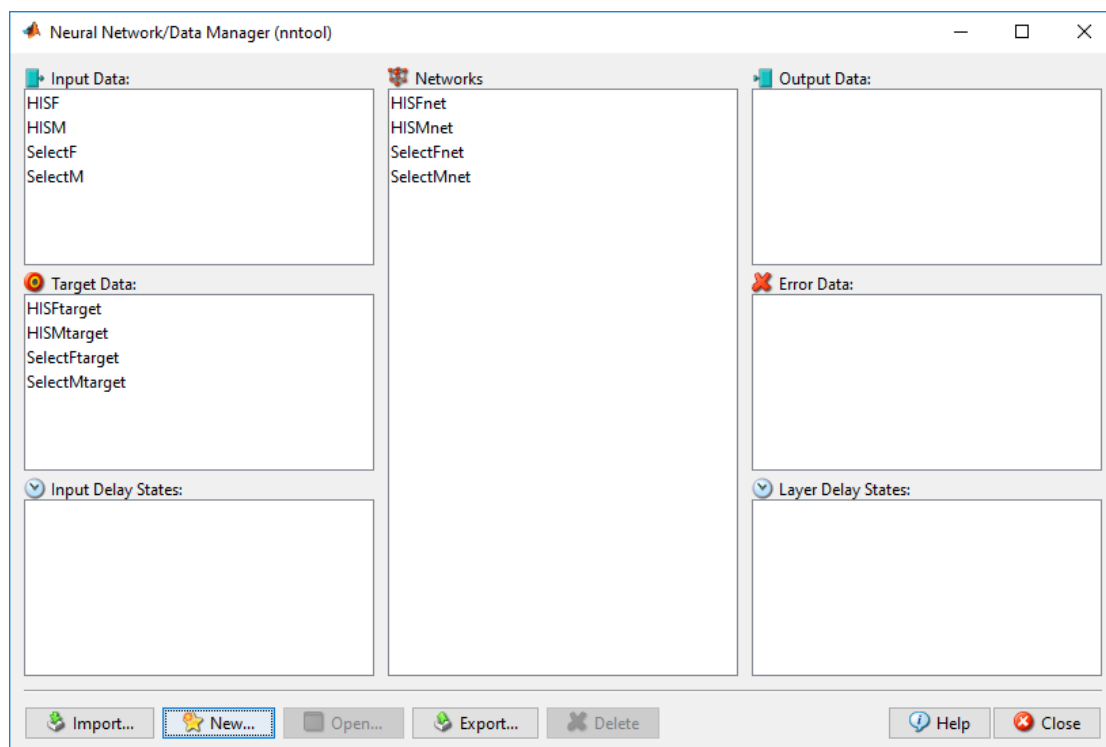


Figure 4.5: The Network/Data Manager which creates neural network, imports, exports, and uses the neural networks and data

All the created networks are exported to the workspace in Matlab.

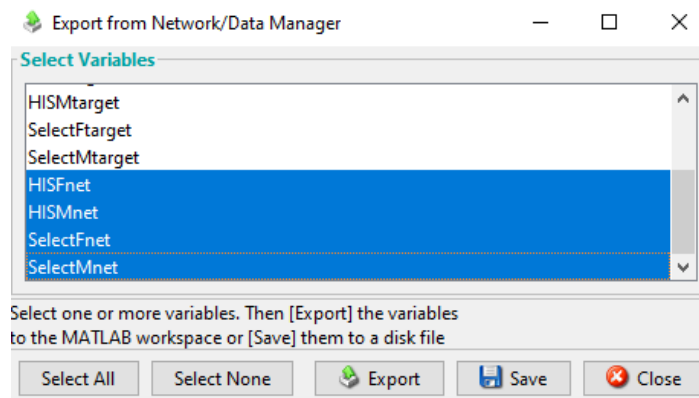


Figure 4.6: Example of exporting the neural networks to workspace of Matlab

Figure 4.6 shows that HISFnet, HISMnet, SelectFnet and SelectMnet are going to be exported to workspace in Matlab.

Below is the command to randomly generate column indices to be used to divide the data into 70% training data and 30% testing data respectively.

```
>> [trainInd, testInd, valInd] = dividerand(183, 0.7, 0.3, 0);
```

Figure 4.7: The command to randomly generate column indices for the division of the data into 70% training data and 30% testing data respectively

The command in Figure 4.7 is entered in the command window in Matlab to randomly generate column indices for the division of the data into 70% training data and 30% testing data respectively. The number of data, which is the number of individuals will be specified first before the percentages of the divided data. For the female and male networks, the number will be 183 and 182 respectively. The generated column indices will be applied on those variables which are formed by the data of input features and target features. Since one column represents one individual in the variable, column indices are used in the division of the data.

Below is the command to change the divide function in order to enable dividing data with indices.

```
>> HISFnet.dividefcn = 'divideind';
```

Figure 4.8: The command to change the divide function in order to enable dividing data with indices

The command in Figure 4.8 is entered in the command window in Matlab to change the divide function in order to enable dividing data with indices.

Below are the commands to set the generated indices to a particular neural network.

```
>> HISFnet.divideParam.trainInd = trainInd;  
>> HISFnet.divideParam.testInd = testInd;  
>> HISFnet.divideParam.valInd = valInd;
```

Figure 4.9: The commands to set the generated indices to a particular neural network

The command in Figure 4.9 is entered in the command window in Matlab to set the generated indices to a particular neural network. All networks will have their respective input data divided in this method.

Below is the neural network design for HISFnet and HISMnet.

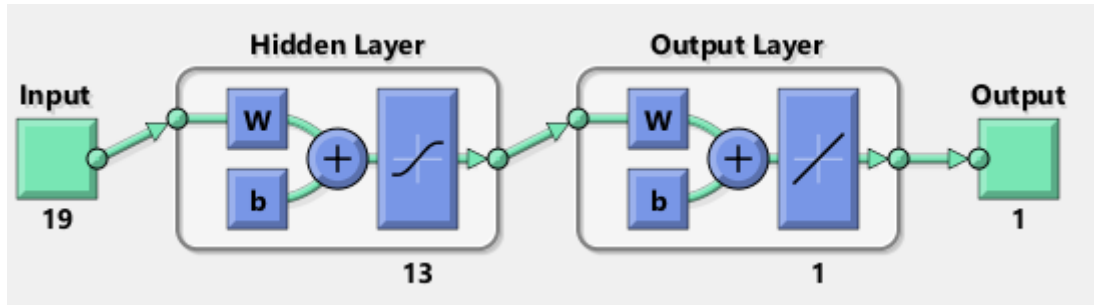


Figure 4.10: The neural network design for HISFnet and HISMnet

Figure 4.10 shows the neural network design for HISFnet and HISMnet. There are 19 input features, which are the 19 bones while the hidden layer consists of the weights and biases and also 13 neurons. There are weights and biases and also one neuron for the output layer and one output feature will be produced.

Below is the neural network design for SelectFnet.

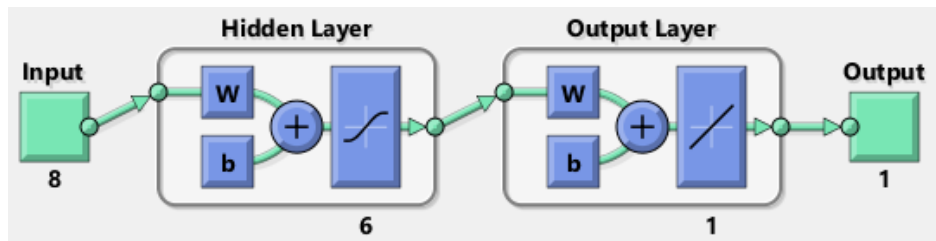


Figure 4.11: The neural network design for SelectFnet

Figure 4.11 shows the neural network design for SelectFnet. There are eight input features, which are the eight selected bones using GA approach while the hidden layer consists of the weights and biases and also six neurons. There are weights and biases and also one neuron for the output layer and one output feature will be produced.

Below is the neural network design for SelectMnet.

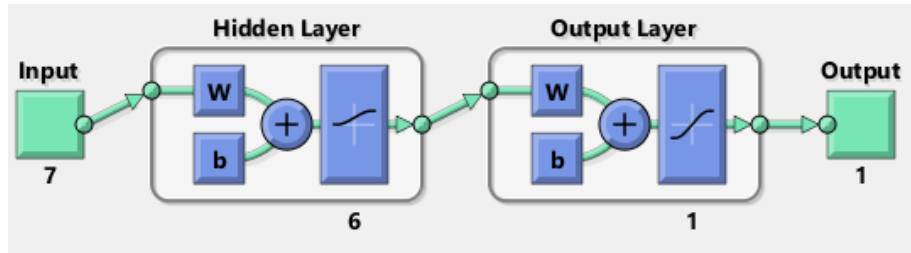


Figure 4.12: The neural network design for SelectMnet

Figure 4.12 shows the neural network design for SelectMnet. There are seven input features, which are the seven selected bones using GA approach while the hidden layer consists of the weights and biases and also six neurons. There are weights and biases and also one neuron for the output layer and one output feature will be produced.

After training all of the networks, the outputs and errors for each network are exported to the workspace in Matlab.

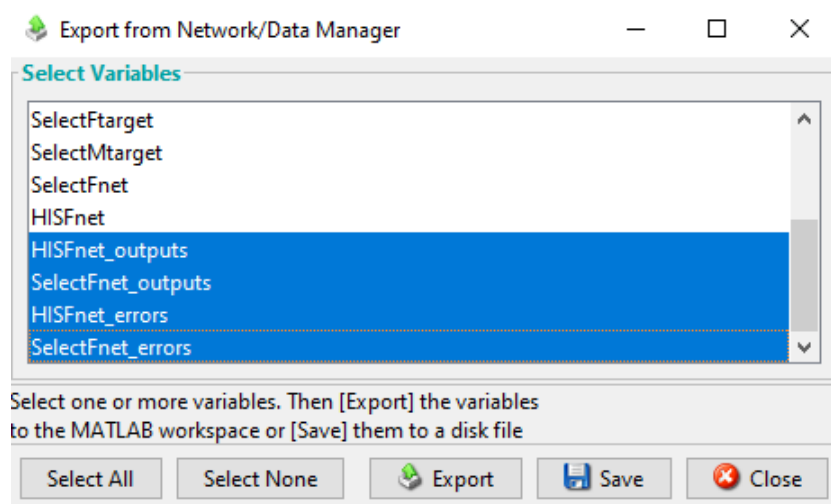


Figure 4.13: Example of exporting the outputs and errors of one of the neural networks

Figure 4.13 shows that HISFnet_outputs, SelectFnet_outputs, HISFnet_errors and SelectFnet_errors are going to be exported to workspace in Matlab. The errors and

outputs of the training for all networks will be exported to the workspace in Matlab as it contains the results of the training.

Below are the commands to extract the data based on the generated indices.

```
>> i = 1
```

Figure 4.14: The command to initialize the variable of a for-loop in order to extract the data based on generated indices

The command in Figure 4.14 is entered in the command window in Matlab to initialize the variable of a for-loop in order to extract the data based on generated indices.

```
>> for i = 1:19  
HISFtrain(i,:) = HISF(i, [trainInd]);  
end
```

Figure 4.15: The for-loop command and also command to extract the data based on generated indices from a specific data

The commands in Figure 4.15 are entered in the command window in Matlab to execute the for-loop and to extract the data based on generated indices from a specific data.

4.2.2 HYBRID GENETIC ALGORITHM-ARTIFICIAL NEURAL NETWORK

Below is the command to activate the Miniconda environment to enable the use of Python.

```
>conda activate yeemayfypenv
```

Figure 4.16: Example of command to activate Miniconda environment

The command in Figure 4.16 is entered in the Command Prompt to activate Miniconda environment named 'yeemayfypenv'.

Below is the command to run the Python script to select features for female Hispanic population.

```
>python GAffemale.py
```

Figure 4.17: Example of command to run Python script

The command in Figure 4.17 is entered in the Command Prompt to run the Python script named 'GAffemale.py' in order to carry out the process of feature selection on the data of input features and target feature of female Hispanic population using GA approach. Another Python script named 'GAmale.py' is run to select features for male Hispanic population.

The script will generate a graph showing the best scores and the average scores throughout 200 generations.

Below is the graph for the results of GA approach for 200 generations.

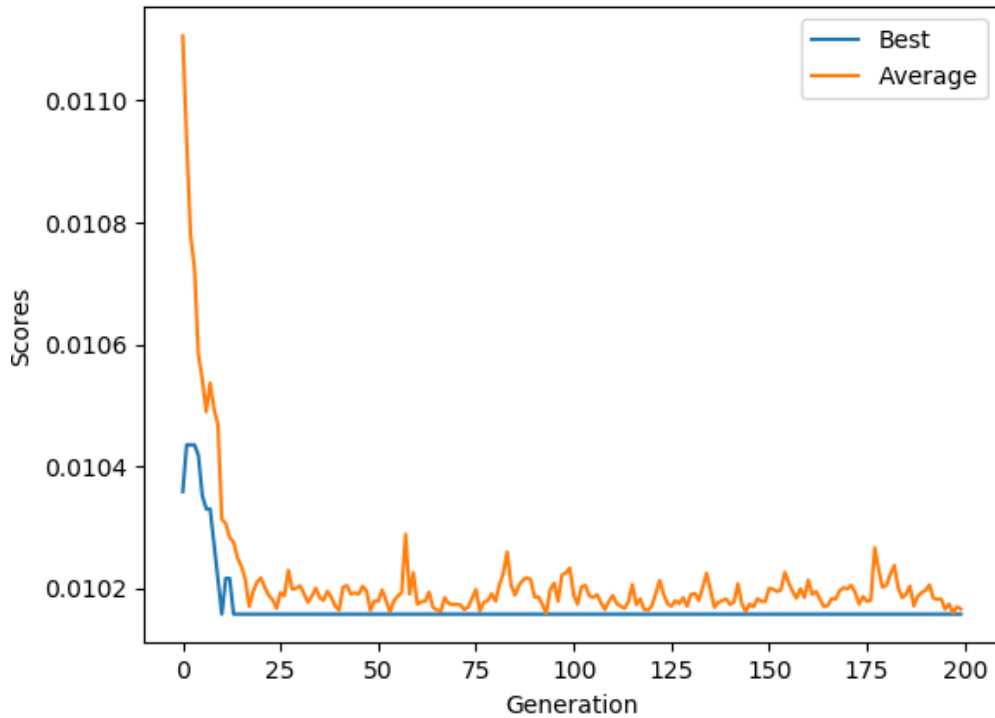


Figure 4.18: A generated graph of the best scores and the average scores throughout 200 generations

The x-axis represents the generation during the training of the GA approach while the y-axis represents the cross-validation MSE score. The blue line represents the best cross-validation MSE score that was obtained in that particular generation while the orange line represents the mean cross-validation MSE score among 200 chromosomes in that generation. As observed, the cross-validation MSE scores are decreasing with each generation, which implies an increase in performance for the GA. The 200th generation, which is the last generation, has displayed similar scores between the best and mean scores. This means that the majority of the chromosomes in the 200th generation have high fitness scores.

The script will generate the cross-validation MSE values before feature selection and after feature selection.


```
(yeemayfypenv) C:\Users\MS\Documents\FYP_YM\GA_dataset>python GAfemale.py
CV MSE before feature selection: 0.01111391
CV MSE after feature selection: 0.01015822
```

Figure 4.19: The generated cross-validation MSE values before and after the feature selection throughout 200 generations

Figure 4.19 shows that the generated cross-validation MSE values before and after the feature selection throughout 200 generations for female Hispanic population.

A CSV file of the data of latest population will be generated by running the script.

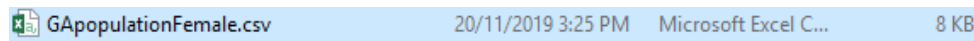


Figure 4.20: A generated CSV file of the data of latest population

The CSV file contains 200 chromosomes in the last generation, which are the fittest individuals in the GA approach. Each chromosome will have value of 0 or 1 for each bone, where 1 represents the bone has been selected after the process of feature selection while 0 represents the bone is not being selected. This CSV file is used to analyse which features are selected during the execution of the GA approach.

The implementation of part of the ANN approach of GA-ANN hybrid approach is the same as the ANN approach discussed in Section 4.2.1.

4.3 RESULTS

Below are the cross-validation (CV) MSE results of female and male Hispanic population before feature selection and after feature selection for 200 generations.

	Female		Male	
	Before feature selection	After feature selection	Before feature selection	After feature selection
CV MSE	0.01111391	0.01015822	0.00896640	0.00836916

Table 4.3: The CV MSE results of female and male Hispanic population before and after feature selection for 200 generations

Before the feature selection, the CV MSE value of female Hispanic population is 0.01111391 and after using the GA approach for feature selection, the CV MSE value has dropped to 0.01015822. For male Hispanic population, the CV MSE value before feature selection is 0.00896640 while after the feature selection using GA approach, the CV MSE value is dropped to 0.00836916. The smaller the MSE, the more preferable or desirable selection it is as it shows the data values are dispersed closely to its central moment. Therefore, GA approach has improved the performance of the CV MSE of both genders of Hispanic population.

There are 200 individuals for both female and male Hispanic population after feature selection. Each individual will have a chromosome consisting of 0 and 1. The 0 implies the feature is not selected while the 1 implies the feature is selected. Below is the total number of 1 (s) for each bone across 200 individuals for female and male Hispanic population.

Female		Male	
Bone1	0	Bone1	200
Bone2	0	Bone2	0
Bone3	0	Bone3	0
Bone4	190	Bone4	0
Bone5	200	Bone5	0
Bone6	200	Bone6	0
Bone7	0	Bone7	190
Bone8	0	Bone8	0
Bone9	0	Bone9	0
Bone10	0	Bone10	0
Bone11	200	Bone11	0
Bone12	0	Bone12	0
Bone13	0	Bone13	200
Bone14	0	Bone14	200
Bone15	180	Bone15	200
Bone16	200	Bone16	200
Bone17	0	Bone17	0

Bone18	200	Bone18	0
Bone19	195	Bone19	175

Table 4.4: The sum of 1 (s) for each bone across 200 individuals for female and male Hispanic population

The features which consists more than 85% of the sum of 1 (s) after will be selected. Therefore, Bone 4, Bone 5, Bone 6, Bone 11, Bone 15, Bone 16, Bone 18, and Bone 19 are the selected features across 200 individuals for female Hispanic population. Bone 1, Bone 7, Bone 13, Bone 14, Bone 15, Bone 16, and Bone 19 are the selected features across 200 individuals for male Hispanic population.

After exporting the outputs of the all networks, the errors are used to confirm the validity of the testing target data and the testing output data. When both data are correct, the testing target data and the testing output data are denormalized using the age range of zero to 18. The denormalization formula is the inverse of the normalization formula. The test errors are calculated for each individual by subtracting the testing target data with the testing output data. Table 4.5 records the testing target data, testing output data and the test set MSE for all four networks.

Female						Male					
ANN			GA-ANN			ANN			GA-ANN		
Target	Output	Error	Target	Output	Error	Target	Output	Error	Target	Output	Error
1	1.35528	-0.35528	1	1.602258	-0.60226	0	2.279791	-2.27979	0	1.880664	-1.88066
1	1.93271	-0.93271	1	2.00224	-1.00224	2	2.187777	-0.18778	2	2.587926	-0.58793
2	2.651318	-0.65132	2	3.131187	-1.13119	2	3.003875	-1.00387	2	2.576813	-0.57681
2	2.450083	-0.45008	2	2.376769	-0.37677	3	4.046295	-1.04629	3	4.491861	-1.49186
2	2.032009	-0.03201	2	1.650577	0.349423	3	2.507708	0.492292	3	2.696342	0.303658
5	4.825922	0.174078	5	4.905369	0.094631	4	5.990798	-1.9908	4	4.981924	-0.98192
6	5.177644	0.822356	6	5.339039	0.660961	4	2.276225	1.723775	4	2.432137	1.567863
6	4.960624	1.039376	6	4.966668	1.033332	5	5.596441	-0.59644	5	6.465045	-1.46505
6	5.264418	0.735582	6	5.669032	0.330968	5	7.464246	-2.46425	5	5.459937	-0.45994
6	5.869192	0.130808	6	5.740944	0.259056	5	5.223394	-0.22339	5	6.074604	-1.0746
7	5.891033	1.108967	7	5.283911	1.716089	5	5.56253	-0.56253	5	6.148144	-1.14814
7	8.978694	-1.97869	7	7.027829	-0.02783	6	5.728094	0.271906	6	6.050651	-0.05065
7	6.808137	0.191862	7	7.618726	-0.61873	6	5.842269	0.157731	6	6.625014	-0.62501
7	5.812521	1.187479	7	6.843336	0.156664	7	8.583953	-1.58395	7	7.879906	-0.87991
7	8.446568	-1.44657	7	8.578115	-1.57811	7	7.168705	-0.16871	7	6.36083	0.63917
8	7.162575	0.837425	8	6.572998	1.427002	7	8.315154	-1.31515	7	7.965232	-0.96523
8	7.815088	0.184912	8	8.388447	-0.38845	7	7.174108	-0.17411	7	7.029521	-0.02952
9	9.86298	-0.86298	9	9.440739	-0.44074	8	8.313136	-0.31314	8	9.095267	-1.09527
9	12.58331	-3.58331	9	11.50172	-2.50172	8	6.807768	1.192232	8	8.806426	-0.80643
9	11.63552	-2.63552	9	11.55004	-2.55004	8	5.420717	2.579283	8	7.087681	0.912319
9	12.23898	-3.23898	9	12.28615	-3.28615	8	9.409963	-1.40996	8	8.851337	-0.85134
9	7.797965	1.202035	9	9.090821	-0.09082	9	9.895757	-0.89576	9	9.615734	-0.61573
10	10.26316	-0.26316	10	10.80231	-0.80231	9	12.21161	-3.21161	9	10.72366	-1.72366
10	12.17328	-2.17328	10	13.02223	-3.02223	9	11.80746	-2.80746	9	11.43072	-2.43072
10	9.582207	0.417793	10	9.864999	0.135001	10	8.486977	1.513023	10	8.458489	1.541511
11	9.241172	1.758828	11	9.40432	1.59568	10	10.39876	-0.39876	10	10.69318	-0.69318
11	12.98901	-1.98901	11	12.20051	-1.20051	10	11.58187	-1.58187	10	11.03624	-1.03624
11	12.97893	-1.97893	11	13.41669	-2.41669	11	9.799663	1.200337	11	10.67472	0.325278
11	13.73625	-2.73625	11	13.32999	-2.32999	11	11.36402	-0.36402	11	11.90709	-0.90709
11	12.65805	-1.65805	11	12.87533	-1.87533	12	14.71938	-2.71938	12	14.66741	-2.66741
11	11.94189	-0.94189	11	12.37348	-1.37348	12	12.37906	-0.37906	12	11.87477	0.12523

11	12.25132	-1.25132	11	12.49909	-1.49909	12	13.07592	-1.07592	12	12.85053	-0.85053
12	14.92424	-2.92424	12	15.19039	-3.19039	12	12.56777	-0.56777	12	13.00823	-1.00823
12	11.99764	0.002363	12	10.99238	1.007624	12	11.74689	0.253114	12	11.80488	0.195123
13	13.74639	-0.74639	13	12.98272	0.017276	12	13.51462	-1.51462	12	13.68316	-1.68316
13	13.99543	-0.99543	13	13.50413	-0.50413	13	12.88582	0.114182	13	12.79072	0.209278
13	13.33478	-0.33478	13	13.19567	-0.19567	13	14.9704	-1.9704	13	15.13407	-2.13407
13	14.15341	-1.15341	13	14.43896	-1.43896	13	12.01156	0.988439	13	13.97713	-0.97713
13	14.30844	-1.30844	13	14.5066	-1.5066	13	14.08809	-1.08809	13	14.01194	-1.01194
14	14.56199	-0.56199	14	14.57401	-0.57401	13	13.42562	-0.42562	13	12.08177	0.918231
14	15.04137	-1.04137	14	14.66595	-0.66595	13	15.50421	-2.50421	13	14.16714	-1.16714
15	12.80126	2.198745	15	12.34801	2.651988	13	14.07654	-1.07654	13	13.41905	-0.41905
15	15.24468	-0.24468	15	15.21399	-0.21399	14	15.93439	-1.93439	14	15.26242	-1.26242
15	14.48925	0.51075	15	13.73032	1.269677	14	14.01759	-0.01759	14	14.37134	-0.37134
16	14.72155	1.278452	16	14.18035	1.819646	14	13.30549	0.694506	14	13.44556	0.554436
17	13.68745	3.312554	17	13.41085	3.589153	15	15.21143	-0.21143	15	14.95085	0.049152
17	14.49464	2.505358	17	14.78106	2.218936	15	15.54973	-0.54973	15	15.39551	-0.39551
17	12.86728	4.132715	17	12.58131	4.418687	16	14.93511	1.064889	16	14.93419	1.065813
17	14.60787	2.392127	17	14.31431	2.685692	16	14.5612	1.4388	16	14.40863	1.591374
18	14.61873	3.381266	18	14.70434	3.295661	17	16.32826	0.671738	17	15.4769	1.523102
18	15.29742	2.70258	18	15.24403	2.755965	17	15.10961	1.89039	17	15.07638	1.923624
18	13.63488	4.365124	18	14.26258	3.737422	17	14.92314	2.076855	17	15.29023	1.709767
18	15.1634	2.836596	18	15.45068	2.549322	17	15.26041	1.739593	17	15.30366	1.696342
18	13.89982	4.100182	18	14.28753	3.712467	18	14.75171	3.248289	18	14.36874	3.631262
18	12.83249	5.16751	18	12.67444	5.325557	18	13.81731	4.182687	18	15.30516	2.694836

Table 4.5: The testing target data, testing output data and the test set MSE for all four ANNs

The test set MSE value, which is the mean of the test errors for that particular network, is then calculated for all four networks. The formula for the calculation of the test set MSE can be found in Equation 2.12 in chapter 2.

Below are the results of age estimation models using ANN approach and GA-ANN hybrid approach for both female and male Hispanic population.

	Female		Male	
	ANN	GA-ANN	ANN	GA-ANN
Test Regression (r value)	0.90686	0.9086	0.9426	0.96226
Test set MSE	4.096813	4.060201	2.429385	1.702814

Table 4.6: The results of age estimation models using ANN approach and GA-ANN hybrid approach for both female and male Hispanic population

The test regression value and the test set MSE are used as metrics for evaluation. The training regression value will always improve as the number of the data used for training increases, which makes the training regression value not a reliable metric. The overall regression value is also not selected as it is calculated to be the mean between the training and test regression values. The test set MSE is the MSE calculated on the test data only, without regard to the training data.

In the results shown in Table 4.6, the performance of the hybrid GA-ANN approach is better than the ANN approach. For the female networks, the test regression value for hybrid GA-ANN approach, 0.9086 is slightly higher than the test regression value for ANN approach, 0.90686. For the male networks, the test regression value for hybrid GA-ANN approach, 0.96226 is higher than the test regression value for ANN approach, 0.9426. The lower the test set MSE value, the less errors, which implies a better performance of the network. For the female networks, the test set MSE for hybrid GA-ANN approach, 4.060201 years is slightly lower than the ANN approach, 4.096813 years. For the male networks, the test set MSE for hybrid GA-ANN approach, 1.702814 years is significantly lower than the ANN approach,

2.429385 years. Overall, there is a better performance in the hybrid GA-ANN approach compared to the ANN approach for networks of both gender, which produces a higher test regression value and a lower test set MSE.

4.4 DISCUSSION AND JUSTIFICATION

Adaption learning function, transfer function, cross-validation, and test set MSE are discussed in the following sections.

4.4.1 ADAPTION LEARNING FUNCTION

Learning function	Layer 1	Layer 2	Female		Male	
			Test regression (R)	MSE	Test regression (R)	MSE
learnGD	purelin	tansig	0.94419	0.00595	0.91708	0.00514
learnGDM	purelin	tansig	0.94419	0.00595	0.91708	0.00514

Table 4.7: The comparison results of using different learning functions for neural networks

The learning function will compute the learning rate for different parameters, calculate the weight change during training, ultimately improving the ANN. The comparison above is done after dividing the dataset into 70% training data and 30% testing data, in which the female neural networks use the same training and testing data and the male neural networks use the same training and testing data. The performance of the female neural networks are the same whereas the performance of the male neural networks are the same too, despite the differences of the adaption learning function. Since most researches uses learnGDM, the learnGDM learning function is chosen for this paper.

4.4.2 TRANSFER FUNCTION

Layer 1	Layer 2	Female		Male	
		Test regression (R)	MSE	Test regression (R)	MSE
tansig	logsig	0.5005	0.0143	0.64528	0.0146
tansig	tansig	0.38641	1.21e-10	0.69575	1.61e-10
tansig	purelin	0.56364	1.11e-23	0.78547	2.49e-22
logsig	logsig	0.42071	0.0143	0.58935	0.0146
logsig	tansig	0.16697	2.87e-11	0.476	6.05e-11
logsig	purelin	0.67053	7.79e-17	0.81512	9.96e-16
purelin	logsig	0.72728	0.0183	0.75804	0.0162
purelin	tansig	0.9152	0.00549	0.91076	0.00490
purelin	purelin	0.91394	0.00534	0.90863	0.00431

Table 4.8: The comparison results of different combinations of transfer functions for Layer 1 and Layer 2 for several tests of neural networks

Transfer function, also known as activation functions help neural networks to learn non-linear relationships by applying the transfer function to produce the final output. In Table 4.8, the combination of purelin and tansig of transfer functions has the highest performance with the highest test regression values, which are 0.9152 for female and 0.91076 for male. Since any MSE value is considered to be valid when it is less than 2, the MSE value of 0.00549 years for female and 0.00490 years for male are acceptable. Even though there are MSE values that are even lower than the purelin and tansig combination, the difference between the test regression values can be quite large, which becomes the reason for elimination.

4.4.3 CROSS-VALIDATION

Cross-validation aims to test the ability of the result to generalize to an independent dataset. With cross-validation, a validation data set can be used to test the model during training. Problems such as underfitting and overfitting can be limited through cross-validation (scikit-learn developers, 2014). In the GA approach, cross-validation is used when selecting the input features to generate the best features. Five-fold cross validation has been used in the GA approach, where the training dataset is

divided into groups of five chromosomes. The first four chromosomes are used for training, while the fifth chromosome will be used for validation. This process repeats for all chromosome groups of five.

4.4.4 TEST SET MEAN SQUARED ERROR (MSE)

Errors are the difference between the expected output and the actual output, which have been one of the metrics to evaluate the accuracy of the data. The theory is that the less errors are generated, the more accurate the model is. The test set MSE is the average of the test errors for that particular network (Rowe, 2015). It is calculated after denormalization of the test target data and test output data, where the actual age predicted by the ANN can be obtained. The test set errors is then calculated by using the denormalized test target data and denormalized test output data. The average of the sum of test errors will then be calculated, obtaining the test set MSE. Microsoft Excel has been used for the denormalization until the calculation of the test set MSE.

4.5 SUMMARY

In this Chapter 4, the overall performance of the GA-ANN hybrid approach is better comparative to the ANN approach, with the GA-ANN hybrid approach producing a higher test regression value and a lower test set MSE value. The feature selection introduced in the GA-ANN hybrid approach will have two main advantages, which are a better performance and the results is obtained and less data is required. An increase in performance will enable the neural network to produce more accurate results. The feature selection for this GA-ANN hybrid approach has reduced the input features required from 19 bones to seven or eight, which is more than 50% of the data, resulting in an optimization of the effort in procuring training data.

CHAPTER 5

CONCLUSION AND RECOMMENDATION

5.1 INTRODUCTION

This chapter concludes the paper, where the project constraints and the future work of this research is discussed.

5.2 PROJECT CONSTRAINT

Below are the project constraints faced during this research:

- Lack of readily available data

The training dataset used for the training of both ANN approach and GA-ANN hybrid approach originated from X-ray images that does not have any measurements of the 19 bones. The measurements were obtained manually by using Photo Pos Pro, which is a tiring process. Too much time was used to acquire the measurements.

- Lack of good hardware

The research was carried out on a laptop with 8 GB of RAM. However, the laptop does not have the suitable technical specifications to support Photo Pos Pro, and the application would have latency during use. There are times when the laptop has shut down on its own, causing loss of measurement data during the procurement of measurements of the 19 bones.

5.3 RECOMMENDATION AND FUTURE WORK

For this research, more training data can be used to produce more accurate results, as the data will not too biased since there are more variety of data. The increase in the amount of training data can also evaluate the performance of the feature selection.

Other bones, such as dental and right hand bones can be used as a combination with the left hand bones for the Hispanic population to produce even more accurate age estimations. This can also increase the robustness of the GA-ANN hybrid approach as it receives more bones to estimate the age.

Other ethnicity should also be used, such as Caucasian or Native Americans. The results of the feature selection for the 19 bones should be different due to biological factors, which means that it is possible that the weightage for the 19 bones may be different compared to another ethnic group during age estimation.

More neural networks can be integrated with the approach. For example, two GA approaches can be integrated with the ANN approach to possibly improve the feature selection. The possibility of an approach with a comparatively more robust feature selection should be tested.

Another approach for the feature selection other than the GA approach can be tested to see if it performs well. There is a possibility where another approach can produce better results during feature selection.

5.4 CLOSING NOTE

In this paper, the measurements of the 19 bones for the Hispanic population have been made to form the general dataset. The dataset is then normalized to avoid computing errors when fed into the ANN.

The ANN approach involves using all 19 bones as input features to estimate the age of the Hispanic population. Since the age estimation will be different based on the gender of the individual, two ANN approaches are created for male and female.

The GA-ANN hybrid approach will use the GA approach to perform feature selection, which reduces the number of input features. The selected input features will then be fed into the ANN approach to produce the expected output data and actual output data.

The expected output data and the actual output data are denormalized to obtain the estimated age and the actual age. The difference between the two ages, which are known as errors are then computed to evaluate the performances of both approaches.

The future work and recommendations are then proposed at the end of this paper.

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APPENDIX A

ORIGINAL DATASET OF FEMALE HISPANIC POPULATION

Image ID	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	Age
5584	1.46	2.48	2.33	1.98	1.83	1.13	1.56	1.73	1.61	1.29	0.81	1.15	1.03	0.64	0.76	0.5	0.59	0.67	0.57	0
5068	1.69	3.16	2.93	2.49	2.13	1.3	1.75	1.96	1.88	1.49	1.07	1.23	1.2	0.92	1.05	0.62	0.7	0.77	0.57	1
5070	1.48	2.35	2.28	2	1.74	1.11	1.51	1.82	1.76	1.29	0.87	1.18	1.23	0.76	0.81	0.64	0.68	0.73	0.57	1
5102	1.67	2.59	2.35	2.47	2.08	1.09	1.75	1.9	1.76	1.5	0.78	1.19	1.01	0.67	0.87	0.52	0.67	0.68	0.55	1
5111	1.39	2.2	2.1	1.86	1.62	1.09	1.51	1.68	1.65	1.23	0.85	1.07	1.03	0.63	0.81	0.63	0.67	0.67	0.57	1
5585	1.65	2.97	2.78	2.48	2.2	1.21	1.88	1.99	1.98	1.58	0.84	1.12	1.17	0.65	1.01	0.58	0.65	0.74	0.49	1
5017	1.93	3.45	3.3	2.92	2.72	1.44	2.04	2.28	2.38	1.6	1.29	1.55	1.48	0.89	1.2	0.9	1.01	0.99	0.72	2
5036	1.68	3.08	2.84	2.59	2.27	1.22	1.68	1.94	1.83	1.35	0.99	1.11	1.13	0.6	0.86	0.66	0.7	0.74	0.57	2
5107	1.56	2.99	2.88	2.35	2.11	1.18	1.83	2.09	2	1.29	0.89	1.19	1.23	0.65	1	0.57	0.61	0.57	0.56	2
5117	1.88	3.08	3	2.74	2.29	1.29	1.81	2.07	1.96	1.48	1.15	1.42	1.41	0.79	0.93	0.73	0.77	0.87	0.7	2
5588	1.68	3.14	3.02	2.61	2.34	1.32	2	2.26	1.96	1.51	1.18	1.49	1.37	0.85	0.97	0.65	0.94	0.8	0.63	2
5061	2.19	3.65	3.47	3.04	2.68	1.6	1.94	2.26	2.33	1.63	1.3	1.6	1.53	0.97	1.15	0.62	0.89	0.72	0.64	3
5065	2.67	3.81	3.64	3.36	3.07	1.55	2.19	2.5	2.34	1.69	1.24	1.52	1.49	0.85	1.29	0.78	0.83	0.87	0.8	3
5067	2.59	3.9	3.67	3.3	3.05	1.73	2.18	2.47	2.57	1.74	1.3	1.64	1.55	0.86	1.46	0.98	1.13	1.14	0.78	3
5090	2.06	3.96	3.82	3.38	3	1.41	2.01	2.31	2.18	1.56	1.36	1.65	1.53	1.04	1.28	0.66	0.84	0.84	0.66	3
6137	2.15	3.84	3.74	3.39	3.21	1.59	2.27	2.55	2.39	1.84	1.23	1.68	1.55	0.91	1.22	0.59	0.77	0.8	0.71	3
5109	2.33	4.04	3.92	3.56	3.09	1.66	2.27	2.88	2.72	2.11	1.33	1.71	1.56	1.25	1.3	0.76	0.79	1.07	0.69	4
5120	2.35	3.91	3.92	3.45	3.2	1.63	2.24	2.58	2.36	1.75	1.22	1.57	1.49	0.82	1.43	0.91	0.98	1.04	0.85	4
5140	2.54	3.83	3.6	3.09	2.96	1.44	2.02	2.34	2.21	1.63	1.3	1.46	1.57	0.99	1.21	0.79	0.78	1.02	0.71	4
5269	2.14	3.84	3.7	3.33	3.02	1.6	2.16	2.46	2.56	1.95	1.42	1.66	1.44	1.01	1.32	1.01	1.13	1.16	0.81	4
6130	2.32	4.06	3.88	3.51	3.16	1.66	2.28	2.61	2.78	2.16	1.33	1.71	1.68	1.18	1.32	0.72	0.85	0.88	0.73	4
5050	2.59	4.21	4.16	3.8	3.45	1.68	2.4	2.87	2.61	1.94	1.3	1.74	1.61	0.98	1.23	0.96	1.06	1.12	0.96	5
5133	3.08	4.43	4.43	3.87	3.59	2.05	2.57	2.88	2.74	2.15	1.37	1.72	1.68	1.25	1.51	0.85	0.97	1.01	0.85	5
5141	2.48	4.49	4.24	3.81	3.57	1.64	2.42	2.71	2.56	1.86	1.18	1.57	1.45	0.84	1.08	0.81	0.87	0.94	0.81	5
5143	2.94	4.41	4.36	3.95	3.54	1.64	2.29	2.97	2.81	2.17	1.47	1.82	1.75	1.11	1.18	0.88	1	0.99	0.81	5
5156	2.2	3.92	3.86	3.38	3	1.93	2.18	2.45	2.59	1.78	1.47	1.56	1.46	1.15	1.45	0.84	0.89	0.9	0.8	5

7038	2.16	4.17	4.08	3.54	3.15	1.36	2.46	2.7	2.5	1.75	1.38	1.57	1.57	1.01	1.43	0.8	0.85	0.9	0.76	5
7039	2.87	4.32	4.38	3.88	3.5	1.79	2.59	3.05	2.9	2.13	1.29	1.76	1.89	1.28	1.59	0.87	0.94	0.96	0.88	5
7040	2.76	4.29	4.23	3.71	3.19	1.57	2.53	2.87	2.71	2.01	1.41	1.74	1.64	1.02	1.4	0.88	1.08	0.97	0.77	5
7121	2.66	4.01	3.93	3.51	3.17	1.54	2.26	2.58	2.44	1.82	1.25	1.53	1.54	1.03	1.22	0.78	1	1.06	0.81	5
7244	2.2	4.08	4.01	3.51	3.23	1.56	2.23	2.63	2.72	2.1	1.4	1.71	1.64	1.03	1.3	0.79	0.8	1.01	0.69	5
5056	2.51	4.39	4.33	3.82	3.41	1.77	2.59	2.9	2.8	1.92	1.61	1.94	1.81	1.15	1.49	1.07	1.11	1.16	0.85	6
5057	3.21	4.84	4.75	4.15	3.75	1.93	2.79	3.57	3.29	2.49	1.66	1.94	2.07	1.32	1.57	0.87	1.01	1.23	0.72	6
5079	3.04	4.54	4.5	3.85	3.55	1.87	2.71	3.09	2.86	1.95	1.47	1.67	1.56	1.14	1.47	0.87	1.16	1.05	0.86	6
5142	3.05	4.37	4.24	3.85	3.56	1.89	2.54	2.84	2.74	2.07	1.55	1.71	1.75	1.18	1.2	0.93	1.2	1.23	1.06	6
5343	3.13	4.44	4.26	3.81	3.47	1.83	2.51	3.11	2.94	2.19	1.47	1.9	1.81	1.16	1.56	0.9	1.05	1.06	0.89	6
7041	2.97	4.42	4.13	3.8	3.55	1.64	2.56	2.94	2.79	1.95	1.46	1.77	1.67	1.15	1.41	0.92	0.98	1.23	0.9	6
7042	2.54	4.5	4.3	4.01	3.65	1.75	2.62	3.24	2.97	2.31	1.55	1.97	1.84	1.11	1.57	0.94	1.05	1.09	0.91	6
7043	2.97	4.15	3.93	3.57	3.25	1.99	2.65	2.98	2.86	2.17	1.31	1.69	1.65	1.25	1.48	0.91	1.05	1.06	0.92	6
7254	3.03	4.4	4.26	3.81	3.6	1.82	2.5	3.16	3.02	2.25	1.53	1.93	1.87	1.16	1.48	0.98	1.1	1.11	1.07	6
7278	2.97	4.5	4.34	3.83	3.56	1.99	2.76	3.15	2.98	2.1	1.46	1.86	1.91	1.29	1.64	0.96	1.09	1.2	0.91	6
5136	3.49	5.06	5.04	4.41	3.93	1.87	2.67	3.5	3.15	2.31	1.63	2.02	1.59	0.98	1.28	0.91	0.99	1.27	0.87	7
5137	3.37	5.33	5.22	4.69	4.18	2.2	2.92	3.24	3	2.35	1.86	1.99	2.18	1.47	1.76	1.03	1.11	1.19	0.87	7
5145	3.48	4.91	4.76	4.33	3.92	1.92	2.97	3.4	3.13	2.38	1.79	2.08	1.93	1.32	1.66	0.93	1.02	1	1.07	7
5153	3.51	4.79	4.6	4.06	3.81	2.44	2.98	3.45	3.31	2.35	1.94	2.2	2.2	1.54	1.82	1.35	1.19	1.5	1.31	7
5926	2.89	5	4.89	4.03	3.96	2.21	2.9	3.28	2.81	2.12	1.69	2.06	1.91	1.29	1.62	0.99	1.31	1.33	1.16	7
7044	3.37	5.05	4.83	4.3	3.94	2.22	3.04	3.22	3.14	2.12	1.79	2.22	1.98	1.32	1.74	1.15	1.31	1.33	0.94	7
7082	3.26	4.98	4.79	4.34	4.02	2.19	3.02	3.39	3.16	2.41	1.78	2.14	2.02	1.41	1.63	0.91	1.02	1.03	0.87	7
7139	3.01	4.63	4.52	4.04	3.66	2.14	2.58	3.09	2.96	2.09	1.62	2.01	1.89	1.42	1.61	1.03	1.2	1.25	0.87	7
7207	3.14	4.85	4.7	4.11	3.84	1.81	2.96	3.29	3.09	2.05	1.76	2.1	1.93	1.25	1.66	1.24	1.23	1.28	0.88	7
7285	3.47	4.95	4.86	4.25	3.8	2.14	2.85	3.19	2.99	2.33	1.43	2.02	1.95	1.23	1.63	1.04	1.27	1.29	1.17	7
5054	3.77	5.59	5.28	4.64	4.08	2.32	3.23	3.51	3.23	2.34	1.79	2.18	2.12	1.45	1.75	1.2	1.3	1.13	1.01	8
5062	3.53	5.02	4.92	4.44	4.08	2.09	3.19	3.47	3.35	2.62	1.87	2.31	2.26	1.44	1.82	1.34	1.24	1.52	1.32	8
5761	3.3	4.96	4.92	4.31	3.82	2.24	2.98	3.41	3.19	2.41	1.77	2.09	2.09	1.38	1.77	1.09	1.17	1.43	1.25	8
5762	3.5	5.08	4.97	4.39	3.87	2.28	3.06	3.45	3.28	2.46	1.89	2.23	2.12	1.44	1.74	1.18	1.24	1.5	1.05	8
5776	2.82	4.83	4.56	4.08	3.8	2.14	3.03	3.29	3.15	2.49	1.79	2.12	1.98	1.35	1.63	0.95	1.26	1.32	1.18	8
7112	3.31	4.59	4.56	4.07	3.81	2.19	2.77	3.14	2.95	2.3	1.59	1.9	1.8	1.27	1.66	1.2	1.27	1.29	1.22	8
7115	3.39	5.1	5.04	4.53	4.2	2.46	3.26	3.52	3.36	2.65	1.93	2.34	2.19	1.56	1.76	1.13	1.2	1.46	1.1	8
7226	3.48	5.36	5.34	4.66	4.19	2.24	3.07	3.51	3.3	2.41	1.85	2.29	2.17	1.39	1.76	1.04	1.16	1.17	0.98	8
7249	3.6	5.5	5.36	4.82	4.38	2.34	3.09	3.32	3.25	2.59	1.76	2.14	2.1	1.6	1.78	1.07	1.39	1.45	1.2	8

5045	3.69	5.43	5.2	4.54	4.13	2.41	3.28	3.7	3.53	2.77	1.88	2.35	2.26	1.62	1.77	1.07	1.42	1.5	1.11	9
6022	3.93	5.83	5.66	5.05	4.67	2.53	3.4	3.89	3.7	2.77	2.07	2.64	2.48	1.78	1.87	1.37	1.61	1.64	1.41	9
6023	3.7	5.28	5.36	4.9	4.34	2.34	3.15	3.67	3.42	2.69	1.83	2.21	2.12	1.4	1.89	1.42	1.49	1.26	1.36	9
6024	3.4	5.21	5.3	4.69	4.35	2.12	3.37	3.81	3.53	2.77	1.97	2.41	2.23	1.56	1.83	1.4	1.51	1.49	1.34	9
6025	3.95	6	5.82	5	4.59	2.68	3.59	4.13	3.78	2.87	2	2.48	2.38	1.65	1.71	1.3	1.42	1.2	1.27	9
7075	3.35	5	4.74	4.05	3.74	2.1	2.88	3.18	2.92	2.26	1.63	1.98	1.86	1.16	1.54	1.19	1.24	1.23	1.14	9
7081	3.66	5.24	5.32	4.58	4.3	2.37	3.13	3.6	3.31	2.43	1.82	2.27	2.13	1.33	1.82	1.5	1.57	1.61	1.41	9
7100	3.82	5.46	5.56	4.94	4.48	2.01	3.2	3.71	3.64	2.73	1.72	2.34	2.25	1.57	1.97	1.41	1.28	1.4	1.42	9
7224	3.9	5.92	5.7	5.01	4.69	2.65	3.54	3.93	3.74	2.72	2.01	2.47	2.32	1.7	1.94	1.49	1.59	1.53	1.35	9
7284	3.33	4.92	4.95	4.58	4.28	2.22	3.01	3.42	3.26	2.52	1.77	2.2	2.1	1.37	1.73	1.23	1.3	1.36	1.15	9
5702	3.61	5.52	5.61	4.89	4.54	2.48	3.49	3.95	3.67	2.92	2.12	2.51	2.35	1.62	1.99	1.5	1.53	1.51	1.38	10
5707	3.79	5.47	5.36	4.72	4.33	2.48	3.32	3.77	3.55	2.74	1.92	2.34	2.21	1.39	1.94	1.28	1.12	1.51	1.3	10
5717	3.54	5.51	5.4	4.87	4.49	2.37	3.26	3.71	3.52	2.78	1.82	2.15	2.08	1.59	1.78	1.19	0.97	1.37	1.23	10
5725	4.12	5.83	5.55	4.88	4.47	2.79	3.62	4.05	3.83	2.98	2.15	2.68	2.57	1.74	2.08	1.47	1.56	1.62	1.42	10
5726	3.75	5.38	5.35	4.73	4.42	2.47	3.06	3.51	3.23	2.53	1.86	2.27	2.14	1.43	1.82	1.33	1.44	1.43	1.3	10
5728	3.82	5.65	5.5	4.95	4.66	2.58	3.43	3.91	3.72	2.88	1.92	2.43	2.3	1.55	1.84	1.44	1.56	1.56	1.38	10
5754	3.66	5.32	5.12	4.59	4.18	2.21	3.17	3.61	3.45	2.7	1.86	2.37	2.2	1.53	1.83	1.08	1.48	1.5	1.14	10
5763	3.39	5.06	4.95	4.38	3.89	2.14	2.86	3.39	3.19	2.37	1.64	2.09	2.01	1.37	1.77	1.06	1.44	1.45	1.04	10
5773	4.1	5.75	5.63	5.03	4.58	2.65	3.44	3.94	3.78	2.89	2.03	2.53	2.51	1.8	2.01	1.58	1.6	1.66	1.48	10
5777	3.66	5.27	5.04	4.54	4.15	2.36	3.28	3.71	3.48	2.71	1.97	2.39	2.3	1.57	1.77	1.32	1.44	1.5	1.24	10
7134	3.94	5.66	5.71	5.02	4.72	2.58	3.52	3.77	3.66	2.78	2.03	2.45	2.39	1.58	2.13	1.41	1.51	1.62	1.53	10
7189	3.88	5.6	5.45	4.72	4.33	2.59	3.58	3.96	3.66	2.94	2.15	2.55	2.3	1.71	2.07	1.4	1.51	1.53	1.34	10
7237	3.71	5.47	5.33	4.73	4.24	2.48	3.29	3.81	3.44	2.64	1.79	2.3	2.08	1.47	1.88	1.34	1.5	1.55	1.37	10
7283	3.69	5.77	5.61	4.93	4.5	2.49	3.31	3.82	3.56	2.66	1.85	2.36	2.29	1.48	1.9	1.41	1.47	1.43	1.27	10
5703	3.6	5.13	5.04	4.44	4.02	2.36	3.41	3.75	3.56	2.61	1.99	2.43	2.3	1.63	1.89	1.35	1.45	1.53	1.23	11
5708	3.88	5.75	5.38	4.91	4.52	2.57	3.49	3.96	3.75	2.87	1.95	2.43	2.3	1.51	1.7	1.39	1.49	1.59	1.35	11
5709	4	5.9	5.72	4.96	4.61	2.72	3.63	4.15	3.91	2.93	1.94	2.46	2.36	1.48	2.05	1.36	1.49	1.63	1.45	11
5718	4.41	6.08	6	5.45	5.02	3.01	3.94	4.26	4.01	3.15	2.27	2.73	2.73	1.77	1.99	1.4	1.59	1.54	1.43	11
5719	4.22	6.07	5.85	4.94	4.63	2.73	3.59	4.06	3.87	3.1	1.99	2.43	2.39	1.58	1.76	1.46	1.55	1.73	1.37	11
5729	3.97	5.9	5.71	5.11	4.86	2.67	3.55	4.07	3.83	3.01	2.05	2.53	2.38	1.63	1.87	1.49	1.57	1.6	1.45	11
5733	3.94	5.61	5.46	4.75	4.46	2.55	3.52	4	3.79	2.94	1.94	2.49	2.25	1.22	1.99	1.47	1.58	1.71	1.38	11
5741	3.88	5.97	5.77	5.33	4.84	2.54	3.45	3.9	3.76	2.96	1.98	2.39	2.28	1.66	1.94	1.4	1.54	1.51	1.37	11
5745	4.13	5.86	5.62	4.77	4.64	2.84	3.58	3.95	3.69	3.06	1.99	2.49	2.44	1.68	1.88	1.45	1.46	1.55	1.26	11
5747	4.26	6.02	6.05	5.26	4.92	2.87	3.7	4.2	3.88	3.04	2.1	2.57	2.45	1.69	2.11	1.49	1.61	1.64	1.57	11

5908	3.98	5.76	5.62	5.05	4.78	2.61	3.5	4.01	3.8	2.96	2	2.52	2.39	1.61	1.82	1.47	1.54	1.61	1.4	11
7068	4.09	5.87	5.71	5.07	4.63	2.89	3.71	4.14	3.88	3.02	2.14	2.65	2.45	1.81	1.92	1.33	1.49	1.51	1.41	11
7072	3.95	6.18	5.99	5.31	5.01	2.92	3.95	4.4	4.14	3.35	2.21	2.8	2.76	1.87	2.02	1.62	1.74	1.68	1.55	11
7073	4.2	5.81	5.7	5.15	4.76	2.55	3.46	3.86	3.66	2.9	2.15	2.61	2.42	1.73	2.08	1.52	1.68	1.69	1.47	11
7227	3.87	5.62	5.43	4.89	4.44	2.61	3.32	3.74	3.49	2.81	1.97	2.31	2.18	1.63	1.82	1.36	1.52	1.53	1.37	11
5710	4.18	6.05	5.84	5	4.71	2.76	3.7	4.15	3.98	3.02	2.01	2.55	2.38	1.54	1.85	1.42	1.51	1.66	1.39	12
5716	4.09	5.97	5.65	4.94	4.5	2.88	3.73	4.29	3.97	3.09	2.04	2.68	2.47	1.67	1.83	1.4	1.47	1.39	1.38	12
5723	4.21	6.31	6.13	5.61	5.1	2.99	3.98	4.43	4.16	3.21	2.33	2.85	2.68	1.76	2.21	1.6	1.77	1.82	1.28	12
5727	3.77	5.83	5.89	5.11	4.56	2.75	3.59	4.1	3.89	2.98	2.15	2.63	2.59	1.68	1.97	1.43	1.63	1.72	1.45	12
5734	4.02	5.7	5.62	4.65	4.51	2.62	3.54	4.02	3.8	2.96	1.94	2.48	2.35	1.27	2.07	1.45	1.6	1.65	1.38	12
5744	4.15	6.06	5.91	5.37	4.86	2.89	3.71	4.1	3.75	3	2.09	2.64	2.46	1.76	2.11	1.58	1.74	1.74	1.57	12
5746	4.27	6.04	5.78	4.99	4.71	2.86	3.67	4.06	3.78	3.21	2.11	2.6	2.51	1.74	2.09	1.52	1.61	1.62	1.35	12
5758	4.24	6.26	6.16	5.57	4.93	2.98	3.76	4.23	4.09	3.23	2.16	2.76	2.71	1.94	1.99	1.52	1.64	1.62	1.54	12
5765	4.01	6.07	5.83	5.2	4.56	2.76	3.61	4.08	3.82	2.99	1.93	2.48	2.32	1.5	1.77	1.4	1.64	1.53	1.45	12
5769	4.13	6.36	6.24	5.45	5	2.83	3.87	4.35	4.13	3.16	2.12	2.57	2.38	1.75	1.89	1.45	1.57	1.71	1.39	12
5774	4.32	6.56	6.21	5.75	5.24	3.03	3.82	4.25	4.09	3.17	2.1	2.56	2.48	1.69	2.07	1.57	1.74	1.79	1.53	12
7107	4.65	7.15	6.85	6.28	5.77	3.38	4.29	4.68	4.47	3.53	2.59	3.01	2.93	1.95	2.36	1.63	1.83	1.89	1.49	12
7200	3.8	5.56	5.3	4.44	4.08	2.54	3.32	3.94	3.64	2.82	1.92	2.44	2.27	1.35	1.89	1.44	1.5	1.52	1.4	12
7210	3.91	5.98	5.84	5.29	4.66	2.8	3.65	4.17	3.91	3.05	1.96	2.53	2.28	1.6	2.11	1.47	1.61	1.6	1.42	12
7218	4.17	5.87	5.81	5.07	4.8	2.82	3.56	3.96	3.73	2.93	2.09	2.44	2.34	1.69	2.1	1.47	1.6	1.54	1.52	12
5701	3.88	6.05	5.92	5.13	4.58	2.82	3.76	4.24	3.92	3.05	2.33	2.68	2.53	1.78	1.88	1.49	1.63	1.63	1.43	13
5705	4.18	5.94	5.64	5.05	4.36	2.68	3.59	3.94	3.72	2.83	2.09	2.44	2.33	1.47	2.07	1.62	1.71	1.72	1.61	13
5706	4.28	6.13	5.96	5.18	4.91	2.97	3.96	4.45	4.14	3.26	2.36	2.78	2.67	1.97	2.12	1.63	1.79	1.77	1.61	13
5714	3.99	5.85	5.74	5.1	4.7	2.82	3.55	4.14	3.76	2.88	1.89	2.46	2.32	1.51	1.74	1.35	1.44	1.43	1.31	13
5721	4.2	6.01	5.7	4.99	4.65	2.85	3.76	4.11	3.88	3	2.14	2.55	2.42	1.62	2.03	1.7	1.87	1.78	1.6	13
5735	4.02	5.55	5.58	4.77	4.49	2.73	3.56	4.07	3.79	3.05	2.01	2.48	2.33	1.24	2.02	1.52	1.65	1.71	1.42	13
5748	4.28	6.17	6.04	5.19	5.04	2.91	3.84	4.29	3.95	3.08	2.23	2.7	2.52	1.76	2.06	1.61	1.68	1.72	1.51	13
5756	4.24	6.37	5.99	5.35	4.85	2.78	3.81	4.3	4.05	3.2	2.06	2.43	2.4	1.67	2.12	1.63	1.69	1.75	1.51	13
5759	4.2	6.34	6.13	5.43	4.86	3.02	3.81	4.29	4.08	3.25	2.17	2.76	2.71	1.88	1.91	1.49	1.64	1.63	1.46	13
5770	4.16	6.46	6.26	5.51	5.06	2.94	3.81	4.36	4.14	3.17	2.13	2.65	2.49	1.71	1.97	1.49	1.62	1.7	1.39	13
5902	4.11	6.36	6.22	5.66	5.1	2.9	3.8	4.18	3.99	3.07	2.22	2.62	2.51	1.71	2.06	1.6	1.66	1.7	1.51	13
7071	4.23	6.11	5.97	5.25	4.62	2.71	3.58	4.12	3.66	2.82	2.2	2.62	2.39	1.72	2.04	1.48	1.58	1.59	1.39	13
7077	4.05	5.84	5.5	4.81	4.51	2.67	3.72	4.14	3.84	2.9	2	2.46	2.3	1.02	1.85	1.49	1.54	1.6	1.42	13
7247	4.42	6.28	6.08	5.39	4.9	2.86	3.74	4.17	3.89	2.95	2.16	2.55	2.5	1.67	2.06	1.56	1.67	1.67	1.52	13

7252	4.06	6.24	5.95	5.09	4.8	2.93	3.81	4.26	3.96	3.14	2.15	2.66	2.48	1.73	2.11	1.73	1.82	1.85	1.55	13
5715	4	5.84	5.66	5.14	4.69	2.87	3.55	4.17	3.8	2.89	1.95	2.47	2.38	1.51	1.73	1.36	1.4	1.47	1.3	14
5737	4.71	6.59	6.27	5.51	5.22	3.02	3.93	4.28	4.01	3.24	2.3	2.77	2.68	1.78	2.17	1.68	1.77	1.77	1.69	14
5743	4.2	6.18	6.16	5.71	5.14	2.82	3.68	4.13	3.95	3.17	2.15	2.57	2.34	1.8	1.98	1.53	1.62	1.63	1.53	14
5755	3.94	5.92	5.88	5.28	4.78	2.82	3.53	3.96	3.72	2.95	2.04	2.47	2.37	1.75	1.4	1.53	1.7	1.73	1.52	14
5760	4.31	6.37	6.24	5.63	5.09	3.05	3.72	4.2	4.07	3.28	2.2	2.82	2.72	1.86	1.96	1.49	1.61	1.62	1.5	14
5900	4.18	6.08	5.94	5.2	4.77	2.72	3.78	4.25	4.04	3.11	2.08	2.57	2.44	1.65	2.02	1.44	1.58	1.67	1.44	14
5901	4.01	5.76	5.6	5.07	4.54	2.69	3.49	3.91	3.68	2.95	1.89	2.44	2.47	1.7	1.78	1.38	1.37	1.51	1.38	14
5910	4.36	6.4	6.2	5.57	5.13	3	3.92	4.39	4.05	3.17	2.31	2.81	2.63	1.87	2.06	1.68	1.84	1.84	1.62	14
5914	4.58	6.42	6.13	5.36	5.08	3.19	4.11	4.55	4.2	3.31	2.32	2.68	2.56	1.79	2.09	1.66	1.73	1.76	1.51	14
5917	4.44	6.17	6.14	5.45	5.08	2.94	3.87	4.27	3.97	3.12	2.23	2.7	2.51	1.71	2.09	1.61	1.68	1.74	1.52	14
5925	4.74	6.65	6.16	5.32	5.12	3.25	4.24	4.75	4.39	3.37	2.51	3.05	2.9	1.98	1.86	1.56	1.72	1.75	1.43	14
7209	4.22	6.38	6.36	5.66	5.29	2.87	3.71	4.14	3.89	3.04	2.1	2.64	2.45	1.66	2.17	1.55	1.76	1.71	1.57	14
7216	3.86	5.99	5.92	5.41	5	2.8	3.75	4.13	3.8	3.02	2.19	2.61	2.48	1.66	2.01	1.49	1.72	1.62	1.52	14
7264	4.39	6.65	6.44	5.79	5.42	3.15	4.03	4.54	4.29	3.23	2.31	2.78	2.51	1.68	2.24	1.68	1.85	1.93	1.66	14
5039	4.6	6.42	6.49	5.66	5.15	3.01	3.84	4.32	4.06	3.19	2.3	2.66	2.56	1.95	2.07	1.77	1.79	1.82	1.64	15
5905	4.5	6.44	6.47	5.78	5.32	2.92	3.95	4.49	4.09	3.26	2.2	2.67	2.5	1.85	2.04	1.62	1.69	1.75	1.58	15
5909	4.14	6.1	5.89	5.22	5.03	2.74	3.67	4.18	3.94	3.09	2.06	2.59	2.46	1.67	1.83	1.58	1.61	1.66	1.45	15
5911	4.03	5.75	5.54	4.85	4.5	2.67	3.56	4.1	3.85	3.04	2.01	2.5	2.36	1.28	2.04	1.5	1.65	1.72	1.41	15
5913	4.74	6.65	6.41	5.65	5.28	3.06	4.01	4.31	4.02	3.28	2.16	2.7	2.64	1.77	2.14	1.68	1.81	1.76	1.7	15
5915	4.3	6.39	6.2	5.71	5.19	2.82	3.7	4.16	3.98	3.21	2.09	2.57	2.37	1.78	1.95	1.59	1.7	1.69	1.54	15
5919	4.08	5.81	5.27	4.49	4.23	2.73	3.49	3.94	3.59	2.7	2.01	2.39	2.13	1.18	1.76	1.45	1.49	1.57	1.4	15
5922	4.54	6.47	6.33	5.7	5.29	2.99	3.98	4.45	4.18	3.24	2.34	2.83	2.66	1.68	2.02	1.69	1.79	1.86	1.57	15
5923	4.2	6.52	6.35	5.54	5.07	2.85	3.84	4.42	4.17	3.25	2.2	2.58	2.48	1.7	2.04	1.5	1.62	1.68	1.41	15
6026	3.92	5.87	5.63	5.12	4.72	2.98	3.59	4.18	3.96	3.15	2.08	2.58	2.47	1.74	1.97	1.57	1.69	1.69	1.54	15
5030	3.79	5.72	5.57	4.93	4.56	2.84	3.65	4.1	3.8	2.9	2.1	2.59	2.39	1.61	2.05	1.57	1.6	1.57	1.5	16
5058	4.71	6.82	6.68	5.9	5.46	3.15	4.07	4.52	4.19	3.25	2.32	2.78	2.59	1.67	2.23	1.69	1.84	1.78	1.62	16
5103	4.27	6.12	5.98	5.38	4.9	2.81	3.84	4.19	3.93	3.05	2.21	2.62	2.4	1.59	2.07	1.54	1.73	1.73	1.42	16
5139	4.31	6.48	6.42	5.82	5.42	3.02	3.88	4.31	4.12	3.21	2.05	2.62	2.49	1.74	2.06	1.63	1.8	1.84	1.55	16
5906	4.55	6.48	6.5	5.77	5.39	2.92	4.01	4.54	4.1	3.24	2.2	2.73	2.52	1.9	2.08	1.63	1.66	1.81	1.59	16
5916	4.01	6.11	6.07	5.39	4.95	2.59	3.57	4.08	3.86	2.91	1.92	2.33	2.24	1.34	1.9	1.43	1.6	1.66	1.51	16
5918	4.18	6	6.05	5.29	4.95	2.84	3.71	4.3	3.96	2.98	2.07	2.5	2.42	1.51	2.05	1.67	1.73	1.72	1.63	16
5924	4.36	6.47	6.46	5.67	5.15	2.83	4.03	4.62	4.26	3.33	2.22	2.86	2.69	1.75	1.73	1.33	1.74	1.81	1.32	16
6027	4.26	6.17	5.87	5.28	4.73	2.87	3.79	4.19	3.92	3.05	2.15	2.58	2.46	1.66	2.02	1.71	1.84	1.82	1.69	16

6028	4.43	6.34	6.2	5.58	5.11	2.98	3.92	4.42	4.12	3.16	2.34	2.84	2.67	1.86	2.07	1.77	1.85	1.88	1.59	16
5055	4.09	5.68	5.58	4.77	4.67	2.79	3.67	4.2	3.84	3.14	2.02	2.44	2.38	1.84	1.91	1.45	1.65	1.71	1.57	17
5071	3.97	6.24	6.28	5.54	5.03	2.77	3.72	4.17	4	3.1	2.09	2.58	2.52	1.68	1.85	1.52	1.59	1.67	1.46	17
5081	4.23	6.15	6.01	5.38	4.85	2.85	3.72	4.07	3.84	2.99	2.21	2.59	2.54	1.73	1.89	1.49	1.58	1.59	1.46	17
5099	3.98	5.97	5.82	5.23	4.87	2.75	3.64	4.07	3.76	2.89	2.15	2.48	2.46	1.76	2.01	1.57	1.71	1.68	1.54	17
5100	4.27	6.3	5.95	5.39	5.01	3.08	3.89	4.34	3.99	3.25	2.2	2.75	2.59	1.89	1.95	1.58	1.7	1.73	1.55	17
5116	4.21	6.3	6.13	5.69	5.26	3.07	3.84	4.29	4.08	3.16	2.34	2.77	2.65	1.94	2.27	1.59	1.82	1.9	1.61	17
5118	4.08	6	5.66	4.91	4.44	2.7	3.47	4.02	3.63	2.8	1.87	2.43	2.25	1.4	1.87	1.49	1.61	1.62	1.45	17
5126	4.63	6.64	6.56	5.56	5.19	3.27	4.14	4.68	4.29	3.46	2.29	2.95	2.76	2.06	2.42	1.73	1.87	1.92	1.7	17
5268	3.84	5.69	5.61	4.91	4.55	2.76	3.59	4.1	3.82	2.97	2.14	2.54	2.4	1.6	2.04	1.58	1.62	1.56	1.53	17
5920	4.13	6.38	6.17	5.59	4.88	2.8	3.68	4.25	3.98	3.04	2.02	2.56	2.46	1.6	2.14	1.55	1.71	1.69	1.51	17
5086	4.28	6.34	6.31	5.72	5.13	2.82	3.81	4.4	4.11	3.25	2.3	2.69	2.63	1.95	2.31	1.78	1.84	1.91	1.63	18
5092	4.59	6.75	6.47	5.71	5.39	3.27	4.21	4.72	4.34	3.57	2.58	3.07	2.86	2.11	2.38	1.81	1.96	1.94	1.8	18
5104	4.01	5.88	5.83	5.19	4.66	2.78	3.79	4.11	3.88	2.98	2.09	2.62	2.57	1.73	1.98	1.51	1.51	1.61	1.44	18
5110	3.98	5.98	5.77	5.11	4.52	2.52	3.47	4.03	3.73	2.7	1.97	2.46	2.31	1.44	1.92	1.39	1.5	1.53	1.36	18
5112	4.09	5.87	5.86	5.32	4.79	2.93	3.65	4.14	3.85	3.04	2.12	2.63	2.49	1.79	2.26	1.63	1.8	1.86	1.56	18
5271	4.09	6.2	6.01	5.41	4.91	2.85	3.72	4.13	3.83	3.02	2.17	2.73	2.45	1.74	2.19	1.62	1.78	1.73	1.59	18
5273	4.32	6.04	5.79	4.98	4.78	2.92	3.72	4.11	3.86	3.19	2.09	2.59	2.51	1.76	1.81	1.55	1.61	1.65	1.35	18
5274	4.68	6.54	6.43	5.7	5.18	3.29	4.1	4.68	4.34	3.42	2.34	3	2.78	2.04	2.41	1.75	1.87	1.87	1.75	18
5294	4.14	6.18	6.02	5.36	5.08	2.8	3.7	4.22	3.99	3.06	2.02	2.62	2.42	1.74	1.88	1.58	1.63	1.68	1.49	18
5304	3.83	5.77	5.66	4.99	4.6	2.75	3.65	4.12	3.83	2.97	2.14	2.6	2.41	1.58	2.02	1.56	1.58	1.58	1.48	18

APPENDIX B

ORIGINAL DATASET OF MALE HISPANIC POPULATION

Image ID	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	Age
5589	1	1.59	1.52	1.33	1.22	0.79	1.1	1.24	1.19	0.9	0.63	0.82	0.81	0.48	0.57	0.42	0.51	0.47	0.39	0
5590	1.25	1.87	1.87	1.63	1.59	1.04	1.35	1.53	1.47	1.19	0.8	1.04	0.96	0.6	0.75	0.55	0.66	0.67	0.51	0
5591	1.26	2.13	2.01	1.86	1.57	1.01	1.4	1.59	1.5	1.13	0.81	1.01	0.97	0.65	0.77	0.56	0.6	0.68	0.55	0
5592	1.37	2.3	2.13	1.84	1.69	1.02	1.51	1.72	1.67	1.28	0.81	1.02	0.99	0.65	0.7	0.46	0.56	0.51	0.41	0
5272	2.05	3.49	3.27	2.89	2.74	1.51	2.15	2.6	2.44	1.97	1.29	1.62	1.57	1.07	1.04	0.72	0.85	0.85	0.72	1
5538	0.87	1.4	1.34	1.2	1.05	0.68	0.96	1.12	1.02	0.81	0.53	0.67	0.64	0.42	0.46	0.34	0.4	0.41	0.37	1
5540	1.58	2.78	2.47	2.25	1.98	1.2	1.81	2.05	1.81	1.43	0.9	1.12	1.09	0.8	1.06	0.63	0.7	0.72	0.63	1
5593	1.53	2.66	2.45	2.14	1.95	1.18	1.86	2.07	1.99	1.54	0.98	1.26	1.16	0.61	1.07	0.64	0.75	0.78	0.57	1
5594	1.57	2.55	2.45	2.26	1.96	1.28	1.78	1.95	1.86	1.49	0.95	1.19	1.18	0.78	0.91	0.68	0.73	0.76	0.56	1
5542	1.87	2.99	2.73	2.33	2.16	1.4	2.07	2.32	2.21	1.73	1.19	1.46	1.41	0.95	1.12	0.69	0.85	0.87	0.67	2
5543	1.89	3.18	3.04	2.72	2.51	1.49	2.17	2.47	2.36	1.77	1.3	1.61	1.57	1.02	1.16	0.82	0.9	0.96	0.76	2
5544	1.84	3.13	2.98	2.64	2.39	1.34	2.06	2.35	2.21	1.71	1.11	1.37	1.33	0.85	1.2	0.75	0.96	0.93	0.73	2
5596	2.07	3.4	3.31	2.87	2.69	1.41	2.16	2.43	2.33	1.8	1.26	1.54	1.51	0.95	1.12	0.73	0.91	0.93	0.69	2
5598	1.94	3.25	2.94	2.62	2.37	1.42	2.09	2.29	2.19	1.67	1.11	1.37	1.35	0.77	1.21	0.74	0.93	0.94	0.72	2
5270	2.21	3.48	3.24	2.97	2.63	1.49	2.09	2.4	2.29	1.71	1.26	1.47	1.43	0.89	1.17	0.9	0.78	0.94	0.68	3
5545	2.03	3.61	3.49	3.08	2.77	1.48	2.37	2.74	2.65	1.99	1.27	1.72	1.63	1.07	1.24	0.82	0.8	0.89	0.72	3
5546	0.97	1.76	1.68	1.47	1.31	0.83	1.15	1.28	1.21	0.91	0.61	0.81	0.78	0.5	0.62	0.41	0.49	0.5	0.41	3
6118	2.35	3.65	3.38	3.06	2.62	1.76	2.44	2.73	2.56	2	1.29	1.64	1.58	0.97	1.29	0.82	1	1.04	0.92	3
6148	1.99	3.62	3.44	3	2.58	1.79	2.41	2.62	2.43	1.87	1.42	1.66	1.59	0.91	1.33	0.91	1.02	1.03	0.87	3
5038	2.32	3.85	3.8	3.35	3.01	1.65	2.49	2.8	2.59	1.99	1.4	1.69	1.63	1.12	1.37	0.99	1.04	1.03	0.96	4
5105	2.1	3.71	3.55	3.23	2.85	1.58	2.43	2.81	2.66	2.01	1.48	1.86	1.75	1.1	1.35	1.08	1.11	1.19	1	4
5177	2.24	3.84	3.63	3.26	2.98	1.6	2.34	2.62	2.46	1.91	1.34	1.52	1.43	0.94	1.4	1.06	1.14	1.14	1.01	4
5316	2.73	4.63	4.61	4.13	3.73	2.12	2.71	3.17	3.03	2.26	1.48	1.86	1.8	1.17	1.53	1.01	1.21	1.2	0.93	4
6047	1.85	3.42	3.22	2.85	2.44	1.3	1.99	2.23	2.07	1.55	1.18	1.43	1.3	0.69	1.13	0.73	0.82	1	0.66	4
5053	2.77	4.43	4.17	3.73	3.29	1.76	2.54	2.82	2.62	2.03	1.48	1.76	1.7	1.14	1.43	1.06	1.14	1.21	1	5
5085	2.44	3.85	3.66	3.15	2.95	1.58	2.38	2.8	2.62	1.77	1.46	1.8	1.74	1.22	1.39	0.88	0.93	1.19	1.05	5
5106	2.8	4.3	4.08	3.66	3.24	1.83	2.79	3.14	3.01	2.25	1.69	2.09	1.97	1.32	1.63	1.22	1.25	1.3	1.1	5

5165	2.65	3.9	3.6	3.2	2.71	1.78	2.37	2.62	2.46	1.81	1.34	1.61	1.48	0.88	1.37	1.03	1.08	1.1	0.94	5
5166	2.91	4.36	4.27	3.73	3.45	1.83	2.74	3	2.8	2.15	1.58	1.94	1.88	1.34	1.53	1.07	1.21	1.25	1.03	5
7045	2.62	4.49	4.28	4	3.63	2.06	2.94	3.2	3.01	2.36	1.69	2.06	1.98	1.49	1.47	1.04	1.14	1.2	0.98	5
7104	2.82	4.37	4.17	3.67	3.46	1.94	2.71	2.98	2.8	2.24	1.49	1.76	1.73	1.15	1.43	1.07	1.14	1.16	0.96	5
7208	2.52	3.95	3.71	3.2	3.03	1.77	2.3	2.71	2.53	1.89	1.25	1.66	1.61	1.07	1.33	0.93	1.02	1.09	0.91	5
7215	2.38	4.21	3.96	3.47	3.12	1.66	2.58	2.84	2.71	1.99	1.35	1.68	1.61	0.9	1.37	0.94	1.11	1.13	0.95	5
5060	2.66	4.86	4.61	4.12	3.71	2.2	2.87	3.19	3.02	2.34	1.69	2.06	2.07	1.42	1.69	1.22	1.3	1.35	1.19	6
5063	3.31	4.86	4.66	4.12	3.85	2.14	3	3.34	3.19	2.5	1.69	2.04	1.96	1.37	1.7	1.26	1.35	1.37	1.12	6
5098	3.01	4.62	4.67	4.08	3.68	1.87	2.95	3.3	3.1	2.31	1.72	2.09	2	1.43	1.7	1.26	1.36	1.34	1.17	6
5134	2.79	4.52	4.5	3.99	3.54	1.86	2.75	3.09	3.03	2.25	1.56	1.9	1.89	1.29	1.46	1.07	1.16	1.14	1.12	6
5144	2.71	4.02	4	3.53	3.16	1.8	2.38	2.69	2.56	1.97	1.31	1.7	1.68	1.03	1.31	1	1.09	1.11	0.94	6
7046	2.92	4.43	4.35	3.84	3.45	2.03	2.75	3	2.82	2.14	1.5	1.78	1.71	1.16	1.49	1.06	1.22	1.28	1.07	6
7047	2.6	3.93	3.92	3.46	3.08	1.73	2.57	2.91	2.65	2.02	1.48	1.73	1.66	1.08	1.33	0.97	1.08	1.14	1.01	6
7048	2.92	4.63	4.38	3.82	3.46	1.97	2.6	2.93	2.76	2.17	1.54	1.97	1.86	1.29	1.51	1.07	1.18	1.23	1	6
7049	2.97	4.72	4.58	4.09	3.77	2.07	2.78	3.15	2.92	2.41	1.61	1.94	1.86	1.25	1.7	1.17	1.2	1.29	1.12	6
5029	3.31	5.15	5.02	4.44	4.11	2.32	3.04	3.42	3.19	2.48	1.9	2.28	2.08	1.4	1.71	1.23	1.36	1.38	1.1	7
5069	2.86	4.45	4.29	3.81	3.42	2.01	2.75	3.03	2.86	2.17	1.71	2.05	1.89	1.2	1.57	1.05	1.15	1.23	0.99	7
5091	3.1	4.68	4.45	4.05	3.65	2.06	2.75	3.16	2.96	2.2	1.54	1.88	1.85	1.2	1.5	1.04	1.16	1.17	1.07	7
5093	3.01	4.64	4.5	4.06	3.77	2.05	2.83	3.32	3.11	2.33	1.77	2.24	2.12	1.25	1.79	1.27	1.44	1.46	1.22	7
5108	3.03	4.65	4.5	3.89	3.59	2.16	2.84	3.07	2.83	2.29	1.76	2.15	1.96	1.36	1.65	1.18	1.26	1.36	1.19	7
7050	3.2	4.96	4.76	4.26	3.81	2.33	2.95	3.24	3.05	2.38	1.81	2.17	2.04	1.5	1.7	1.34	1.36	1.4	1.22	7
7051	3.21	4.63	4.41	3.92	3.56	2.21	2.66	3	2.89	2.23	1.63	1.94	1.91	1.39	1.64	1.21	1.26	1.3	1.15	7
7099	3.43	4.98	4.95	4.54	4	2.15	2.83	3.29	3.13	2.37	1.64	1.99	1.97	1.31	1.74	1.29	1.4	1.48	1.27	7
7133	3.16	4.75	4.61	3.94	3.66	2.26	3.02	3.32	3.12	2.33	1.74	2.14	2.04	1.46	1.73	1.22	1.35	1.41	1.2	7
7211	2.9	4.39	4.33	3.86	3.32	1.92	2.61	2.96	2.73	2.02	1.49	1.76	1.67	1.03	1.53	1.1	1.17	1.19	1.05	7
5035	3.05	4.73	4.5	4.03	3.68	2.02	2.7	2.99	2.88	2.16	1.59	1.87	1.73	1.18	1.59	1.11	1.24	1.27	1.02	8
5041	3.09	4.84	4.8	4.45	4.13	2.12	2.85	3.16	3.03	2.35	1.62	1.91	1.8	1.13	1.69	1.29	1.32	1.39	1.23	8
5064	3.46	4.9	4.88	4.43	4	2.16	3.12	3.47	3.37	2.53	1.72	2.23	2.1	1.41	1.78	1.23	1.35	1.39	1.23	8
5500	3.45	4.95	4.85	4.27	3.82	2.29	3.16	3.51	3.3	2.52	1.75	2.16	2.1	1.38	1.79	1.31	1.43	1.44	1.24	8
5547	3.15	4.63	4.67	4.17	3.9	2.04	2.99	3.3	3.21	2.5	1.74	2.08	1.99	1.41	1.63	1.25	1.3	1.32	1.16	8
5548	3.42	5.13	4.88	4.09	3.95	2.31	3.09	3.39	3.24	2.58	1.83	2.27	2.11	1.28	1.9	1.32	1.4	1.47	1.3	8
7052	3.09	4.52	4.52	4.04	3.65	2.18	2.88	3.21	3.04	2.35	1.69	2.2	2.06	1.39	1.66	1.23	1.31	1.34	1.17	8
7053	3.42	5.13	5.14	4.48	4.01	2.31	2.97	3.34	3.15	2.39	1.76	2.04	1.84	1.2	1.7	1.31	1.46	1.45	1.23	8
7097	3.12	4.72	4.6	3.96	3.57	2.27	2.89	3.14	3.09	2.39	1.81	2.09	2.02	1.4	1.83	1.26	1.37	1.37	1.15	8

7119	3.39	5	4.89	4.38	4.07	2.12	2.97	3.48	3.27	2.45	1.62	2.02	1.93	1.24	1.78	1.36	1.35	1.41	1.17	8
5023	3.39	5.01	4.99	4.42	3.93	2.21	2.99	3.34	3.17	2.4	1.83	2.1	1.95	1.49	1.77	1.3	1.33	1.38	1.19	9
5119	4.09	5.95	5.66	5.11	4.82	2.68	3.5	3.95	3.68	2.81	2.04	2.51	2.32	1.49	1.94	1.49	1.66	1.63	1.3	9
5135	3.27	4.67	4.54	3.96	3.74	2.06	2.83	3.24	3	2.26	1.55	1.94	1.82	1.07	1.64	1.2	1.3	1.35	1.17	9
5147	3.47	5	5.02	4.56	4.09	2.26	3.05	3.45	3.3	2.48	1.74	2.15	2.06	1.41	2.04	1.32	1.57	1.62	1.34	9
5549	3.84	5.71	5.61	5.06	4.49	2.4	3.08	3.73	3.41	2.42	1.5	2.04	1.89	0.9	1.78	1.27	1.48	1.57	1.18	9
7054	3.55	5.19	5.34	4.65	4.11	2.35	3.3	3.59	3.53	2.66	1.87	2.36	2.43	1.63	1.79	1.34	1.55	1.6	1.36	9
7055	3.25	4.98	4.88	4.2	3.8	2.27	3.05	3.42	3.15	2.42	1.63	2.11	1.92	1.24	1.75	1.2	1.34	1.3	1.16	9
7056	3.61	5.41	5.16	4.54	4.32	2.54	3.31	3.64	3.4	2.67	2.05	2.29	2.17	1.45	2.08	1.53	1.64	1.68	1.49	9
7070	3.66	5.28	5.19	4.48	4.04	2.57	3.18	3.6	3.3	2.63	1.82	2.25	2.16	1.68	1.86	1.34	1.44	1.49	1.35	9
7111	3.34	5.03	5.12	4.47	4.02	2.36	3.03	3.46	3.23	2.58	1.79	2.19	2.1	1.54	1.83	1.34	1.46	1.48	1.29	9
5043	3.15	4.75	4.68	4.19	3.77	2.2	2.86	3.26	3.07	2.38	1.59	1.92	1.87	1.25	1.83	1.38	1.37	1.43	1.26	10
5044	3.74	5.47	5.3	4.73	4.29	2.21	3.27	3.72	3.54	2.72	1.83	2.32	2.27	1.56	1.88	1.38	1.54	1.56	1.37	10
5089	3.7	5.5	5.44	4.78	4.27	2.36	3.29	3.64	3.39	2.59	1.89	2.35	2.21	1.44	1.75	1.21	1.38	1.46	0.98	10
5161	4.11	5.51	5.32	4.61	4.24	2.59	3.34	3.66	3.51	2.79	1.95	2.34	2.25	1.66	1.82	1.36	1.46	1.45	1.35	10
5164	3.43	4.9	4.84	4.41	3.75	2.22	3.09	3.46	3.26	2.32	1.77	2.15	2.09	1.35	1.78	1.36	1.53	1.57	1.3	10
5501	3.65	5.41	5.37	4.78	4.19	2.4	3.33	3.73	3.48	2.62	1.84	2.21	2.09	1.35	2.03	1.39	1.5	1.56	1.42	10
5502	3.14	4.93	4.78	4.26	3.8	2.14	3.04	3.49	3.25	2.48	1.6	2.05	1.92	1.23	1.77	1.21	1.36	1.39	1.17	10
5503	3.5	5.22	4.93	4.34	4	2.34	3.09	3.48	3.22	2.45	1.74	2.18	2.07	1.43	1.76	1.3	1.34	1.48	1.21	10
5504	3.72	5.11	4.87	4.28	4.09	2.62	3.16	3.37	3.22	2.48	1.95	2.27	2.19	1.42	1.71	1.38	1.48	1.57	1.39	10
5505	3.56	5.7	5.65	5	4.53	2.5	3.28	3.84	3.65	2.7	2.04	2.5	2.37	1.66	1.98	1.55	1.64	1.65	1.43	10
7057	3.53	5.44	5.33	4.71	4.37	2.53	3.06	3.56	3.36	2.7	1.81	2.28	2.19	1.64	1.83	1.39	1.5	1.58	1.46	10
7241	3.69	5.54	5.42	4.89	4.28	2.39	3.38	3.96	3.65	2.78	1.89	2.47	2.28	1.63	1.89	1.47	1.57	1.59	1.41	10
5082	3.58	5.07	4.96	4.41	3.97	2.32	3.03	3.42	3.16	2.35	1.82	2.26	2.08	1.4	1.88	1.41	1.55	1.62	1.35	11
5101	3.47	5.28	5.26	4.56	4.27	2.26	3.05	3.5	3.3	2.66	1.81	2.23	2.15	1.59	1.94	1.43	1.57	1.54	1.43	11
5211	3.59	5.47	5.45	4.8	4.46	2.56	3.33	3.78	3.58	2.78	1.96	2.47	2.38	1.66	2.06	1.6	1.59	1.63	1.53	11
5318	4.06	5.85	5.65	5.06	4.54	2.8	3.72	4.14	3.85	2.9	1.98	2.44	2.24	1.36	1.99	1.42	1.67	1.65	1.32	11
5320	3.89	5.6	5.46	4.84	4.4	2.42	3.29	3.6	3.46	2.82	2	2.46	2.3	1.72	1.96	1.5	1.61	1.6	1.36	11
5340	4.62	6.69	6.47	5.78	5.41	3.1	3.98	4.46	4.22	3.22	2.3	2.81	2.69	1.66	2.21	1.7	1.9	1.82	1.57	11
5506	3.54	5.34	5.26	4.72	4.15	2.42	3.2	3.61	3.37	2.63	1.91	2.24	2.17	1.58	1.85	1.47	1.64	1.67	1.52	11
5507	2.02	2.89	2.79	2.48	2.33	1.35	1.75	1.94	1.86	1.42	0.99	1.16	1.1	0.77	0.98	0.7	0.73	0.72	0.65	11
5508	2.09	3	2.87	2.47	2.32	1.41	1.81	2.06	1.93	1.5	1.1	1.37	1.35	0.88	1.11	0.81	0.9	0.96	0.8	11
5550	4.69	6.42	6.41	5.62	5.08	2.89	3.74	4.24	3.99	3.1	2.2	2.61	2.55	1.82	2.22	1.58	1.6	1.79	1.54	11
7058	3.71	5.59	5.28	4.65	4.43	2.33	3.2	3.68	3.48	2.63	1.8	2.2	2.12	1.5	1.93	1.44	1.5	1.52	1.43	11

7059	3.37	4.8	4.89	4.32	3.88	2.17	2.86	3.29	3.1	2.35	1.67	2.08	2.03	1.32	1.66	1.27	1.32	1.39	1.17	11
7106	3.89	5.63	5.45	4.79	4.35	2.34	3.09	3.47	3.29	2.53	1.83	2.28	2.24	1.61	1.88	1.4	1.55	1.57	1.44	11
7135	3.91	5.64	5.46	4.8	4.58	2.48	3.35	3.64	3.53	2.63	1.97	2.35	2.27	1.57	1.99	1.54	1.58	1.59	1.42	11
5509	3.79	6.03	5.91	5.35	4.77	2.75	3.4	3.94	3.78	2.9	2.13	2.65	2.51	1.84	2.12	1.63	1.72	1.78	1.59	12
5510	3.98	5.86	5.86	5.2	4.62	2.64	3.53	4.02	3.71	2.87	2.19	2.67	2.55	1.8	2.11	1.6	1.73	1.76	1.57	12
5511	4.53	6.59	6.12	5.51	5.04	3.18	3.88	4.26	4.1	3.19	2.29	2.78	2.53	1.59	2.05	1.66	1.75	1.73	1.6	12
5512	4.81	6.93	6.86	6.03	5.46	3.25	4.1	4.63	4.37	3.35	2.31	2.76	2.72	1.94	2.46	1.61	1.83	1.97	1.66	12
5513	3.85	6.16	6.06	5.44	4.78	2.86	3.47	4.04	3.8	2.96	2.2	2.65	2.6	1.87	2.15	1.64	1.76	1.82	1.64	12
5514	4.2	6.07	6.16	5.41	4.88	2.83	3.68	4.15	3.9	3	2.33	2.76	2.67	1.88	2.26	1.66	1.78	1.81	1.64	12
5516	4.5	6.69	6.22	5.37	5.15	3.23	3.92	4.32	4.15	3.19	2.28	2.78	2.62	1.64	2.08	1.71	1.81	1.78	1.59	12
5551	4.22	6.06	6.08	5.47	5	2.64	3.78	4.17	3.92	3.02	2.2	2.78	2.61	1.79	2.24	1.68	1.81	1.64	1.65	12
5552	4.39	6.68	6.53	5.83	5.43	3.05	4.05	4.6	4.22	3.4	1.97	2.71	2.52	1.63	2.27	1.46	1.47	1.63	1.45	12
6116	4.34	6.26	6.07	5.26	4.93	3.2	3.88	4.11	3.93	3.18	2.11	2.7	2.59	1.89	2.21	1.63	1.76	1.84	1.62	12
7060	3.77	5.72	5.47	4.72	4.34	2.5	3.24	3.73	3.4	2.65	1.82	2.23	2.14	1.31	1.95	1.54	1.72	1.75	1.47	12
7061	4.3	6.2	6.13	5.33	4.93	2.95	3.87	4.32	4.01	3.31	2.19	2.7	2.62	2.02	2.24	1.72	1.79	1.8	1.72	12
7223	3.67	5.34	5.2	4.65	4.3	2.45	3.29	3.63	3.38	2.72	1.85	2.26	2.02	1.44	1.98	1.56	1.6	1.66	1.46	12
7236	4.36	6.17	6.15	5.32	5.11	2.99	3.84	4.39	4.21	3.26	2.2	2.7	2.58	1.86	2.13	1.71	1.78	1.84	1.67	12
7277	4.29	6.07	5.99	5.27	4.78	2.9	3.72	4.13	3.89	3.07	2.11	2.5	2.38	1.76	2.15	1.53	1.61	1.69	1.53	12
5517	4.09	6.22	6.04	5.39	4.76	2.83	3.72	4.19	3.93	3.06	2.1	2.56	2.45	1.86	2.2	1.72	1.83	1.87	1.72	13
5518	4.19	6.24	6.09	5.34	4.92	2.81	3.54	3.98	3.69	2.86	2.07	2.47	2.43	1.72	2.15	1.53	1.64	1.68	1.48	13
5519	3.98	5.73	5.58	4.89	4.62	2.56	3.39	3.75	3.55	2.75	2.04	2.35	2.19	1.5	2.28	1.54	1.52	1.64	1.46	13
5520	4.36	5.92	5.56	4.93	4.7	2.86	3.65	4.15	3.93	3.02	2.16	2.62	2.49	1.67	2.04	1.46	1.67	1.69	1.54	13
5521	4.98	7.14	7.02	6.23	5.66	3.29	4.17	4.71	4.45	3.49	2.43	2.84	2.84	2.04	2.49	1.65	1.87	2.05	1.7	13
5522	4.11	5.77	5.74	5.06	4.57	2.76	3.57	4.07	3.74	2.89	2.13	2.61	2.38	1.59	2	1.45	1.49	1.54	1.43	13
5553	4.44	6.5	6.26	5.46	4.99	3.1	4.03	4.43	4.02	3.2	2.25	2.85	2.61	1.8	2.38	1.72	1.94	1.93	1.64	13
5554	4.39	6.6	6.53	5.81	5.15	3.09	3.88	4.18	4.14	3.3	2.37	2.72	2.65	1.95	2.31	1.72	1.8	1.84	1.7	13
5555	4.54	6.54	6.22	5.46	5.07	3.27	3.92	4.34	4.14	3.23	2.35	2.84	2.63	1.66	2.09	1.68	1.61	1.77	1.6	13
5556	4.16	6.64	6.45	5.7	5.25	2.88	3.72	4.26	4.03	3.13	1.88	2.47	2.47	1.55	1.98	1.63	1.94	1.9	1.71	13
7062	4.7	6.86	6.67	5.72	5.34	3.18	4.02	4.45	4.18	3.3	2.27	2.88	2.68	1.79	2.4	1.71	1.81	1.83	1.7	13
7063	3.92	5.54	5.53	4.94	4.7	2.81	3.54	4.08	3.83	3.04	2.04	2.53	2.46	1.8	2.21	1.61	1.69	1.72	1.59	13
7064	4.59	6.66	6.36	5.65	5.34	3.16	3.88	4.33	4.05	3.28	2.27	2.73	2.7	1.82	2.16	1.69	1.87	1.85	1.76	13
7201	4.4	6.6	6.26	5.6	5.32	3.27	4.08	4.59	4.24	3.31	2.22	2.78	2.56	1.92	2.26	1.71	1.84	1.93	1.69	13
7219	4.17	5.76	5.75	5.13	4.89	2.78	3.55	3.95	3.75	2.94	2.06	2.52	2.4	1.68	2.19	1.69	1.71	1.78	1.61	13
5523	4.58	6.6	6.37	5.54	5.14	3.16	4.04	4.5	4.12	3.28	2.12	2.8	2.69	1.82	2.39	1.67	1.95	1.94	1.7	14

5524	4.51	6.55	6.37	5.62	5.25	3.24	4.01	4.53	4.28	3.34	2.26	2.66	2.63	1.93	2.39	1.68	1.94	2	1.63	14
5525	4.91	7.25	6.89	6.13	5.66	3.57	4.38	4.86	4.62	3.49	2.66	3.24	2.78	1.96	2.39	1.92	2	2.08	1.77	14
5526	4.26	6.52	6.34	5.68	5.07	3	3.94	4.36	4.11	3.25	2.3	2.73	2.59	1.95	2.26	1.79	1.91	1.97	1.82	14
5527	4.24	6.46	6.3	5.58	5.11	3.15	3.96	4.3	4.04	3.24	2.32	2.62	2.46	1.85	2.2	1.6	1.82	1.77	1.6	14
5557	4.7	6.96	6.88	6.13	5.63	3.24	4.09	4.68	4.38	3.37	2.48	2.9	2.78	1.98	2.25	1.8	1.91	1.92	1.79	14
5558	2.14	3.2	3.13	2.84	2.63	1.46	1.92	2.2	2.17	1.6	1.19	1.41	1.32	0.96	1.06	0.79	0.86	0.87	0.8	14
5559	4.81	6.96	6.92	6.22	5.71	3.21	4.24	4.75	4.5	3.52	2.48	3.11	2.9	1.84	2.44	1.84	1.99	1.96	1.85	14
5560	4.23	6.62	6.42	5.68	5.21	2.94	3.72	4.3	4.04	3.17	1.82	2.47	2.43	1.55	2.21	1.64	1.91	1.91	1.76	14
5561	4.63	6.33	5.92	5.33	4.98	3.04	3.85	4.34	4.08	3.18	2.27	2.75	2.6	1.75	2.07	1.57	1.75	1.77	1.56	14
7065	4.8	6.73	6.51	5.88	5.51	3.35	4.2	4.68	4.42	3.6	2.38	2.86	2.86	2	2.2	1.6	1.7	1.83	1.59	14
7066	5.15	7.44	7.47	6.63	6.25	3.7	4.37	4.81	4.62	3.73	2.46	2.99	2.89	2.02	2.36	1.79	1.91	1.9	1.78	14
7067	4.76	6.55	6.36	5.7	5.4	3.4	4.09	4.56	4.31	3.35	2.44	3.11	2.79	1.99	2.45	1.77	1.91	1.92	1.68	14
7212	4.79	6.75	6.54	5.63	5.24	3.23	4.03	4.52	4.29	3.38	2.34	2.87	2.63	1.9	2.07	1.63	1.84	1.81	1.77	14
5528	4.62	6.65	6.4	5.53	5.13	3.2	4.15	4.48	4.14	3.28	2.18	2.74	2.61	1.82	2.37	1.73	1.98	2	1.61	15
5529	4.63	6.86	6.81	6.08	5.57	3.36	4.47	5.02	4.71	3.82	2.51	2.95	2.9	2	2.42	1.84	1.97	1.91	1.89	15
5530	4.82	6.31	6.14	5.6	5.06	3.12	3.89	4.26	4.07	3.23	2.36	2.83	2.62	1.88	1.99	1.43	1.69	1.84	1.45	15
5531	4.66	6.32	5.97	5.31	5.03	3.08	3.87	4.43	4.1	3.21	2.29	2.77	2.62	1.75	2.14	1.55	1.79	1.75	1.58	15
5532	4.57	6.74	6.5	5.7	5.5	3.17	4.24	4.67	4.54	3.61	2.6	3.09	2.97	2.25	2.19	1.75	1.91	1.92	1.74	15
5533	4.82	6.43	6.17	5.65	5.1	3.2	3.94	4.29	4.13	3.3	2.38	2.84	2.65	1.87	1.96	1.4	1.7	1.83	1.49	15
5534	4.19	6.45	6.31	5.61	5.19	3.18	3.95	4.34	4.08	3.28	2.37	2.64	2.52	1.88	2.21	1.67	1.83	1.84	1.64	15
5562	4.9	7.55	7.35	6.66	6.12	3.09	4.28	4.88	4.44	3.37	2.17	2.83	2.59	1.52	2.58	1.98	2.06	2.09	1.87	15
5563	4.73	6.98	6.59	6.05	5.47	3.2	4.12	4.66	4.22	3.35	2.37	2.72	2.58	1.82	2.49	1.87	2.04	1.98	1.86	15
6110	4.42	6.46	6.31	5.61	5.19	3.16	4.06	4.64	4.39	3.39	2.37	2.99	2.94	1.78	2.45	1.9	2.09	2.12	1.92	15
5013	4.99	7.13	6.81	6.13	5.81	3.42	4.23	4.62	4.39	3.52	2.6	3.06	2.97	2.06	2.25	1.76	1.94	1.97	1.65	16
5332	4.71	7.13	7.27	6.44	5.61	3.32	4.25	4.83	4.55	3.48	2.41	2.98	2.95	1.99	2.14	1.82	1.98	2.06	1.87	16
5351	4.42	6.59	6.38	5.67	5.19	3	3.74	4.52	4.28	3.31	2.25	2.63	2.43	1.81	2.24	1.59	1.68	1.67	1.66	16
5352	4.87	7.03	6.87	5.97	5.66	3.24	4.15	4.73	4.4	3.45	2.38	2.96	2.84	2.07	2.52	1.89	2.09	2.11	1.94	16
5535	4.35	6.35	6.16	5.44	5.24	2.98	3.75	4.16	3.97	3.1	2.29	2.69	2.52	1.72	2.39	1.69	1.71	1.79	1.67	16
5536	4.67	6.86	6.8	6.15	5.54	3.41	4.51	5.06	4.71	3.82	2.49	2.93	2.89	2.02	2.42	1.84	1.97	1.97	1.94	16
5564	4.78	6.93	6.96	6.22	5.7	3.22	4.28	4.82	4.52	3.54	2.58	3.16	2.96	1.9	2.18	1.84	1.97	1.97	1.84	16
5565	4.98	7.6	7.36	6.69	6.17	3.21	4.16	4.87	4.47	3.35	2.31	2.84	2.56	1.49	2.49	2.05	2.13	2.12	1.96	16
5566	4.19	6.04	6.02	5.25	4.75	2.8	3.66	4.16	3.81	2.97	2.16	2.67	2.44	1.64	1.99	1.57	1.56	1.59	1.46	16
6132	4.74	7.32	7.23	6.4	5.8	3.47	4.3	4.89	4.57	3.68	2.56	3.08	2.89	2.13	2.54	1.95	1.99	2.06	1.77	16
5083	4.61	6.51	6.53	5.86	5.25	3.17	3.94	4.44	4.22	3.22	2.31	2.83	2.62	1.73	2.26	1.63	1.87	1.86	1.69	17

5148	5.1	7.04	7.04	6.31	5.59	3.34	4.38	4.9	4.62	3.54	2.53	3.01	2.87	2.09	2.51	1.69	2	2.05	1.83	17
5292	4.68	6.42	6.35	5.68	5.28	3.18	4	4.58	4.24	3.27	2.26	2.74	2.6	1.95	2.25	1.79	1.92	1.92	1.77	17
5306	4.72	7.16	7.1	6.32	5.77	3.25	4.16	4.77	4.52	3.5	2.51	3.07	2.93	2.05	2.53	1.96	2.15	2.15	1.95	17
5315	4.82	6.8	6.81	6.02	5.55	3.26	4.17	4.61	4.28	3.42	2.41	2.87	2.84	2.03	2.44	1.92	1.99	2.15	1.79	17
5341	4.79	6.83	6.7	5.95	5.65	3.16	4.02	4.59	4.33	3.4	2.35	2.84	2.66	1.87	2.31	1.68	1.79	1.81	1.68	17
5344	4.73	6.62	6.64	5.84	5.33	3.12	3.83	4.45	4.14	3.3	2.13	2.71	2.64	1.96	2.12	1.56	1.61	1.73	1.53	17
5360	4.68	6.75	6.49	5.75	5.28	2.97	3.82	4.33	4.03	3.12	2.19	2.66	2.57	1.69	2.18	1.78	1.93	1.96	1.74	17
5390	4.61	6.54	6.49	5.75	5.31	3.01	3.66	4.23	4.04	3.16	2.12	2.58	2.45	1.76	2.14	1.77	1.94	1.92	1.66	17
5567	4.73	7.09	6.64	6.16	5.55	3.24	4.08	4.68	4.2	3.35	2.35	2.8	2.62	1.85	2.47	1.83	1.99	1.97	1.83	17
5014	4.82	7	6.76	5.97	5.53	3.18	4.03	4.48	4.13	3.35	2.39	2.85	2.72	1.94	2.22	1.68	1.77	1.78	1.61	18
5059	5.01	7.08	6.73	6.06	5.67	3.78	4.35	4.9	4.58	3.62	2.46	3.19	3.04	2.07	2.34	1.86	1.96	2.05	1.83	18
5066	4.44	6.48	6.3	5.67	5.18	2.97	3.79	4.39	4.01	3.21	2.23	2.82	2.57	1.75	2.39	1.88	2.01	2.06	1.78	18
5084	4.58	6.77	6.38	5.69	5.48	3.13	3.92	4.54	4.33	3.4	2.24	2.68	2.72	2.1	2.19	1.75	1.87	1.94	1.81	18
5300	4.97	7.09	6.78	6.15	5.81	3.48	4.24	4.68	4.45	3.55	2.6	3.01	2.95	2.06	2.26	1.8	1.86	1.93	1.66	18
5342	4.64	6.76	6.52	5.81	5.23	3.17	3.85	4.28	3.99	3.21	2.18	2.77	2.51	1.83	2.16	1.69	1.71	1.78	1.67	18
5345	4.62	6.66	6.51	5.65	5.06	3.17	3.86	4.34	4.13	3.34	2.25	2.62	2.58	1.89	2.35	1.81	1.89	1.86	1.8	18
5346	4.62	6.89	6.58	6.08	5.5	3.1	4.1	4.55	4.4	3.38	2.39	2.82	2.73	1.84	2.3	1.93	2.07	2.13	1.81	18
5359	4.86	7.49	6.99	6.11	5.93	3.23	4.15	4.78	4.37	3.42	2.48	3.11	2.82	1.84	2.39	1.79	1.91	1.99	1.8	18
5537	4.73	7.06	6.67	5.98	5.52	3.41	4.34	4.89	4.42	3.56	2.64	3.06	2.84	2.03	2.39	1.89	1.99	1.97	1.86	18

APPENDIX C

NORMALIZED TRAINING DATASET OF INPUT FEATURES AND TARGET FEATURES FOR FEMALE HISPANIC POPULATION USING ANN APPROACH

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	Age
0.116 716	0.145 253	0.138 737	0.121 719	0.140 482	0.113 974	0.114 388	0.113 029	0.1 0.1	0.120 513	0.113 26	0.1 32	0.108 333	0.121 192	0.1 0.1	0.1 0.1	0.1 0.1	0.158 394	0.148 855	0.1 0.1
0.171 642	0.255 152	0.239 789	0.214 027	0.198 313	0.173 362	0.169 065	0.172 964	0.175 524	0.188 889	0.228 177	0.1 64	0.179 167	0.269 536	0.239 759	0.173 2824	0.164 234	0.216 788	0.148 855	0.144 444
0.121 493	0.124 242	0.130 316	0.125 339	0.123 133	0.106 987	0.1 0.1	0.136 482	0.141 958	0.120 513	0.139 779	0.1 44	0.191 667	0.184 768	0.124 096	0.185 4962	0.152 555	0.193 431	0.148 855	0.144 444
0.166 866	0.163 03	0.142 105	0.210 407	0.188 675	0.1 0.1	0.169 065	0.157 329	0.141 958	0.192 308	0.1 0.1	0.1 48	0.108 0.1	0.137 086	0.153 012	0.112 2137	0.146 715	0.164 234	0.136 641	0.144 444
0.217 015	0.242 222	0.251 579	0.259 276	0.229 157	0.169 869	0.186 331	0.201 629	0.197 902	0.185 47	0.263 536	0.2 4	0.266 667	0.200 662	0.181 928	0.240 458	0.205 109	0.275 182	0.228 244	0.188 889
0.169 254	0.251 919	0.254 947	0.235 747	0.238 795	0.180 349	0.241 007	0.251 14	0.197 902	0.195 726	0.276 796	0.2 68	0.25 0.25	0.232 45	0.201 205	0.191 6031	0.304 38	0.234 307	0.185 496	0.188 889
0.291 045	0.334 343	0.330 737	0.313 575	0.304 337	0.278 166	0.223 741	0.251 14	0.301 399	0.236 752	0.329 834	0.3 12	0.316 667	0.296 026	0.287 952	0.173 2824	0.275 182	0.187 591	0.191 603	0.233 333
0.405 672	0.360 202	0.359 368	0.371 493	0.379 518	0.260 699	0.295 683	0.313 681	0.304 196	0.257 265	0.303 315	0.2 8	0.3 0.3	0.232 45	0.355 422	0.270 9924	0.240 146	0.275 182	0.289 313	0.233 333
0.386 567	0.374 747	0.364 421	0.360 633	0.375 663	0.323 581	0.292 806	0.305 863	0.368 531	0.274 359	0.329 834	0.3 28	0.325 0.325	0.237 748	0.437 349	0.393 1298	0.415 328	0.432 847	0.277 099	0.233 333
0.26	0.384 444	0.389 684	0.375 113	0.366 024	0.211 79	0.243 885	0.264 169	0.259 441	0.212 821	0.356 354	0.3 32	0.316 667	0.333 113	0.350 602	0.197 7099	0.245 985	0.257 664	0.203 817	0.233 333
0.281 493	0.365 051	0.376 211	0.376 923	0.406 506	0.274 672	0.318 705	0.326 71	0.318 182	0.308 547	0.298 895	0.3 44	0.325 0.325	0.264 238	0.321 687	0.154 9618	0.205 109	0.234 307	0.234 351	0.233 333
0.324 478	0.397 374	0.406 526	0.407 692	0.383 373	0.299 127	0.318 705	0.412 704	0.410 49	0.400 855	0.343 094	0.3 56	0.329 167	0.444 371	0.360 241	0.258 7786	0.216 788	0.391 971	0.222 137	0.277 778
0.329 254	0.376 364	0.406 526	0.387 783	0.404 578	0.288 646	0.310 072	0.334 528	0.309 79	0.277 778	0.294 475	0.3 0.3	0.3 0.3	0.216 556	0.422 892	0.350 3817	0.327 737	0.374 453	0.319 847	0.277 778

0.374 627	0.363 434	0.352 632	0.322 624	0.358 313	0.222 271	0.246 763	0.271 987	0.267 832	0.236 752	0.329 834	0.2 56	0.333 333	0.306 623	0.316 867	0.277 0992	0.210 949	0.362 774	0.234 351	0.277 778
0.279 104	0.365 051	0.369 474	0.366 063	0.369 88	0.278 166	0.287 05	0.303 257	0.365 734	0.346 154	0.382 873	0.3 36	0.279 167	0.317 219	0.369 88	0.411 4504	0.415 328	0.444 526	0.295 42	0.277 778
0.322 09	0.400 606	0.399 789	0.398 643	0.396 867	0.299 127	0.321 583	0.342 345	0.427 273	0.417 949	0.343 094	0.3 56	0.379 167	0.407 285	0.369 88	0.234 3511	0.251 825	0.281 022	0.246 565	0.277 778
0.386 567	0.424 848	0.446 947	0.451 131	0.452 771	0.306 114	0.356 115	0.410 098	0.379 72	0.342 735	0.329 834	0.3 68	0.35 325	0.301 325	0.326 506	0.380 916	0.374 453	0.421 168	0.387 023	0.322 222
0.503 582	0.460 404	0.492 421	0.463 801	0.479 759	0.435 371	0.405 036	0.412 704	0.416 084	0.414 53	0.360 773	0.3 6	0.379 167	0.444 371	0.461 446	0.313 7405	0.321 898	0.356 934	0.319 847	0.322 222
0.360 299	0.470 101	0.460 421	0.452 941	0.475 904	0.292 14	0.361 871	0.368 404	0.365 734	0.315 385	0.276 796	0.3 0.3	0.283 333	0.227 152	0.254 217	0.289 313	0.263 504	0.316 058	0.295 42	0.322 222
0.470 149	0.457 172	0.480 632	0.478 281	0.470 12	0.292 14	0.324 46	0.436 156	0.435 664	0.421 368	0.404 972	0.4 0.4	0.408 333	0.370 199	0.302 41	0.332 0611	0.339 416	0.345 255	0.295 42	0.322 222
0.293 433	0.377 98	0.396 421	0.375 113	0.366 024	0.393 45	0.292 806	0.300 651	0.374 126	0.288 034	0.404 972	0.2 96	0.287 5	0.391 391	0.432 53	0.307 6336	0.275 182	0.292 701	0.289 313	0.322 222
0.283 881	0.418 384	0.433 474	0.404 072	0.394 94	0.194 323	0.373 381	0.365 798	0.348 951	0.277 778	0.365 193	0.3 0.3	0.333 333	0.317 219	0.422 892	0.283 2061	0.251 825	0.292 701	0.264 885	0.322 222
0.427 164	0.437 778	0.458 737	0.434 842	0.402 651	0.267 686	0.393 525	0.410 098	0.407 692	0.366 667	0.378 453	0.3 68	0.362 5	0.322 517	0.408 434	0.332 0611	0.386 131	0.333 577	0.270 992	0.322 222
0.403 284	0.392 525	0.408 211	0.398 643	0.398 795	0.257 205	0.315 827	0.334 528	0.332 168	0.301 709	0.307 735	0.2 84	0.320 833	0.327 815	0.321 687	0.270 9924	0.339 416	0.386 131	0.295 42	0.322 222
0.293 433	0.403 838	0.421 684	0.398 643	0.410 361	0.264 192	0.307 194	0.347 557	0.410 49	0.397 436	0.374 033	0.3 56	0.362 5	0.327 815	0.360 241	0.277 0992	0.222 628	0.356 934	0.222 137	0.322 222
0.534 627	0.526 667	0.546 316	0.514 48	0.510 602	0.393 45	0.468 345	0.592 508	0.569 93	0.530 769	0.488 95	0.4 48	0.541 667	0.481 457	0.490 361	0.325 9542	0.345 255	0.485 401	0.240 458	0.366 667
0.494 03	0.478 182	0.504 211	0.460 181	0.472 048	0.372 489	0.445 324	0.467 427	0.449 65	0.346 154	0.404 972	0.3 4	0.329 167	0.386 093	0.442 169	0.325 9542	0.432 847	0.380 292	0.325 954	0.366 667
0.496 418	0.450 707	0.460 421	0.460 181	0.473 976	0.379 476	0.396 403	0.402 28	0.416 084	0.387 179	0.440 331	0.3 56	0.408 333	0.407 285	0.312 048	0.362 5954	0.456 204	0.485 401	0.448 092	0.366 667
0.515 522	0.462 02	0.463 789	0.452 941	0.456 627	0.358 515	0.387 77	0.472 638	0.472 028	0.428 205	0.404 972	0.4 32	0.433 333	0.396 689	0.485 542	0.344 2748	0.368 613	0.386 131	0.344 275	0.366 667
0.374 627	0.471 717	0.470 526	0.489 14	0.491 325	0.330 568	0.419 424	0.506 515	0.480 42	0.469 231	0.440 331	0.4 6	0.445 833	0.370 199	0.490 361	0.368 7023	0.368 613	0.403 65	0.356 489	0.366 667

0.491 642	0.455 556	0.463 789	0.452 941	0.481 687	0.355 022	0.384 892	0.485 668	0.494 406	0.448 718	0.431 492	0.4 44	0.458 333	0.396 689	0.446 988	0.393 1298	0.397 81	0.415 328	0.454 198	0.366 667
0.601 493	0.562 222	0.595 158	0.561 538	0.545 301	0.372 489	0.433 813	0.574 267	0.530 769	0.469 231	0.475 691	0.4 8	0.341 667	0.301 325	0.350 602	0.350 3817	0.333 577	0.508 759	0.332 061	0.411 111
0.572 836	0.605 859	0.625 474	0.612 217	0.593 494	0.487 773	0.505 755	0.506 515	0.488 811	0.482 906	0.577 348	0.4 68	0.587 5	0.560 927	0.581 928	0.423 6641	0.403 65	0.462 044	0.332 061	0.411 111
0.606 269	0.518 586	0.521 053	0.498 19	0.522 169	0.571 616	0.523 022	0.561 238	0.575 524	0.482 906	0.612 707	0.5 52	0.595 833	0.598 013	0.610 843	0.619 084	0.450 365	0.643 066	0.600 763	0.411 111
0.546 567	0.549 293	0.553 053	0.548 869	0.562 651	0.484 279	0.534 532	0.545 603	0.533 566	0.503 419	0.541 989	0.5 28	0.520 833	0.529 139	0.519 277	0.350 3817	0.351 095	0.368 613	0.332 061	0.411 111
0.517 91	0.528 283	0.537 895	0.507 24	0.527 952	0.351 528	0.517 266	0.519 544	0.513 986	0.380 342	0.533 149	0.5 12	0.483 333	0.444 371	0.533 735	0.551 9084	0.473 723	0.514 599	0.338 168	0.411 111
0.668 358	0.647 879	0.635 579	0.603 167	0.574 217	0.529 694	0.594 964	0.576 873	0.553 147	0.479 487	0.546 409	0.5 44	0.562 5	0.550 331	0.577 108	0.527 4809	0.514 599	0.427 007	0.417 557	0.455 556
0.611 045	0.555 758	0.574 947	0.566 968	0.574 217	0.449 345	0.583 453	0.566 45	0.586 713	0.575 214	0.581 768	0.5 96	0.620 833	0.545 033	0.610 843	0.612 9771	0.479 562	0.654 745	0.606 87	0.455 556
0.556 119	0.546 061	0.574 947	0.543 439	0.524 096	0.501 747	0.523 022	0.550 814	0.541 958	0.503 419	0.537 569	0.5 08	0.55 0.55	0.513 245	0.586 747	0.460 3053	0.438 686	0.602 19	0.564 122	0.455 556
0.603 881	0.565 455	0.583 368	0.557 919	0.533 735	0.515 721	0.546 043	0.561 238	0.567 133	0.520 513	0.590 608	0.5 64	0.562 5	0.545 033	0.572 289	0.515 2672	0.479 562	0.643 066	0.441 985	0.455 556
0.577 612	0.568 687	0.595 158	0.583 258	0.597 349	0.578 603	0.603 597	0.579 479	0.589 51	0.585 47	0.608 287	0.6 08	0.591 667	0.608 609	0.581 928	0.484 7328	0.456 204	0.619 708	0.472 519	0.455 556
0.599 104	0.610 707	0.645 684	0.606 787	0.595 422	0.501 747	0.548 921	0.576 873	0.572 727	0.503 419	0.572 928	0.5 88	0.583 333	0.518 543	0.581 928	0.429 771	0.432 847	0.450 365	0.399 237	0.455 556
0.627 761	0.633 333	0.649 053	0.635 747	0.632 048	0.536 681	0.554 676	0.527 362	0.558 741	0.564 957	0.533 149	0.5 28	0.554 167	0.629 801	0.591 566	0.448 0916	0.567 153	0.613 869	0.533 588	0.455 556
0.649 254	0.622 02	0.622 105	0.585 068	0.583 855	0.561 135	0.609 353	0.626 384	0.637 063	0.626 496	0.586 188	0.6 12	0.620 833	0.640 397	0.586 747	0.448 0916	0.584 672	0.643 066	0.478 626	0.5 0.5
0.706 567	0.686 667	0.699 579	0.677 376	0.687 952	0.603 057	0.643 885	0.675 896	0.684 615	0.626 496	0.670 166	0.7 28	0.712 5	0.725 166	0.634 94	0.631 2977	0.695 62	0.724 818	0.661 832	0.5 0.5
0.651 642	0.597 778	0.649 053	0.650 226	0.624 337	0.536 681	0.571 942	0.618 567	0.606 294	0.599 145	0.564 088	0.5 56	0.562 5	0.523 841	0.644 578	0.661 8321	0.625 547	0.502 92	0.631 298	0.5 0.5
0.568 06	0.552 525	0.544 632	0.496 38	0.508 675	0.452 838	0.494 245	0.490 879	0.466 434	0.452 137	0.475 691	0.4 64	0.454 167	0.396 689	0.475 904	0.521 374	0.479 562	0.485 401	0.496 947	0.5 0.5

0.680 299	0.626 869	0.682 737	0.657 466	0.651 325	0.421 397	0.586 331	0.628 99	0.667 832	0.612 821	0.515 47	0.6 08	0.616 667	0.613 907	0.683 133	0.655 7252	0.502 92	0.584 672	0.667 939	0.5
0.630 149	0.636 566	0.691 158	0.648 416	0.662 892	0.585 59	0.669 784	0.691 531	0.676 224	0.677 778	0.692 265	0.6 76	0.658 333	0.640 397	0.692 771	0.710 687	0.648 905	0.648 905	0.643 511	0.544 444
0.673 134	0.628 485	0.649 053	0.617 647	0.622 41	0.585 59	0.620 863	0.644 625	0.642 657	0.616 239	0.603 867	0.6 08	0.6 0.6	0.518 543	0.668 675	0.576 3359	0.409 489	0.648 905	0.594 656	0.544 444
0.613 433	0.634 949	0.655 789	0.644 796	0.653 253	0.547 162	0.603 597	0.628 99	0.634 266	0.629 915	0.559 669	0.5 32	0.545 833	0.624 503	0.591 566	0.521 374	0.321 898	0.567 153	0.551 908	0.544 444
0.751 94	0.686 667	0.681 053	0.646 606	0.649 398	0.693 886	0.707 194	0.717 59	0.720 979	0.698 291	0.705 525	0.7 44	0.75 0.75	0.703 974	0.736 145	0.692 3664	0.666 423	0.713 139	0.667 939	0.544 444
0.680 299	0.657 576	0.672 632	0.659 276	0.686 024	0.620 524	0.652 518	0.681 107	0.690 21	0.664 103	0.603 867	0.6 44	0.637 5	0.603 311	0.620 482	0.674 0458	0.666 423	0.678 102	0.643 511	0.544 444
0.642 09	0.604 242	0.608 632	0.594 118	0.593 494	0.491 266	0.577 698	0.602 932	0.614 685	0.602 564	0.577 348	0.6 2	0.595 833	0.592 715	0.615 663	0.454 1985	0.619 708	0.643 066	0.496 947	0.544 444
0.577 612	0.562 222	0.58	0.556 109	0.537 59	0.466 812	0.488 489	0.545 603	0.541 958	0.489 744	0.480 11	0.5 08	0.516 667	0.507 947	0.586 747	0.441 9847	0.596 35	0.613 869	0.435 878	0.544 444
0.708 955	0.659 192	0.708	0.671 946	0.697 59	0.620 524	0.678 417	0.644 625	0.673 427	0.629 915	0.652 486	0.6 52	0.675	0.619 205	0.760 241	0.655 7252	0.637 226	0.713 139	0.735 115	0.544 444
0.694 627	0.649 495	0.664 211	0.617 647	0.622 41	0.624 017	0.695 683	0.694 137	0.673 427	0.684 615	0.705 525	0.6 92	0.637 5	0.688 079	0.731 325	0.649 6183	0.637 226	0.660 584	0.619 084	0.544 444
0.654 03	0.628 485	0.644	0.619 457	0.605 06	0.585 59	0.612 23	0.655 049	0.611 888	0.582 051	0.546 409	0.5 92	0.545 833	0.560 927	0.639 759	0.612 9771	0.631 387	0.672 263	0.637 405	0.544 444
0.649 254	0.676 97	0.691 158	0.655 656	0.655 181	0.589 083	0.617 986	0.657 655	0.645 455	0.588 889	0.572 928	0.6 16	0.633 333	0.566 225	0.649 398	0.655 7252	0.613 869	0.602 19	0.576 336	0.544 444
0.694 627	0.673 737	0.652 421	0.652 036	0.659 036	0.617 031	0.669 784	0.694 137	0.698 601	0.660 684	0.617 127	0.6 44	0.637 5	0.582 119	0.553 012	0.643 5115	0.625 547	0.695 62	0.625 191	0.588 889
0.708 955	0.651 111	0.665 895	0.623 077	0.647 47	0.610 044	0.678 417	0.704 56	0.709 79	0.684 615	0.612 707	0.6 68	0.616 667	0.428 477	0.692 771	0.692 3664	0.678 102	0.765 693	0.643 511	0.588 889
0.754 328	0.691 515	0.692 842	0.626 697	0.682 169	0.711 354	0.695 683	0.691 531	0.681 818	0.725 641	0.634 807	0.6 68	0.695 833	0.672 185	0.639 759	0.680 1527	0.608 029	0.672 263	0.570 229	0.588 889
0.785 373	0.717 374	0.765 263	0.715 385	0.736 145	0.721 834	0.730 216	0.756 678	0.734 965	0.718 803	0.683 425	0.7 0.7	0.7 0.7	0.677 483	0.750 602	0.704 5802	0.695 62	0.724 818	0.759 542	0.588 889
0.718 507	0.675 354	0.692 842	0.677 376	0.709 157	0.631 004	0.672 662	0.707 166	0.712 587	0.691 453	0.639 227	0.6 8	0.675	0.635 099	0.610 843	0.692 3664	0.654 745	0.707 299	0.655 725	0.588 889

0.711 343	0.743 232	0.755 158	0.724 434	0.753 494	0.739 301	0.802 158	0.808 795	0.807 692	0.824 786	0.732 044	0.7 92	0.829 167	0.772 848	0.707 229	0.783 9695	0.771 533	0.748 175	0.747 328	0.588 889
0.771 045	0.683 434	0.706 316	0.695 475	0.705 301	0.610 044	0.661 151	0.668 078	0.673 427	0.670 94	0.705 525	0.7 16	0.687 5	0.698 675	0.736 145	0.722 9008	0.736 496	0.754 015	0.698 473	0.588 889
0.692 239	0.652 727	0.660 842	0.648 416	0.643 614	0.631 004	0.620 863	0.636 808	0.625 874	0.640 171	0.625 967	0.5 96	0.587 5	0.645 695	0.610 843	0.625 1908	0.643 066	0.660 584	0.637 405	0.588 889
0.766 269	0.722 222	0.729 895	0.668 326	0.695 663	0.683 406	0.730 216	0.743 648	0.762 937	0.711 966	0.643 646	0.6 92	0.670 833	0.598 013	0.625 301	0.661 8321	0.637 226	0.736 496	0.649 618	0.633 333
0.744 776	0.709 293	0.697 895	0.657 466	0.655 181	0.725 328	0.738 849	0.780 13	0.760 14	0.735 897	0.656 906	0.7 44	0.708 333	0.666 887	0.615 663	0.649 6183	0.613 869	0.578 832	0.643 511	0.633 333
0.773 433	0.764 242	0.778 737	0.778 733	0.770 843	0.763 755	0.810 791	0.816 612	0.813 287	0.776 923	0.785 083	0.8 12	0.795 833	0.714 57	0.798 795	0.771 7557	0.789 051	0.829 927	0.582 443	0.633 333
0.668 358	0.686 667	0.738 316	0.688 235	0.666 747	0.679 913	0.698 561	0.730 619	0.737 762	0.698 291	0.705 525	0.7 24	0.758 333	0.672 185	0.683 133	0.667 9389	0.707 299	0.771 533	0.686 26	0.633 333
0.728 06	0.665 657	0.692 842	0.604 977	0.657 108	0.634 498	0.684 173	0.709 772	0.712 587	0.691 453	0.612 707	0.6 64	0.658 333	0.454 967	0.731 325	0.680 1527	0.689 781	0.730 657	0.643 511	0.633 333
0.759 104	0.723 838	0.741 684	0.735 294	0.724 578	0.728 821	0.733 094	0.730 619	0.698 601	0.705 128	0.679 006	0.7 28	0.704 167	0.714 57	0.750 602	0.759 542	0.771 533	0.783 212	0.759 542	0.633 333
0.787 761	0.720 606	0.719 789	0.666 516	0.695 663	0.718 341	0.721 583	0.720 195	0.706 993	0.776 923	0.687 845	0.7 12	0.725 0.725	0.703 974	0.740 964	0.722 9008	0.695 62	0.713 139	0.625 191	0.633 333
0.780 597	0.756 162	0.783 789	0.771 493	0.738 072	0.760 262	0.747 482	0.764 495	0.793 706	0.783 761	0.709 945	0.7 76	0.808 333	0.809 934	0.692 771	0.722 9008	0.713 139	0.713 139	0.741 221	0.633 333
0.725 672	0.725 455	0.728 211	0.704 525	0.666 747	0.683 406	0.704 317	0.725 407	0.718 182	0.701 709	0.608 287	0.6 64	0.645 833	0.576 821	0.586 747	0.649 6183	0.713 139	0.660 584	0.686 26	0.633 333
0.754 328	0.772 323	0.797 263	0.749 774	0.751 566	0.707 86	0.779 137	0.795 765	0.804 895	0.759 829	0.692 265	0.7 0.7	0.670 833	0.709 272	0.644 578	0.680 1527	0.672 263	0.765 693	0.649 618	0.633 333
0.878 507	0.9 0.9	0.9 0.9	0.9 0.9	0.9 0.9	0.9 0.9	0.9 0.9	0.881 759	0.9 0.9	0.886 325	0.9 0.9	0.8 76	0.9 0.9	0.815 232	0.871 084	0.790 0763	0.824 088	0.870 803	0.710 687	0.633 333
0.701 791	0.710 909	0.729 895	0.720 814	0.686 024	0.697 38	0.715 827	0.748 86	0.743 357	0.722 222	0.621 547	0.6 84	0.629 167	0.629 801	0.750 602	0.692 3664	0.695 62	0.701 46	0.667 939	0.633 333
0.763 881	0.693 131	0.724 842	0.680 995	0.713 012	0.704 367	0.689 928	0.694 137	0.693 007	0.681 197	0.679 006	0.6 48	0.654 167	0.677 483	0.745 783	0.692 3664	0.689 781	0.666 423	0.729 008	0.633 333
0.790 149	0.735 152	0.750 105	0.700 905	0.734 217	0.756 769	0.805 036	0.821 824	0.807 692	0.794 017	0.798 343	0.7 84	0.791 667	0.825 828	0.755 422	0.790 0763	0.800 73	0.800 73	0.783 969	0.677 778

0.771 045	0.715 758	0.706 316	0.666 516	0.684 096	0.714 847	0.747 482	0.733 225	0.734 965	0.705 128	0.701 105	0.6 92	0.687 5	0.640 397	0.712 048	0.832 8244	0.847 445	0.806 569	0.777 863	0.677 778
0.728 06	0.641 414	0.686 105	0.626 697	0.653 253	0.672 926	0.689 928	0.722 801	0.709 79	0.722 222	0.643 646	0.6 64	0.65 073	0.439 229	0.707 229	0.722 9008	0.718 978	0.765 693	0.667 939	0.677 778
0.790 149	0.741 616	0.763 579	0.702 715	0.759 277	0.735 808	0.770 504	0.780 13	0.754 545	0.732 479	0.740 884	0.7 52	0.729 167	0.714 57	0.726 506	0.777 8626	0.736 496	0.771 533	0.722 901	0.677 778
0.780 597	0.773 939	0.755 158	0.731 674	0.722 651	0.690 393	0.761 871	0.782 736	0.782 517	0.773 504	0.665 746	0.6 44	0.679 167	0.666 887	0.755 422	0.790 0763	0.742 336	0.789 051	0.722 901	0.677 778
0.761 493	0.788 485	0.800 632	0.760 633	0.763 133	0.746 288	0.761 871	0.798 371	0.807 692	0.763 248	0.696 685	0.7 32	0.716 667	0.688 079	0.683 133	0.704 5802	0.701 46	0.759 854	0.649 618	0.677 778
0.778 209	0.731 919	0.751 789	0.713 575	0.678 313	0.665 939	0.695 683	0.735 831	0.673 427	0.643 59	0.727 624	0.7 2	0.675 377	0.693 867	0.716 867	0.698 4733	0.678 102	0.695 62	0.649 618	0.677 778
0.735 224	0.688 283	0.672 632	0.633 937	0.657 108	0.651 965	0.735 971	0.741 042	0.723 776	0.670 94	0.639 227	0.6 56	0.637 5	0.322 517	0.625 301	0.704 5802	0.654 745	0.701 46	0.667 939	0.677 778
0.823 582	0.759 394	0.770 316	0.738 914	0.732 289	0.718 341	0.741 727	0.748 86	0.737 762	0.688 034	0.709 945	0.6 92	0.720 833	0.666 887	0.726 506	0.747 3282	0.730 657	0.742 336	0.729 008	0.677 778
0.737 612	0.752 929	0.748 421	0.684 615	0.713 012	0.742 795	0.761 871	0.772 313	0.757 343	0.752 991	0.705 525	0.7 36	0.712 5	0.698 675	0.750 602	0.851 145	0.818 248	0.847 445	0.747 328	0.677 778
0.723 284	0.688 283	0.699 579	0.693 665	0.691 807	0.721 834	0.687 05	0.748 86	0.712 587	0.667 521	0.617 127	0.6 6	0.670 833	0.582 119	0.567 47	0.625 1908	0.572 993	0.625 547	0.594 656	0.722 222
0.771 045	0.743 232	0.783 789	0.796 833	0.778 554	0.704 367	0.724 46	0.738 436	0.754 545	0.763 248	0.705 525	0.7 0.7	0.654 167	0.735 762	0.687 952	0.729 0076	0.701 46	0.718 978	0.735 115	0.722 222
0.708 955	0.701 212	0.736 632	0.719 005	0.709 157	0.704 367	0.681 295	0.694 137	0.690 21	0.688 034	0.656 906	0.6 6	0.666 667	0.709 272	0.408 434	0.729 0076	0.748 175	0.777 372	0.729 008	0.722 222
0.797 313	0.773 939	0.797 263	0.782 353	0.768 916	0.784 716	0.735 971	0.756 678	0.788 112	0.800 855	0.727 624	0.8 0.8	0.812 5	0.767 55	0.678 313	0.704 5802	0.695 62	0.713 139	0.716 794	0.722 222
0.766 269	0.727 071	0.746 737	0.704 525	0.707 229	0.669 432	0.753 237	0.769 707	0.779 72	0.742 735	0.674 586	0.7 0.7	0.695 833	0.656 291	0.707 229	0.674 0458	0.678 102	0.742 336	0.680 153	0.722 222
0.725 672	0.675 354	0.689 474	0.680 995	0.662 892	0.658 952	0.669 784	0.681 107	0.679 021	0.688 034	0.590 608	0.6 48	0.708 333	0.682 781	0.591 566	0.637 4046	0.555 474	0.648 905	0.643 511	0.722 222
0.809 254	0.778 788	0.790 526	0.771 493	0.776 627	0.767 249	0.793 525	0.806 189	0.782 517	0.763 248	0.776 243	0.7 96	0.775 0775	0.772 848	0.726 506	0.820 6107	0.829 927	0.841 606	0.790 076	0.722 222
0.828 358	0.741 616	0.780 421	0.749 774	0.766 988	0.746 288	0.779 137	0.774 919	0.760 14	0.746 154	0.740 884	0.7 52	0.725 0725	0.688 079	0.740 964	0.777 8626	0.736 496	0.783 212	0.729 008	0.722 222

0.9	0.819 192	0.783 789	0.726 244	0.774 699	0.854 585	0.885 612	0.9	0.877 622	0.831 624	0.864 641	0.8 92	0.887 5	0.831 126	0.630 12	0.747 3282	0.759 854	0.789 051	0.674 046	0.722 222
0.775 821	0.775 556	0.817 474	0.787 783	0.807 47	0.721 834	0.733 094	0.741 042	0.737 762	0.718 803	0.683 425	0.7 28	0.7 5	0.661 589	0.779 518	0.741 2214	0.783 212	0.765 693	0.759 542	0.722 222
0.689 851	0.712 525	0.743 368	0.742 534	0.751 566	0.697 38	0.744 604	0.738 436	0.712 587	0.711 966	0.723 204	0.7 16	0.712 5	0.661 589	0.702 41	0.704 5802	0.759 854	0.713 139	0.729 008	0.722 222
0.816 418	0.819 192	0.830 947	0.811 312	0.832 53	0.819 651	0.825 18	0.845 277	0.849 65	0.783 761	0.776 243	0.7 84	0.725 185	0.672 253	0.813 253	0.820 6107	0.835 766	0.894 161	0.814 504	0.722 222
0.866 567	0.782 02	0.839 368	0.787 783	0.780 482	0.770 742	0.770 504	0.787 948	0.785 315	0.770 085	0.771 823	0.7 36	0.745 833	0.815 232	0.731 325	0.875 5725	0.800 73	0.829 927	0.802 29	0.766 667
0.842 687	0.785 253	0.836 502	0.809 253	0.813 301	0.739 158	0.802 248	0.832 706	0.793 017	0.794 624	0.727 4	0.7 833	0.720 252	0.762 867	0.716 9695	0.783 336	0.742 051	0.789 649	0.765 667	0.766 667
0.756 716	0.730 303	0.738 316	0.708 145	0.757 349	0.676 419	0.721 583	0.751 466	0.751 748	0.735 897	0.665 746	0.7 08	0.704 167	0.666 887	0.615 663	0.759 542	0.695 62	0.736 496	0.686 26	0.766 667
0.794 925	0.777 172	0.790 526	0.796 833	0.788 193	0.704 367	0.730 216	0.746 254	0.762 937	0.776 923	0.679 006	0.7 0.7	0.666 667	0.725 166	0.673 494	0.765 6489	0.748 175	0.754 015	0.741 221	0.766 667
0.742 388	0.683 434	0.633 895	0.576 018	0.603 133	0.672 926	0.669 784	0.688 925	0.653 846	0.602 564	0.643 646	0.6 28	0.566 667	0.407 285	0.581 928	0.680 1527	0.625 547	0.683 942	0.655 725	0.766 667
0.852 239	0.790 101	0.812 421	0.795 023	0.807 47	0.763 755	0.810 791	0.821 824	0.818 881	0.787 179	0.789 503	0.8 04	0.787 5	0.672 185	0.707 229	0.826 7176	0.800 73	0.853 285	0.759 542	0.766 667
0.704 179	0.693 131	0.694 526	0.690 045	0.697 59	0.760 262	0.698 561	0.751 466	0.757 343	0.756 41	0.674 586	0.7 04	0.708 333	0.703 974	0.683 133	0.753 4351	0.742 336	0.754 015	0.741 221	0.766 667
0.673 134	0.668 889	0.684 421	0.655 656	0.666 747	0.711 354	0.715 827	0.730 619	0.712 587	0.670 94	0.683 425	0.7 08	0.675 099	0.635 687	0.721 4351	0.753 4351	0.689 781	0.683 942	0.716 794	0.811 111
0.892 836	0.846 667	0.871 368	0.831 222	0.840 241	0.819 651	0.836 691	0.840 065	0.821 678	0.790 598	0.780 663	0.7 84	0.758 333	0.666 887	0.808 434	0.826 7176	0.829 927	0.806 569	0.790 076	0.811 111
0.787 761	0.733 535	0.753 474	0.737 104	0.732 289	0.700 873	0.770 504	0.754 072	0.748 951	0.722 222	0.732 044	0.7 2	0.679 167	0.624 503	0.731 325	0.735 1145	0.765 693	0.777 372	0.667 939	0.811 111
0.797 313	0.791 717	0.827 579	0.816 742	0.832 53	0.774 236	0.782 014	0.785 342	0.802 098	0.776 923	0.661 326	0.7 2	0.716 667	0.703 974	0.726 506	0.790 0763	0.806 569	0.841 606	0.747 328	0.811 111
0.854 627	0.791 717	0.841 053	0.807 692	0.826 747	0.739 301	0.819 424	0.845 277	0.796 503	0.787 179	0.727 624	0.7 64	0.729 167	0.788 742	0.736 145	0.790 0763	0.724 818	0.824 088	0.771 756	0.811 111
0.725 672	0.731 919	0.768 632	0.738 914	0.741 928	0.624 017	0.692 806	0.725 407	0.729 371	0.674 359	0.603 867	0.6 04	0.612 5	0.492 053	0.649 398	0.667 9389	0.689 781	0.736 496	0.722 901	0.811 111

0.766 269	0.714 141	0.765 263	0.720 814	0.741 928	0.711 354	0.733 094	0.782 736	0.757 343	0.698 291	0.670 166	0.6 72	0.687 5	0.582 119	0.721 687	0.814 5038	0.765 693	0.771 533	0.796 183	0.811 111
0.785 373	0.741 616	0.734 947	0.719 005	0.699 518	0.721 834	0.756 115	0.754 072	0.746 154	0.722 222	0.705 525	0.7 04	0.704 167	0.661 589	0.707 229	0.838 9313	0.829 927	0.829 927	0.832 824	0.811 111
0.825 97	0.769 091	0.790 526	0.773 303	0.772 771	0.760 262	0.793 525	0.814 007	0.802 098	0.759 829	0.789 503	0.8 08	0.791 667	0.767 55	0.731 325	0.875 5725	0.835 766	0.864 964	0.771 756	0.811 111
0.744 776	0.662 424	0.686 105	0.626 697	0.687 952	0.693 886	0.721 583	0.756 678	0.723 776	0.752 991	0.648 066	0.6 48	0.670 833	0.756 954	0.654 217	0.680 1527	0.718 978	0.765 693	0.759 542	0.855 556
0.716 119	0.752 929	0.804	0.766 063	0.757 349	0.686 9	0.735 971	0.748 86	0.768 531	0.739 316	0.679 006	0.7 04	0.729 167	0.672 185	0.625 301	0.722 9008	0.683 942	0.742 336	0.692 366	0.855 556
0.778 209	0.738 384	0.758 526	0.737 104	0.722 651	0.714 847	0.735 971	0.722 801	0.723 776	0.701 709	0.732 044	0.7 08	0.737 5	0.698 675	0.644 578	0.704 5802	0.678 102	0.695 62	0.692 366	0.855 556
0.773 433	0.762 626	0.778 737	0.793 213	0.801 687	0.791 703	0.770 504	0.780 13	0.790 909	0.759 829	0.789 503	0.7 8	0.783 333	0.809 934	0.827 711	0.765 6489	0.818 248	0.876 642	0.783 969	0.855 556
0.742 388	0.714 141	0.699 579	0.652 036	0.643 614	0.662 445	0.664 029	0.709 772	0.665 035	0.636 752	0.581 768	0.6 44	0.616 667	0.523 841	0.634 94	0.704 5802	0.695 62	0.713 139	0.686 26	0.855 556
0.873 731	0.817 576	0.851 158	0.769 683	0.788 193	0.861 572	0.856 835	0.881 759	0.849 65	0.862 393	0.767 403	0.8 52	0.829 167	0.873 51	0.9	0.851 145	0.847 445	0.888 321	0.838 931	0.855 556
0.725 672	0.694 747	0.728 211	0.702 715	0.686 024	0.690 393	0.756 115	0.733 225	0.734 965	0.698 291	0.679 006	0.7 2	0.75	0.698 675	0.687 952	0.716 7939	0.637 226	0.707 299	0.680 153	0.9
0.718 507	0.710 909	0.718 105	0.688 235	0.659 036	0.599 563	0.664 029	0.712 378	0.693 007	0.602 564	0.625 967	0.6 56	0.641 667	0.545 033	0.659 036	0.643 5115	0.631 387	0.660 584	0.631 298	0.9
0.744 776	0.746 465	0.758 526	0.742 534	0.734 217	0.714 847	0.735 971	0.738 436	0.720 979	0.711 966	0.714 365	0.7 64	0.7	0.703 974	0.789 157	0.783 9695	0.794 891	0.777 372	0.771 756	0.9
0.799 701	0.720 606	0.721 474	0.664 706	0.709 157	0.739 301	0.735 971	0.733 225	0.729 371	0.770 085	0.679 006	0.7 08	0.725	0.714 57	0.606 024	0.741 2214	0.695 62	0.730 657	0.625 191	0.9

APPENDIX D

NORMALIZED TEST DATASET OF INPUT FEATURES AND TARGET FEATURES FOR FEMALE HISPANIC POPULATION USING ANN APPROACH

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	Age
0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.111 189	0.1	0.130 939	0.1	0.108 333	0.115 894	0.124 096	0.179 3893	0.146 715	0.158 394	0.148 855	0.144 444
0.162 09	0.224 444	0.214 526	0.212 217	0.211 807	0.141 921	0.206 475	0.180 782	0.203 497	0.219 658	0.126 519	0.1	0.166 667	0.126 49	0.220 482	0.148 855	0.135 036	0.199 27	0.1	0.144 444
0.228 955	0.302 02	0.302 105	0.291 855	0.312 048	0.222 271	0.252 518	0.256 352	0.315 385	0.226 496	0.325 414	0.2	0.295 833	0.253 642	0.312 048	0.344 2748	0.345 255	0.345 255	0.240 458	0.188 889
0.169 254	0.242 222	0.224 632	0.232 127	0.225 301	0.145 415	0.148 921	0.167 752	0.161 538	0.141 026	0.192 818	0.1	0.15	0.1	0.148 193	0.197 7099	0.164 234	0.199 27	0.148 855	0.188 889
0.140 597	0.227 677	0.231 368	0.188 688	0.194 458	0.131 441	0.192 086	0.206 84	0.209 091	0.120 513	0.148 619	0.1	0.191 667	0.126 49	0.215 663	0.142 7481	0.111 679	0.1	0.142 748	0.188 889
0.453 433	0.442 626	0.484	0.465 611	0.462 41	0.344 541	0.410 791	0.457 003	0.460 839	0.407 692	0.325 414	0.3	0.466 667	0.460 265	0.5	0.325 9542	0.304 38	0.327 737	0.338 168	0.322 222
0.367 463	0.453 939	0.475 579	0.454 751	0.445 06	0.337 555	0.410 791	0.417 915	0.432 867	0.335 897	0.466 851	0.4	0.433 333	0.391 391	0.451 807	0.448 0916	0.403 65	0.444 526	0.319 847	0.366 667
0.477 313	0.458 788	0.441 895	0.451 131	0.472 048	0.292 14	0.402 158	0.428 339	0.430 07	0.346 154	0.400 552	0.3	0.375	0.391 391	0.413 253	0.356 4886	0.327 737	0.485 401	0.350 382	0.366 667
0.477 313	0.415 152	0.408 211	0.409 502	0.414 217	0.414 41	0.428 058	0.438 762	0.449 65	0.421 368	0.334 254	0.3	0.366 667	0.444 371	0.446 988	0.350 3817	0.368 613	0.386 131	0.362 595	0.366 667
0.477 313	0.471 717	0.477 263	0.456 561	0.473 976	0.414 41	0.459 712	0.483 062	0.483 217	0.397 436	0.400 552	0.4	0.475	0.465 563	0.524 096	0.380 916	0.391 971	0.467 883	0.356 489	0.366 667
0.599 104	0.537 98	0.548	0.547 059	0.543 373	0.389 956	0.520 144	0.548 208	0.525 175	0.493 162	0.546 409	0.5	0.483 333	0.481 457	0.533 735	0.362 5954	0.351 095	0.351 095	0.454 198	0.411 111
0.458 209	0.552 525	0.569 895	0.492 76	0.551 084	0.491 266	0.5	0.516 938	0.435 664	0.404 274	0.502 21	0.4	0.475	0.465 563	0.514 458	0.399 2366	0.520 438	0.543 796	0.509 16	0.411 111
0.572 836	0.560 606	0.559 789	0.541 629	0.547 229	0.494 76	0.540 288	0.501 303	0.527 972	0.404 274	0.546 409	0.5	0.504 167	0.481 457	0.572 289	0.496 9466	0.520 438	0.543 796	0.374 809	0.411 111

0.486 866	0.492 727	0.507 579	0.494 57	0.493 253	0.466 812	0.407 914	0.467 427	0.477 622	0.394 017	0.471 271	0.4 76	0.466 667	0.534 437	0.509 639	0.423 6641	0.456 204	0.497 08	0.332 061	0.411 111
0.596 716	0.544 444	0.564 842	0.532 579	0.520 241	0.466 812	0.485 612	0.493 485	0.486 014	0.476 068	0.387 293	0.4 8	0.491 667	0.433 775	0.519 277	0.429 771	0.497 08	0.520 438	0.515 267	0.411 111
0.441 493	0.525 051	0.514 316	0.501 81	0.520 241	0.466 812	0.537 41	0.519 544	0.530 769	0.530 769	0.546 409	0.5 2	0.504 167	0.497 351	0.519 277	0.374 8092	0.491 241	0.537 956	0.521 374	0.455 556
0.558 507	0.486 263	0.514 316	0.5 0.5	0.522 169	0.484 279	0.462 59	0.480 456	0.474 825	0.465 812	0.458 011	0.4 32	0.429 167	0.454 967	0.533 735	0.527 4809	0.497 08	0.520 438	0.545 802	0.455 556
0.58 465	0.586 947	0.638 947	0.612 217	0.626 265	0.459 825	0.635 252	0.655 049	0.637 063	0.626 496	0.625 967	0.6 36	0.608 333	0.608 609	0.615 663	0.649 6183	0.637 226	0.637 226	0.619 084	0.5 0.5
0.711 343	0.714 141	0.726 526	0.668 326	0.672 53	0.655 459	0.698 561	0.738 436	0.706 993	0.660 684	0.639 227	0.6 64	0.670 833	0.656 291	0.557 831	0.588 5496	0.584 672	0.467 883	0.576 336	0.5 0.5
0.642 09	0.591 313	0.642 316	0.592 308	0.616 627	0.547 162	0.566 187	0.600 326	0.575 524	0.510 256	0.559 669	0.5 8	0.566 667	0.486 755	0.610 843	0.710 687	0.672 263	0.707 299	0.661 832	0.5 0.5
0.699 403	0.701 212	0.706 316	0.670 136	0.691 807	0.644 978	0.684 173	0.686 319	0.695 804	0.609 402	0.643 646	0.6 6	0.645 833	0.682 781	0.668 675	0.704 5802	0.683 942	0.660 584	0.625 191	0.5 0.5
0.563 284	0.539 596	0.58 0.58	0.592 308	0.612 771	0.494 76	0.531 655	0.553 42	0.561 538	0.541 026	0.537 569	0.5 52	0.554 167	0.507 947	0.567 47	0.545 8015	0.514 599	0.561 314	0.503 053	0.5 0.5
0.663 582	0.613 939	0.647 368	0.619 457	0.639 759	0.582 096	0.546 043	0.576 873	0.553 147	0.544 444	0.577 348	0.5 8	0.570 833	0.539 735	0.610 843	0.606 8702	0.596 35	0.602 19	0.594 656	0.544 444
0.747 164	0.673 737	0.694 526	0.673 756	0.670 602	0.644 978	0.655 396	0.688 925	0.706 993	0.667 521	0.652 486	0.6 84	0.725 0.725	0.735 762	0.702 41	0.759 542	0.689 781	0.736 496	0.704 58	0.544 444
0.642 09	0.596 162	0.595 158	0.585 068	0.587 711	0.543 668	0.609 353	0.628 99	0.623 077	0.605 983	0.625 967	0.6 28	0.637 5	0.613 907	0.586 747	0.600 7634	0.596 35	0.643 066	0.558 015	0.544 444
0.627 761	0.573 535	0.595 158	0.566 968	0.562 651	0.543 668	0.646 763	0.639 414	0.645 455	0.571 795	0.634 807	0.6 44	0.637 5	0.645 695	0.644 578	0.619 084	0.602 19	0.660 584	0.551 908	0.588 889
0.723 284	0.697 98	0.709 684	0.661 086	0.676 386	0.669 432	0.710 072	0.743 648	0.743 357	0.681 197	0.612 707	0.6 56	0.662 5	0.566 225	0.721 687	0.625 1908	0.625 547	0.718 978	0.686 26	0.588 889
0.821 194	0.727 071	0.756 842	0.749 774	0.755 422	0.770 742	0.799 281	0.772 313	0.771 329	0.756 41	0.758 564	0.7 64	0.816 667	0.719 868	0.692 771	0.649 6183	0.683 942	0.666 423	0.674 046	0.588 889
0.775 821	0.725 455	0.731 579	0.657 466	0.680 241	0.672 926	0.698 561	0.720 195	0.732 168	0.739 316	0.634 807	0.6 44	0.675 0.675	0.619 205	0.581 928	0.686 2595	0.660 584	0.777 372	0.637 405	0.588 889
0.716 119	0.697 98	0.708 0.708	0.688 235	0.724 578	0.651 965	0.687 05	0.722 801	0.720 979	0.708 547	0.661 326	0.6 84	0.670 833	0.645 695	0.634 94	0.704 5802	0.672 263	0.701 46	0.686 26	0.588 889

0.694 627	0.709 293	0.718 105	0.728 054	0.720 723	0.606 55	0.658 273	0.678 502	0.701 399	0.691 453	0.630 387	0.6 28	0.629 167	0.661 589	0.668 675	0.649 6183	0.654 745	0.648 905	0.637 405	0.588 889
0.744 776	0.693 131	0.708	0.680 995	0.680 241	0.728 821	0.733 094	0.741 042	0.734 965	0.711 966	0.701 105	0.7 32	0.7 06	0.741 036	0.659 036	0.606 8702	0.625 547	0.648 905	0.661 832	0.588 889
0.799 701	0.804 646	0.792 211	0.804 072	0.797 831	0.777 729	0.764 748	0.769 707	0.793 706	0.763 248	0.683 425	0.6 96	0.712 5	0.677 483	0.731 325	0.753 4351	0.771 533	0.812 409	0.735 115	0.633 333
0.675 522	0.643 03	0.638 947	0.566 968	0.574 217	0.606 55	0.620 863	0.688 925	0.667 832	0.643 59	0.603 867	0.6 48	0.625	0.497 351	0.644 578	0.674 0458	0.631 387	0.654 745	0.655 725	0.633 333
0.694 627	0.722 222	0.743 368	0.691 855	0.670 602	0.704 367	0.747 482	0.767 101	0.746 154	0.722 222	0.785 083	0.7 44	0.733 333	0.725 166	0.639 759	0.704 5802	0.707 299	0.718 978	0.674 046	0.677 778
0.766 269	0.704 444	0.696 211	0.677 376	0.628 193	0.655 459	0.698 561	0.688 925	0.690 21	0.647 009	0.679 006	0.6 48	0.65	0.560 927	0.731 325	0.783 9695	0.754 015	0.771 533	0.783 969	0.677 778
0.720 896	0.689 899	0.713 053	0.686 425	0.693 735	0.704 367	0.687 05	0.741 042	0.701 399	0.664 103	0.590 608	0.6 56	0.645 833	0.582 119	0.572 289	0.619 084	0.596 35	0.602 19	0.600 763	0.677 778
0.771 045	0.769 091	0.778 737	0.746 154	0.724 578	0.774 236	0.761 871	0.780 13	0.790 909	0.790 598	0.714 365	0.7 76	0.808 333	0.778 146	0.654 217	0.704 5802	0.713 139	0.718 978	0.692 366	0.677 778
0.749 552	0.772 323	0.793 895	0.787 783	0.770 843	0.732 314	0.758 993	0.751 466	0.765 734	0.729 06	0.736 464	0.7 2	0.725	0.688 079	0.726 506	0.771 7557	0.724 818	0.759 854	0.722 901	0.677 778
0.892 836	0.809 495	0.802 316	0.760 633	0.793 976	0.774 236	0.796 403	0.777 524	0.771 329	0.787 179	0.771 823	0.7 8	0.795 833	0.725 166	0.779 518	0.820 6107	0.789 051	0.800 73	0.832 824	0.722 222
0.861 791	0.782 02	0.778 737	0.733 484	0.766 988	0.833 624	0.848 201	0.847 883	0.824 476	0.811 111	0.780 663	0.7 44	0.745 833	0.730 464	0.740 964	0.808 3969	0.765 693	0.794 891	0.722 901	0.722 222
0.730 448	0.673 737	0.679 368	0.641 176	0.655 181	0.651 965	0.689 928	0.730 619	0.726 573	0.718 803	0.643 646	0.6 72	0.662 5	0.460 265	0.716 867	0.710 687	0.718 978	0.771 533	0.661 832	0.766 667
0.9	0.819 192	0.825 895	0.785 973	0.805 542	0.788 21	0.819 424	0.785 342	0.774 126	0.800 855	0.709 945	0.7 52	0.779 167	0.719 868	0.765 06	0.820 6107	0.812 409	0.794 891	0.838 931	0.766 667
0.771 045	0.798 182	0.815 789	0.766 063	0.765 06	0.714 847	0.770 504	0.814 007	0.816 084	0.790 598	0.727 624	0.7 04	0.712 5	0.682 781	0.716 867	0.710 687	0.701 46	0.748 175	0.661 832	0.766 667
0.809 254	0.790 101	0.834 316	0.789 593	0.780 482	0.707 86	0.825 18	0.866 124	0.841 259	0.817 949	0.736 464	0.8 16	0.8	0.709 272	0.567 47	0.606 8702	0.771 533	0.824 088	0.606 87	0.811 111
0.718 507	0.709 293	0.726 526	0.709 955	0.726 506	0.679 913	0.712 95	0.722 801	0.701 399	0.667 521	0.705 525	0.6 64	0.704 167	0.714 57	0.702 41	0.753 4351	0.754 015	0.748 175	0.741 221	0.855 556
0.787 761	0.762 626	0.748 421	0.738 914	0.753 494	0.795 197	0.784 892	0.793 16	0.765 734	0.790 598	0.727 624	0.7 72	0.758 333	0.783 444	0.673 494	0.759 542	0.748 175	0.777 372	0.747 328	0.855 556

0.685 075	0.664 04	0.691 158	0.652 036	0.664 819	0.683 406	0.698 561	0.730 619	0.718 182	0.694 872	0.701 105	0.6 88	0.679 167	0.629 801	0.716 867	0.759 542	0.701 46	0.678 102	0.735 115	0.855 556
0.754 328	0.775 556	0.785 474	0.775 113	0.728 434	0.697 38	0.724 46	0.769 707	0.762 937	0.718 803	0.648 066	0.6 96	0.704 167	0.629 801	0.765 06	0.741 2214	0.754 015	0.754 015	0.722 901	0.855 556
0.790 149	0.769 091	0.809 053	0.798 643	0.776 627	0.704 367	0.761 871	0.808 795	0.799 301	0.790 598	0.771 823	0.7 48	0.775 232	0.815 232	0.846 988	0.881 6794	0.829 927	0.882 482	0.796 183	0.9
0.864 179	0.835 354	0.836	0.796 833	0.826 747	0.861 572	0.876 978	0.892 182	0.863 636	0.9	0.895 58	0.9	0.870 833	0.9	0.880 723	0.9	0.9	0.9	0.9	0.9
0.744 776	0.693 131	0.733 263	0.726 244	0.711 084	0.742 795	0.715 827	0.741 042	0.726 573	0.718 803	0.692 265	0.7 24	0.716 667	0.730 464	0.822 892	0.790 0763	0.806 569	0.853 285	0.753 435	0.9
0.885 672	0.801 414	0.829 263	0.795 023	0.786 265	0.868 559	0.845 324	0.881 759	0.863 636	0.848 718	0.789 503	0.8 72	0.837 5	0.862 914	0.895 181	0.863 3588	0.847 445	0.859 124	0.869 466	0.9
0.756 716	0.743 232	0.760 211	0.733 484	0.766 988	0.697 38	0.730 216	0.761 889	0.765 734	0.725 641	0.648 066	0.7 2	0.687 5	0.703 974	0.639 759	0.759 542	0.707 299	0.748 175	0.710 687	0.9
0.682 687	0.676 97	0.699 579	0.666 516	0.674 458	0.679 913	0.715 827	0.735 831	0.720 979	0.694 872	0.701 105	0.7 12	0.683 333	0.619 205	0.707 229	0.747 3282	0.678 102	0.689 781	0.704 58	0.9

APPENDIX E

NORMALIZED TRAINING DATASET OF INPUT FEATURES AND TARGET FEATURES FOR MALE HISPANIC POPULATION USING ANN APPROACH

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	Age
0.124 299	0.124 516	0.123 491	0.118 944	0.126 154	0.128 387	0.131 549	0.124 365	0.136 856	0.123 92	0.137 559	0.146 693	0.156 667	0.126 23	0.141 5094	0.137 427	0.150 286	0.127 586	0.110 063	0.1
0.172 897	0.194 194	0.187 439	0.196 175	0.18 161	0.185 155	0.199 431	0.195 065	0.204 05	0.185 164	0.205 837	0.205 0.21	0.200 546	0.216 9811	0.202 924	0.191 429	0.224 138	0.190 566	0.1	
0.193 458	0.216 129	0.203 1	0.193 26	0.198 462	0.187 742	0.223 944	0.221 827	0.240 921	0.224 917	0.205 164	0.208 949	0.216 667	0.200 546	0.190 566	0.156 14	0.173 143	0.145 977	0.120 126	0.1
0.320 561	0.369 677	0.351 876	0.346 266	0.36 0.36	0.314 194	0.368 169	0.400 508	0.407 859	0.408 306	0.385 446	0.395 72	0.41 0.41	0.384 153	0.318 8679	0.277 778	0.305 714	0.302 299	0.276 101	0.144 444
0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.144 444
0.232 71	0.278 065	0.247 471	0.253 005	0.243 077	0.234 194	0.291 549	0.288 832	0.271 274	0.264 784	0.238 967	0.240 078	0.25 0.25	0.266 12	0.326 4151	0.235 673	0.237 143	0.242 529	0.230 818	0.144 444
0.223 364	0.262 581	0.244 861	0.236 976	0.238 462	0.229 032	0.302 817	0.292 893	0.310 298	0.294 02	0.269 014	0.283 658	0.273 333	0.183 06	0.330 1887	0.240 351	0.26 0.26	0.270 115	0.200 629	0.144 444
0.230 841	0.248 387	0.244 861	0.254 463	0.24 0.24	0.254 839	0.284 789	0.268 528	0.282 114	0.280 731	0.257 746	0.261 868	0.28 0.28	0.257 377	0.269 8113	0.259 064	0.250 857	0.260 92	0.195 597	0.144 444
0.281 308	0.323 226	0.314 029	0.309 836	0.306 154	0.270 323	0.347 887	0.349 746	0.357 995	0.339 203	0.317 84	0.317 899	0.33 0.33	0.287 978	0.379 2453	0.291 813	0.356 0.356	0.339 08	0.281 132	0.188 889
0.324 299	0.358 065	0.357 096	0.343 352	0.352 308	0.288 387	0.370 423	0.365 99	0.384 011	0.363 123	0.374 178	0.370 817	0.39 0.39	0.331 694	0.349 0566	0.282 456	0.333 143	0.339 08	0.261 006	0.188 889
0.3	0.338 71	0.308 809	0.306 922	0.303 077	0.290 968	0.354 648	0.337 563	0.353 659	0.328 571	0.317 84	0.317 899	0.336 667	0.253 005	0.383 0189	0.287 135	0.342 286	0.343 678	0.276 101	0.188 889
0.350 467	0.368 387	0.347 961	0.357 923	0.343 077	0.309 032	0.354 648	0.359 898	0.375 339	0.339 203	0.374 178	0.349 027	0.363 333	0.305 464	0.367 9245	0.361 988	0.273 714	0.343 678	0.255 975	0.233 333
0.316 822	0.385 161	0.380 587	0.373 953	0.364 615	0.306 452	0.417 746	0.428 934	0.453 388	0.413 621	0.377 934	0.426 848	0.43 0.43	0.384 153	0.394 3396	0.324 561	0.282 857	0.320 69	0.276 101	0.233 333

0.118	0.146	0.144	0.139		0.138	0.142	0.132	0.141	0.126	0.130	0.143	0.146	0.134	0.160	0.132	0.141	0.141	0.120	0.233
692	452	372	344	0.14	71	817	487	192	578	047	58	667	973	3774	749	143	379	126	333
0.371	0.416	0.421	0.413	0.401	0.350	0.444	0.441	0.440	0.413	0.426	0.417		0.406	0.443	0.404	0.392	0.385	0.396	0.277
028	129	044	297	538	323	789	117	379	621	761	51	0.43	011	3962	094	571	057	855	778
0.329	0.398	0.388	0.395	0.376	0.332	0.431	0.443	0.455	0.418	0.456	0.470		0.397	0.435	0.446	0.424	0.458	0.416	0.277
907	065	418	811	923	258	268	147	556	937	808	428	0.47	268	8491	199	571	621	981	778
0.356	0.414	0.398	0.400	0.396	0.337	0.410	0.404	0.412	0.392	0.404	0.364	0.363	0.327	0.454	0.436	0.438	0.435	0.422	0.277
075	839	858	182	923	419	986	569	195	359	225	591	333	322	717	842	286	632	013	778
0.460	0.474	0.457	0.458	0.436	0.396	0.512	0.510	0.531	0.482	0.535	0.542	0.543	0.493	0.541	0.511	0.488	0.509	0.467	0.322
748	194	586	47	923	774	394	152	436	724	681	023	333	443	5094	696	571	195	296	222
0.481	0.481	0.482	0.468	0.469	0.396	0.501	0.481	0.485	0.456	0.494	0.495	0.513	0.502	0.503	0.441	0.470	0.486	0.432	0.322
308	935	382	67	231	774	127	726	908	146	366	331	333	186	7736	52	286	207	075	222
0.427	0.498	0.483	0.508	0.496	0.456	0.546	0.522	0.531	0.511	0.535	0.532	0.546	0.567	0.481	0.427	0.438	0.463	0.406	0.322
103	71	687	015	923	129	197	335	436	96	681	685	667	76	1321	485	286	218	918	222
0.408	0.429	0.409	0.391	0.404	0.381	0.401	0.422	0.427	0.387	0.370	0.408	0.423	0.384	0.428	0.376	0.383	0.412	0.371	0.322
411	032	299	439	615	29	972	843	371	043	423	171	333	153	3019	023	429	644	698	222
0.382	0.462	0.441	0.430	0.418	0.352	0.465	0.449	0.466	0.413	0.407	0.414	0.423	0.309	0.443	0.380	0.424	0.431	0.391	0.322
243	581	925	783	462	903	07	239	396	621	981	397	333	836	3962	702	571	034	824	222
0.434	0.546	0.526	0.525	0.509	0.492	0.530	0.520	0.533	0.506	0.535	0.532	0.576	0.537	0.564	0.511	0.511	0.532	0.512	0.366
579	452	754	501	231	258	423	305	604	645	681	685	667	158	1509	696	429	184	579	667
0.556	0.546	0.533	0.525	0.530	0.476	0.559	0.550	0.570	0.549	0.535	0.526		0.515	0.567	0.530	0.534	0.541	0.477	0.366
075	452	279	501	769	774	718	761	461	169	681	459	0.54	301	9245	409	286	379	358	667
0.5	0.515	0.534	0.519	0.504	0.407	0.548	0.542	0.550	0.498	0.546	0.542	0.553	0.541	0.567	0.530	0.538	0.527	0.502	0.366
	484	584	672	615	097	451	64	949	671	948	023	333	53	9245	409	857	586	516	667
0.458	0.502	0.512	0.506	0.483	0.404	0.503		0.535	0.482	0.486	0.482	0.516	0.480	0.477	0.441	0.447	0.435	0.477	0.366
879	581	398	557	077	516	38	0.5	772	724	854	879	667	328	3585	52	429	632	358	667
0.483	0.490	0.492	0.484	0.469	0.448	0.503	0.481	0.490	0.453	0.464	0.445	0.456	0.423	0.488	0.436	0.474		0.452	0.366
178	968	822	699	231	387	38	726	244	488	319	525	667	497	6792	842	857	0.5	201	667
0.423	0.426	0.436	0.429	0.412	0.370	0.462	0.463	0.453	0.421	0.456	0.429		0.388	0.428	0.394	0.410	0.435	0.422	0.366
364	452	705	326	308	968	817	452	388	595	808	961	0.44	525	3019	737	857	632	013	667
0.492	0.528	0.522	0.521	0.518	0.458	0.510	0.512	0.511	0.525	0.505	0.495	0.506	0.462	0.567	0.488	0.465	0.504	0.477	0.366
523	387	838	129	462	71	141	183	924	249	634	331	667	842	9245	304	714	598	358	667
0.556	0.583	0.580	0.572	0.570	0.523	0.568	0.567	0.570	0.543	0.614	0.601		0.528	0.571	0.516	0.538	0.545	0.467	0.411
075	871	261	131	769	226	732	005	461	854	554	167	0.58	415	6981	374	857	977	296	111

0.471 963	0.493 548	0.484 992	0.480 328	0.464 615	0.443 226	0.503 38	0.487 817	0.498 916	0.461 462	0.543 192	0.529 572	0.516 667	0.440 984	0.518 8679	0.432 164	0.442 857	0.477 011	0.411 95	0.411 111
0.503 738	0.519 355	0.512 398	0.491 985	0.490 769	0.481 935	0.523 662	0.495 939	0.492 412	0.493 355	0.561 972	0.560 7	0.54 0.54	0.510 929	0.549 0566	0.492 982	0.493 143	0.536 782	0.512 579	0.411 111
0.535 514	0.559 355	0.546 33	0.545 902	0.524 615	0.525 806	0.548 451	0.530 457	0.540 108	0.517 276	0.580 751	0.566 926	0.566 667	0.572 131	0.567 9245	0.567 836	0.538 857	0.555 172	0.527 673	0.411 111
0.537 383	0.516 774	0.500 653	0.496 357	0.486 154	0.494 839	0.483 099	0.481 726	0.505 42	0.477 409	0.513 146	0.495 331	0.523 333	0.524 044	0.545 283	0.507 018	0.493 143	0.509 195	0.492 453	0.411 111
0.578 505	0.561 935	0.571 126	0.586 703	0.553 846	0.479 355	0.521 408	0.540 609	0.557 453	0.514 618	0.516 901	0.510 895	0.543 333	0.489 071	0.583 0189	0.544 444	0.557 143	0.591 954	0.552 83	0.411 111
0.507 477	0.529 677	0.512 398	0.512 386	0.504 615	0.445 806	0.492 113	0.479 695	0.503 252	0.458 804	0.498 122	0.473 541	0.463 333	0.432 24	0.526 4151	0.460 234	0.484 0.484	0.495 402	0.427 044	0.455 556
0.514 953	0.543 871	0.551 55	0.573 588	0.573 846	0.471 613	0.525 915	0.514 213	0.535 772	0.509 302	0.509 39	0.485 992	0.486 667	0.410 383	0.564 1509	0.544 444	0.520 571	0.550 575	0.532 704	0.455 556
0.584 112	0.551 613	0.561 99	0.570 674	0.553 846	0.481 935	0.586 761	0.577 157	0.609 485	0.557 143	0.546 948	0.585 603	0.586 667	0.532 787	0.598 1132	0.516 374	0.534 286	0.550 575	0.532 704	0.455 556
0.526 168	0.516 774	0.534 584	0.532 787	0.538 462	0.450 968	0.557 465	0.542 64	0.574 797	0.549 169	0.554 46	0.538 911	0.55 0.55	0.532 787	0.541 5094	0.525 731	0.511 429	0.518 391	0.497 484	0.455 556
0.514 953	0.502 581	0.515 008	0.513 843	0.5 0.5	0.487 097	0.532 676	0.524 365	0.537 94	0.509 302	0.535 681	0.576 265	0.573 333	0.524 044	0.552 8302	0.516 374	0.516 0.516	0.527 586	0.502 516	0.455 556
0.576 636	0.581 29	0.595 922	0.577 96	0.555 385	0.520 645	0.552 958	0.550 761	0.561 789	0.519 934	0.561 972	0.526 459	0.5 0.5	0.440 984	0.567 9245	0.553 801	0.584 571	0.578 161	0.532 704	0.455 556
0.571 028	0.565 806	0.576 346	0.569 217	0.543 077	0.494 839	0.557 465	0.550 761	0.566 125	0.522 591	0.588 263	0.545 136	0.536 667	0.567 76	0.594 3396	0.549 123	0.525 143	0.545 977	0.512 579	0.5 0.5
0.701 869	0.687 097	0.663 785	0.669 763	0.68 0.68	0.616 129	0.672 394	0.674 619	0.676 694	0.631 561	0.667 136	0.672 763	0.66 0.66	0.567 76	0.658 4906	0.638 012	0.676 0.676	0.660 92	0.567 925	0.5 0.5
0.548 598	0.521 935	0.517 618	0.502 186	0.513 846	0.456 129	0.521 408	0.530 457	0.529 268	0.485 382	0.483 099	0.495 331	0.493 333	0.384 153	0.545 283	0.502 339	0.511 429	0.532 184	0.502 516	0.5 0.5
0.600 935	0.589 032	0.622 023	0.602 732	0.570 769	0.530 968	0.627 324	0.601 523	0.644 173	0.591 694	0.603 286	0.626 07	0.696 667	0.628 962	0.601 8868	0.567 836	0.625 714	0.647 126	0.598 113	0.5 0.5
0.544 86	0.561 935	0.561 99	0.537 158	0.523 077	0.510 323	0.570 986	0.567 005	0.561 789	0.527 907	0.513 146	0.548 249	0.526 667	0.458 47	0.586 7925	0.502 339	0.529 714	0.509 195	0.497 484	0.5 0.5
0.612 15	0.617 419	0.598 532	0.586 703	0.603 077	0.58 0.58	0.629 577	0.611 675	0.615 989	0.594 352	0.670 892	0.604 28	0.61 0.61	0.550 273	0.711 3208	0.656 725	0.666 857	0.683 908	0.663 522	0.5 0.5

0.561 682	0.568 387	0.593 312	0.576 503	0.556 923	0.533 548	0.566 479	0.575 127	0.579 133	0.570 432	0.573 239	0.573 152	0.586 667	0.589 617	0.616 9811	0.567 836	0.584 571	0.591 954	0.562 893	0.5
0.636 449	0.625 161	0.616 803	0.614 39	0.598 462	0.494 839	0.620 563	0.627 919	0.646 341	0.607 641	0.588 263	0.613 619	0.643 333	0.598 361	0.635 8491	0.586 55	0.621 143	0.628 736	0.603 145	0.544 444
0.628 972	0.629 032	0.635 073	0.621 676	0.595 385	0.533 548	0.625 07	0.611 675	0.613 821	0.573 09	0.610 798	0.622 957	0.623 333	0.545 902	0.586 7925	0.507 018	0.548	0.582 759	0.406 918	0.544 444
0.705 607	0.630 323	0.619 413	0.596 903	0.590 769	0.592 903	0.636 338	0.615 736	0.639 837	0.626 246	0.633 333	0.619 844	0.636 667	0.642 077	0.613 2075	0.577 193	0.584 571	0.578 161	0.593 082	0.544 444
0.578 505	0.551 613	0.556 77	0.567 76	0.515 385	0.497 419	0.58	0.575 127	0.585 637	0.501 329	0.565 728	0.560 7	0.583 333	0.506 557	0.598 1132	0.577 193	0.616 571	0.633 333	0.567 925	0.544 444
0.524 299	0.555 484	0.548 94	0.545 902	0.523 077	0.476 774	0.568 732	0.581 218	0.583 469	0.543 854	0.501 878	0.529 572	0.526 667	0.454 098	0.594 3396	0.507 018	0.538 857	0.550 575	0.502 516	0.544 444
0.591 589	0.592 903	0.568 515	0.557 559	0.553 846	0.528 387	0.58	0.579 188	0.576 965	0.535 88	0.554 46	0.570 039	0.576 667	0.541 53	0.590 566	0.549 123	0.529 714	0.591 954	0.522 642	0.544 444
0.632 71	0.578 71	0.560 685	0.548 816	0.567 692	0.600 645	0.595 775	0.556 853	0.576 965	0.543 854	0.633 333	0.598 054	0.616 667	0.537 158	0.571 6981	0.586 55	0.593 714	0.633 333	0.613 208	0.544 444
0.602 804	0.654 839	0.662 48	0.653 734	0.635 385	0.569 677	0.622 817	0.652 284	0.670 19	0.602 326	0.667 136	0.669 65	0.676 667	0.642 077	0.673 5849	0.666 082	0.666 857	0.670 115	0.633 333	0.544 444
0.597 196	0.621 29	0.620 718	0.611 475	0.610 769	0.577 419	0.573 239	0.595 431	0.607 317	0.602 326	0.580 751	0.601 167	0.616 667	0.633 333	0.616 9811	0.591 228	0.602 857	0.637 931	0.648 428	0.544 444
0.606 542	0.573 548	0.572 431	0.567 76	0.549 231	0.523 226	0.566 479	0.567 005	0.563 957	0.509 302	0.584 507	0.594 942	0.58	0.528 415	0.635 8491	0.600 585	0.625 714	0.656 322	0.593 082	0.588 889
0.608 411	0.625 161	0.636 378	0.624 59	0.624 615	0.585 161	0.634 085	0.640 102	0.655 014	0.623 588	0.637 089	0.660 311	0.68	0.642 077	0.703 7736	0.689 474	0.644	0.660 92	0.683 648	0.588 889
0.696 262	0.674 194	0.662 48	0.662 477	0.636 923	0.647 097	0.721 972	0.713 198	0.713 55	0.655 482	0.644 601	0.650 973	0.633 333	0.510 929	0.677 3585	0.605 263	0.680 571	0.670 115	0.577 987	0.588 889
0.664 486	0.641 935	0.637 684	0.630 419	0.615 385	0.549 032	0.625 07	0.603 553	0.628 997	0.634 219	0.652 113	0.657 198	0.653 333	0.668 306	0.666 0377	0.642 69	0.653 143	0.647 126	0.598 113	0.588 889
0.800 935	0.782 581	0.769 494	0.767 395	0.770 769	0.724 516	0.780 563	0.778 173	0.793 767	0.740 532	0.764 789	0.766 148	0.783 333	0.642 077	0.760 3774	0.736 257	0.785 714	0.748 276	0.703 774	0.588 889
0.599 065	0.608 387	0.611 582	0.612 933	0.576 923	0.549 032	0.604 789	0.605 584	0.609 485	0.583 721	0.618 31	0.588 716	0.61	0.607 104	0.624 5283	0.628 655	0.666 857	0.679 31	0.678 616	0.588 889
0.314 953	0.292 258	0.289 233	0.286 521	0.296 923	0.272 903	0.278 028	0.266 497	0.282 114	0.262 126	0.272 77	0.252 529	0.253 333	0.253 005	0.296 2264	0.268 421	0.250 857	0.242 529	0.240 881	0.588 889

0.328 037	0.306 452	0.299 674	0.285 064	0.295 385	0.288 387	0.291 549	0.290 863	0.297 29	0.283 389	0.314 085	0.317 899	0.336 667	0.301 093	0.345 283	0.319 883	0.328 571	0.352 874	0.316 352	0.588 889
0.814 019	0.747 742	0.761 664	0.744 08	0.72	0.670 323	0.726 479	0.733 503	0.743 902	0.708 638	0.727 23	0.703 891	0.736 667	0.712 022	0.764 1509	0.680 117	0.648 571	0.734 483	0.688 679	0.588 889
0.630 841	0.640 645	0.614 192	0.602 732	0.62	0.525 806	0.604 789	0.619 797	0.633 333	0.583 721	0.576 995	0.576 265	0.593 333	0.572 131	0.654 717	0.614 62	0.602 857	0.610 345	0.633 333	0.588 889
0.567 29	0.538 71	0.563 295	0.554 645	0.535 385	0.484 516	0.528 169	0.540 609	0.550 949	0.509 302	0.528 169	0.538 911	0.563 333	0.493 443	0.552 8302	0.535 088	0.520 571	0.550 575	0.502 516	0.588 889
0.664 486	0.645 806	0.636 378	0.623 133	0.607 692	0.528 387	0.58	0.577 157	0.592 141	0.557 143	0.588 263	0.601 167	0.633 333	0.620 219	0.635 8491	0.595 906	0.625 714	0.633 333	0.638 365	0.588 889
0.645 794	0.697 419	0.696 411	0.704 736	0.672 308	0.634 194	0.649 859	0.672 589	0.698 374	0.655 482	0.700 939	0.716 342	0.723 333	0.720 765	0.726 4151	0.703 509	0.703 429	0.729 885	0.713 836	0.633 333
0.681 308	0.675 484	0.689 886	0.682 878	0.649 231	0.605 806	0.679 155	0.688 832	0.683 198	0.647 508	0.723 474	0.722 568	0.736 667	0.703 279	0.722 6415	0.689 474	0.708	0.720 69	0.703 774	0.633 333
0.784 112	0.769 677	0.723 817	0.728 051	0.713 846	0.745 161	0.758 028	0.737 563	0.767 751	0.732 558	0.761 033	0.756 809	0.73	0.611 475	0.7	0.717 544	0.717 143	0.706 897	0.718 868	0.633 333
0.778 505	0.782 581	0.736 868	0.707 65	0.730 769	0.758 065	0.767 042	0.749 746	0.778 591	0.732 558	0.757 277	0.756 809	0.76	0.633 333	0.711 3208	0.740 936	0.744 571	0.729 885	0.713 836	0.633 333
0.757 944	0.781 29	0.777 325	0.774 681	0.773 846	0.711 613	0.796 338	0.806 599	0.793 767	0.788 372	0.640 845	0.735 019	0.726 667	0.628 962	0.783 0189	0.623 977	0.589 143	0.660 92	0.643 396	0.633 333
0.748 598	0.727 097	0.717 292	0.691 621	0.696 923	0.750 323	0.758 028	0.707 107	0.730 894	0.729 9	0.693 427	0.731 907	0.75	0.742 623	0.760 3774	0.703 509	0.721 714	0.757 471	0.728 931	0.633 333
0.642 056	0.657 419	0.638 989	0.612 933	0.606 154	0.569 677	0.613 803	0.629 949	0.615 989	0.589 037	0.584 507	0.585 603	0.6	0.489 071	0.662 2642	0.661 404	0.703 429	0.716 092	0.653 459	0.633 333
0.741 121	0.719 355	0.725 122	0.701 821	0.696 923	0.685 806	0.755 775	0.749 746	0.748 238	0.764 452	0.723 474	0.731 907	0.76	0.799 454	0.771 6981	0.745 614	0.735 429	0.739 08	0.779 245	0.633 333
0.752 336	0.715 484	0.727 732	0.700 364	0.724 615	0.696 129	0.749 014	0.763 959	0.791 599	0.751 163	0.727 23	0.731 907	0.746 667	0.729 508	0.730 1887	0.740 936	0.730 857	0.757 471	0.754 088	0.633 333
0.701 869	0.721 935	0.713 377	0.710 565	0.670 769	0.654 839	0.721 972	0.723 35	0.730 894	0.698 007	0.689 671	0.688 327	0.703 333	0.729 508	0.756 6038	0.745 614	0.753 714	0.771 264	0.779 245	0.677 778
0.681 308	0.658 71	0.653 344	0.637 705	0.649 231	0.585 161	0.647 606	0.634 01	0.648 509	0.615 615	0.667 136	0.622 957	0.616 667	0.572 131	0.786 7925	0.661 404	0.612	0.665 517	0.648 428	0.677 778
0.752 336	0.683 226	0.650 734	0.643 534	0.661 538	0.662 581	0.706 197	0.715 228	0.730 894	0.687 375	0.712 207	0.707 004	0.716 667	0.646 448	0.696 2264	0.623 977	0.680 571	0.688 506	0.688 679	0.677 778

0.705 607	0.663 871	0.674 225	0.662 477	0.641 538	0.636 774	0.688 169	0.698 985	0.689 702	0.652 824	0.700 939	0.703 891	0.68	0.611 475	0.681 1321	0.619 298	0.598 286	0.619 54	0.633 333	0.677 778
0.767 29	0.758 065	0.742 088	0.720 765	0.706 154	0.724 516	0.791 831	0.772 081	0.750 407	0.735 216	0.746 009	0.778 599	0.756 667	0.703 279	0.824 5283	0.745 614	0.804	0.798 851	0.738 994	0.677 778
0.714 953	0.776 129	0.766 884	0.755 738	0.746 154	0.667 742	0.721 972	0.737 563	0.752 575	0.716 611	0.607 042	0.660 311	0.71	0.593 989	0.673 5849	0.703 509	0.804	0.785 057	0.774 214	0.677 778
0.815 888	0.804 516	0.795 595	0.758 652	0.76	0.745 161	0.789 577	0.776 142	0.785 095	0.761 794	0.753 521	0.787 938	0.78	0.698 907	0.832 0755	0.740 936	0.744 571	0.752 874	0.769 182	0.677 778
0.795 327	0.778 71	0.755 139	0.748 452	0.76	0.74	0.758 028	0.751 777	0.756 911	0.756 478	0.753 521	0.741 245	0.786 667	0.712 022	0.741 5094	0.731 579	0.772	0.762 069	0.799 371	0.677 778
0.793 458	0.770 968	0.756 444	0.732 423	0.729 231	0.74	0.794 085	0.786 294	0.772 087	0.756 478	0.697 183	0.763 035	0.783 333	0.712 022	0.828 3019	0.722 222	0.808	0.803 448	0.769 182	0.722 222
0.780 374	0.764 516	0.756 444	0.744 08	0.746 154	0.760 645	0.787 324	0.792 386	0.806 775	0.772 425	0.749 765	0.719 455	0.763 333	0.760 109	0.828 3019	0.726 901	0.804	0.831 034	0.733 962	0.722 222
0.815 888	0.817 419	0.823 002	0.818 397	0.804 615	0.760 645	0.805 352	0.822 843	0.828 455	0.780 399	0.832 394	0.794 163	0.813 333	0.781 967	0.775 4717	0.783 041	0.790 286	0.794 253	0.814 465	0.722 222
0.337 383	0.332 258	0.333 605	0.338 98	0.343 077	0.301 29	0.316 338	0.319 289	0.349 322	0.309 967	0.347 887	0.330 35	0.326 667	0.336 066	0.326 4151	0.310 526	0.310 286	0.311 494	0.316 352	0.722 222
0.836 449	0.817 419	0.828 222	0.831 512	0.816 923	0.752 903	0.839 155	0.837 056	0.854 472	0.820 266	0.832 394	0.859 533	0.853 333	0.720 765	0.847 1698	0.801 754	0.826 857	0.812 644	0.844 654	0.722 222
0.728 037	0.773 548	0.762 969	0.752 823	0.74	0.683 226	0.721 972	0.745 685	0.754 743	0.727 243	0.584 507	0.660 311	0.696 667	0.593 989	0.760 3774	0.708 187	0.790 286	0.789 655	0.799 371	0.722 222
0.802 804	0.736 129	0.697 716	0.701 821	0.704 615	0.709 032	0.751 268	0.753 807	0.763 415	0.729 9	0.753 521	0.747 471	0.753 333	0.681 421	0.707 5472	0.675 439	0.717 143	0.725 287	0.698 742	0.722 222
0.834 579	0.787 742	0.774 715	0.781 967	0.786 154	0.789 032	0.830 141	0.822 843	0.837 127	0.841 528	0.794 836	0.781 712	0.84	0.790 71	0.756 6038	0.689 474	0.694 286	0.752 874	0.713 836	0.722 222
0.9	0.879 355	0.9	0.891 257	0.9	0.879 355	0.868 451	0.849 239	0.880 488	0.876 08	0.824 883	0.822 179	0.85	0.799 454	0.816 9811	0.778 363	0.790 286	0.785 057	0.809 434	0.722 222
0.827 103	0.764 516	0.755 139	0.755 738	0.769 231	0.801 935	0.805 352	0.798 477	0.813 279	0.775 083	0.817 371	0.859 533	0.816 667	0.786 339	0.850 9434	0.769 006	0.790 286	0.794 253	0.759 119	0.722 222
0.832 71	0.790 323	0.778 63	0.745 537	0.744 615	0.758 065	0.791 831	0.790 355	0.808 943	0.783 056	0.779 812	0.784 825	0.763 333	0.746 995	0.707 5472	0.703 509	0.758 286	0.743 678	0.804 403	0.722 222
0.800 935	0.777 419	0.760 359	0.730 965	0.727 692	0.750 323	0.818 873	0.782 234	0.776 423	0.756 478	0.719 718	0.744 358	0.756 667	0.712 022	0.820 7547	0.750 292	0.822 286	0.831 034	0.723 899	0.766 667

0.802	0.804	0.813	0.811	0.795	0.791	0.890	0.891			0.843	0.809	0.853	0.790	0.839	0.801	0.817	0.789	0.864	0.766
804	516	866	111	385	613	986	878	0.9	0.9	662	728	333	71	6226	754	714	655	78	667
0.808	0.734	0.704	0.698	0.712	0.719	0.755	0.772	0.767	0.737	0.761	0.753		0.681	0.733	0.666	0.735	0.716	0.708	0.766
411	839	241	907	308	355	775	081	751	874	033	696	0.76	421	9623	082	429	092	805	667
0.791	0.789	0.773	0.755	0.784	0.742	0.839	0.820	0.863	0.844	0.877	0.853	0.876		0.752	0.759	0.790	0.794	0.789	0.766
589	032	409	738	615	581	155	812	144	186	465	307	667	0.9	8302	649	286	253	308	667
0.838	0.749	0.730	0.748	0.723	0.750	0.771	0.743	0.774	0.761	0.794	0.775		0.733	0.666	0.595	0.694	0.752	0.663	0.766
318	032	343	452	077	323	549	655	255	794	836	486	0.77	88	0377	906	286	874	522	667
0.720	0.751	0.748	0.742	0.736	0.745	0.773	0.753	0.763	0.756	0.791	0.713	0.726	0.738	0.760	0.722	0.753	0.757	0.738	0.766
561	613	613	623	923	161	803	807	415	478	08	23	667	251	3774	222	714	471	994	667
0.853	0.893	0.884	0.895		0.721	0.848	0.863	0.841	0.780	0.715	0.772		0.580		0.867	0.858	0.872	0.854	0.766
271	548	339	628	0.88	935	169	452	463	399	962	374	0.75	874	0.9	251	857	414	717	667
0.763	0.752	0.748	0.742	0.736		0.798	0.814	0.830	0.785	0.791	0.822	0.866	0.694	0.850	0.829	0.872	0.886	0.879	0.766
551	903	613	623	923	0.74	592	721	623	714	08	179	667	536	9434	825	571	207	874	667
0.870	0.839	0.813	0.818	0.832	0.807	0.836	0.810	0.830	0.820	0.877	0.843	0.876	0.816	0.775	0.764		0.817	0.744	0.811
093	355	866	397	308	097	901	66	623	266	465	969	667	94	4717	327	0.804	241	025	111
0.817	0.839	0.873	0.863	0.801	0.781	0.841	0.853	0.865	0.809	0.806	0.819		0.786	0.733	0.792	0.822	0.858	0.854	0.811
757	355	899	57	538	29	408	299	312	635	103	066	0.87	339	9623	398	286	621	717	111
0.763	0.769	0.757	0.751	0.736	0.698	0.726	0.790	0.806	0.764	0.746	0.710	0.696	0.707	0.771	0.684	0.685	0.679	0.749	0.811
551	677	749	366	923	71	479	355	775	452	009	117	667	65	6981	795	143	31	057	111
0.847	0.826	0.821	0.795	0.809	0.760	0.818	0.832	0.832	0.801	0.794	0.812	0.833	0.821	0.877	0.825	0.872	0.881	0.889	0.811
664	452	697	082	231	645	873	995	791	661	836	84	333	311	3585	146	571	609	937	111
0.750	0.738	0.729	0.717	0.744	0.693	0.728	0.717	0.739	0.708	0.761	0.728	0.726	0.668	0.828	0.731	0.698	0.734	0.754	0.811
467	71	038	851	615	548	732	259	566	638	033	794	667	306	3019	579	857	483	088	111
0.830	0.813	0.833	0.831	0.815	0.755	0.848	0.851	0.858	0.825	0.869	0.875	0.873	0.746	0.749	0.801	0.817	0.817	0.839	0.811
841	548	442	512	385	484	169	269	808	581	953	097	333	995	0566	754	714	241	623	111
0.868		0.885		0.887	0.752	0.821	0.861	0.847	0.775	0.768	0.775		0.567	0.866		0.890	0.886		0.811
224	0.9	644	0.9	692	903	127	421	967	083	545	486	0.74	76	0377	0.9	857	207	0.9	111
0.720	0.698	0.710	0.690	0.669	0.647	0.708	0.717	0.704	0.674	0.712	0.722		0.633	0.677	0.675	0.630	0.642	0.648	0.811
561	71	767	164	231	097	451	259	878	086	207	568	0.7	333	3585	439	286	529	428	111
0.799	0.759	0.777	0.779	0.746	0.742	0.771	0.774	0.793	0.740	0.768	0.772		0.672	0.779	0.703		0.766	0.764	0.855
065	355	325	053	154	581	549	112	767	532	545	374	0.76	678	2453	509	0.772	667	151	556
0.890	0.827	0.843	0.844	0.798	0.786	0.870	0.867	0.880	0.825	0.851	0.828	0.843	0.830	0.873	0.731	0.831	0.854	0.834	0.855
654	742	883	627	462	452	704	513	488	581	174	405	333	055	5849	579	429	023	591	556

0.838 318	0.796 774	0.813 866	0.802 368	0.792 308	0.765 806	0.823 38	0.808 629	0.806 775	0.793 688	0.806 103	0.784 825	0.833 333	0.803 825	0.847 1698	0.839 181	0.826 857	0.9	0.814 465	0.855 556
0.832 71	0.800 645	0.799 511	0.792 168	0.807 692	0.74 577	0.789 569	0.804 615	0.817 372	0.788 568	0.783 486	0.775 333	0.773 88	0.733 1132	0.798 901	0.726 429	0.735 678	0.743 119	0.759 556	0.855
0.821 495	0.773 548	0.791 68	0.776 138	0.758 462	0.729 677	0.746 761	0.776 142	0.776 423	0.761 794	0.700 939	0.735 019	0.766 667	0.773 224	0.726 4151	0.670 76	0.653 143	0.706 897	0.683 648	0.855 556
0.821 495	0.834 194	0.791 68	0.822 769	0.792 308	0.760 645	0.803 099	0.822 843	0.789 431	0.775 083	0.783 568	0.763 035	0.76 137	0.725 4906	0.858 076	0.797 857	0.826 241	0.817 591	0.834 556	0.855
0.838 318	0.822 581	0.807 341	0.795 082	0.789 231	0.745 161	0.791 831	0.782 234	0.774 255	0.775 083	0.798 592	0.778 599	0.793 333	0.764 481	0.764 1509	0.726 901	0.726 286	0.729 885	0.723 899	0.9
0.873 832	0.832 903	0.803 426	0.808 197	0.810 769	0.9 0.9	0.863 944	0.867 513	0.871 816	0.846 844	0.824 883	0.884 436	0.9 0.9	0.821 311	0.809 434	0.811 111	0.813 143	0.854 023	0.834 591	0.9
0.793 458	0.792 903	0.757 749	0.754 281	0.781 538	0.732 258	0.767 042	0.794 416	0.817 615	0.788 372	0.742 254	0.725 681	0.793 333	0.834 426	0.752 8302	0.759 649	0.772	0.803 448	0.824 528	0.9
0.866 355	0.834 194	0.809 951	0.821 311	0.832 308	0.822 581	0.839 155	0.822 843	0.843 631	0.828 239	0.877 465	0.828 405	0.87 0.87	0.816 94	0.779 2453	0.783 041	0.767 429	0.798 851	0.749 057	0.9
0.804 673	0.791 613	0.776 02	0.771 767	0.743 077	0.742 581	0.751 268	0.741 624	0.743 902	0.737 874	0.719 718	0.753 696	0.723 333	0.716 393	0.741 5094	0.731 579	0.698 857	0.729 885	0.754 088	0.9
0.800 935	0.808 387	0.783 85	0.811 111	0.784 615	0.724 516	0.807 606	0.796 447	0.832 791	0.783 056	0.798 592	0.769 261	0.796 667	0.720 765	0.794 3396	0.843 86	0.863 429	0.890 805	0.824 528	0.9
0.845 794	0.885 806	0.837 357	0.815 483	0.850 769	0.758 065	0.818 873	0.843 147	0.826 287	0.793 688	0.832 394	0.859 533	0.826 667	0.720 765	0.828 3019	0.778 363	0.790 286	0.826 437	0.819 497	0.9
0.821 495	0.830 323	0.795 595	0.796 539	0.787 692	0.804 516	0.861 69	0.865 482	0.837 127	0.830 897	0.892 488	0.843 969	0.833 333	0.803 825	0.828 3019	0.825 146	0.826 857	0.817 241	0.849 686	0.9

APPENDIX F

NORMALIZED TEST DATASET OF INPUT FEATURES AND TARGET FEATURES FOR MALE HISPANIC POPULATION USING ANN APPROACH

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	Age
0.171 028	0.160 645	0.169 168	0.162 659	0.183 077	0.192 903	0.187 887	0.183 249	0.197 561	0.200 997	0.201 408	0.215 175	0.206 667	0.178 689	0.209 434	0.198 246	0.218 857	0.219 54	0.170 44	0.1
0.286 916	0.305 161	0.281 403	0.264 663	0.270 769	0.285 806	0.350 141	0.343 655	0.357 995	0.344 518	0.347 887	0.345 914	0.356 667	0.331 694	0.349 0566	0.263 743	0.305 714	0.311 494	0.250 943	0.188 889
0.290 654	0.329 677	0.321 86	0.321 494	0.324 615	0.309 032	0.372 676	0.374 112	0.390 515	0.355 15	0.389 202	0.392 607	0.41 0.41	0.362 295	0.364 1509	0.324 561	0.328 571	0.352 874	0.296 226	0.188 889
0.376 636	0.390 323	0.366 232	0.371 038	0.341 538	0.378 71	0.433 521	0.426 904	0.433 875	0.416 279	0.385 446	0.401 946	0.413 333	0.340 437	0.413 2075	0.324 561	0.374 286	0.389 655	0.376 73	0.233 333
0.309 346	0.386 452	0.374 062	0.362 295	0.335 385	0.386 452	0.426 761	0.404 569	0.405 691	0.381 728	0.434 272	0.408 171	0.416 667	0.314 208	0.428 3019	0.366 667	0.383 429	0.385 057	0.351 572	0.233 333
0.447 664	0.516 774	0.526 754	0.526 958	0.512 308	0.471 613	0.494 366	0.516 244	0.535 772	0.485 382	0.456 808	0.470 428	0.486 667	0.427 869	0.503 7736	0.413 45	0.470 286	0.463 218	0.381 761	0.277 778
0.283 178	0.360 645	0.345 351	0.340 437	0.313 846	0.26 0.26	0.332 113	0.325 381	0.327 642	0.296 678	0.344 131	0.336 576	0.32 0.32	0.218 033	0.352 8302	0.282 456	0.292 0.292	0.371 264	0.245 912	0.277 778
0.455 14	0.490 968	0.469 331	0.468 67	0.444 615	0.378 71	0.456 056	0.445 178	0.446 883	0.424 252	0.456 808	0.439 3	0.453 333	0.414 754	0.466 0377	0.436 842	0.438 286	0.467 816	0.416 981	0.322 222
0.393 458	0.416 129	0.402 773	0.384 153	0.392 308	0.332 258	0.42 0.42	0.441 117	0.446 883	0.355 15	0.449 296	0.451 751	0.466 667	0.449 727	0.450 9434	0.352 632	0.342 286	0.458 621	0.442 138	0.322 222
0.432 71	0.422 581	0.394 943	0.391 439	0.355 385	0.383 871	0.417 746	0.404 569	0.412 195	0.365 781	0.404 225	0.392 607	0.38 0.38	0.301 093	0.443 3962	0.422 807	0.410 857	0.417 241	0.386 792	0.322 222
0.464 486	0.483 226	0.469 331	0.459 927	0.470 769	0.425 161	0.494 366	0.477 665	0.485 908	0.480 066	0.460 563	0.439 3	0.463 333	0.419 126	0.466 0377	0.441 52	0.438 286	0.444 828	0.396 855	0.322 222
0.443 925	0.438 065	0.447 145	0.439 526	0.424 615	0.389 032	0.42 0.42	0.418 782	0.433 875	0.408 306	0.392 958	0.420 623	0.446 667	0.366 667	0.420 7547	0.408 772	0.415 429	0.421 839	0.386 792	0.366 667
0.483 178	0.516 774	0.496 737	0.481 785	0.470 769	0.432 903	0.469 577	0.467 513	0.477 236	0.461 462	0.479 343	0.504 669	0.506 667	0.480 328	0.496 2264	0.441 52	0.456 571	0.477 011	0.416 981	0.366 667

0.516 822	0.523 226	0.505 873	0.515 301	0.5 0.5	0.456 129	0.503 38	0.514 213	0.520 596	0.469 435	0.479 343	0.476 654	0.503 333	0.440 984	0.492 4528	0.427 485	0.447 429	0.449 425	0.452 201	0.411 111
0.5 065	0.518 065	0.512 398	0.516 758	0.518 462	0.453 548	0.521 408	0.546 701	0.553 117	0.503 987	0.565 728	0.588 716	0.593 333	0.462 842	0.601 8868	0.535 088	0.575 429	0.582 759	0.527 673	0.411 111
0.528 037	0.532 258	0.526 754	0.499 271	0.501 538	0.507 742	0.564 225	0.546 701	0.555 285	0.503 987	0.554 46	0.557 588	0.566 667	0.554 645	0.579 2453	0.511 696	0.534 286	0.559 77	0.517 61	0.411 111
0.479 439	0.485 806	0.490 212	0.487 614	0.449 231	0.42 0.42	0.471 831	0.473 604	0.470 732	0.421 595	0.460 563	0.439 3	0.443 333	0.366 667	0.503 7736	0.455 556	0.452 0.452	0.458 621	0.442 138	0.411 111
0.582 243	0.558 065	0.558 075	0.547 359	0.526 154	0.515 484	0.595 775	0.585 279	0.594 309	0.554 485	0.558 216	0.563 813	0.586 667	0.519 672	0.601 8868	0.553 801	0.570 857	0.573 563	0.537 736	0.455 556
0.576 636	0.581 29	0.561 99	0.521 129	0.546 154	0.520 645	0.58 0.58	0.560 914	0.581 301	0.570 432	0.588 263	0.598 054	0.59 0.59	0.475 956	0.643 3962	0.558 48	0.557 143	0.587 356	0.567 925	0.455 556
0.520 561	0.528 387	0.525 449	0.502 186	0.487 692	0.510 323	0.534 93	0.510 152	0.548 78	0.519 934	0.580 751	0.542 023	0.56 0.56	0.528 415	0.616 9811	0.530 409	0.543 429	0.541 379	0.492 453	0.455 556
0.571 028	0.564 516	0.563 295	0.563 388	0.564 615	0.471 613	0.552 958	0.579 188	0.587 805	0.535 88	0.509 39	0.520 233	0.53 0.53	0.458 47	0.598 1132	0.577 193	0.534 286	0.559 77	0.502 516	0.455 556
0.585 981	0.564 516	0.580 261	0.589 617	0.567 692	0.507 742	0.570 986	0.573 096	0.594 309	0.543 854	0.554 46	0.560 7	0.573 333	0.532 787	0.696 2264	0.558 48	0.634 857	0.656 322	0.588 05	0.5 0.5
0.655 14	0.656 129	0.657 259	0.662 477	0.629 231	0.543 871	0.577 746	0.629 949	0.618 157	0.527 907	0.464 319	0.526 459	0.516 667	0.309 836	0.598 1132	0.535 088	0.593 714	0.633 333	0.507 547	0.5 0.5
0.621 495	0.600 645	0.602 447	0.577 96	0.56 0.56	0.587 742	0.600 282	0.603 553	0.594 309	0.583 721	0.584 507	0.591 829	0.606 667	0.650 82	0.628 3019	0.567 836	0.575 429	0.596 552	0.593 082	0.5 0.5
0.526 168	0.532 258	0.535 889	0.535 701	0.518 462	0.492 258	0.528 169	0.534 518	0.544 444	0.517 276	0.498 122	0.489 105	0.51 0.51	0.462 842	0.616 9811	0.586 55	0.543 429	0.568 966	0.547 799	0.544 444
0.619 626	0.617 419	0.625 938	0.621 676	0.583 077	0.543 871	0.634 085	0.629 949	0.633 333	0.581 063	0.592 019	0.579 377	0.583 333	0.506 557	0.692 4528	0.591 228	0.602 857	0.628 736	0.628 302	0.544 444
0.627 103	0.634 194	0.632 463	0.637 705	0.596 923	0.541 29	0.645 352	0.676 65	0.670 19	0.623 588	0.610 798	0.660 311	0.646 667	0.628 962	0.639 6226	0.628 655	0.634 857	0.642 529	0.623 27	0.544 444
0.585 981	0.600 645	0.611 582	0.589 617	0.595 385	0.507 742	0.570 986	0.583 249	0.594 309	0.591 694	0.580 751	0.585 603	0.603 333	0.611 475	0.658 4906	0.609 942	0.634 857	0.619 54	0.633 333	0.588 889
0.668 224	0.647 097	0.637 684	0.624 59	0.643 077	0.564 516	0.638 592	0.611 675	0.644 173	0.583 721	0.640 845	0.622 957	0.643 333	0.602 732	0.677 3585	0.661 404	0.639 429	0.642 529	0.628 302	0.588 889
0.836 449	0.813 548	0.820 392	0.803 825	0.778 462	0.763 226	0.807 606	0.812 69	0.826 287	0.775 083	0.768 545	0.750 584	0.793 333	0.764 481	0.854 717	0.694 152	0.753 714	0.817 241	0.749 057	0.633 333

0.657 009	0.714 194	0.715 987	0.717 851	0.673 846	0.662 581	0.665 634	0.692 893	0.702 71	0.671 429	0.727 23	0.716 342	0.753 333	0.733 88	0.737 7358	0.708 187	0.721 714	0.748 276	0.738 994	0.633 333
0.722 43	0.702 581	0.729 038	0.713 479	0.689 231	0.654 839	0.712 958	0.715 228	0.724 39	0.682 06	0.776 056	0.750 584	0.776 667	0.738 251	0.779 2453	0.717 544	0.730 857	0.743 678	0.738 994	0.633 333
0.726 168	0.701 29	0.718 597	0.722 222	0.707 692	0.605 806	0.735 493	0.719 289	0.728 726	0.687 375	0.727 23	0.756 809	0.756 667	0.698 907	0.771 6981	0.726 901	0.744 571	0.665 517	0.744 025	0.633 333
0.623 364	0.608 387	0.603 752	0.602 732	0.6 774	0.556 774	0.625 07	0.609 645	0.611 653	0.607 641	0.595 775	0.594 942	0.56 0.56	0.545 902	0.673 5849	0.670 76	0.648 571	0.674 713	0.648 428	0.633 333
0.739 252	0.702 581	0.706 852	0.693 078	0.673 846	0.672 903	0.721 972	0.711 168	0.722 222	0.700 664	0.693 427	0.669 65	0.68 0.68	0.685 792	0.737 7358	0.656 725	0.653 143	0.688 506	0.683 648	0.633 333
0.720 561	0.724 516	0.719 902	0.703 279	0.695 385	0.649 677	0.681 408	0.680 711	0.678 862	0.644 85	0.678 404	0.660 311	0.696 667	0.668 306	0.737 7358	0.656 725	0.666 857	0.683 908	0.658 491	0.677 778
0.868 224	0.840 645	0.841 272	0.832 969	0.809 231	0.773 548	0.823 38	0.828 934	0.843 631	0.812 292	0.813 615	0.775 486	0.833 333	0.808 197	0.866 0377	0.712 865	0.772 0.772	0.854 023	0.769 182	0.677 778
0.757 944	0.770 968	0.777 325	0.771 767	0.730 769	0.721 935	0.758 028	0.721 32	0.776 423	0.761 794	0.791 08	0.738 132	0.77 0.77	0.768 852	0.798 1132	0.745 614	0.74 0.74	0.757 471	0.769 182	0.677 778
0.785 981	0.763 226	0.736 868	0.720 765	0.718 462	0.768 387	0.767 042	0.753 807	0.776 423	0.743 189	0.783 568	0.775 486	0.763 333	0.642 077	0.715 0943	0.726 901	0.653 143	0.725 287	0.718 868	0.677 778
0.670 093	0.634 194	0.646 819	0.644 991	0.661 538	0.649 677	0.681 408	0.701 015	0.709 214	0.692 691	0.667 136	0.678 988	0.706 667	0.703 279	0.760 3774	0.694 152	0.689 714	0.702 299	0.713 836	0.677 778
0.759 813	0.770 968	0.742 088	0.741 166	0.756 923	0.768 387	0.803 099	0.804 569	0.798 103	0.764 452	0.734 742	0.756 809	0.74 0.74	0.755 738	0.779 2453	0.740 936	0.758 286	0.798 851	0.764 151	0.677 778
0.716 822	0.662 581	0.675 53	0.672 678	0.690 769	0.641 935	0.683 662	0.674 619	0.691 87	0.666 113	0.674 648	0.675 875	0.686 667	0.650 82	0.752 8302	0.731 579	0.698 857	0.729 885	0.723 899	0.677 778
0.855 14	0.854 839	0.824 307	0.818 397	0.809 231	0.845 806	0.870 704	0.859 391	0.880 488	0.812 292	0.9 0.9	0.9 0.9	0.813 333	0.773 224	0.828 3019	0.839 181	0.831 429	0.867 816	0.804 403	0.722 222
0.733 645	0.760 645	0.752 529	0.752 823	0.718 462	0.698 71	0.771 549	0.757 868	0.769 919	0.748 505	0.764 789	0.741 245	0.75 0.75	0.768 852	0.779 2453	0.778 363	0.790 286	0.817 241	0.829 56	0.722 222
0.729 907	0.752 903	0.747 308	0.738 251	0.724 615	0.737 419	0.776 056	0.745 685	0.754 743	0.745 847	0.772 3	0.707 004	0.706 667	0.725 137	0.756 6038	0.689 474	0.749 143	0.725 287	0.718 868	0.722 222
0.838 318	0.733 548	0.726 427	0.741 166	0.716 923	0.729 677	0.760 282	0.737 563	0.761 247	0.743 189	0.787 324	0.772 374	0.76 0.76	0.738 251	0.677 3585	0.609 942	0.689 714	0.757 471	0.643 396	0.766 667
0.821 495	0.82 0.82	0.785 155	0.806 74	0.78 0.78	0.750 323	0.812 113	0.818 782	0.793 767	0.775 083	0.791 08	0.738 132	0.746 667	0.712 022	0.866 0377	0.815 789	0.849 714	0.821 839	0.849 686	0.766 667

0.810 28	0.804 516	0.812 561	0.821 311	0.790 769	0.804 516	0.9	0.9	0.9	0.9	0.836 15	0.803 502	0.85	0.799 454	0.839 6226	0.801 754	0.817 714	0.817 241	0.889 937	0.811 111
0.823 364	0.863 871	0.868 679	0.857 741	0.830 769	0.82 676	0.852 676	0.865 482	0.869 648	0.862 791	0.862 441	0.850 195	0.85	0.847 541	0.884 9057	0.853 216	0.826 857	0.858 621	0.804 403	0.811 111
0.812 15	0.747 742	0.753 834	0.752 823	0.750 769	0.745 161	0.785 07	0.802 538	0.798 103	0.753 821	0.749 765	0.744 358	0.753 333	0.768 852	0.775 4717	0.778 363	0.794 857	0.794 253	0.804 403	0.855 556
0.819 626	0.843 226	0.851 713	0.846 084	0.826 154	0.763 226	0.821 127	0.841 117	0.858 808	0.814 95	0.843 662	0.847 082	0.863 333	0.812 568	0.881 1321	0.857 895	0.9	0.9	0.894 969	0.855 556
0.812 15	0.790 323	0.772 104	0.763 024	0.750 769	0.690 968	0.744 507	0.751 777	0.752 575	0.713 953	0.723 474	0.719 455	0.743 333	0.655 191	0.749 0566	0.773 684	0.799 429	0.812 644	0.789 308	0.855 556
0.799 065	0.763 226	0.772 104	0.763 024	0.755 385	0.701 29	0.708 451	0.731 472	0.754 743	0.724 585	0.697 183	0.694 553	0.703 333	0.685 792	0.733 9623	0.769 006	0.804	0.794 253	0.749 057	0.855 556
0.767 29	0.755 484	0.747 308	0.751 366	0.735 385	0.690 968	0.737 746	0.763 959	0.748 238	0.737 874	0.738 498	0.769 261	0.743 333	0.681 421	0.828 3019	0.820 468	0.836	0.858 621	0.809 434	0.9
0.800 935	0.778 71	0.774 715	0.748 452	0.716 923	0.742 581	0.753 521	0.753 807	0.774 255	0.772 425	0.746 009	0.707 004	0.746 667	0.742 623	0.813 2075	0.787 719	0.781 143	0.766 667	0.819 497	0.9

APPENDIX G

NORMALIZED DATASET OF INPUT FEATURES AND TARGET FEATURES FOR FEMALE HISPANIC POPULATION FOR FEATURE SELECTION USING GA APPROACH

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	Age
0.116 716	0.145 253	0.138 737	0.121 719	0.140 482	0.113 974	0.114 388	0.113 029	0.1	0.120 513	0.113 26	0.1 32	0.108 333	0.121 192	0.1	0.1	0.1	0.158 394	0.148 855	0.1
0.171 642	0.255 152	0.239 789	0.214 027	0.198 313	0.173 362	0.169 065	0.172 964	0.175 524	0.188 889	0.228 177	0.1 64	0.179 167	0.269 536	0.239 759	0.173 282	0.164 234	0.216 788	0.148 855	0.144 444
0.121 493	0.124 242	0.130 316	0.125 339	0.123 133	0.106 987	0.1	0.136 482	0.141 958	0.120 513	0.139 779	0.1 44	0.191 667	0.184 768	0.124 096	0.185 496	0.152 555	0.193 431	0.148 855	0.144 444
0.166 866	0.163 03	0.142 105	0.210 407	0.188 675	0.1	0.169 065	0.157 329	0.141 958	0.192 308	0.1	0.1 48	0.108 0.1	0.137 086	0.153 012	0.112 214	0.146 715	0.164 234	0.136 641	0.144 444
0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.111 189	0.1	0.130 939	0.1	0.108 333	0.115 894	0.124 096	0.179 389	0.146 715	0.158 394	0.148 855	0.144 444
0.162 09	0.224 444	0.214 526	0.212 217	0.211 807	0.141 921	0.206 475	0.180 782	0.203 497	0.219 658	0.126 519	0.1 2	0.166 667	0.126 49	0.220 482	0.148 855	0.135 036	0.199 27	0.1	0.144 444
0.228 955	0.302 02	0.302 105	0.291 855	0.312 048	0.222 271	0.252 518	0.256 352	0.315 385	0.226 496	0.325 414	0.2 92	0.295 833	0.253 642	0.312 048	0.344 275	0.345 255	0.345 255	0.240 458	0.188 889
0.169 254	0.242 222	0.224 632	0.232 127	0.225 301	0.145 415	0.148 921	0.167 752	0.161 538	0.141 026	0.192 818	0.1 16	0.15	0.1	0.148 193	0.197 71	0.164 234	0.199 27	0.148 855	0.188 889
0.140 597	0.227 677	0.231 368	0.188 688	0.194 458	0.131 441	0.192 086	0.206 84	0.209 091	0.120 513	0.148 619	0.1 48	0.191 667	0.126 49	0.215 663	0.142 748	0.111 679	0.1	0.142 748	0.188 889
0.217 015	0.242 222	0.251 579	0.259 276	0.229 157	0.169 869	0.186 331	0.201 629	0.197 902	0.185 47	0.263 536	0.2 4	0.266 667	0.200 662	0.181 928	0.240 458	0.205 109	0.275 182	0.228 244	0.188 889
0.169 254	0.251 919	0.254 947	0.235 747	0.238 795	0.180 349	0.241 007	0.251 14	0.197 902	0.195 726	0.276 796	0.2 68	0.25	0.232 45	0.201 205	0.191 603	0.304 38	0.234 307	0.185 496	0.188 889
0.291 045	0.334 343	0.330 737	0.313 575	0.304 337	0.278 166	0.223 741	0.251 14	0.301 399	0.236 752	0.329 834	0.3 12	0.316 667	0.296 026	0.287 952	0.173 282	0.275 182	0.187 591	0.191 603	0.233 333
0.405 672	0.360 202	0.359 368	0.371 493	0.379 518	0.260 699	0.295 683	0.313 681	0.304 196	0.257 265	0.303 315	0.2 8	0.3	0.232 45	0.355 422	0.270 992	0.240 146	0.275 182	0.289 313	0.233 333

0.386 567	0.374 747	0.364 421	0.360 633	0.375 663	0.323 581	0.292 806	0.305 863	0.368 531	0.274 359	0.329 834	0.3 28	0.325	0.237 748	0.437 349	0.393 13	0.415 328	0.432 847	0.277 099	0.233 333
0.26	0.384 444	0.389 684	0.375 113	0.366 024	0.211 79	0.243 885	0.264 169	0.259 441	0.212 821	0.356 354	0.3 32	0.316 667	0.333 113	0.350 602	0.197 71	0.245 985	0.257 664	0.203 817	0.233 333
0.281 493	0.365 051	0.376 211	0.376 923	0.406 506	0.274 672	0.318 705	0.326 71	0.318 182	0.308 547	0.298 895	0.3 44	0.325	0.264 238	0.321 687	0.154 962	0.205 109	0.234 307	0.234 351	0.233 333
0.324 478	0.397 374	0.406 526	0.407 692	0.383 373	0.299 127	0.318 705	0.412 704	0.410 49	0.400 855	0.343 094	0.3 56	0.329 167	0.444 371	0.360 241	0.258 779	0.216 788	0.391 971	0.222 137	0.277 778
0.329 254	0.376 364	0.406 526	0.387 783	0.404 578	0.288 646	0.310 072	0.334 528	0.309 79	0.277 778	0.294 475	0.3 0.3	0.3	0.216 556	0.422 892	0.350 382	0.327 737	0.374 453	0.319 847	0.277 778
0.374 627	0.363 434	0.352 632	0.322 624	0.358 313	0.222 271	0.246 763	0.271 987	0.267 832	0.236 752	0.329 834	0.2 56	0.333 333	0.306 623	0.316 867	0.277 099	0.210 949	0.362 774	0.234 351	0.277 778
0.279 104	0.365 051	0.369 474	0.366 063	0.369 88	0.278 166	0.287 05	0.303 257	0.365 734	0.346 154	0.382 873	0.3 36	0.279 167	0.317 219	0.369 88	0.411 45	0.415 328	0.444 526	0.295 42	0.277 778
0.322 09	0.400 606	0.399 789	0.398 643	0.396 867	0.299 127	0.321 583	0.342 345	0.427 273	0.417 949	0.343 094	0.3 56	0.379 167	0.407 285	0.369 88	0.234 351	0.251 825	0.281 022	0.246 565	0.277 778
0.386 567	0.424 848	0.446 947	0.451 131	0.452 771	0.306 114	0.356 115	0.410 098	0.379 72	0.342 735	0.329 834	0.3 68	0.35	0.301 325	0.326 506	0.380 916	0.374 453	0.421 168	0.387 023	0.322 222
0.503 582	0.460 404	0.492 421	0.463 801	0.479 759	0.435 371	0.405 036	0.412 704	0.416 084	0.414 53	0.360 773	0.3 6	0.379 167	0.444 371	0.461 446	0.313 74	0.321 898	0.356 934	0.319 847	0.322 222
0.360 299	0.470 101	0.460 421	0.452 941	0.475 904	0.292 14	0.361 871	0.368 404	0.365 734	0.315 385	0.276 796	0.3 0.3	0.283 333	0.227 152	0.254 217	0.289 313	0.263 504	0.316 058	0.295 42	0.322 222
0.470 149	0.457 172	0.480 632	0.478 281	0.470 12	0.292 14	0.324 46	0.436 156	0.435 664	0.421 368	0.404 972	0.4 0.4	0.408 333	0.370 199	0.302 41	0.332 061	0.339 416	0.345 255	0.295 42	0.322 222
0.293 433	0.377 98	0.396 421	0.375 113	0.366 024	0.393 45	0.292 806	0.300 651	0.374 126	0.288 034	0.404 972	0.2 96	0.287 5	0.391 391	0.432 53	0.307 634	0.275 182	0.292 701	0.289 313	0.322 222
0.283 881	0.418 384	0.433 474	0.404 072	0.394 94	0.194 323	0.373 381	0.365 798	0.348 951	0.277 778	0.365 193	0.3 0.3	0.333 333	0.317 219	0.422 892	0.283 206	0.251 825	0.292 701	0.264 885	0.322 222
0.453 433	0.442 626	0.484	0.465 611	0.462 41	0.344 541	0.410 791	0.457 003	0.460 839	0.407 692	0.325 414	0.3 76	0.466 667	0.460 265	0.5	0.325 954	0.304 38	0.327 737	0.338 168	0.322 222
0.427 164	0.437 778	0.458 737	0.434 842	0.402 651	0.267 686	0.393 525	0.410 098	0.407 692	0.366 667	0.378 453	0.3 68	0.362 5	0.322 517	0.408 434	0.332 061	0.386 131	0.333 577	0.270 992	0.322 222
0.403 284	0.392 525	0.408 211	0.398 643	0.398 795	0.257 205	0.315 827	0.334 528	0.332 168	0.301 709	0.307 735	0.2 84	0.320 833	0.327 815	0.321 687	0.270 992	0.339 416	0.386 131	0.295 42	0.322 222

0.293 433	0.403 838	0.421 684	0.398 643	0.410 361	0.264 192	0.307 194	0.347 557	0.410 49	0.397 436	0.374 033	0.3 56	0.362 5	0.327 815	0.360 241	0.277 099	0.222 628	0.356 934	0.222 137	0.322 222
0.367 463	0.453 939	0.475 579	0.454 751	0.445 06	0.337 555	0.410 791	0.417 915	0.432 867	0.335 897	0.466 851	0.4 48	0.433 333	0.391 391	0.451 807	0.448 092	0.403 65	0.444 526	0.319 847	0.366 667
0.534 627	0.526 667	0.546 316	0.514 48	0.510 602	0.393 45	0.468 345	0.592 508	0.569 93	0.530 769	0.488 95	0.4 48	0.541 667	0.481 457	0.490 361	0.325 954	0.345 255	0.485 401	0.240 458	0.366 667
0.494 03	0.478 182	0.504 211	0.460 181	0.472 048	0.372 489	0.445 324	0.467 427	0.449 65	0.346 154	0.404 972	0.3 4	0.329 167	0.386 093	0.442 169	0.325 954	0.432 847	0.380 292	0.325 954	0.366 667
0.496 418	0.450 707	0.460 421	0.460 181	0.473 976	0.379 476	0.396 403	0.402 28	0.416 084	0.387 179	0.440 331	0.3 56	0.408 333	0.407 285	0.312 048	0.362 595	0.456 204	0.485 401	0.448 092	0.366 667
0.515 522	0.462 02	0.463 789	0.452 941	0.456 627	0.358 515	0.387 77	0.472 638	0.472 028	0.428 205	0.404 972	0.4 32	0.433 333	0.396 689	0.485 542	0.344 275	0.368 613	0.386 131	0.344 275	0.366 667
0.477 313	0.458 788	0.441 895	0.451 131	0.472 048	0.292 14	0.402 158	0.428 339	0.430 07	0.346 154	0.400 552	0.3 8	0.375 391	0.391 253	0.413 489	0.356 489	0.327 737	0.485 401	0.350 382	0.366 667
0.374 627	0.471 717	0.470 526	0.489 14	0.491 325	0.330 568	0.419 424	0.506 515	0.480 42	0.469 231	0.440 331	0.4 6	0.445 833	0.370 199	0.490 361	0.368 702	0.368 613	0.403 65	0.356 489	0.366 667
0.477 313	0.415 152	0.408 211	0.409 502	0.414 217	0.414 41	0.428 058	0.438 762	0.449 65	0.421 368	0.334 254	0.3 48	0.366 667	0.444 371	0.446 988	0.350 382	0.368 613	0.386 131	0.362 595	0.366 667
0.491 642	0.455 556	0.463 789	0.452 941	0.481 687	0.355 022	0.384 892	0.485 668	0.494 406	0.448 718	0.431 492	0.4 44	0.458 333	0.396 689	0.446 988	0.393 13	0.397 81	0.415 328	0.454 198	0.366 667
0.477 313	0.471 717	0.477 263	0.456 561	0.473 976	0.414 41	0.459 712	0.483 062	0.483 217	0.397 436	0.400 552	0.4 16	0.475 563	0.465 096	0.524 916	0.380 916	0.391 971	0.467 883	0.356 489	0.366 667
0.601 493	0.562 222	0.595 158	0.561 538	0.545 301	0.372 489	0.433 813	0.574 267	0.530 769	0.469 231	0.475 691	0.4 8	0.341 667	0.301 325	0.350 602	0.350 382	0.333 577	0.508 759	0.332 061	0.411 111
0.572 836	0.605 859	0.625 474	0.612 217	0.593 494	0.487 773	0.505 755	0.506 515	0.488 811	0.482 906	0.577 348	0.4 68	0.587 5	0.560 927	0.581 928	0.423 664	0.403 65	0.462 044	0.332 061	0.411 111
0.599 104	0.537 98	0.548	0.547 059	0.543 373	0.389 956	0.520 144	0.548 208	0.525 175	0.493 162	0.546 409	0.5 04	0.483 333	0.481 457	0.533 735	0.362 595	0.351 095	0.351 095	0.454 198	0.411 111
0.606 269	0.518 586	0.521 053	0.498 19	0.522 169	0.571 616	0.523 022	0.561 238	0.575 524	0.482 906	0.612 707	0.5 52	0.595 833	0.598 013	0.610 843	0.619 084	0.450 365	0.643 066	0.600 763	0.411 111
0.458 209	0.552 525	0.569 895	0.492 76	0.551 084	0.491 266	0.5	0.516 938	0.435 664	0.404 274	0.502 21	0.4 96	0.475	0.465 563	0.514 458	0.399 237	0.520 438	0.543 796	0.509 16	0.411 111
0.572 836	0.560 606	0.559 789	0.541 629	0.547 229	0.494 76	0.540 288	0.501 303	0.527 972	0.404 274	0.546 409	0.5 6	0.504 167	0.481 457	0.572 289	0.496 947	0.520 438	0.543 796	0.374 809	0.411 111

0.546 567	0.549 293	0.553 053	0.548 869	0.562 651	0.484 279	0.534 532	0.545 603	0.533 566	0.503 419	0.541 989	0.5 28	0.520 833	0.529 139	0.519 277	0.350 382	0.351 095	0.368 613	0.332 061	0.411 111
0.486 866	0.492 727	0.507 579	0.494 57	0.493 253	0.466 812	0.407 914	0.467 427	0.477 622	0.394 017	0.471 271	0.4 76	0.466 667	0.534 437	0.509 639	0.423 664	0.456 204	0.497 08	0.332 061	0.411 111
0.517 91	0.528 283	0.537 895	0.507 24	0.527 952	0.351 528	0.517 266	0.519 544	0.513 986	0.380 342	0.533 149	0.5 12	0.483 333	0.444 371	0.533 735	0.551 908	0.473 723	0.514 599	0.338 168	0.411 111
0.596 716	0.544 444	0.564 842	0.532 579	0.520 241	0.466 812	0.485 612	0.493 485	0.486 014	0.476 068	0.387 293	0.4 8	0.491 667	0.433 775	0.519 277	0.429 771	0.497 08	0.520 438	0.515 267	0.411 111
0.668 358	0.647 879	0.635 579	0.603 167	0.574 217	0.529 694	0.594 964	0.576 873	0.553 147	0.479 487	0.546 409	0.5 44	0.562 5	0.550 331	0.577 108	0.527 481	0.514 599	0.427 007	0.417 557	0.455 556
0.611 045	0.555 758	0.574 947	0.566 968	0.574 217	0.449 345	0.583 453	0.566 45	0.586 713	0.575 214	0.581 768	0.5 96	0.620 833	0.545 033	0.610 843	0.612 977	0.479 562	0.654 745	0.606 87	0.455 556
0.556 119	0.546 061	0.574 947	0.543 439	0.524 096	0.501 747	0.523 022	0.550 814	0.541 958	0.503 419	0.537 569	0.5 08	0.55 245	0.513 245	0.586 747	0.460 305	0.438 686	0.602 19	0.564 122	0.455 556
0.603 881	0.565 455	0.583 368	0.557 919	0.533 735	0.515 721	0.546 043	0.561 238	0.567 133	0.520 513	0.590 608	0.5 64	0.562 5	0.545 033	0.572 289	0.515 267	0.479 562	0.643 066	0.441 985	0.455 556
0.441 493	0.525 051	0.514 316	0.501 81	0.520 241	0.466 812	0.537 41	0.519 544	0.530 769	0.530 769	0.546 409	0.5 2	0.504 167	0.497 351	0.519 277	0.374 809	0.491 241	0.537 956	0.521 374	0.455 556
0.558 507	0.486 263	0.514 316	0.5 0.5	0.522 169	0.484 279	0.462 59	0.480 456	0.474 825	0.465 812	0.458 011	0.4 32	0.429 167	0.454 967	0.533 735	0.527 481	0.497 08	0.520 438	0.545 802	0.455 556
0.577 612	0.568 687	0.595 158	0.583 258	0.597 349	0.578 603	0.603 597	0.579 479	0.589 51	0.585 47	0.608 287	0.6 08	0.591 667	0.608 609	0.581 928	0.484 733	0.456 204	0.619 708	0.472 519	0.455 556
0.599 104	0.610 707	0.645 684	0.606 787	0.595 422	0.501 747	0.548 921	0.576 873	0.572 727	0.503 419	0.572 928	0.5 88	0.583 333	0.518 543	0.581 928	0.429 771	0.432 847	0.450 365	0.399 237	0.455 556
0.627 761	0.633 333	0.649 053	0.635 747	0.632 048	0.536 681	0.554 676	0.527 362	0.558 741	0.564 957	0.533 149	0.5 28	0.554 167	0.629 801	0.591 566	0.448 092	0.567 153	0.613 869	0.533 588	0.455 556
0.649 254	0.622 02	0.622 105	0.585 068	0.583 855	0.561 135	0.609 353	0.626 384	0.637 063	0.626 496	0.586 188	0.6 12	0.620 833	0.640 397	0.586 747	0.448 092	0.584 672	0.643 066	0.478 626	0.5 0.5
0.706 567	0.686 667	0.699 579	0.677 376	0.687 952	0.603 057	0.643 885	0.675 896	0.684 615	0.626 496	0.670 166	0.7 28	0.712 5	0.725 166	0.634 94	0.631 298	0.695 62	0.724 818	0.661 832	0.5 0.5
0.651 642	0.597 778	0.649 053	0.650 226	0.624 337	0.536 681	0.571 942	0.618 567	0.606 294	0.599 145	0.564 088	0.5 56	0.562 5	0.523 841	0.644 578	0.661 832	0.625 547	0.502 92	0.631 298	0.5 0.5
0.58 0.58	0.586 465	0.638 947	0.612 217	0.626 265	0.459 825	0.635 252	0.655 049	0.637 063	0.626 496	0.625 967	0.6 36	0.608 333	0.608 609	0.615 663	0.649 618	0.637 226	0.637 226	0.619 084	0.5 0.5

0.711 343	0.714 141	0.726 526	0.668 326	0.672 53	0.655 459	0.698 561	0.738 436	0.706 993	0.660 684	0.639 227	0.6 64	0.670 833	0.656 291	0.557 831	0.588 55	0.584 672	0.467 883	0.576 336	0.5
0.568 06	0.552 525	0.544 632	0.496 38	0.508 675	0.452 838	0.494 245	0.490 879	0.466 434	0.452 137	0.475 691	0.4 64	0.454 167	0.396 689	0.475 904	0.521 374	0.479 562	0.485 401	0.496 947	0.5
0.642 09	0.591 313	0.642 316	0.592 308	0.616 627	0.547 162	0.566 187	0.600 326	0.575 524	0.510 256	0.559 669	0.5 8	0.566 667	0.486 755	0.610 843	0.710 687	0.672 263	0.707 299	0.661 832	0.5
0.680 299	0.626 869	0.682 737	0.657 466	0.651 325	0.421 397	0.586 331	0.628 99	0.667 832	0.612 821	0.515 47	0.6 08	0.616 667	0.613 907	0.683 133	0.655 725	0.502 92	0.584 672	0.667 939	0.5
0.699 403	0.701 212	0.706 316	0.670 136	0.691 807	0.644 978	0.684 173	0.686 319	0.695 804	0.609 402	0.643 646	0.6 6	0.645 833	0.682 781	0.668 675	0.704 58	0.683 942	0.660 584	0.625 191	0.5
0.563 284	0.539 596	0.58 308	0.592 771	0.612 76	0.494 76	0.531 655	0.553 42	0.561 538	0.541 026	0.537 569	0.5 52	0.554 167	0.507 947	0.567 47	0.545 802	0.514 599	0.561 314	0.503 053	0.5
0.630 149	0.636 566	0.691 158	0.648 416	0.662 892	0.585 59	0.669 784	0.691 531	0.676 224	0.677 778	0.692 265	0.6 76	0.658 333	0.640 397	0.692 771	0.710 687	0.648 905	0.648 905	0.643 511	0.544 444
0.673 134	0.628 485	0.649 053	0.617 647	0.622 41	0.585 59	0.620 863	0.644 625	0.642 657	0.616 239	0.603 867	0.6 08	0.6 0.6	0.518 543	0.668 675	0.576 336	0.409 489	0.648 905	0.594 656	0.544 444
0.613 433	0.634 949	0.655 789	0.644 796	0.653 253	0.547 162	0.603 597	0.628 99	0.634 266	0.629 915	0.559 669	0.5 32	0.545 833	0.624 503	0.591 566	0.521 374	0.321 898	0.567 153	0.551 908	0.544 444
0.751 94	0.686 667	0.681 053	0.646 606	0.649 398	0.693 886	0.707 194	0.717 59	0.720 979	0.698 291	0.705 525	0.7 44	0.75 0.75	0.703 974	0.736 145	0.692 366	0.666 423	0.713 139	0.667 939	0.544 444
0.663 582	0.613 939	0.647 368	0.619 457	0.639 759	0.582 096	0.546 043	0.576 873	0.553 147	0.544 444	0.577 348	0.5 8	0.570 833	0.539 735	0.610 843	0.606 87	0.596 35	0.602 19	0.594 656	0.544 444
0.680 299	0.657 576	0.672 632	0.659 276	0.686 024	0.620 524	0.652 518	0.681 107	0.690 21	0.664 103	0.603 867	0.6 44	0.637 5	0.603 311	0.620 482	0.674 046	0.666 423	0.678 102	0.643 511	0.544 444
0.642 09	0.604 242	0.608 632	0.594 118	0.593 494	0.491 266	0.577 698	0.602 932	0.614 685	0.602 564	0.577 348	0.6 2	0.595 833	0.592 715	0.615 663	0.454 198	0.619 708	0.643 066	0.496 947	0.544 444
0.577 612	0.562 222	0.58 109	0.556 109	0.537 59	0.466 812	0.488 489	0.545 603	0.541 958	0.489 744	0.480 11	0.5 08	0.516 667	0.507 947	0.586 747	0.441 985	0.596 35	0.613 869	0.435 878	0.544 444
0.747 164	0.673 737	0.694 526	0.673 756	0.670 602	0.644 978	0.655 396	0.688 925	0.706 993	0.667 521	0.652 486	0.6 84	0.725 0.725	0.735 762	0.702 41	0.759 542	0.689 781	0.736 496	0.704 58	0.544 444
0.642 09	0.596 162	0.595 158	0.585 068	0.587 711	0.543 668	0.609 353	0.628 99	0.623 077	0.605 983	0.625 967	0.6 28	0.637 5	0.613 907	0.586 747	0.600 763	0.596 35	0.643 066	0.558 015	0.544 444
0.708 955	0.659 192	0.708 0.708	0.671 946	0.697 59	0.620 524	0.678 417	0.644 625	0.673 427	0.629 915	0.652 486	0.6 52	0.675 0.675	0.619 205	0.760 241	0.655 725	0.637 226	0.713 139	0.735 115	0.544 444

0.694 627	0.649 495	0.664 211	0.617 647	0.622 41	0.624 017	0.695 683	0.694 137	0.673 427	0.684 615	0.705 525	0.6 92	0.637 5	0.688 079	0.731 325	0.649 618	0.637 226	0.660 584	0.619 084	0.544 444
0.654 03	0.628 485	0.644	0.619 457	0.605 06	0.585 59	0.612 23	0.655 049	0.611 888	0.582 051	0.546 409	0.5 92	0.545 833	0.560 927	0.639 759	0.612 977	0.631 387	0.672 263	0.637 405	0.544 444
0.649 254	0.676 97	0.691 158	0.655 656	0.655 181	0.589 083	0.617 986	0.657 655	0.645 455	0.588 889	0.572 928	0.6 16	0.633 333	0.566 225	0.649 398	0.655 725	0.613 869	0.602 19	0.576 336	0.544 444
0.627 761	0.573 535	0.595 158	0.566 968	0.562 651	0.543 668	0.646 763	0.639 414	0.645 455	0.571 795	0.634 807	0.6 44	0.637 5	0.645 695	0.644 578	0.619 084	0.602 19	0.660 584	0.551 908	0.588 889
0.694 627	0.673 737	0.652 421	0.652 036	0.659 036	0.617 031	0.669 784	0.694 137	0.698 601	0.660 684	0.617 127	0.6 44	0.637 5	0.582 119	0.553 012	0.643 511	0.625 547	0.695 62	0.625 191	0.588 889
0.723 284	0.697 98	0.709 684	0.661 086	0.676 386	0.669 432	0.710 072	0.743 648	0.743 357	0.681 197	0.612 707	0.6 56	0.662 5	0.566 225	0.721 687	0.625 191	0.625 547	0.718 978	0.686 26	0.588 889
0.821 194	0.727 071	0.756 842	0.749 774	0.755 422	0.770 742	0.799 281	0.772 313	0.771 329	0.756 41	0.758 564	0.7 64	0.816 667	0.719 868	0.692 771	0.649 618	0.683 942	0.666 423	0.674 046	0.588 889
0.775 821	0.725 455	0.731 579	0.657 466	0.680 241	0.672 926	0.698 561	0.720 195	0.732 168	0.739 316	0.634 807	0.6 44	0.675	0.619 205	0.581 928	0.686 26	0.660 584	0.777 372	0.637 405	0.588 889
0.716 119	0.697 98	0.708	0.688 235	0.724 578	0.651 965	0.687 05	0.722 801	0.720 979	0.708 547	0.661 326	0.6 84	0.670 833	0.645 695	0.634 94	0.704 58	0.672 263	0.701 46	0.686 26	0.588 889
0.708 955	0.651 111	0.665 895	0.623 077	0.647 47	0.610 044	0.678 417	0.704 56	0.709 79	0.684 615	0.612 707	0.6 68	0.616 667	0.428 477	0.692 771	0.692 366	0.678 102	0.765 693	0.643 511	0.588 889
0.694 627	0.709 293	0.718 105	0.728 054	0.720 723	0.606 55	0.658 273	0.678 502	0.701 399	0.691 453	0.630 387	0.6 28	0.629 167	0.661 589	0.668 675	0.649 618	0.654 745	0.648 905	0.637 405	0.588 889
0.754 328	0.691 515	0.692 842	0.626 697	0.682 169	0.711 354	0.695 683	0.691 531	0.681 818	0.725 641	0.634 807	0.6 68	0.695 833	0.672 185	0.639 759	0.680 153	0.608 029	0.672 263	0.570 229	0.588 889
0.785 373	0.717 374	0.765 263	0.715 385	0.736 145	0.721 834	0.730 216	0.756 678	0.734 965	0.718 803	0.683 425	0.7 0.7	0.677 483	0.750 602	0.704 58	0.695 62	0.724 818	0.759 542	0.588 889	
0.718 507	0.675 354	0.692 842	0.677 376	0.709 157	0.631 004	0.672 662	0.707 166	0.712 587	0.691 453	0.639 227	0.6 8	0.675	0.635 099	0.610 843	0.692 366	0.654 745	0.707 299	0.655 725	0.588 889
0.744 776	0.693 131	0.708	0.680 995	0.680 241	0.728 821	0.733 094	0.741 042	0.734 965	0.711 966	0.701 105	0.7 32	0.7	0.741 06	0.659 036	0.606 87	0.625 547	0.648 905	0.661 832	0.588 889
0.711 343	0.743 232	0.755 158	0.724 434	0.753 494	0.739 301	0.802 158	0.808 795	0.807 692	0.824 786	0.732 044	0.7 92	0.829 167	0.772 848	0.707 229	0.783 969	0.771 533	0.748 175	0.747 328	0.588 889
0.771 045	0.683 434	0.706 316	0.695 475	0.705 301	0.610 044	0.661 151	0.668 078	0.673 427	0.670 94	0.705 525	0.7 16	0.687 5	0.698 675	0.736 145	0.722 901	0.736 496	0.754 015	0.698 473	0.588 889

0.692 239	0.652 727	0.660 842	0.648 416	0.643 614	0.631 004	0.620 863	0.636 808	0.625 874	0.640 171	0.625 967	0.5 96	0.587 5	0.645 695	0.610 843	0.625 191	0.643 066	0.660 584	0.637 405	0.588 889
0.766 269	0.722 222	0.729 895	0.668 326	0.695 663	0.683 406	0.730 216	0.743 648	0.762 937	0.711 966	0.643 646	0.6 92	0.670 833	0.598 013	0.625 301	0.661 832	0.637 226	0.736 496	0.649 618	0.633 333
0.744 776	0.709 293	0.697 895	0.657 466	0.655 181	0.725 328	0.738 849	0.780 13	0.760 14	0.735 897	0.656 906	0.7 44	0.708 333	0.666 887	0.615 663	0.649 618	0.613 869	0.578 832	0.643 511	0.633 333
0.773 433	0.764 242	0.778 737	0.778 733	0.770 843	0.763 755	0.810 791	0.816 612	0.813 287	0.776 923	0.785 083	0.8 12	0.795 833	0.714 57	0.798 795	0.771 756	0.789 051	0.829 927	0.582 443	0.633 333
0.668 358	0.686 667	0.738 316	0.688 235	0.666 747	0.679 913	0.698 561	0.730 619	0.737 762	0.698 291	0.705 525	0.7 24	0.758 333	0.672 185	0.683 133	0.667 939	0.707 299	0.771 533	0.686 26	0.633 333
0.728 06	0.665 657	0.692 842	0.604 977	0.657 108	0.634 498	0.684 173	0.709 772	0.712 587	0.691 453	0.612 707	0.6 64	0.658 333	0.454 967	0.731 325	0.680 153	0.689 781	0.730 657	0.643 511	0.633 333
0.759 104	0.723 838	0.741 684	0.735 294	0.724 578	0.728 821	0.733 094	0.730 619	0.698 601	0.705 128	0.679 006	0.7 28	0.704 167	0.714 57	0.750 602	0.759 542	0.771 533	0.783 212	0.759 542	0.633 333
0.787 761	0.720 606	0.719 789	0.666 516	0.695 663	0.718 341	0.721 583	0.720 195	0.706 993	0.776 923	0.687 845	0.7 12	0.725	0.703 974	0.740 964	0.722 901	0.695 62	0.713 139	0.625 191	0.633 333
0.780 597	0.756 162	0.783 789	0.771 493	0.738 072	0.760 262	0.747 482	0.764 495	0.793 706	0.783 761	0.709 945	0.7 76	0.808 333	0.809 934	0.692 771	0.722 901	0.713 139	0.713 139	0.741 221	0.633 333
0.725 672	0.725 455	0.728 211	0.704 525	0.666 747	0.683 406	0.704 317	0.725 407	0.718 182	0.701 709	0.608 287	0.6 64	0.645 833	0.576 821	0.586 747	0.649 618	0.713 139	0.660 584	0.686 26	0.633 333
0.754 328	0.772 323	0.797 263	0.749 774	0.751 566	0.707 86	0.779 137	0.795 765	0.804 895	0.759 829	0.692 265	0.7 0.7	0.670 833	0.709 272	0.644 578	0.680 153	0.672 263	0.765 693	0.649 618	0.633 333
0.799 701	0.804 646	0.792 211	0.804 072	0.797 831	0.777 729	0.764 748	0.769 707	0.793 706	0.763 248	0.683 425	0.6 96	0.712 5	0.677 483	0.731 325	0.753 435	0.771 533	0.812 409	0.735 115	0.633 333
0.878 507	0.9	0.9	0.9	0.9	0.9	0.9	0.881 759	0.9	0.886 325	0.9	0.8 76	0.9	0.815 232	0.871 084	0.790 076	0.824 088	0.870 803	0.710 687	0.633 333
0.675 522	0.643 03	0.638 947	0.566 968	0.574 217	0.606 55	0.620 863	0.688 925	0.667 832	0.643 59	0.603 867	0.6 48	0.625	0.497 351	0.644 578	0.674 046	0.631 387	0.654 745	0.655 725	0.633 333
0.701 791	0.710 909	0.729 895	0.720 814	0.686 024	0.697 38	0.715 827	0.748 86	0.743 357	0.722 222	0.621 547	0.6 84	0.629 167	0.629 801	0.750 602	0.692 366	0.695 62	0.701 46	0.667 939	0.633 333
0.763 881	0.693 131	0.724 842	0.680 995	0.713 012	0.704 367	0.689 928	0.694 137	0.693 007	0.681 197	0.679 006	0.6 48	0.654 167	0.677 483	0.745 783	0.692 366	0.689 781	0.666 423	0.729 008	0.633 333
0.694 627	0.722 222	0.743 368	0.691 855	0.670 602	0.704 367	0.747 482	0.767 101	0.746 154	0.722 222	0.785 083	0.7 44	0.733 333	0.725 166	0.639 759	0.704 58	0.707 299	0.718 978	0.674 046	0.677 778

0.766 269	0.704 444	0.696 211	0.677 376	0.628 193	0.655 459	0.698 561	0.688 925	0.690 21	0.647 009	0.679 006	0.6 48	0.65	0.560 927	0.731 325	0.783 969	0.754 015	0.771 533	0.783 969	0.677 778
0.790 149	0.735 152	0.750 105	0.700 905	0.734 217	0.756 769	0.805 036	0.821 824	0.807 692	0.794 017	0.798 343	0.7 84	0.791 667	0.825 828	0.755 422	0.790 076	0.800 73	0.800 73	0.783 969	0.677 778
0.720 896	0.689 899	0.713 053	0.686 425	0.693 735	0.704 367	0.687 05	0.741 042	0.701 399	0.664 103	0.590 608	0.6 56	0.645 833	0.582 119	0.572 289	0.619 084	0.596 35	0.602 19	0.600 763	0.677 778
0.771 045	0.715 758	0.706 316	0.666 516	0.684 096	0.714 847	0.747 482	0.733 225	0.734 965	0.705 128	0.701 105	0.6 92	0.687 5	0.640 397	0.712 048	0.832 824	0.847 445	0.806 569	0.777 863	0.677 778
0.728 06	0.641 414	0.686 105	0.626 697	0.653 253	0.672 926	0.689 928	0.722 801	0.709 79	0.722 222	0.643 646	0.6 64	0.65	0.439 073	0.707 229	0.722 901	0.718 978	0.765 693	0.667 939	0.677 778
0.790 149	0.741 616	0.763 579	0.702 715	0.759 277	0.735 808	0.770 504	0.780 13	0.754 545	0.732 479	0.740 884	0.7 52	0.729 167	0.714 57	0.726 506	0.777 863	0.736 496	0.771 533	0.722 901	0.677 778
0.780 597	0.773 939	0.755 158	0.731 674	0.722 651	0.690 393	0.761 871	0.782 736	0.782 517	0.773 504	0.665 746	0.6 44	0.679 167	0.666 887	0.755 422	0.790 076	0.742 336	0.789 051	0.722 901	0.677 778
0.771 045	0.769 091	0.778 737	0.746 154	0.724 578	0.774 236	0.761 871	0.780 13	0.790 909	0.790 598	0.714 365	0.7 76	0.808 333	0.778 146	0.654 217	0.704 58	0.713 139	0.718 978	0.692 366	0.677 778
0.761 493	0.788 485	0.800 632	0.760 633	0.763 133	0.746 288	0.761 871	0.798 371	0.807 692	0.763 248	0.696 685	0.7 32	0.716 667	0.688 079	0.683 133	0.704 58	0.701 46	0.759 854	0.649 618	0.677 778
0.749 552	0.772 323	0.793 895	0.787 783	0.770 843	0.732 314	0.758 993	0.751 466	0.765 734	0.729 06	0.736 464	0.7 2	0.725	0.688 079	0.726 506	0.771 756	0.724 818	0.759 854	0.722 901	0.677 778
0.778 209	0.731 919	0.751 789	0.713 575	0.678 313	0.665 939	0.695 683	0.735 831	0.673 427	0.643 59	0.727 624	0.7 2	0.675	0.693 377	0.716 867	0.698 473	0.678 102	0.695 62	0.649 618	0.677 778
0.735 224	0.688 283	0.672 632	0.633 937	0.657 108	0.651 965	0.735 971	0.741 042	0.723 776	0.670 94	0.639 227	0.6 56	0.637 5	0.322 517	0.625 301	0.704 58	0.654 745	0.701 46	0.667 939	0.677 778
0.823 582	0.759 394	0.770 316	0.738 914	0.732 289	0.718 341	0.741 727	0.748 86	0.737 762	0.688 034	0.709 945	0.6 92	0.720 833	0.666 887	0.726 506	0.747 328	0.730 657	0.742 336	0.729 008	0.677 778
0.737 612	0.752 929	0.748 421	0.684 615	0.713 012	0.742 795	0.761 871	0.772 313	0.757 343	0.752 991	0.705 525	0.7 36	0.712 5	0.698 675	0.750 602	0.851 145	0.818 248	0.847 445	0.747 328	0.677 778
0.723 284	0.688 283	0.699 579	0.693 665	0.691 807	0.721 834	0.687 05	0.748 86	0.712 587	0.667 521	0.617 127	0.6 6	0.670 833	0.582 119	0.567 47	0.625 191	0.572 993	0.625 547	0.594 656	0.722 222
0.892 836	0.809 495	0.802 316	0.760 633	0.793 976	0.774 236	0.796 403	0.777 524	0.771 329	0.787 179	0.771 823	0.7 8	0.795 833	0.725 166	0.779 518	0.820 611	0.789 051	0.800 73	0.832 824	0.722 222
0.771 045	0.743 232	0.783 789	0.796 833	0.778 554	0.704 367	0.724 46	0.738 436	0.754 545	0.763 248	0.705 525	0.7 0	0.654 167	0.735 762	0.687 952	0.729 008	0.701 46	0.718 978	0.735 115	0.722 222

0.708 955	0.701 212	0.736 632	0.719 005	0.709 157	0.704 367	0.681 295	0.694 137	0.690 21	0.688 034	0.656 906	0.6 6	0.666 667	0.709 272	0.408 434	0.729 008	0.748 175	0.777 372	0.729 008	0.722 222
0.797 313	0.773 939	0.797 263	0.782 353	0.768 916	0.784 716	0.735 971	0.756 678	0.788 112	0.800 855	0.727 624	0.8 0.8	0.812 5	0.767 55	0.678 313	0.704 58	0.695 62	0.713 139	0.716 794	0.722 222
0.766 269	0.727 071	0.746 737	0.704 525	0.707 229	0.669 432	0.753 237	0.769 707	0.779 72	0.742 735	0.674 586	0.7 0.7	0.695 833	0.656 291	0.707 229	0.674 046	0.678 102	0.742 336	0.680 153	0.722 222
0.725 672	0.675 354	0.689 474	0.680 995	0.662 892	0.658 952	0.669 784	0.681 107	0.679 021	0.688 034	0.590 608	0.6 48	0.708 333	0.682 781	0.591 566	0.637 405	0.555 474	0.648 905	0.643 511	0.722 222
0.809 254	0.778 788	0.790 526	0.771 493	0.776 627	0.767 249	0.793 525	0.806 189	0.782 517	0.763 248	0.776 243	0.7 96	0.775 0.775	0.772 848	0.726 506	0.820 611	0.829 927	0.841 606	0.790 076	0.722 222
0.861 791	0.782 02	0.778 737	0.733 484	0.766 988	0.833 624	0.848 201	0.847 883	0.824 476	0.811 111	0.780 663	0.7 44	0.745 833	0.730 464	0.740 964	0.808 397	0.765 693	0.794 891	0.722 901	0.722 222
0.828 358	0.741 616	0.780 421	0.749 774	0.766 988	0.746 288	0.779 137	0.774 919	0.760 14	0.746 154	0.740 884	0.7 52	0.725 0.725	0.688 079	0.740 964	0.777 863	0.736 496	0.783 212	0.729 008	0.722 222
0.9 0.9	0.819 192	0.783 789	0.726 244	0.774 699	0.854 585	0.885 612	0.9 0.9	0.877 622	0.831 624	0.864 641	0.8 92	0.887 5	0.831 126	0.630 12	0.747 328	0.759 854	0.789 051	0.674 046	0.722 222
0.775 821	0.775 556	0.817 474	0.787 783	0.807 47	0.721 834	0.733 094	0.741 042	0.737 762	0.718 803	0.683 425	0.7 28	0.7 0.7	0.661 589	0.779 518	0.741 221	0.783 212	0.765 693	0.759 542	0.722 222
0.689 851	0.712 525	0.743 368	0.742 534	0.751 566	0.697 38	0.744 604	0.738 436	0.712 587	0.711 966	0.723 204	0.7 16	0.712 5	0.661 589	0.702 41	0.704 58	0.759 854	0.713 139	0.729 008	0.722 222
0.816 418	0.819 192	0.830 947	0.811 312	0.832 53	0.819 651	0.825 18	0.845 277	0.849 65	0.783 761	0.776 243	0.7 84	0.725 0.725	0.672 185	0.813 253	0.820 611	0.835 766	0.894 161	0.814 504	0.722 222
0.866 567	0.782 02	0.839 368	0.787 783	0.780 482	0.770 742	0.770 504	0.787 948	0.785 315	0.770 085	0.771 823	0.7 36	0.745 833	0.815 232	0.731 325	0.875 573	0.800 73	0.829 927	0.802 29	0.766 667
0.842 687	0.785 253	0.836 0.836	0.809 502	0.813 253	0.739 301	0.802 158	0.832 248	0.793 706	0.794 017	0.727 624	0.7 4	0.720 833	0.762 252	0.716 867	0.783 969	0.742 336	0.789 051	0.765 649	0.766 667
0.756 716	0.730 303	0.738 316	0.708 145	0.757 349	0.676 419	0.721 583	0.751 466	0.751 748	0.735 897	0.665 746	0.7 08	0.704 167	0.666 887	0.615 663	0.759 542	0.695 62	0.736 496	0.686 26	0.766 667
0.730 448	0.673 737	0.679 368	0.641 176	0.655 181	0.651 965	0.689 928	0.730 619	0.726 573	0.718 803	0.643 646	0.6 72	0.662 5	0.460 265	0.716 867	0.710 687	0.718 978	0.771 533	0.661 832	0.766 667
0.9 0.9	0.819 192	0.825 895	0.785 973	0.805 542	0.788 21	0.819 424	0.785 342	0.774 126	0.800 855	0.709 945	0.7 52	0.779 167	0.719 868	0.765 06	0.820 611	0.812 409	0.794 891	0.838 931	0.766 667
0.794 925	0.777 172	0.790 526	0.796 833	0.788 193	0.704 367	0.730 216	0.746 254	0.762 937	0.776 923	0.679 006	0.7 0.7	0.666 667	0.725 166	0.673 494	0.765 649	0.748 175	0.754 015	0.741 221	0.766 667

0.742 388	0.683 434	0.633 895	0.576 018	0.603 133	0.672 926	0.669 784	0.688 925	0.653 846	0.602 564	0.643 646	0.6 28	0.566 667	0.407 285	0.581 928	0.680 153	0.625 547	0.683 942	0.655 725	0.766 667
0.852 239	0.790 101	0.812 421	0.795 023	0.807 47	0.763 755	0.810 791	0.821 824	0.818 881	0.787 179	0.789 503	0.8 04	0.787 5	0.672 185	0.707 229	0.826 718	0.800 73	0.853 285	0.759 542	0.766 667
0.771 045	0.798 182	0.815 789	0.766 063	0.765 06	0.714 847	0.770 504	0.814 007	0.816 084	0.790 598	0.727 624	0.7 04	0.712 5	0.682 781	0.716 867	0.710 687	0.701 46	0.748 175	0.661 832	0.766 667
0.704 179	0.693 131	0.694 526	0.690 045	0.697 59	0.760 262	0.698 561	0.751 466	0.757 343	0.756 41	0.674 586	0.7 04	0.708 333	0.703 974	0.683 133	0.753 435	0.742 336	0.754 015	0.741 221	0.766 667
0.673 134	0.668 889	0.684 421	0.655 656	0.666 747	0.711 354	0.715 827	0.730 619	0.712 587	0.670 94	0.683 425	0.7 08	0.675 099	0.635 099	0.721 687	0.753 435	0.689 781	0.683 942	0.716 794	0.811 111
0.892 836	0.846 667	0.871 368	0.831 222	0.840 241	0.819 651	0.836 691	0.840 065	0.821 678	0.790 598	0.780 663	0.7 84	0.758 333	0.666 887	0.808 434	0.826 718	0.829 927	0.806 569	0.790 076	0.811 111
0.787 761	0.733 535	0.753 474	0.737 104	0.732 289	0.700 873	0.770 504	0.754 072	0.748 951	0.722 222	0.732 044	0.7 2	0.679 167	0.624 503	0.731 325	0.735 115	0.765 693	0.777 372	0.667 939	0.811 111
0.797 313	0.791 717	0.827 579	0.816 742	0.832 53	0.774 236	0.782 014	0.785 342	0.802 098	0.776 923	0.661 326	0.7 2	0.716 667	0.703 974	0.726 506	0.790 076	0.806 569	0.841 606	0.747 328	0.811 111
0.854 627	0.791 717	0.841 053	0.807 692	0.826 747	0.739 301	0.819 424	0.845 277	0.796 503	0.787 179	0.727 624	0.7 64	0.729 167	0.788 742	0.736 145	0.790 076	0.724 818	0.824 088	0.771 756	0.811 111
0.725 672	0.731 919	0.768 632	0.738 914	0.741 928	0.624 017	0.692 806	0.725 407	0.729 371	0.674 359	0.603 867	0.6 04	0.612 5	0.492 053	0.649 398	0.667 939	0.689 781	0.736 496	0.722 901	0.811 111
0.766 269	0.714 141	0.765 263	0.720 814	0.741 928	0.711 354	0.733 094	0.782 736	0.757 343	0.698 291	0.670 166	0.6 72	0.687 5	0.582 119	0.721 687	0.814 504	0.765 693	0.771 533	0.796 183	0.811 111
0.809 254	0.790 101	0.834 316	0.789 593	0.780 482	0.707 86	0.825 18	0.866 124	0.841 259	0.817 949	0.736 464	0.8 16	0.8 0.8	0.709 272	0.567 47	0.606 87	0.771 533	0.824 088	0.606 87	0.811 111
0.785 373	0.741 616	0.734 947	0.719 005	0.699 518	0.721 834	0.756 115	0.754 072	0.746 154	0.722 222	0.705 525	0.7 04	0.704 167	0.661 589	0.707 229	0.838 931	0.829 927	0.829 927	0.832 824	0.811 111
0.825 97	0.769 091	0.790 526	0.773 303	0.772 771	0.760 262	0.793 525	0.814 007	0.802 098	0.759 829	0.789 503	0.8 08	0.791 667	0.767 55	0.731 325	0.875 573	0.835 766	0.864 964	0.771 756	0.811 111
0.744 776	0.662 424	0.686 105	0.626 697	0.687 952	0.693 886	0.721 583	0.756 678	0.723 776	0.752 991	0.648 066	0.6 48	0.670 833	0.756 954	0.654 217	0.680 153	0.718 978	0.765 693	0.759 542	0.855 556
0.716 119	0.752 929	0.804	0.766 063	0.757 349	0.686 9	0.735 971	0.748 86	0.768 531	0.739 316	0.679 006	0.7 04	0.729 167	0.672 185	0.625 301	0.722 901	0.683 942	0.742 336	0.692 366	0.855 556
0.778 209	0.738 384	0.758 526	0.737 104	0.722 651	0.714 847	0.735 971	0.722 801	0.723 776	0.701 709	0.732 044	0.7 08	0.737 5	0.698 675	0.644 578	0.704 58	0.678 102	0.695 62	0.692 366	0.855 556

0.718 507	0.709 293	0.726 526	0.709 955	0.726 506	0.679 913	0.712 95	0.722 801	0.701 399	0.667 521	0.705 525	0.6 64	0.704 167	0.714 57	0.702 41	0.753 435	0.754 015	0.748 175	0.741 221	0.855 556
0.787 761	0.762 626	0.748 421	0.738 914	0.753 494	0.795 197	0.784 892	0.793 16	0.765 734	0.790 598	0.727 624	0.7 72	0.758 333	0.783 444	0.673 494	0.759 542	0.748 175	0.777 372	0.747 328	0.855 556
0.773 433	0.762 626	0.778 737	0.793 213	0.801 687	0.791 703	0.770 504	0.780 13	0.790 909	0.759 829	0.789 503	0.7 8	0.783 333	0.809 934	0.827 711	0.765 649	0.818 248	0.876 642	0.783 969	0.855 556
0.742 388	0.714 141	0.699 579	0.652 036	0.643 614	0.662 445	0.664 029	0.709 772	0.665 035	0.636 752	0.581 768	0.6 44	0.616 667	0.523 841	0.634 94	0.704 58	0.695 62	0.713 139	0.686 26	0.855 556
0.873 731	0.817 576	0.851 158	0.769 683	0.788 193	0.861 572	0.856 835	0.881 759	0.849 65	0.862 393	0.767 403	0.8 52	0.829 167	0.873 51	0.9 0.9	0.851 145	0.847 445	0.888 321	0.838 931	0.855 556
0.685 075	0.664 04	0.691 158	0.652 036	0.664 819	0.683 406	0.698 561	0.730 619	0.718 182	0.694 872	0.701 105	0.6 88	0.679 167	0.629 801	0.716 867	0.759 542	0.701 46	0.678 102	0.735 115	0.855 556
0.754 328	0.775 556	0.785 474	0.775 113	0.728 434	0.697 38	0.724 46	0.769 707	0.762 937	0.718 803	0.648 066	0.6 96	0.704 167	0.629 801	0.765 06	0.741 221	0.754 015	0.754 015	0.722 901	0.855 556
0.790 149	0.769 091	0.809 053	0.798 643	0.776 627	0.704 367	0.761 871	0.808 795	0.799 301	0.790 598	0.771 823	0.7 48	0.775 0.775	0.815 232	0.846 988	0.881 679	0.829 927	0.882 482	0.796 183	0.9
0.864 179	0.835 354	0.836 0.836	0.796 833	0.826 747	0.861 572	0.876 978	0.892 182	0.863 636	0.9 0.9	0.895 58	0.9 0.9	0.870 833	0.9 0.9	0.880 723	0.9 0.9	0.9 0.9	0.9 0.9	0.9 0.9	0.9
0.725 672	0.694 747	0.728 211	0.702 715	0.686 024	0.690 393	0.756 115	0.733 225	0.734 965	0.698 291	0.679 006	0.7 2	0.75 0.75	0.698 675	0.687 952	0.716 794	0.637 226	0.707 299	0.680 153	0.9
0.718 507	0.710 909	0.718 105	0.688 235	0.659 036	0.599 563	0.664 029	0.712 378	0.693 007	0.602 564	0.625 967	0.6 56	0.641 667	0.545 033	0.659 036	0.643 511	0.631 387	0.660 584	0.631 298	0.9
0.744 776	0.693 131	0.733 263	0.726 244	0.711 084	0.742 795	0.715 827	0.741 042	0.726 573	0.718 803	0.692 265	0.7 24	0.716 667	0.730 464	0.822 892	0.790 076	0.806 569	0.853 285	0.753 435	0.9
0.744 776	0.746 465	0.758 526	0.742 534	0.734 217	0.714 847	0.735 971	0.738 436	0.720 979	0.711 966	0.714 365	0.7 64	0.7 0.7	0.703 974	0.789 157	0.783 969	0.794 891	0.777 372	0.771 756	0.9
0.799 701	0.720 606	0.721 474	0.664 706	0.709 157	0.739 301	0.735 971	0.733 225	0.729 371	0.770 085	0.679 006	0.7 08	0.725 0.725	0.714 57	0.606 024	0.741 221	0.695 62	0.730 657	0.625 191	0.9
0.885 672	0.801 414	0.829 263	0.795 023	0.786 265	0.868 559	0.845 324	0.881 759	0.863 636	0.848 718	0.789 503	0.8 72	0.837 5	0.862 914	0.895 181	0.863 359	0.847 445	0.859 124	0.869 466	0.9
0.756 716	0.743 232	0.760 211	0.733 484	0.766 988	0.697 38	0.730 216	0.761 889	0.765 734	0.725 641	0.648 066	0.7 2	0.687 5	0.703 974	0.639 759	0.759 542	0.707 299	0.748 175	0.710 687	0.9
0.682 687	0.676 97	0.699 579	0.666 516	0.674 458	0.679 913	0.715 827	0.735 831	0.720 979	0.694 872	0.701 105	0.7 12	0.683 333	0.619 205	0.707 229	0.747 328	0.678 102	0.689 781	0.704 58	0.9

APPENDIX H

NORMALIZED DATASET OF INPUT FEATURES AND TARGET FEATURES FOR MALE HISPANIC POPULATION FOR FEATURE SELECTION USING GA APPROACH

1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	Age
0.124 299	0.124 516	0.123 491	0.118 944	0.126 154	0.128 387	0.131 549	0.124 365	0.136 856	0.123 92	0.137 559	0.146 693	0.156 667	0.126 23	0.141 509	0.137 427	0.150 286	0.127 586	0.110 063	0.1
0.171 028	0.160 645	0.169 168	0.162 659	0.183 077	0.192 903	0.187 887	0.183 249	0.197 561	0.200 997	0.201 408	0.215 175	0.206 667	0.178 689	0.209 434	0.198 246	0.218 857	0.219 54	0.170 44	0.1
0.172 897	0.194 194	0.187 439	0.196 175	0.18 018	0.185 161	0.199 155	0.195 431	0.204 065	0.185 05	0.205 164	0.205 837	0.205 0.21	0.200 546	0.216 981	0.202 924	0.191 429	0.224 138	0.190 566	0.1
0.193 458	0.216 129	0.203 1	0.193 26	0.198 462	0.187 742	0.223 944	0.221 827	0.240 921	0.224 917	0.205 164	0.208 949	0.216 667	0.200 546	0.190 566	0.156 14	0.173 143	0.145 977	0.120 126	0.1
0.320 561	0.369 677	0.351 876	0.346 266	0.36 036	0.314 194	0.368 169	0.400 508	0.407 859	0.408 306	0.385 446	0.395 72	0.395 0.41	0.384 153	0.318 868	0.277 778	0.305 714	0.302 299	0.276 101	0.144 444
0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.144 444
0.232 71	0.278 065	0.247 471	0.253 005	0.243 077	0.234 194	0.291 549	0.288 832	0.271 274	0.264 784	0.238 967	0.240 078	0.25 0.25	0.266 12	0.326 415	0.235 673	0.237 143	0.242 529	0.230 818	0.144 444
0.223 364	0.262 581	0.244 861	0.236 976	0.238 462	0.229 032	0.302 817	0.292 893	0.310 298	0.294 02	0.269 014	0.283 658	0.273 333	0.183 06	0.330 189	0.240 351	0.26 0.26	0.270 115	0.200 629	0.144 444
0.230 841	0.248 387	0.244 861	0.254 463	0.24 0.24	0.254 839	0.284 789	0.268 528	0.282 114	0.280 731	0.257 746	0.261 868	0.28 0.28	0.257 377	0.269 811	0.259 064	0.250 857	0.260 92	0.195 597	0.144 444
0.286 916	0.305 161	0.281 403	0.264 663	0.270 769	0.285 806	0.350 141	0.343 655	0.357 995	0.344 518	0.347 887	0.345 914	0.356 667	0.331 694	0.349 057	0.263 743	0.305 714	0.311 494	0.250 943	0.188 889
0.290 654	0.329 677	0.321 86	0.321 494	0.324 615	0.309 032	0.372 676	0.374 112	0.390 515	0.355 15	0.389 202	0.392 607	0.41 0.41	0.362 295	0.364 151	0.324 561	0.328 571	0.352 874	0.296 226	0.188 889
0.281 308	0.323 226	0.314 029	0.309 836	0.306 154	0.270 323	0.347 887	0.349 746	0.357 995	0.339 203	0.317 84	0.317 899	0.33 0.33	0.287 978	0.379 245	0.291 813	0.356 0.356	0.339 08	0.281 132	0.188 889
0.324 299	0.358 065	0.357 096	0.343 352	0.352 308	0.288 387	0.370 423	0.365 99	0.384 011	0.363 123	0.374 178	0.370 817	0.39 0.39	0.331 694	0.349 057	0.282 456	0.333 143	0.339 08	0.261 006	0.188 889

0.3	0.338 71	0.308 809	0.306 922	0.303 077	0.290 968	0.354 648	0.337 563	0.353 659	0.328 571	0.317 84	0.317 899	0.336 667	0.253 005	0.383 019	0.287 135	0.342 286	0.343 678	0.276 101	0.188 889
0.350 467	0.368 387	0.347 961	0.357 923	0.343 077	0.309 032	0.354 648	0.359 898	0.375 339	0.339 203	0.374 178	0.349 027	0.363 333	0.305 464	0.367 925	0.361 988	0.273 714	0.343 678	0.255 975	0.233 333
0.316 822	0.385 161	0.380 587	0.373 953	0.364 615	0.306 452	0.417 746	0.428 934	0.453 388	0.413 621	0.377 934	0.426 848	0.43 0.43	0.384 153	0.394 34	0.324 561	0.282 857	0.320 69	0.276 101	0.233 333
0.118 692	0.146 452	0.144 372	0.139 344	0.14 0.14	0.138 71	0.142 817	0.132 487	0.141 192	0.126 578	0.130 047	0.143 58	0.146 667	0.134 973	0.160 377	0.132 749	0.141 143	0.141 379	0.120 126	0.233 333
0.376 636	0.390 323	0.366 232	0.371 038	0.341 538	0.378 71	0.433 521	0.426 904	0.433 875	0.416 279	0.385 446	0.401 946	0.413 333	0.340 437	0.413 208	0.324 561	0.374 286	0.389 655	0.376 73	0.233 333
0.309 346	0.386 452	0.374 062	0.362 295	0.335 385	0.386 452	0.426 761	0.404 569	0.405 691	0.381 728	0.434 272	0.408 171	0.416 667	0.314 208	0.428 302	0.366 667	0.383 429	0.385 057	0.351 572	0.233 333
0.371 028	0.416 129	0.421 044	0.413 297	0.401 538	0.350 323	0.444 789	0.441 117	0.440 379	0.413 621	0.426 761	0.417 51	0.43 0.43	0.406 011	0.443 396	0.404 094	0.392 571	0.385 057	0.396 855	0.277 778
0.329 907	0.398 065	0.388 418	0.395 811	0.376 923	0.332 258	0.431 268	0.443 147	0.455 556	0.418 937	0.456 808	0.470 428	0.47 0.47	0.397 268	0.435 849	0.446 199	0.424 571	0.458 621	0.416 981	0.277 778
0.356 075	0.414 839	0.398 858	0.400 182	0.396 923	0.337 419	0.410 986	0.404 569	0.412 195	0.392 359	0.404 225	0.364 591	0.363 333	0.327 322	0.454 717	0.436 842	0.438 286	0.435 632	0.422 013	0.277 778
0.447 664	0.516 774	0.526 754	0.526 958	0.512 308	0.471 613	0.494 366	0.516 244	0.535 772	0.485 382	0.456 808	0.470 428	0.486 667	0.427 869	0.503 774	0.413 45	0.470 286	0.463 218	0.381 761	0.277 778
0.283 178	0.360 645	0.345 351	0.340 437	0.313 846	0.26 0.26	0.332 113	0.325 381	0.327 642	0.296 678	0.344 131	0.336 576	0.32 0.32	0.218 033	0.352 83	0.282 456	0.292 0.292	0.371 264	0.245 912	0.277 778
0.455 14	0.490 968	0.469 331	0.468 67	0.444 615	0.378 71	0.456 056	0.445 178	0.446 883	0.424 252	0.456 808	0.439 3	0.453 333	0.414 754	0.466 038	0.436 842	0.438 286	0.467 816	0.416 981	0.322 222
0.393 458	0.416 129	0.402 773	0.384 153	0.392 308	0.332 258	0.42 0.42	0.441 117	0.446 883	0.355 15	0.449 296	0.451 751	0.466 667	0.449 727	0.450 943	0.352 632	0.342 286	0.458 621	0.442 138	0.322 222
0.460 748	0.474 194	0.457 586	0.458 47	0.436 923	0.396 774	0.512 394	0.510 152	0.531 436	0.482 724	0.535 681	0.542 023	0.543 333	0.493 443	0.541 509	0.511 696	0.488 571	0.509 195	0.467 296	0.322 222
0.432 71	0.422 581	0.394 943	0.391 439	0.355 385	0.383 871	0.417 746	0.404 569	0.412 195	0.365 781	0.404 225	0.392 607	0.38 0.38	0.301 093	0.443 396	0.422 807	0.410 857	0.417 241	0.386 792	0.322 222
0.481 308	0.481 935	0.482 382	0.468 67	0.469 231	0.396 774	0.501 127	0.481 726	0.485 908	0.456 146	0.494 366	0.495 331	0.513 333	0.502 186	0.503 774	0.441 52	0.470 286	0.486 207	0.432 075	0.322 222
0.427 103	0.498 71	0.483 687	0.508 015	0.496 923	0.456 129	0.546 197	0.522 335	0.531 436	0.511 96	0.535 681	0.532 685	0.546 667	0.567 76	0.481 132	0.427 485	0.438 286	0.463 218	0.406 918	0.322 222

0.464 486	0.483 226	0.469 331	0.459 927	0.470 769	0.425 161	0.494 366	0.477 665	0.485 908	0.480 066	0.460 563	0.439 3	0.463 333	0.419 126	0.466 038	0.441 52	0.438 286	0.444 828	0.396 855	0.322 222
0.408 411	0.429 032	0.409 299	0.391 439	0.404 615	0.381 29	0.401 972	0.422 843	0.427 371	0.387 043	0.370 423	0.408 171	0.423 333	0.384 153	0.428 302	0.376 023	0.383 429	0.412 644	0.371 698	0.322 222
0.382 243	0.462 581	0.441 925	0.430 783	0.418 462	0.352 903	0.465 07	0.449 239	0.466 396	0.413 621	0.407 981	0.414 397	0.423 333	0.309 836	0.443 396	0.380 702	0.424 571	0.431 034	0.391 824	0.322 222
0.434 579	0.546 452	0.526 754	0.525 501	0.509 231	0.492 258	0.530 423	0.520 305	0.533 604	0.506 645	0.535 681	0.532 685	0.576 667	0.537 158	0.564 151	0.511 696	0.511 429	0.532 184	0.512 579	0.366 667
0.556 075	0.546 452	0.533 279	0.525 501	0.530 769	0.476 774	0.559 718	0.550 761	0.570 461	0.549 169	0.535 681	0.526 459	0.54 0.54	0.515 301	0.567 925	0.530 409	0.534 286	0.541 379	0.477 358	0.366 667
0.5 0.5	0.515 484	0.534 584	0.519 672	0.504 615	0.407 097	0.548 451	0.542 64	0.550 949	0.498 671	0.546 948	0.542 023	0.553 333	0.541 53	0.567 925	0.530 409	0.538 857	0.527 586	0.502 516	0.366 667
0.458 879	0.502 581	0.512 398	0.506 557	0.483 077	0.404 516	0.503 38	0.5 0.5	0.535 772	0.482 724	0.486 854	0.482 879	0.516 667	0.480 328	0.477 358	0.441 52	0.447 429	0.435 632	0.477 358	0.366 667
0.443 925	0.438 065	0.447 145	0.439 526	0.424 615	0.389 032	0.42 0.42	0.418 782	0.433 875	0.408 306	0.392 958	0.420 623	0.446 667	0.366 667	0.420 755	0.408 772	0.415 429	0.421 839	0.386 792	0.366 667
0.483 178	0.490 968	0.492 822	0.484 699	0.469 231	0.448 387	0.503 38	0.481 726	0.490 244	0.453 488	0.464 319	0.445 525	0.456 667	0.423 497	0.488 679	0.436 842	0.474 857	0.5 0.5	0.452 201	0.366 667
0.423 364	0.426 452	0.436 705	0.429 326	0.412 308	0.370 968	0.462 817	0.463 452	0.453 388	0.421 595	0.456 808	0.429 961	0.44 0.44	0.388 525	0.428 302	0.394 737	0.410 857	0.435 632	0.422 013	0.366 667
0.483 178	0.516 774	0.496 737	0.481 785	0.470 769	0.432 903	0.469 577	0.467 513	0.477 236	0.461 462	0.479 343	0.504 669	0.506 667	0.480 328	0.496 226	0.441 52	0.456 571	0.477 011	0.416 981	0.366 667
0.492 523	0.528 387	0.522 838	0.521 129	0.518 462	0.458 71	0.510 141	0.512 183	0.511 924	0.525 249	0.505 634	0.495 331	0.506 667	0.462 842	0.567 925	0.488 304	0.465 714	0.504 598	0.477 358	0.366 667
0.556 075	0.583 871	0.580 261	0.572 131	0.570 769	0.523 226	0.568 732	0.567 005	0.570 461	0.543 854	0.614 554	0.601 167	0.58 0.58	0.528 415	0.571 698	0.516 374	0.538 857	0.545 977	0.467 296	0.411 111
0.471 963	0.493 548	0.484 992	0.480 328	0.464 615	0.443 226	0.503 38	0.487 817	0.498 916	0.461 462	0.543 192	0.529 572	0.516 667	0.440 984	0.518 868	0.432 164	0.442 857	0.477 011	0.411 95	0.411 111
0.516 822	0.523 226	0.505 873	0.515 301	0.5 0.5	0.456 129	0.503 38	0.514 213	0.520 596	0.469 435	0.479 343	0.476 654	0.503 333	0.440 984	0.492 453	0.427 485	0.447 429	0.449 425	0.452 201	0.411 111
0.5 0.5	0.518 065	0.512 398	0.516 758	0.518 462	0.453 548	0.521 408	0.546 701	0.553 117	0.503 987	0.565 728	0.588 716	0.593 333	0.462 842	0.601 887	0.535 088	0.575 429	0.582 759	0.527 673	0.411 111
0.503 738	0.519 355	0.512 398	0.491 985	0.490 769	0.481 935	0.523 662	0.495 939	0.492 412	0.493 355	0.561 972	0.560 7	0.54 0.54	0.510 929	0.549 057	0.492 982	0.493 143	0.536 782	0.512 579	0.411 111

0.535	0.559	0.546	0.545	0.524	0.525	0.548	0.530	0.540	0.517	0.580	0.566	0.566	0.572	0.567	0.567	0.538	0.555	0.527	0.411
514	355	33	902	615	806	451	457	108	276	751	926	667	131	925	836	857	172	673	111
0.537	0.516	0.500	0.496	0.486	0.494	0.483	0.481	0.505	0.477	0.513	0.495	0.523	0.524	0.545	0.507	0.493	0.509	0.492	0.411
383	774	653	357	154	839	099	726	42	409	146	331	333	044	283	018	143	195	453	111
0.578	0.561	0.571	0.586	0.553	0.479	0.521	0.540	0.557	0.514	0.516	0.510	0.543	0.489	0.583	0.544	0.557	0.591	0.552	0.411
505	935	126	703	846	355	408	609	453	618	901	895	333	071	019	444	143	954	83	111
0.528	0.532	0.526	0.499	0.501	0.507	0.564	0.546	0.555	0.503	0.554	0.557	0.566	0.554	0.579	0.511	0.534	0.559	0.517	0.411
037	258	754	271	538	742	225	701	285	987	46	588	667	645	245	696	286	77	61	111
0.479	0.485	0.490	0.487	0.449		0.471	0.473	0.470	0.421	0.460	0.439	0.443	0.366	0.503	0.455		0.458	0.442	0.411
439	806	212	614	231	0.42	831	604	732	595	563	3	333	667	774	556	0.452	621	138	111
0.507	0.529	0.512	0.512	0.504	0.445	0.492	0.479	0.503	0.458	0.498	0.473	0.463	0.432	0.526	0.460		0.495	0.427	0.455
477	677	398	386	615	806	113	695	252	804	122	541	333	24	415	234	0.484	402	044	556
0.514	0.543	0.551	0.573	0.573	0.471	0.525	0.514	0.535	0.509	0.509	0.485	0.486	0.410	0.564	0.544	0.520	0.550	0.532	0.455
953	871	55	588	846	613	915	213	772	302	39	992	667	383	151	444	571	575	704	556
0.584	0.551	0.561	0.570	0.553	0.481	0.586	0.577	0.609	0.557	0.546	0.585	0.586	0.532	0.598	0.516	0.534	0.550	0.532	0.455
112	613	99	674	846	935	761	157	485	143	948	603	667	787	113	374	286	575	704	556
0.582	0.558	0.558	0.547	0.526	0.515	0.595	0.585	0.594	0.554	0.558	0.563	0.586	0.519	0.601	0.553	0.570	0.573	0.537	0.455
243	065	075	359	154	484	775	279	309	485	216	813	667	672	887	801	857	563	736	556
0.526	0.516	0.534	0.532	0.538	0.450	0.557	0.542	0.574	0.549	0.554	0.538		0.532	0.541	0.525	0.511	0.518	0.497	0.455
168	774	584	787	462	968	465	64	797	169	46	911	0.55	787	509	731	429	391	484	556
0.576	0.581	0.561	0.521	0.546	0.520		0.560	0.581	0.570	0.588	0.598		0.475	0.643	0.558	0.557	0.587	0.567	0.455
636	29	99	129	154	645	0.58	914	301	432	263	054	0.59	956	396	48	143	356	925	556
0.514	0.502	0.515	0.513		0.487	0.532	0.524	0.537	0.509	0.535	0.576	0.573	0.524	0.552	0.516		0.527	0.502	0.455
953	581	008	843	0.5	097	676	365	94	302	681	265	333	044	83	374	0.516	586	516	556
0.576	0.581	0.595	0.577	0.555	0.520	0.552	0.550	0.561	0.519	0.561	0.526		0.440	0.567	0.553	0.584	0.578	0.532	0.455
636	29	922	96	385	645	958	761	789	934	972	459	0.5	984	925	801	571	161	704	556
0.520	0.528	0.525	0.502	0.487	0.510	0.534	0.510	0.548	0.519	0.580	0.542		0.528	0.616	0.530	0.543	0.541	0.492	0.455
561	387	449	186	692	323	93	152	78	934	751	023	0.56	415	981	409	429	379	453	556
0.571	0.564	0.563	0.563	0.564	0.471	0.552	0.579	0.587	0.535	0.509	0.520		0.458	0.598	0.577	0.534	0.559	0.502	0.455
028	516	295	388	615	613	958	188	805	88	39	233	0.53	47	113	193	286	77	516	556
0.571	0.565	0.576	0.569	0.543	0.494	0.557	0.550	0.566	0.522	0.588	0.545	0.536	0.567	0.594	0.549	0.525	0.545	0.512	
028	806	346	217	077	839	465	761	125	591	263	136	667	76	34	123	143	977	579	0.5
0.701	0.687	0.663	0.669		0.616	0.672	0.674	0.676	0.631	0.667	0.672		0.567	0.658	0.638		0.660	0.567	
869	097	785	763	0.68	129	394	619	694	561	136	763	0.66	76	491	012	0.676	92	925	0.5

0.548 598	0.521 935	0.517 618	0.502 186	0.513 846	0.456 129	0.521 408	0.530 457	0.529 268	0.485 382	0.483 099	0.495 331	0.493 333	0.384 153	0.545 283	0.502 339	0.511 429	0.532 184	0.502 516	0.5
0.585 981	0.564 516	0.580 261	0.589 617	0.567 692	0.507 742	0.570 986	0.573 096	0.594 309	0.543 854	0.554 46	0.560 7	0.573 333	0.532 787	0.696 226	0.558 48	0.634 857	0.656 322	0.588 05	0.5
0.655 14	0.656 129	0.657 259	0.662 477	0.629 231	0.543 871	0.577 746	0.629 949	0.618 157	0.527 907	0.464 319	0.526 459	0.516 667	0.309 836	0.598 113	0.535 088	0.593 714	0.633 333	0.507 547	0.5
0.600 935	0.589 032	0.622 023	0.602 732	0.570 769	0.530 968	0.627 324	0.601 523	0.644 173	0.591 694	0.603 286	0.626 07	0.696 667	0.628 962	0.601 887	0.567 836	0.625 714	0.647 126	0.598 113	0.5
0.544 86	0.561 935	0.561 99	0.537 158	0.523 077	0.510 323	0.570 986	0.567 005	0.561 789	0.527 907	0.513 146	0.548 249	0.526 667	0.458 47	0.586 792	0.502 339	0.529 714	0.509 195	0.497 484	0.5
0.612 15	0.617 419	0.598 532	0.586 703	0.603 077	0.58	0.629 577	0.611 675	0.615 989	0.594 352	0.670 892	0.604 28	0.61	0.550 273	0.711 321	0.656 725	0.666 857	0.683 908	0.663 522	0.5
0.621 495	0.600 645	0.602 447	0.577 96	0.587 0.56	0.587 742	0.600 282	0.603 553	0.594 309	0.583 721	0.584 507	0.591 829	0.606 667	0.650 82	0.628 302	0.567 836	0.575 429	0.596 552	0.593 082	0.5
0.561 682	0.568 387	0.593 312	0.576 503	0.556 923	0.533 548	0.566 479	0.575 127	0.579 133	0.570 432	0.573 239	0.573 152	0.586 667	0.589 617	0.616 981	0.567 836	0.584 571	0.591 954	0.562 893	0.5
0.526 168	0.532 258	0.535 889	0.535 701	0.518 462	0.492 258	0.528 169	0.534 518	0.544 444	0.517 276	0.498 122	0.489 105	0.51	0.462 842	0.616 981	0.586 55	0.543 429	0.568 966	0.547 799	0.544 444
0.636 449	0.625 161	0.616 803	0.614 39	0.598 462	0.494 839	0.620 563	0.627 919	0.646 341	0.607 641	0.588 263	0.613 619	0.643 333	0.598 361	0.635 849	0.586 55	0.621 143	0.628 736	0.603 145	0.544 444
0.628 972	0.629 032	0.635 073	0.621 676	0.595 385	0.533 548	0.625 07	0.611 675	0.613 821	0.573 09	0.610 798	0.622 957	0.623 333	0.545 902	0.586 792	0.507 018	0.548	0.582 759	0.406 918	0.544 444
0.705 607	0.630 323	0.619 413	0.596 903	0.590 769	0.592 903	0.636 338	0.615 736	0.639 837	0.626 246	0.633 333	0.619 844	0.636 667	0.642 077	0.613 208	0.577 193	0.584 571	0.578 161	0.593 082	0.544 444
0.578 505	0.551 613	0.556 77	0.567 76	0.515 385	0.497 419	0.58	0.575 127	0.585 637	0.501 329	0.565 728	0.560 7	0.583 333	0.506 557	0.598 113	0.577 193	0.616 571	0.633 333	0.567 925	0.544 444
0.619 626	0.617 419	0.625 938	0.621 676	0.583 077	0.543 871	0.634 085	0.629 949	0.633 333	0.581 063	0.592 019	0.579 377	0.583 333	0.506 557	0.692 453	0.591 228	0.602 857	0.628 736	0.628 302	0.544 444
0.524 299	0.555 484	0.548 94	0.545 902	0.523 077	0.476 774	0.568 732	0.581 218	0.583 469	0.543 854	0.501 878	0.529 572	0.526 667	0.454 098	0.594 34	0.507 018	0.538 857	0.550 575	0.502 516	0.544 444
0.591 589	0.592 903	0.568 515	0.557 559	0.553 846	0.528 387	0.58	0.579 188	0.576 965	0.535 88	0.554 46	0.570 039	0.576 667	0.541 53	0.590 566	0.549 123	0.529 714	0.591 954	0.522 642	0.544 444
0.632 71	0.578 71	0.560 685	0.548 816	0.567 692	0.600 645	0.595 775	0.556 853	0.576 965	0.543 854	0.633 333	0.598 054	0.616 667	0.537 158	0.571 698	0.586 55	0.593 714	0.633 333	0.613 208	0.544 444

0.602 804	0.654 839	0.662 48	0.653 734	0.635 385	0.569 677	0.622 817	0.652 284	0.670 19	0.602 326	0.667 136	0.669 65	0.676 667	0.642 077	0.673 585	0.666 082	0.666 857	0.670 115	0.633 333	0.544 444
0.597 196	0.621 29	0.620 718	0.611 475	0.610 769	0.577 419	0.573 239	0.595 431	0.607 317	0.602 326	0.580 751	0.601 167	0.616 667	0.633 333	0.616 981	0.591 228	0.602 857	0.637 931	0.648 428	0.544 444
0.627 103	0.634 194	0.632 463	0.637 705	0.596 923	0.541 29	0.645 352	0.676 65	0.670 19	0.623 588	0.610 798	0.660 311	0.646 667	0.628 962	0.639 623	0.628 655	0.634 857	0.642 529	0.623 27	0.544 444
0.606 542	0.573 548	0.572 431	0.567 76	0.549 231	0.523 226	0.566 479	0.567 005	0.563 957	0.509 302	0.584 507	0.594 942	0.58 0.58	0.528 415	0.635 849	0.600 585	0.625 714	0.656 322	0.593 082	0.588 889
0.585 981	0.600 645	0.611 582	0.589 617	0.595 385	0.507 742	0.570 986	0.583 249	0.594 309	0.591 694	0.580 751	0.585 603	0.603 333	0.611 475	0.658 491	0.609 942	0.634 857	0.619 54	0.633 333	0.588 889
0.608 411	0.625 161	0.636 378	0.624 59	0.624 615	0.585 161	0.634 085	0.640 102	0.655 014	0.623 588	0.637 089	0.660 311	0.68 0.68	0.642 077	0.703 774	0.689 474	0.644 0.644	0.660 92	0.683 648	0.588 889
0.696 262	0.674 194	0.662 48	0.662 477	0.636 923	0.647 097	0.721 972	0.713 198	0.713 55	0.655 482	0.644 601	0.650 973	0.633 333	0.510 929	0.677 358	0.605 263	0.680 571	0.670 115	0.577 987	0.588 889
0.664 486	0.641 935	0.637 684	0.630 419	0.615 385	0.549 032	0.625 07	0.603 553	0.628 997	0.634 219	0.652 113	0.657 198	0.653 333	0.668 306	0.666 038	0.642 69	0.653 143	0.647 126	0.598 113	0.588 889
0.800 935	0.782 581	0.769 494	0.767 395	0.770 769	0.724 516	0.780 563	0.778 173	0.793 767	0.740 532	0.764 789	0.766 148	0.783 333	0.642 077	0.760 377	0.736 257	0.785 714	0.748 276	0.703 774	0.588 889
0.599 065	0.608 387	0.611 582	0.612 933	0.576 923	0.549 032	0.604 789	0.605 584	0.609 485	0.583 721	0.618 31	0.588 716	0.61 0.61	0.607 104	0.624 528	0.628 655	0.666 857	0.679 31	0.678 616	0.588 889
0.314 953	0.292 258	0.289 233	0.286 521	0.296 923	0.272 903	0.278 028	0.266 497	0.282 114	0.262 126	0.272 77	0.252 529	0.253 333	0.253 005	0.296 226	0.268 421	0.250 857	0.242 529	0.240 881	0.588 889
0.328 037	0.306 452	0.299 674	0.285 064	0.295 385	0.288 387	0.291 549	0.290 863	0.297 29	0.283 389	0.314 085	0.317 899	0.336 667	0.301 093	0.345 283	0.319 883	0.328 571	0.352 874	0.316 352	0.588 889
0.814 019	0.747 742	0.761 664	0.744 08	0.72 0.72	0.670 323	0.726 479	0.733 503	0.743 902	0.708 638	0.727 23	0.703 891	0.736 667	0.712 022	0.764 151	0.680 117	0.648 571	0.734 483	0.688 679	0.588 889
0.630 841	0.640 645	0.614 192	0.602 732	0.62 0.62	0.525 806	0.604 789	0.619 797	0.633 333	0.583 721	0.576 995	0.576 265	0.593 333	0.572 131	0.654 717	0.614 62	0.602 857	0.610 345	0.633 333	0.588 889
0.567 29	0.538 71	0.563 295	0.554 645	0.535 385	0.484 516	0.528 169	0.540 609	0.550 949	0.509 302	0.528 169	0.538 911	0.563 333	0.493 443	0.552 83	0.535 088	0.520 571	0.550 575	0.502 516	0.588 889
0.664 486	0.645 806	0.636 378	0.623 133	0.607 692	0.528 387	0.58 0.58	0.577 157	0.592 141	0.557 143	0.588 263	0.601 167	0.633 333	0.620 219	0.635 849	0.595 906	0.625 714	0.633 333	0.638 365	0.588 889
0.668 224	0.647 097	0.637 684	0.624 59	0.643 077	0.564 516	0.638 592	0.611 675	0.644 173	0.583 721	0.640 845	0.622 957	0.643 333	0.602 732	0.677 358	0.661 404	0.639 429	0.642 529	0.628 302	0.588 889

0.645 794	0.697 419	0.696 411	0.704 736	0.672 308	0.634 194	0.649 859	0.672 589	0.698 374	0.655 482	0.700 939	0.716 342	0.723 333	0.720 765	0.726 415	0.703 509	0.703 429	0.729 885	0.713 836	0.633 333
0.681 308	0.675 484	0.689 886	0.682 878	0.649 231	0.605 806	0.679 155	0.688 832	0.683 198	0.647 508	0.723 474	0.722 568	0.736 667	0.703 279	0.722 642	0.689 474	0.708 69	0.720 774	0.703 333	0.633 333
0.784 112	0.769 677	0.723 817	0.728 051	0.713 846	0.745 161	0.758 028	0.737 563	0.767 751	0.732 558	0.761 033	0.756 809	0.73 0.73	0.611 475	0.7 0.7	0.717 544	0.717 143	0.706 897	0.718 868	0.633 333
0.836 449	0.813 548	0.820 392	0.803 825	0.778 462	0.763 226	0.807 606	0.812 69	0.826 287	0.775 083	0.768 545	0.750 584	0.793 333	0.764 481	0.854 717	0.694 152	0.753 714	0.817 241	0.749 057	0.633 333
0.657 009	0.714 194	0.715 987	0.717 851	0.673 846	0.662 581	0.665 634	0.692 893	0.702 71	0.671 429	0.727 23	0.716 342	0.753 333	0.733 88	0.737 736	0.708 187	0.721 714	0.748 276	0.738 994	0.633 333
0.722 43	0.702 581	0.729 038	0.713 479	0.689 231	0.654 839	0.712 958	0.715 228	0.724 39	0.682 06	0.776 056	0.750 584	0.776 667	0.738 251	0.779 245	0.717 544	0.730 857	0.743 678	0.738 994	0.633 333
0.778 505	0.782 581	0.736 868	0.707 65	0.730 769	0.758 065	0.767 042	0.749 746	0.778 591	0.732 558	0.757 277	0.756 809	0.76 0.76	0.633 333	0.711 321	0.740 936	0.744 571	0.729 885	0.713 836	0.633 333
0.726 168	0.701 29	0.718 597	0.722 222	0.707 692	0.605 806	0.735 493	0.719 289	0.728 726	0.687 375	0.727 23	0.756 809	0.756 667	0.698 907	0.771 698	0.726 901	0.744 571	0.665 517	0.744 025	0.633 333
0.757 944	0.781 29	0.777 325	0.774 681	0.773 846	0.711 613	0.796 338	0.806 599	0.793 767	0.788 372	0.640 845	0.735 019	0.726 667	0.628 962	0.783 019	0.623 977	0.589 143	0.660 92	0.643 396	0.633 333
0.748 598	0.727 097	0.717 292	0.691 621	0.696 923	0.750 323	0.758 028	0.707 107	0.730 894	0.729 9	0.693 427	0.731 907	0.75 0.75	0.742 623	0.760 377	0.703 509	0.721 714	0.757 471	0.728 931	0.633 333
0.642 056	0.657 419	0.638 989	0.612 933	0.606 154	0.569 677	0.613 803	0.629 949	0.615 989	0.589 037	0.584 507	0.585 603	0.6 0.6	0.489 071	0.662 264	0.661 404	0.703 429	0.716 092	0.653 459	0.633 333
0.741 121	0.719 355	0.725 122	0.701 821	0.696 923	0.685 806	0.755 775	0.749 746	0.748 238	0.764 452	0.723 474	0.731 907	0.76 0.76	0.799 454	0.771 698	0.745 614	0.735 429	0.739 08	0.779 245	0.633 333
0.623 364	0.608 387	0.603 752	0.602 732	0.6 0.6	0.556 774	0.625 07	0.609 645	0.611 653	0.607 641	0.595 775	0.594 942	0.56 0.56	0.545 902	0.673 585	0.670 76	0.648 571	0.674 713	0.648 428	0.633 333
0.752 336	0.715 484	0.727 732	0.700 364	0.724 615	0.696 129	0.749 014	0.763 959	0.791 599	0.751 163	0.727 23	0.731 907	0.746 667	0.729 508	0.730 189	0.740 936	0.730 857	0.757 471	0.754 088	0.633 333
0.739 252	0.702 581	0.706 852	0.693 078	0.673 846	0.672 903	0.721 972	0.711 168	0.722 222	0.700 664	0.693 427	0.669 65	0.68 0.68	0.685 792	0.737 736	0.656 725	0.653 143	0.688 506	0.683 648	0.633 333
0.701 869	0.721 935	0.713 377	0.710 565	0.670 769	0.654 839	0.721 972	0.723 35	0.730 894	0.698 007	0.689 671	0.688 327	0.703 333	0.729 508	0.756 604	0.745 614	0.753 714	0.771 264	0.779 245	0.677 778
0.720 561	0.724 516	0.719 902	0.703 279	0.695 385	0.649 677	0.681 408	0.680 711	0.678 862	0.644 85	0.678 404	0.660 311	0.696 667	0.668 306	0.737 736	0.656 725	0.666 857	0.683 908	0.658 491	0.677 778

0.681 308	0.658 71	0.653 344	0.637 705	0.649 231	0.585 161	0.647 606	0.634 01	0.648 509	0.615 615	0.667 136	0.622 957	0.616 667	0.572 131	0.786 792	0.661 404	0.612	0.665 517	0.648 428	0.677 778
0.752 336	0.683 226	0.650 734	0.643 534	0.661 538	0.662 581	0.706 197	0.715 228	0.730 894	0.687 375	0.712 207	0.707 004	0.716 667	0.646 448	0.696 226	0.623 977	0.680 571	0.688 506	0.688 679	0.677 778
0.868 224	0.840 645	0.841 272	0.832 969	0.809 231	0.773 548	0.823 38	0.828 934	0.843 631	0.812 292	0.813 615	0.775 486	0.833 333	0.808 197	0.866 038	0.712 865	0.772	0.854 023	0.769 182	0.677 778
0.705 607	0.663 871	0.674 225	0.662 477	0.641 538	0.636 774	0.688 169	0.698 985	0.689 702	0.652 824	0.700 939	0.703 891	0.68	0.611 475	0.681 132	0.619 298	0.598 286	0.619 54	0.633 333	0.677 778
0.767 29	0.758 065	0.742 088	0.720 765	0.706 154	0.724 516	0.791 831	0.772 081	0.750 407	0.735 216	0.746 009	0.778 599	0.756 667	0.703 279	0.824 528	0.745 614	0.804	0.798 851	0.738 994	0.677 778
0.757 944	0.770 968	0.777 325	0.771 767	0.730 769	0.721 935	0.758 028	0.721 32	0.776 423	0.761 794	0.791 08	0.738 132	0.77	0.768 852	0.798 113	0.745 614	0.74	0.757 471	0.769 182	0.677 778
0.785 981	0.763 226	0.736 868	0.720 765	0.718 462	0.768 387	0.767 042	0.753 807	0.776 423	0.743 189	0.783 568	0.775 486	0.763 333	0.642 077	0.715 094	0.726 901	0.653 143	0.725 287	0.718 868	0.677 778
0.714 953	0.776 129	0.766 884	0.755 738	0.746 154	0.667 742	0.721 972	0.737 563	0.752 575	0.716 611	0.607 042	0.660 311	0.71	0.593 989	0.673 585	0.703 509	0.804	0.785 057	0.774 214	0.677 778
0.815 888	0.804 516	0.795 595	0.758 652	0.76	0.745 161	0.789 577	0.776 142	0.785 095	0.761 794	0.753 521	0.787 938	0.78	0.698 907	0.832 075	0.740 936	0.744 571	0.752 874	0.769 182	0.677 778
0.670 093	0.634 194	0.646 819	0.644 991	0.661 538	0.649 677	0.681 408	0.701 015	0.709 214	0.692 691	0.667 136	0.678 988	0.706 667	0.703 279	0.760 377	0.694 152	0.689 714	0.702 299	0.713 836	0.677 778
0.795 327	0.778 71	0.755 139	0.748 452	0.76	0.74	0.758 028	0.751 777	0.756 911	0.756 478	0.753 521	0.741 245	0.786 667	0.712 022	0.741 509	0.731 579	0.772	0.762 069	0.799 371	0.677 778
0.759 813	0.770 968	0.742 088	0.741 166	0.756 923	0.768 387	0.803 099	0.804 569	0.798 103	0.764 452	0.734 742	0.756 809	0.74	0.755 738	0.779 245	0.740 936	0.758 286	0.798 851	0.764 151	0.677 778
0.716 822	0.662 581	0.675 53	0.672 678	0.690 769	0.641 935	0.683 662	0.674 619	0.691 87	0.666 113	0.674 648	0.675 875	0.686 667	0.650 82	0.752 83	0.731 579	0.698 857	0.729 885	0.723 899	0.677 778
0.793 458	0.770 968	0.756 444	0.732 423	0.729 231	0.74	0.794 085	0.786 294	0.772 087	0.756 478	0.697 183	0.763 035	0.783 333	0.712 022	0.828 302	0.722 222	0.808 571	0.803 448	0.769 182	0.722 222
0.780 374	0.764 516	0.756 444	0.744 08	0.746 154	0.760 645	0.787 324	0.792 386	0.806 775	0.772 425	0.749 765	0.719 455	0.763 333	0.760 109	0.828 302	0.726 901	0.804	0.831 034	0.733 962	0.722 222
0.855 14	0.854 839	0.824 307	0.818 397	0.809 231	0.845 806	0.870 704	0.859 391	0.880 488	0.812 292	0.9	0.9	0.813 333	0.773 224	0.828 302	0.839 181	0.831 429	0.867 816	0.804 403	0.722 222
0.733 645	0.760 645	0.752 529	0.752 823	0.718 462	0.698 71	0.771 549	0.757 868	0.769 919	0.748 505	0.764 789	0.741 245	0.75	0.768 852	0.779 245	0.778 363	0.790 286	0.817 241	0.829 56	0.722 222

0.729 907	0.752 903	0.747 308	0.738 251	0.724 615	0.737 419	0.776 056	0.745 685	0.754 743	0.745 847	0.772 3	0.707 004	0.706 667	0.725 137	0.756 604	0.689 474	0.749 143	0.725 287	0.718 868	0.722 222
0.815 888	0.817 419	0.823 002	0.818 397	0.804 615	0.760 645	0.805 352	0.822 843	0.828 455	0.780 399	0.832 394	0.794 163	0.813 333	0.781 967	0.775 472	0.783 041	0.790 286	0.794 253	0.814 465	0.722 222
0.337 383	0.332 258	0.333 605	0.338 98	0.343 077	0.301 29	0.316 338	0.319 289	0.349 322	0.309 967	0.347 887	0.330 35	0.326 667	0.336 066	0.326 415	0.310 526	0.310 286	0.311 494	0.316 352	0.722 222
0.836 449	0.817 419	0.828 222	0.831 512	0.816 923	0.752 903	0.839 155	0.837 056	0.854 472	0.820 266	0.832 394	0.859 533	0.853 333	0.720 765	0.847 17	0.801 754	0.826 857	0.812 644	0.844 654	0.722 222
0.728 037	0.773 548	0.762 969	0.752 823	0.74 074	0.683 226	0.721 972	0.745 685	0.754 743	0.727 243	0.584 507	0.660 311	0.696 667	0.593 989	0.760 377	0.708 187	0.790 286	0.789 655	0.799 371	0.722 222
0.802 804	0.736 129	0.697 716	0.701 821	0.704 615	0.709 032	0.751 268	0.753 807	0.763 415	0.729 9	0.753 521	0.747 471	0.753 333	0.681 421	0.707 547	0.675 439	0.717 143	0.725 287	0.698 742	0.722 222
0.834 579	0.787 742	0.774 715	0.781 967	0.786 154	0.789 032	0.830 141	0.822 843	0.837 127	0.841 528	0.794 836	0.781 712	0.84 084	0.790 71	0.756 604	0.689 474	0.694 286	0.752 874	0.713 836	0.722 222
0.9 355	0.879 355	0.9 257	0.891 257	0.9 09	0.879 355	0.868 451	0.849 239	0.880 488	0.876 08	0.824 883	0.822 179	0.85 085	0.799 454	0.816 981	0.778 363	0.790 286	0.785 057	0.809 434	0.722 222
0.827 103	0.764 516	0.755 139	0.755 738	0.769 231	0.801 935	0.805 352	0.798 477	0.813 279	0.775 083	0.817 371	0.859 533	0.816 667	0.786 339	0.850 943	0.769 006	0.790 286	0.794 253	0.759 119	0.722 222
0.832 71	0.790 323	0.778 63	0.745 537	0.744 615	0.758 065	0.791 831	0.790 355	0.808 943	0.783 056	0.779 812	0.784 825	0.763 333	0.746 995	0.707 547	0.703 509	0.758 286	0.743 678	0.804 403	0.722 222
0.800 935	0.777 419	0.760 359	0.730 965	0.727 692	0.750 323	0.818 873	0.782 234	0.776 423	0.756 478	0.719 718	0.744 358	0.756 667	0.712 022	0.820 755	0.750 292	0.822 286	0.831 034	0.723 899	0.766 667
0.802 804	0.804 516	0.813 866	0.811 111	0.795 385	0.791 613	0.890 986	0.891 878	0.9 09	0.9 09	0.843 662	0.809 728	0.853 333	0.790 71	0.839 623	0.801 754	0.817 714	0.789 655	0.864 78	0.766 667
0.838 318	0.733 548	0.726 427	0.741 166	0.716 923	0.729 677	0.760 282	0.737 563	0.761 247	0.743 189	0.787 324	0.772 374	0.76 076	0.738 251	0.677 358	0.609 942	0.689 714	0.757 471	0.643 396	0.766 667
0.808 411	0.734 839	0.704 241	0.698 907	0.712 308	0.719 355	0.755 775	0.772 081	0.767 751	0.737 874	0.761 033	0.753 696	0.76 076	0.681 421	0.733 962	0.666 082	0.735 429	0.716 092	0.708 805	0.766 667
0.791 589	0.789 032	0.773 409	0.755 738	0.784 615	0.742 581	0.839 155	0.820 812	0.863 144	0.844 186	0.877 465	0.853 307	0.876 667	0.9 09	0.752 83	0.759 649	0.790 286	0.794 253	0.789 308	0.766 667
0.838 318	0.749 032	0.730 343	0.748 452	0.723 077	0.750 323	0.771 549	0.743 655	0.774 255	0.761 794	0.794 836	0.775 486	0.77 077	0.733 88	0.666 038	0.595 906	0.694 286	0.752 874	0.663 522	0.766 667
0.720 561	0.751 613	0.748 613	0.742 623	0.736 923	0.745 161	0.773 803	0.753 807	0.763 415	0.756 478	0.791 08	0.713 23	0.726 667	0.738 251	0.760 377	0.722 222	0.753 714	0.757 471	0.738 994	0.766 667

0.853 271	0.893 548	0.884 339	0.895 628	0.88	0.721 935	0.848 169	0.863 452	0.841 463	0.780 399	0.715 962	0.772 374	0.75	0.580 874	0.9	0.867 251	0.858 857	0.872 414	0.854 717	0.766 667
0.821 495	0.82 155	0.785 613	0.806 74	0.78	0.750 323	0.812 113	0.818 782	0.793 767	0.775 083	0.791 08	0.738 132	0.746 667	0.712 022	0.866 038	0.815 789	0.849 714	0.821 839	0.849 686	0.766 667
0.763 551	0.752 903	0.748 613	0.742 623	0.736 923	0.74	0.798 592	0.814 721	0.830 623	0.785 714	0.791 08	0.822 179	0.866 667	0.694 536	0.850 943	0.829 825	0.872 571	0.886 207	0.879 874	0.766 667
0.870 093	0.839 355	0.813 866	0.818 397	0.832 308	0.807 097	0.836 901	0.810 66	0.830 623	0.820 266	0.877 465	0.843 969	0.876 667	0.816 94	0.775 472	0.764 327	0.804	0.817 241	0.744 025	0.811 111
0.817 757	0.839 355	0.873 899	0.863 57	0.801 538	0.781 29	0.841 408	0.853 299	0.865 312	0.809 635	0.806 103	0.819 066	0.87	0.786 339	0.733 962	0.792 398	0.822 286	0.858 621	0.854 717	0.811 111
0.763 551	0.769 677	0.757 749	0.751 366	0.736 923	0.698 71	0.726 479	0.790 355	0.806 775	0.764 452	0.746 009	0.710 117	0.696 667	0.707 65	0.771 698	0.684 795	0.685 143	0.679 31	0.749 057	0.811 111
0.847 664	0.826 452	0.821 697	0.795 082	0.809 231	0.760 645	0.818 873	0.832 995	0.832 791	0.801 661	0.794 836	0.812 84	0.833 333	0.821 311	0.877 358	0.825 146	0.872 571	0.881 609	0.889 937	0.811 111
0.750 467	0.738 71	0.729 038	0.717 851	0.744 615	0.693 548	0.728 732	0.717 259	0.739 566	0.708 638	0.761 033	0.728 794	0.726 667	0.668 306	0.828 302	0.731 579	0.698 857	0.734 483	0.754 088	0.811 111
0.810 28	0.804 516	0.812 561	0.821 311	0.790 769	0.804 516	0.9	0.9	0.9	0.9	0.836 15	0.803 502	0.85	0.799 454	0.839 623	0.801 754	0.817 714	0.817 241	0.889 937	0.811 111
0.830 841	0.813 548	0.833 442	0.831 512	0.815 385	0.755 484	0.848 169	0.851 269	0.858 808	0.825 581	0.869 953	0.875 097	0.873 333	0.746 995	0.749 057	0.801 754	0.817 714	0.817 241	0.839 623	0.811 111
0.868 224	0.9	0.885 644	0.9	0.887 692	0.752 903	0.821 127	0.861 421	0.847 967	0.775 083	0.768 545	0.775 486	0.74	0.567 76	0.866 038	0.9	0.890 857	0.886 207	0.9	0.811 111
0.720 561	0.698 71	0.710 767	0.690 164	0.669 231	0.647 097	0.708 451	0.717 259	0.704 878	0.674 086	0.712 207	0.722 568	0.7	0.633 333	0.677 358	0.675 439	0.630 286	0.642 529	0.648 428	0.811 111
0.823 364	0.863 871	0.868 679	0.857 741	0.830 769	0.82	0.852 676	0.865 482	0.869 648	0.862 791	0.862 441	0.850 195	0.85	0.847 541	0.884 906	0.853 216	0.826 857	0.858 621	0.804 403	0.811 111
0.799 065	0.759 355	0.777 325	0.779 053	0.746 154	0.742 581	0.771 549	0.774 112	0.793 767	0.740 532	0.768 545	0.772 374	0.76	0.672 678	0.779 245	0.703 509	0.772	0.766 667	0.764 151	0.855 556
0.890 654	0.827 742	0.843 883	0.844 627	0.798 462	0.786 452	0.870 704	0.867 513	0.880 488	0.825 581	0.851 174	0.828 405	0.843 333	0.830 055	0.873 585	0.731 579	0.831 429	0.854 023	0.834 591	0.855 556
0.812 15	0.747 742	0.753 834	0.752 823	0.750 769	0.745 161	0.785 07	0.802 538	0.798 103	0.753 821	0.749 765	0.744 358	0.753 333	0.768 852	0.775 472	0.778 363	0.794 857	0.794 253	0.804 403	0.855 556
0.819 626	0.843 226	0.851 713	0.846 084	0.826 154	0.763 226	0.821 127	0.841 117	0.858 808	0.814 95	0.843 662	0.847 082	0.863 333	0.812 568	0.881 132	0.857 895	0.9	0.9	0.894 969	0.855 556

0.838 318	0.796 774	0.813 866	0.802 368	0.792 308	0.765 806	0.823 38	0.808 629	0.806 775	0.793 688	0.806 103	0.784 825	0.833 333	0.803 825	0.847 17	0.839 181	0.826 857	0.9 465	0.814 556	0.855
0.832 71	0.800 645	0.799 511	0.792 168	0.807 692	0.74 577	0.789 569	0.804 615	0.817 372	0.788 568	0.783 486	0.775 333	0.773 88	0.733 113	0.798 901	0.726 429	0.735 429	0.743 678	0.759 119	0.855
0.821 495	0.773 548	0.791 68	0.776 138	0.758 462	0.729 677	0.746 761	0.776 142	0.776 423	0.761 794	0.700 939	0.735 019	0.766 667	0.773 224	0.726 415	0.670 76	0.653 143	0.706 897	0.683 648	0.855
0.812 15	0.790 323	0.772 104	0.763 024	0.750 769	0.690 968	0.744 507	0.751 777	0.752 575	0.713 953	0.723 474	0.719 455	0.743 333	0.655 191	0.749 057	0.773 684	0.799 429	0.812 644	0.789 308	0.855
0.799 065	0.763 226	0.772 104	0.763 024	0.755 385	0.701 29	0.708 451	0.731 472	0.754 743	0.724 585	0.697 183	0.694 553	0.703 333	0.685 792	0.733 962	0.769 006	0.804 253	0.794 057	0.749 556	0.855
0.821 495	0.834 194	0.791 68	0.822 769	0.792 308	0.760 645	0.803 099	0.822 843	0.789 431	0.775 083	0.783 568	0.763 035	0.76 137	0.725 491	0.858 076	0.797 857	0.826 241	0.817 591	0.834 556	0.855
0.838 318	0.822 581	0.807 341	0.795 082	0.789 231	0.745 161	0.791 831	0.782 234	0.774 255	0.775 083	0.798 592	0.778 599	0.793 333	0.764 481	0.764 151	0.726 901	0.726 286	0.729 885	0.723 899	0.9
0.873 832	0.832 903	0.803 426	0.808 197	0.810 769	0.9 944	0.863 944	0.867 513	0.871 816	0.846 844	0.824 883	0.884 436	0.9 0.9	0.821 311	0.809 434	0.811 111	0.813 143	0.854 023	0.834 591	0.9
0.767 29	0.755 484	0.747 308	0.751 366	0.735 385	0.690 968	0.737 746	0.763 959	0.748 238	0.737 874	0.738 498	0.769 261	0.743 333	0.681 421	0.828 302	0.820 468	0.836 0.836	0.858 621	0.809 434	0.9
0.793 458	0.792 903	0.757 749	0.754 281	0.781 538	0.732 258	0.767 042	0.794 416	0.817 615	0.788 372	0.742 254	0.725 681	0.793 333	0.834 426	0.752 83	0.759 649	0.772 0.772	0.803 448	0.824 528	0.9
0.866 355	0.834 194	0.809 951	0.821 311	0.832 308	0.822 581	0.839 155	0.822 843	0.843 631	0.828 239	0.877 465	0.828 405	0.87 0.87	0.816 94	0.779 245	0.783 041	0.767 429	0.798 851	0.749 057	0.9
0.804 673	0.791 613	0.776 02	0.771 767	0.743 077	0.742 581	0.751 268	0.741 624	0.743 902	0.737 874	0.719 718	0.753 696	0.723 333	0.716 393	0.741 509	0.731 579	0.698 857	0.729 885	0.754 088	0.9
0.800 935	0.778 71	0.774 715	0.748 452	0.716 923	0.742 581	0.753 521	0.753 807	0.774 255	0.772 425	0.746 009	0.707 004	0.746 667	0.742 623	0.813 208	0.787 719	0.781 143	0.766 667	0.819 497	0.9
0.800 935	0.808 387	0.783 85	0.811 111	0.784 615	0.724 516	0.807 606	0.796 447	0.832 791	0.783 056	0.798 592	0.769 261	0.796 667	0.720 765	0.794 34	0.843 86	0.863 429	0.890 805	0.824 528	0.9
0.845 794	0.885 806	0.837 357	0.815 483	0.850 769	0.758 065	0.818 873	0.843 147	0.826 287	0.793 688	0.832 394	0.859 533	0.826 667	0.720 765	0.828 302	0.778 363	0.790 286	0.826 437	0.819 497	0.9
0.821 495	0.830 323	0.795 595	0.796 539	0.787 692	0.804 516	0.861 69	0.865 482	0.837 127	0.830 897	0.892 488	0.843 969	0.833 333	0.803 825	0.828 302	0.825 146	0.826 857	0.817 241	0.849 686	0.9

APPENDIX I

**GENERATED CSV FILE OF DATASET OF LATEST POPULATION FOR FEMALE HISPANIC POPULATION AFTER
FEATURE SELECTION USING GA APPROACH**

[illegible]

0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	1	0	1	1
0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	1	0	1	1
0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	1	0	1	1
0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	1	0	1	1
0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	1	0	1	1
0	0	0	1	1	1	0	0	0	0	1	0	0	0	1	1	0	1	1

APPENDIX J

GENERATED CSV FILE OF DATASET OF LATEST POPULATION FOR MALE HISPANIC POPULATION AFTER FEATURE SELECTION USING GA APPROACH

[illegible]

1	0	0	0	0	0	1	0	0	0	0	0	1	1	1	1	0	0	1
1	0	0	0	0	0	1	0	0	0	0	0	1	1	1	1	0	0	1
1	0	0	0	0	0	1	0	0	0	0	0	1	1	1	1	0	0	1
1	0	0	0	0	0	1	0	0	0	0	0	1	1	1	1	0	0	1
1	0	0	0	0	0	1	0	0	0	0	0	1	1	1	1	0	0	1
1	0	0	0	0	0	1	0	0	0	0	0	1	1	1	1	0	0	1

APPENDIX K

NORMALIZED TRAINING DATASET OF SELECTED INPUT FEATURES AND SELECTED TARGET FEATURES FOR FEMALE HISPANIC POPULATION USING GA-ANN APPROACH

4	5	6	11	15	16	18	19	Age
0.121719	0.140482	0.113974	0.11326	0.1	0.1	0.158394	0.148855	0.1
0.214027	0.198313	0.173362	0.228177	0.239759	0.173282	0.216788	0.148855	0.144444
0.125339	0.123133	0.106987	0.139779	0.124096	0.185496	0.193431	0.148855	0.144444
0.210407	0.188675	0.1	0.1	0.153012	0.112214	0.164234	0.136641	0.144444
0.259276	0.229157	0.169869	0.263536	0.181928	0.240458	0.275182	0.228244	0.188889
0.235747	0.238795	0.180349	0.276796	0.201205	0.191603	0.234307	0.185496	0.188889
0.313575	0.304337	0.278166	0.329834	0.287952	0.173282	0.187591	0.191603	0.233333
0.371493	0.379518	0.260699	0.303315	0.355422	0.270992	0.275182	0.289313	0.233333
0.360633	0.375663	0.323581	0.329834	0.437349	0.39313	0.432847	0.277099	0.233333
0.375113	0.366024	0.21179	0.356354	0.350602	0.19771	0.257664	0.203817	0.233333
0.376923	0.406506	0.274672	0.298895	0.321687	0.154962	0.234307	0.234351	0.233333
0.407692	0.383373	0.299127	0.343094	0.360241	0.258779	0.391971	0.222137	0.277778
0.387783	0.404578	0.288646	0.294475	0.422892	0.350382	0.374453	0.319847	0.277778
0.322624	0.358313	0.222271	0.329834	0.316867	0.277099	0.362774	0.234351	0.277778
0.366063	0.36988	0.278166	0.382873	0.36988	0.41145	0.444526	0.29542	0.277778
0.398643	0.396867	0.299127	0.343094	0.36988	0.234351	0.281022	0.246565	0.277778
0.451131	0.452771	0.306114	0.329834	0.326506	0.380916	0.421168	0.387023	0.322222
0.463801	0.479759	0.435371	0.360773	0.461446	0.31374	0.356934	0.319847	0.322222
0.452941	0.475904	0.29214	0.276796	0.254217	0.289313	0.316058	0.29542	0.322222
0.478281	0.47012	0.29214	0.404972	0.30241	0.332061	0.345255	0.29542	0.322222
0.375113	0.366024	0.39345	0.404972	0.43253	0.307634	0.292701	0.289313	0.322222
0.404072	0.39494	0.194323	0.365193	0.422892	0.283206	0.292701	0.264885	0.322222
0.434842	0.402651	0.267686	0.378453	0.408434	0.332061	0.333577	0.270992	0.322222
0.398643	0.398795	0.257205	0.307735	0.321687	0.270992	0.386131	0.29542	0.322222
0.398643	0.410361	0.264192	0.374033	0.360241	0.277099	0.356934	0.222137	0.322222
0.51448	0.510602	0.39345	0.48895	0.490361	0.325954	0.485401	0.240458	0.366667

0.460181	0.472048	0.372489	0.404972	0.442169	0.325954	0.380292	0.325954	0.366667
0.460181	0.473976	0.379476	0.440331	0.312048	0.362595	0.485401	0.448092	0.366667
0.452941	0.456627	0.358515	0.404972	0.485542	0.344275	0.386131	0.344275	0.366667
0.48914	0.491325	0.330568	0.440331	0.490361	0.368702	0.40365	0.356489	0.366667
0.452941	0.481687	0.355022	0.431492	0.446988	0.39313	0.415328	0.454198	0.366667
0.561538	0.545301	0.372489	0.475691	0.350602	0.350382	0.508759	0.332061	0.411111
0.612217	0.593494	0.487773	0.577348	0.581928	0.423664	0.462044	0.332061	0.411111
0.49819	0.522169	0.571616	0.612707	0.610843	0.619084	0.643066	0.600763	0.411111
0.548869	0.562651	0.484279	0.541989	0.519277	0.350382	0.368613	0.332061	0.411111
0.50724	0.527952	0.351528	0.533149	0.533735	0.551908	0.514599	0.338168	0.411111
0.603167	0.574217	0.529694	0.546409	0.577108	0.527481	0.427007	0.417557	0.455556
0.566968	0.574217	0.449345	0.581768	0.610843	0.612977	0.654745	0.60687	0.455556
0.543439	0.524096	0.501747	0.537569	0.586747	0.460305	0.60219	0.564122	0.455556
0.557919	0.533735	0.515721	0.590608	0.572289	0.515267	0.643066	0.441985	0.455556
0.583258	0.597349	0.578603	0.608287	0.581928	0.484733	0.619708	0.472519	0.455556
0.606787	0.595422	0.501747	0.572928	0.581928	0.429771	0.450365	0.399237	0.455556
0.635747	0.632048	0.536681	0.533149	0.591566	0.448092	0.613869	0.533588	0.455556
0.585068	0.583855	0.561135	0.586188	0.586747	0.448092	0.643066	0.478626	0.5
0.677376	0.687952	0.603057	0.670166	0.63494	0.631298	0.724818	0.661832	0.5
0.650226	0.624337	0.536681	0.564088	0.644578	0.661832	0.50292	0.631298	0.5
0.49638	0.508675	0.452838	0.475691	0.475904	0.521374	0.485401	0.496947	0.5
0.657466	0.651325	0.421397	0.51547	0.683133	0.655725	0.584672	0.667939	0.5
0.648416	0.662892	0.58559	0.692265	0.692771	0.710687	0.648905	0.643511	0.544444
0.617647	0.62241	0.58559	0.603867	0.668675	0.576336	0.648905	0.594656	0.544444
0.644796	0.653253	0.547162	0.559669	0.591566	0.521374	0.567153	0.551908	0.544444
0.646606	0.649398	0.693886	0.705525	0.736145	0.692366	0.713139	0.667939	0.544444
0.659276	0.686024	0.620524	0.603867	0.620482	0.674046	0.678102	0.643511	0.544444
0.594118	0.593494	0.491266	0.577348	0.615663	0.454198	0.643066	0.496947	0.544444
0.556109	0.53759	0.466812	0.48011	0.586747	0.441985	0.613869	0.435878	0.544444
0.671946	0.69759	0.620524	0.652486	0.760241	0.655725	0.713139	0.735115	0.544444
0.617647	0.62241	0.624017	0.705525	0.731325	0.649618	0.660584	0.619084	0.544444
0.619457	0.60506	0.58559	0.546409	0.639759	0.612977	0.672263	0.637405	0.544444
0.655656	0.655181	0.589083	0.572928	0.649398	0.655725	0.60219	0.576336	0.544444
0.652036	0.659036	0.617031	0.617127	0.553012	0.643511	0.69562	0.625191	0.588889

0.623077	0.64747	0.610044	0.612707	0.692771	0.692366	0.765693	0.643511	0.588889
0.626697	0.682169	0.711354	0.634807	0.639759	0.680153	0.672263	0.570229	0.588889
0.715385	0.736145	0.721834	0.683425	0.750602	0.70458	0.724818	0.759542	0.588889
0.677376	0.709157	0.631004	0.639227	0.610843	0.692366	0.707299	0.655725	0.588889
0.724434	0.753494	0.739301	0.732044	0.707229	0.783969	0.748175	0.747328	0.588889
0.695475	0.705301	0.610044	0.705525	0.736145	0.722901	0.754015	0.698473	0.588889
0.648416	0.643614	0.631004	0.625967	0.610843	0.625191	0.660584	0.637405	0.588889
0.668326	0.695663	0.683406	0.643646	0.625301	0.661832	0.736496	0.649618	0.633333
0.657466	0.655181	0.725328	0.656906	0.615663	0.649618	0.578832	0.643511	0.633333
0.778733	0.770843	0.763755	0.785083	0.798795	0.771756	0.829927	0.582443	0.633333
0.688235	0.666747	0.679913	0.705525	0.683133	0.667939	0.771533	0.68626	0.633333
0.604977	0.657108	0.634498	0.612707	0.731325	0.680153	0.730657	0.643511	0.633333
0.735294	0.724578	0.728821	0.679006	0.750602	0.759542	0.783212	0.759542	0.633333
0.666516	0.695663	0.718341	0.687845	0.740964	0.722901	0.713139	0.625191	0.633333
0.771493	0.738072	0.760262	0.709945	0.692771	0.722901	0.713139	0.741221	0.633333
0.704525	0.666747	0.683406	0.608287	0.586747	0.649618	0.660584	0.68626	0.633333
0.749774	0.751566	0.70786	0.692265	0.644578	0.680153	0.765693	0.649618	0.633333
0.9	0.9	0.9	0.9	0.871084	0.790076	0.870803	0.710687	0.633333
0.720814	0.686024	0.69738	0.621547	0.750602	0.692366	0.70146	0.667939	0.633333
0.680995	0.713012	0.704367	0.679006	0.745783	0.692366	0.666423	0.729008	0.633333
0.700905	0.734217	0.756769	0.798343	0.755422	0.790076	0.80073	0.783969	0.677778
0.666516	0.684096	0.714847	0.701105	0.712048	0.832824	0.806569	0.777863	0.677778
0.626697	0.653253	0.672926	0.643646	0.707229	0.722901	0.765693	0.667939	0.677778
0.702715	0.759277	0.735808	0.740884	0.726506	0.777863	0.771533	0.722901	0.677778
0.731674	0.722651	0.690393	0.665746	0.755422	0.790076	0.789051	0.722901	0.677778
0.760633	0.763133	0.746288	0.696685	0.683133	0.70458	0.759854	0.649618	0.677778
0.713575	0.678313	0.665939	0.727624	0.716867	0.698473	0.69562	0.649618	0.677778
0.633937	0.657108	0.651965	0.639227	0.625301	0.70458	0.70146	0.667939	0.677778
0.738914	0.732289	0.718341	0.709945	0.726506	0.747328	0.742336	0.729008	0.677778
0.684615	0.713012	0.742795	0.705525	0.750602	0.851145	0.847445	0.747328	0.677778
0.693665	0.691807	0.721834	0.617127	0.56747	0.625191	0.625547	0.594656	0.722222
0.796833	0.778554	0.704367	0.705525	0.687952	0.729008	0.718978	0.735115	0.722222
0.719005	0.709157	0.704367	0.656906	0.408434	0.729008	0.777372	0.729008	0.722222
0.782353	0.768916	0.784716	0.727624	0.678313	0.70458	0.713139	0.716794	0.722222

0.704525	0.707229	0.669432	0.674586	0.707229	0.674046	0.742336	0.680153	0.722222
0.680995	0.662892	0.658952	0.590608	0.591566	0.637405	0.648905	0.643511	0.722222
0.771493	0.776627	0.767249	0.776243	0.726506	0.820611	0.841606	0.790076	0.722222
0.749774	0.766988	0.746288	0.740884	0.740964	0.777863	0.783212	0.729008	0.722222
0.726244	0.774699	0.854585	0.864641	0.63012	0.747328	0.789051	0.674046	0.722222
0.787783	0.80747	0.721834	0.683425	0.779518	0.741221	0.765693	0.759542	0.722222
0.742534	0.751566	0.69738	0.723204	0.70241	0.70458	0.713139	0.729008	0.722222
0.811312	0.83253	0.819651	0.776243	0.813253	0.820611	0.894161	0.814504	0.722222
0.787783	0.780482	0.770742	0.771823	0.731325	0.875573	0.829927	0.80229	0.766667
0.809502	0.813253	0.739301	0.727624	0.716867	0.783969	0.789051	0.765649	0.766667
0.708145	0.757349	0.676419	0.665746	0.615663	0.759542	0.736496	0.68626	0.766667
0.796833	0.788193	0.704367	0.679006	0.673494	0.765649	0.754015	0.741221	0.766667
0.576018	0.603133	0.672926	0.643646	0.581928	0.680153	0.683942	0.655725	0.766667
0.795023	0.80747	0.763755	0.789503	0.707229	0.826718	0.853285	0.759542	0.766667
0.690045	0.69759	0.760262	0.674586	0.683133	0.753435	0.754015	0.741221	0.766667
0.655656	0.666747	0.711354	0.683425	0.721687	0.753435	0.683942	0.716794	0.811111
0.831222	0.840241	0.819651	0.780663	0.808434	0.826718	0.806569	0.790076	0.811111
0.737104	0.732289	0.700873	0.732044	0.731325	0.735115	0.777372	0.667939	0.811111
0.816742	0.83253	0.774236	0.661326	0.726506	0.790076	0.841606	0.747328	0.811111
0.807692	0.826747	0.739301	0.727624	0.736145	0.790076	0.824088	0.771756	0.811111
0.738914	0.741928	0.624017	0.603867	0.649398	0.667939	0.736496	0.722901	0.811111
0.720814	0.741928	0.711354	0.670166	0.721687	0.814504	0.771533	0.796183	0.811111
0.719005	0.699518	0.721834	0.705525	0.707229	0.838931	0.829927	0.832824	0.811111
0.773303	0.772771	0.760262	0.789503	0.731325	0.875573	0.864964	0.771756	0.811111
0.626697	0.687952	0.693886	0.648066	0.654217	0.680153	0.765693	0.759542	0.855556
0.766063	0.757349	0.6869	0.679006	0.625301	0.722901	0.742336	0.692366	0.855556
0.737104	0.722651	0.714847	0.732044	0.644578	0.70458	0.69562	0.692366	0.855556
0.793213	0.801687	0.791703	0.789503	0.827711	0.765649	0.876642	0.783969	0.855556
0.652036	0.643614	0.662445	0.581768	0.63494	0.70458	0.713139	0.68626	0.855556
0.769683	0.788193	0.861572	0.767403	0.9	0.851145	0.888321	0.838931	0.855556
0.702715	0.686024	0.690393	0.679006	0.687952	0.716794	0.707299	0.680153	0.9
0.688235	0.659036	0.599563	0.625967	0.659036	0.643511	0.660584	0.631298	0.9
0.742534	0.734217	0.714847	0.714365	0.789157	0.783969	0.777372	0.771756	0.9
0.664706	0.709157	0.739301	0.679006	0.606024	0.741221	0.730657	0.625191	0.9

APPENDIX L

NORMALIZED TEST DATASET OF SELECTED INPUT FEATURES AND SELECTED TARGET FEATURES FOR FEMALE HISPANIC POPULATION USING GA-ANN APPROACH

4	5	6	11	15	16	18	19	Age
0.1	0.1	0.1	0.130939	0.124096	0.179389	0.158394	0.148855	0.144444
0.212217	0.211807	0.141921	0.126519	0.220482	0.148855	0.19927	0.1	0.144444
0.291855	0.312048	0.222271	0.325414	0.312048	0.344275	0.345255	0.240458	0.188889
0.232127	0.225301	0.145415	0.192818	0.148193	0.19771	0.19927	0.148855	0.188889
0.188688	0.194458	0.131441	0.148619	0.215663	0.142748	0.1	0.142748	0.188889
0.465611	0.46241	0.344541	0.325414	0.5	0.325954	0.327737	0.338168	0.322222
0.454751	0.44506	0.337555	0.466851	0.451807	0.448092	0.444526	0.319847	0.366667
0.451131	0.472048	0.29214	0.400552	0.413253	0.356489	0.485401	0.350382	0.366667
0.409502	0.414217	0.41441	0.334254	0.446988	0.350382	0.386131	0.362595	0.366667
0.456561	0.473976	0.41441	0.400552	0.524096	0.380916	0.467883	0.356489	0.366667
0.547059	0.543373	0.389956	0.546409	0.533735	0.362595	0.351095	0.454198	0.411111
0.49276	0.551084	0.491266	0.50221	0.514458	0.399237	0.543796	0.50916	0.411111
0.541629	0.547229	0.49476	0.546409	0.572289	0.496947	0.543796	0.374809	0.411111
0.49457	0.493253	0.466812	0.471271	0.509639	0.423664	0.49708	0.332061	0.411111
0.532579	0.520241	0.466812	0.387293	0.519277	0.429771	0.520438	0.515267	0.411111
0.50181	0.520241	0.466812	0.546409	0.519277	0.374809	0.537956	0.521374	0.455556
0.5	0.522169	0.484279	0.458011	0.533735	0.527481	0.520438	0.545802	0.455556
0.612217	0.626265	0.459825	0.625967	0.615663	0.649618	0.637226	0.619084	0.5
0.668326	0.67253	0.655459	0.639227	0.557831	0.58855	0.467883	0.576336	0.5
0.592308	0.616627	0.547162	0.559669	0.610843	0.710687	0.707299	0.661832	0.5
0.670136	0.691807	0.644978	0.643646	0.668675	0.70458	0.660584	0.625191	0.5
0.592308	0.612771	0.49476	0.537569	0.56747	0.545802	0.561314	0.503053	0.5
0.619457	0.639759	0.582096	0.577348	0.610843	0.60687	0.60219	0.594656	0.544444
0.673756	0.670602	0.644978	0.652486	0.70241	0.759542	0.736496	0.70458	0.544444
0.585068	0.587711	0.543668	0.625967	0.586747	0.600763	0.643066	0.558015	0.544444
0.566968	0.562651	0.543668	0.634807	0.644578	0.619084	0.660584	0.551908	0.588889

0.661086	0.676386	0.669432	0.612707	0.721687	0.625191	0.718978	0.68626	0.588889
0.749774	0.755422	0.770742	0.758564	0.692771	0.649618	0.666423	0.674046	0.588889
0.657466	0.680241	0.672926	0.634807	0.581928	0.68626	0.777372	0.637405	0.588889
0.688235	0.724578	0.651965	0.661326	0.63494	0.70458	0.70146	0.68626	0.588889
0.728054	0.720723	0.60655	0.630387	0.668675	0.649618	0.648905	0.637405	0.588889
0.680995	0.680241	0.728821	0.701105	0.659036	0.60687	0.648905	0.661832	0.588889
0.804072	0.797831	0.777729	0.683425	0.731325	0.753435	0.812409	0.735115	0.633333
0.566968	0.574217	0.60655	0.603867	0.644578	0.674046	0.654745	0.655725	0.633333
0.691855	0.670602	0.704367	0.785083	0.639759	0.70458	0.718978	0.674046	0.677778
0.677376	0.628193	0.655459	0.679006	0.731325	0.783969	0.771533	0.783969	0.677778
0.686425	0.693735	0.704367	0.590608	0.572289	0.619084	0.60219	0.600763	0.677778
0.746154	0.724578	0.774236	0.714365	0.654217	0.70458	0.718978	0.692366	0.677778
0.787783	0.770843	0.732314	0.736464	0.726506	0.771756	0.759854	0.722901	0.677778
0.760633	0.793976	0.774236	0.771823	0.779518	0.820611	0.80073	0.832824	0.722222
0.733484	0.766988	0.833624	0.780663	0.740964	0.808397	0.794891	0.722901	0.722222
0.641176	0.655181	0.651965	0.643646	0.716867	0.710687	0.771533	0.661832	0.766667
0.785973	0.805542	0.78821	0.709945	0.76506	0.820611	0.794891	0.838931	0.766667
0.766063	0.76506	0.714847	0.727624	0.716867	0.710687	0.748175	0.661832	0.766667
0.789593	0.780482	0.70786	0.736464	0.56747	0.60687	0.824088	0.60687	0.811111
0.709955	0.726506	0.679913	0.705525	0.70241	0.753435	0.748175	0.741221	0.855556
0.738914	0.753494	0.795197	0.727624	0.673494	0.759542	0.777372	0.747328	0.855556
0.652036	0.664819	0.683406	0.701105	0.716867	0.759542	0.678102	0.735115	0.855556
0.775113	0.728434	0.69738	0.648066	0.76506	0.741221	0.754015	0.722901	0.855556
0.798643	0.776627	0.704367	0.771823	0.846988	0.881679	0.882482	0.796183	0.9
0.796833	0.826747	0.861572	0.89558	0.880723	0.9	0.9	0.9	0.9
0.726244	0.711084	0.742795	0.692265	0.822892	0.790076	0.853285	0.753435	0.9
0.795023	0.786265	0.868559	0.789503	0.895181	0.863359	0.859124	0.869466	0.9
0.733484	0.766988	0.69738	0.648066	0.639759	0.759542	0.748175	0.710687	0.9
0.666516	0.674458	0.679913	0.701105	0.707229	0.747328	0.689781	0.70458	0.9

APPENDIX M

NORMALIZED TRAINING DATASET OF SELECTED INPUT FEATURES AND SELECTED TARGET FEATURES FOR MALE HISPANIC POPULATION USING GA-ANN APPROACH

1	7	13	14	15	16	19	Age
0.124299	0.131549	0.156667	0.12623	0.141509	0.137427	0.110063	0.1
0.172897	0.199155	0.21	0.200546	0.216981	0.202924	0.190566	0.1
0.193458	0.223944	0.216667	0.200546	0.190566	0.15614	0.120126	0.1
0.320561	0.368169	0.41	0.384153	0.318868	0.277778	0.276101	0.144444
0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.144444
0.23271	0.291549	0.25	0.26612	0.326415	0.235673	0.230818	0.144444
0.223364	0.302817	0.273333	0.18306	0.330189	0.240351	0.200629	0.144444
0.230841	0.284789	0.28	0.257377	0.269811	0.259064	0.195597	0.144444
0.281308	0.347887	0.33	0.287978	0.379245	0.291813	0.281132	0.188889
0.324299	0.370423	0.39	0.331694	0.349057	0.282456	0.261006	0.188889
0.3	0.354648	0.336667	0.253005	0.383019	0.287135	0.276101	0.188889
0.350467	0.354648	0.363333	0.305464	0.367925	0.361988	0.255975	0.233333
0.316822	0.417746	0.43	0.384153	0.39434	0.324561	0.276101	0.233333
0.118692	0.142817	0.146667	0.134973	0.160377	0.132749	0.120126	0.233333
0.371028	0.444789	0.43	0.406011	0.443396	0.404094	0.396855	0.277778
0.329907	0.431268	0.47	0.397268	0.435849	0.446199	0.416981	0.277778
0.356075	0.410986	0.363333	0.327322	0.454717	0.436842	0.422013	0.277778
0.460748	0.512394	0.543333	0.493443	0.541509	0.511696	0.467296	0.322222
0.481308	0.501127	0.513333	0.502186	0.503774	0.44152	0.432075	0.322222
0.427103	0.546197	0.546667	0.56776	0.481132	0.427485	0.406918	0.322222
0.408411	0.401972	0.423333	0.384153	0.428302	0.376023	0.371698	0.322222
0.382243	0.46507	0.423333	0.309836	0.443396	0.380702	0.391824	0.322222
0.434579	0.530423	0.576667	0.537158	0.564151	0.511696	0.512579	0.366667
0.556075	0.559718	0.54	0.515301	0.567925	0.530409	0.477358	0.366667
0.5	0.548451	0.553333	0.54153	0.567925	0.530409	0.502516	0.366667
0.458879	0.50338	0.516667	0.480328	0.477358	0.44152	0.477358	0.366667

0.483178	0.50338	0.456667	0.423497	0.488679	0.436842	0.452201	0.366667
0.423364	0.462817	0.44	0.388525	0.428302	0.394737	0.422013	0.366667
0.492523	0.510141	0.506667	0.462842	0.567925	0.488304	0.477358	0.366667
0.556075	0.568732	0.58	0.528415	0.571698	0.516374	0.467296	0.411111
0.471963	0.50338	0.516667	0.440984	0.518868	0.432164	0.41195	0.411111
0.503738	0.523662	0.54	0.510929	0.549057	0.492982	0.512579	0.411111
0.535514	0.548451	0.566667	0.572131	0.567925	0.567836	0.527673	0.411111
0.537383	0.483099	0.523333	0.524044	0.545283	0.507018	0.492453	0.411111
0.578505	0.521408	0.543333	0.489071	0.583019	0.544444	0.55283	0.411111
0.507477	0.492113	0.463333	0.43224	0.526415	0.460234	0.427044	0.455556
0.514953	0.525915	0.486667	0.410383	0.564151	0.544444	0.532704	0.455556
0.584112	0.586761	0.586667	0.532787	0.598113	0.516374	0.532704	0.455556
0.526168	0.557465	0.55	0.532787	0.541509	0.525731	0.497484	0.455556
0.514953	0.532676	0.573333	0.524044	0.55283	0.516374	0.502516	0.455556
0.576636	0.552958	0.5	0.440984	0.567925	0.553801	0.532704	0.455556
0.571028	0.557465	0.536667	0.56776	0.59434	0.549123	0.512579	0.5
0.701869	0.672394	0.66	0.56776	0.658491	0.638012	0.567925	0.5
0.548598	0.521408	0.493333	0.384153	0.545283	0.502339	0.502516	0.5
0.600935	0.627324	0.696667	0.628962	0.601887	0.567836	0.598113	0.5
0.54486	0.570986	0.526667	0.45847	0.586792	0.502339	0.497484	0.5
0.61215	0.629577	0.61	0.550273	0.711321	0.656725	0.663522	0.5
0.561682	0.566479	0.586667	0.589617	0.616981	0.567836	0.562893	0.5
0.636449	0.620563	0.643333	0.598361	0.635849	0.58655	0.603145	0.544444
0.628972	0.62507	0.623333	0.545902	0.586792	0.507018	0.406918	0.544444
0.705607	0.636338	0.636667	0.642077	0.613208	0.577193	0.593082	0.544444
0.578505	0.58	0.583333	0.506557	0.598113	0.577193	0.567925	0.544444
0.524299	0.568732	0.526667	0.454098	0.59434	0.507018	0.502516	0.544444
0.591589	0.58	0.576667	0.54153	0.590566	0.549123	0.522642	0.544444
0.63271	0.595775	0.616667	0.537158	0.571698	0.58655	0.613208	0.544444
0.602804	0.622817	0.676667	0.642077	0.673585	0.666082	0.633333	0.544444
0.597196	0.573239	0.616667	0.633333	0.616981	0.591228	0.648428	0.544444
0.606542	0.566479	0.58	0.528415	0.635849	0.600585	0.593082	0.588889
0.608411	0.634085	0.68	0.642077	0.703774	0.689474	0.683648	0.588889
0.696262	0.721972	0.633333	0.510929	0.677358	0.605263	0.577987	0.588889

0.664486	0.62507	0.653333	0.668306	0.666038	0.64269	0.598113	0.588889
0.800935	0.780563	0.783333	0.642077	0.760377	0.736257	0.703774	0.588889
0.599065	0.604789	0.61	0.607104	0.624528	0.628655	0.678616	0.588889
0.314953	0.278028	0.253333	0.253005	0.296226	0.268421	0.240881	0.588889
0.328037	0.291549	0.336667	0.301093	0.345283	0.319883	0.316352	0.588889
0.814019	0.726479	0.736667	0.712022	0.764151	0.680117	0.688679	0.588889
0.630841	0.604789	0.593333	0.572131	0.654717	0.61462	0.633333	0.588889
0.56729	0.528169	0.563333	0.493443	0.55283	0.535088	0.502516	0.588889
0.664486	0.58	0.633333	0.620219	0.635849	0.595906	0.638365	0.588889
0.645794	0.649859	0.723333	0.720765	0.726415	0.703509	0.713836	0.633333
0.681308	0.679155	0.736667	0.703279	0.722642	0.689474	0.703774	0.633333
0.784112	0.758028	0.73	0.611475	0.7	0.717544	0.718868	0.633333
0.778505	0.767042	0.76	0.633333	0.711321	0.740936	0.713836	0.633333
0.757944	0.796338	0.726667	0.628962	0.783019	0.623977	0.643396	0.633333
0.748598	0.758028	0.75	0.742623	0.760377	0.703509	0.728931	0.633333
0.642056	0.613803	0.6	0.489071	0.662264	0.661404	0.653459	0.633333
0.741121	0.755775	0.76	0.799454	0.771698	0.745614	0.779245	0.633333
0.752336	0.749014	0.746667	0.729508	0.730189	0.740936	0.754088	0.633333
0.701869	0.721972	0.703333	0.729508	0.756604	0.745614	0.779245	0.677778
0.681308	0.647606	0.616667	0.572131	0.786792	0.661404	0.648428	0.677778
0.752336	0.706197	0.716667	0.646448	0.696226	0.623977	0.688679	0.677778
0.705607	0.688169	0.68	0.611475	0.681132	0.619298	0.633333	0.677778
0.76729	0.791831	0.756667	0.703279	0.824528	0.745614	0.738994	0.677778
0.714953	0.721972	0.71	0.593989	0.673585	0.703509	0.774214	0.677778
0.815888	0.789577	0.78	0.698907	0.832075	0.740936	0.769182	0.677778
0.795327	0.758028	0.786667	0.712022	0.741509	0.731579	0.799371	0.677778
0.793458	0.794085	0.783333	0.712022	0.828302	0.722222	0.769182	0.722222
0.780374	0.787324	0.763333	0.760109	0.828302	0.726901	0.733962	0.722222
0.815888	0.805352	0.813333	0.781967	0.775472	0.783041	0.814465	0.722222
0.337383	0.316338	0.326667	0.336066	0.326415	0.310526	0.316352	0.722222
0.836449	0.839155	0.853333	0.720765	0.84717	0.801754	0.844654	0.722222
0.728037	0.721972	0.696667	0.593989	0.760377	0.708187	0.799371	0.722222
0.802804	0.751268	0.753333	0.681421	0.707547	0.675439	0.698742	0.722222
0.834579	0.830141	0.84	0.79071	0.756604	0.689474	0.713836	0.722222

0.9	0.868451	0.85	0.799454	0.816981	0.778363	0.809434	0.722222
0.827103	0.805352	0.816667	0.786339	0.850943	0.769006	0.759119	0.722222
0.83271	0.791831	0.763333	0.746995	0.707547	0.703509	0.804403	0.722222
0.800935	0.818873	0.756667	0.712022	0.820755	0.750292	0.723899	0.766667
0.802804	0.890986	0.853333	0.79071	0.839623	0.801754	0.86478	0.766667
0.808411	0.755775	0.76	0.681421	0.733962	0.666082	0.708805	0.766667
0.791589	0.839155	0.876667	0.9	0.75283	0.759649	0.789308	0.766667
0.838318	0.771549	0.77	0.73388	0.666038	0.595906	0.663522	0.766667
0.720561	0.773803	0.726667	0.738251	0.760377	0.722222	0.738994	0.766667
0.853271	0.848169	0.75	0.580874	0.9	0.867251	0.854717	0.766667
0.763551	0.798592	0.866667	0.694536	0.850943	0.829825	0.879874	0.766667
0.870093	0.836901	0.876667	0.81694	0.775472	0.764327	0.744025	0.811111
0.817757	0.841408	0.87	0.786339	0.733962	0.792398	0.854717	0.811111
0.763551	0.726479	0.696667	0.70765	0.771698	0.684795	0.749057	0.811111
0.847664	0.818873	0.833333	0.821311	0.877358	0.825146	0.889937	0.811111
0.750467	0.728732	0.726667	0.668306	0.828302	0.731579	0.754088	0.811111
0.830841	0.848169	0.873333	0.746995	0.749057	0.801754	0.839623	0.811111
0.868224	0.821127	0.74	0.56776	0.866038	0.9	0.9	0.811111
0.720561	0.708451	0.7	0.633333	0.677358	0.675439	0.648428	0.811111
0.799065	0.771549	0.76	0.672678	0.779245	0.703509	0.764151	0.855556
0.890654	0.870704	0.843333	0.830055	0.873585	0.731579	0.834591	0.855556
0.838318	0.82338	0.833333	0.803825	0.84717	0.839181	0.814465	0.855556
0.83271	0.789577	0.773333	0.73388	0.798113	0.726901	0.759119	0.855556
0.821495	0.746761	0.766667	0.773224	0.726415	0.67076	0.683648	0.855556
0.821495	0.803099	0.76	0.725137	0.858491	0.797076	0.834591	0.855556
0.838318	0.791831	0.793333	0.764481	0.764151	0.726901	0.723899	0.9
0.873832	0.863944	0.9	0.821311	0.809434	0.811111	0.834591	0.9
0.793458	0.767042	0.793333	0.834426	0.75283	0.759649	0.824528	0.9
0.866355	0.839155	0.87	0.81694	0.779245	0.783041	0.749057	0.9
0.804673	0.751268	0.723333	0.716393	0.741509	0.731579	0.754088	0.9
0.800935	0.807606	0.796667	0.720765	0.79434	0.84386	0.824528	0.9
0.845794	0.818873	0.826667	0.720765	0.828302	0.778363	0.819497	0.9
0.821495	0.86169	0.833333	0.803825	0.828302	0.825146	0.849686	0.9

APPENDIX N

NORMALIZED TEST DATASET OF SELECTED INPUT FEATURES AND SELECTED TARGET FEATURES FOR MALE HISPANIC POPULATION USING GA-ANN APPROACH

1	7	13	14	15	16	19	Age
0.171028	0.187887	0.206667	0.178689	0.209434	0.198246	0.17044	0.1
0.286916	0.350141	0.356667	0.331694	0.349057	0.263743	0.250943	0.188889
0.290654	0.372676	0.41	0.362295	0.364151	0.324561	0.296226	0.188889
0.376636	0.433521	0.413333	0.340437	0.413208	0.324561	0.37673	0.233333
0.309346	0.426761	0.416667	0.314208	0.428302	0.366667	0.351572	0.233333
0.447664	0.494366	0.486667	0.427869	0.503774	0.41345	0.381761	0.277778
0.283178	0.332113	0.32	0.218033	0.35283	0.282456	0.245912	0.277778
0.45514	0.456056	0.453333	0.414754	0.466038	0.436842	0.416981	0.322222
0.393458	0.42	0.466667	0.449727	0.450943	0.352632	0.442138	0.322222
0.43271	0.417746	0.38	0.301093	0.443396	0.422807	0.386792	0.322222
0.464486	0.494366	0.463333	0.419126	0.466038	0.44152	0.396855	0.322222
0.443925	0.42	0.446667	0.366667	0.420755	0.408772	0.386792	0.366667
0.483178	0.469577	0.506667	0.480328	0.496226	0.44152	0.416981	0.366667
0.516822	0.50338	0.503333	0.440984	0.492453	0.427485	0.452201	0.411111
0.5	0.521408	0.593333	0.462842	0.601887	0.535088	0.527673	0.411111
0.528037	0.564225	0.566667	0.554645	0.579245	0.511696	0.51761	0.411111
0.479439	0.471831	0.443333	0.366667	0.503774	0.455556	0.442138	0.411111
0.582243	0.595775	0.586667	0.519672	0.601887	0.553801	0.537736	0.455556
0.576636	0.58	0.59	0.475956	0.643396	0.55848	0.567925	0.455556
0.520561	0.53493	0.56	0.528415	0.616981	0.530409	0.492453	0.455556
0.571028	0.552958	0.53	0.45847	0.598113	0.577193	0.502516	0.455556
0.585981	0.570986	0.573333	0.532787	0.696226	0.55848	0.58805	0.5
0.65514	0.577746	0.516667	0.309836	0.598113	0.535088	0.507547	0.5
0.621495	0.600282	0.606667	0.65082	0.628302	0.567836	0.593082	0.5
0.526168	0.528169	0.51	0.462842	0.616981	0.58655	0.547799	0.544444
0.619626	0.634085	0.583333	0.506557	0.692453	0.591228	0.628302	0.544444

0.627103	0.645352	0.646667	0.628962	0.639623	0.628655	0.62327	0.544444
0.585981	0.570986	0.603333	0.611475	0.658491	0.609942	0.633333	0.588889
0.668224	0.638592	0.643333	0.602732	0.677358	0.661404	0.628302	0.588889
0.836449	0.807606	0.793333	0.764481	0.854717	0.694152	0.749057	0.633333
0.657009	0.665634	0.753333	0.73388	0.737736	0.708187	0.738994	0.633333
0.72243	0.712958	0.776667	0.738251	0.779245	0.717544	0.738994	0.633333
0.726168	0.735493	0.756667	0.698907	0.771698	0.726901	0.744025	0.633333
0.623364	0.62507	0.56	0.545902	0.673585	0.67076	0.648428	0.633333
0.739252	0.721972	0.68	0.685792	0.737736	0.656725	0.683648	0.633333
0.720561	0.681408	0.696667	0.668306	0.737736	0.656725	0.658491	0.677778
0.868224	0.82338	0.833333	0.808197	0.866038	0.712865	0.769182	0.677778
0.757944	0.758028	0.77	0.768852	0.798113	0.745614	0.769182	0.677778
0.785981	0.767042	0.763333	0.642077	0.715094	0.726901	0.718868	0.677778
0.670093	0.681408	0.706667	0.703279	0.760377	0.694152	0.713836	0.677778
0.759813	0.803099	0.74	0.755738	0.779245	0.740936	0.764151	0.677778
0.716822	0.683662	0.686667	0.65082	0.75283	0.731579	0.723899	0.677778
0.85514	0.870704	0.813333	0.773224	0.828302	0.839181	0.804403	0.722222
0.733645	0.771549	0.75	0.768852	0.779245	0.778363	0.82956	0.722222
0.729907	0.776056	0.706667	0.725137	0.756604	0.689474	0.718868	0.722222
0.838318	0.760282	0.76	0.738251	0.677358	0.609942	0.643396	0.766667
0.821495	0.812113	0.746667	0.712022	0.866038	0.815789	0.849686	0.766667
0.81028	0.9	0.85	0.799454	0.839623	0.801754	0.889937	0.811111
0.823364	0.852676	0.85	0.847541	0.884906	0.853216	0.804403	0.811111
0.81215	0.78507	0.753333	0.768852	0.775472	0.778363	0.804403	0.855556
0.819626	0.821127	0.863333	0.812568	0.881132	0.857895	0.894969	0.855556
0.81215	0.744507	0.743333	0.655191	0.749057	0.773684	0.789308	0.855556
0.799065	0.708451	0.703333	0.685792	0.733962	0.769006	0.749057	0.855556
0.76729	0.737746	0.743333	0.681421	0.828302	0.820468	0.809434	0.9
0.800935	0.753521	0.746667	0.742623	0.813208	0.787719	0.819497	0.9

APPENDIX O

NORMALIZED AND DENORMALIZED TRAINING DATASET OF TARGET FEATURE, OUTPUTS AND ERRORS FOR FEMALE HISPANIC POPULATION USING ANN APPROACH AND GA-ANN APPROACH

ANN						GA-ANN					
Normalized			Denormalized			Normalized			Denormalized		
Target	Output	Error	Target	Output	Error	Target	Output	Error	Target	Output	Error
0.1	0.165005	-0.065	0	1.462611	-1.46261	0.1	0.171634	-0.07163	0	1.611771	-1.61177
0.144444	0.190916	-0.04647	1	2.045608	-1.04561	0.144444	0.189427	-0.04498	1	2.012104	-1.0121
0.144444	0.165951	-0.02151	1	1.483898	-0.4839	0.144444	0.179772	-0.03533	1	1.794876	-0.79488
0.144444	0.166355	-0.02191	1	1.49299	-0.49299	0.144444	0.187721	-0.04328	1	1.973727	-0.97373
0.188889	0.210397	-0.02151	2	2.483925	-0.48393	0.188889	0.22601	-0.03712	2	2.835234	-0.83523
0.188889	0.213796	-0.02491	2	2.560415	-0.56042	0.188889	0.198552	-0.00966	2	2.21741	-0.21741
0.233333	0.199852	0.033482	3	2.246662	0.753338	0.233333	0.217863	0.015471	3	2.651913	0.348087
0.233333	0.249125	-0.01579	3	3.355302	-0.3553	0.233333	0.257367	-0.02403	3	3.54076	-0.54076
0.233333	0.258167	-0.02483	3	3.558764	-0.55876	0.233333	0.3014	-0.06807	3	4.531506	-1.53151
0.233333	0.21353	0.019804	3	2.554417	0.445583	0.233333	0.215141	0.018192	3	2.59068	0.40932
0.233333	0.226293	0.007041	3	2.841582	0.158418	0.233333	0.234879	-0.00155	3	3.034768	-0.03477
0.277778	0.272924	0.004854	4	3.890786	0.109214	0.277778	0.288693	-0.01091	4	4.245586	-0.24559
0.277778	0.298052	-0.02027	4	4.456169	-0.45617	0.277778	0.294839	-0.01706	4	4.38387	-0.38387
0.277778	0.233847	0.043931	4	3.011561	0.988439	0.277778	0.233691	0.044087	4	3.008042	0.991958
0.277778	0.259419	0.018359	4	3.586929	0.413071	0.277778	0.298597	-0.02082	4	4.468434	-0.46843
0.277778	0.225638	0.05214	4	2.826858	1.173142	0.277778	0.260345	0.017432	4	3.607773	0.392227
0.322222	0.372989	-0.05077	5	6.142257	-1.14226	0.322222	0.374523	-0.0523	5	6.176775	-1.17678
0.322222	0.328678	-0.00646	5	5.145265	-0.14526	0.322222	0.354026	-0.0318	5	5.715575	-0.71557
0.322222	0.360633	-0.03841	5	5.864244	-0.86424	0.322222	0.34286	-0.02064	5	5.464356	-0.46436
0.322222	0.316239	0.005983	5	4.865374	0.134626	0.322222	0.33135	-0.00913	5	5.205382	-0.20538
0.322222	0.254069	0.068153	5	3.466552	1.533448	0.322222	0.282388	0.039835	5	4.103723	0.896277
0.322222	0.277989	0.044233	5	4.004751	0.995249	0.322222	0.228218	0.094004	5	2.884915	2.115085
0.322222	0.308152	0.01407	5	4.683414	0.316586	0.322222	0.283879	0.038343	5	4.137286	0.862714
0.322222	0.293569	0.028653	5	4.35531	0.64469	0.322222	0.289841	0.032381	5	4.271426	0.728574

0.322222	0.237513	0.084709	5	3.094052	1.905948	0.322222	0.258517	0.063705	5	3.566638	1.433362
0.366667	0.382985	-0.01632	6	6.367157	-0.36716	0.366667	0.33729	0.029377	6	5.339021	0.660979
0.366667	0.385612	-0.01895	6	6.426271	-0.42627	0.366667	0.325615	0.041052	6	5.076329	0.923671
0.366667	0.394984	-0.02832	6	6.637145	-0.63714	0.366667	0.397414	-0.03075	6	6.691818	-0.69182
0.366667	0.308296	0.058371	6	4.686659	1.313341	0.366667	0.314863	0.051804	6	4.834407	1.165593
0.366667	0.332302	0.034365	6	5.226793	0.773207	0.366667	0.319778	0.046889	6	4.944994	1.055006
0.366667	0.336106	0.030561	6	5.312377	0.687623	0.366667	0.340566	0.0261	6	5.412744	0.587256
0.411111	0.461114	-0.05	7	8.125059	-1.12506	0.411111	0.420094	-0.00898	7	7.20212	-0.20212
0.411111	0.431339	-0.02023	7	7.45512	-0.45512	0.411111	0.421975	-0.01086	7	7.244428	-0.24443
0.411111	0.470856	-0.05974	7	8.34426	-1.34426	0.411111	0.507079	-0.09597	7	9.159269	-2.15927
0.411111	0.359218	0.051893	7	5.832397	1.167603	0.411111	0.367116	0.043995	7	6.010118	0.989882
0.411111	0.382208	0.028904	7	6.349669	0.650331	0.411111	0.366599	0.044512	7	5.998486	1.001514
0.455556	0.523232	-0.06768	8	9.522714	-1.52271	0.455556	0.492174	-0.03662	8	8.823917	-0.82392
0.455556	0.461715	-0.00616	8	8.138577	-0.13858	0.455556	0.495258	-0.0397	8	8.893304	-0.8933
0.455556	0.497084	-0.04153	8	8.934396	-0.9344	0.455556	0.471262	-0.01571	8	8.353396	-0.3534
0.455556	0.470471	-0.01492	8	8.335596	-0.3356	0.455556	0.487025	-0.03147	8	8.708065	-0.70807
0.455556	0.456702	-0.00115	8	8.025785	-0.02579	0.455556	0.505609	-0.05005	8	9.126196	-1.1262
0.455556	0.439313	0.016243	8	7.634543	0.365457	0.455556	0.433893	0.021663	8	7.512591	0.487409
0.455556	0.505834	-0.05028	8	9.131257	-1.13126	0.455556	0.532839	-0.07728	8	9.738887	-1.73889
0.5	0.506475	-0.00647	9	9.145679	-0.14568	0.5	0.499801	0.000199	9	8.995521	0.004479
0.5	0.600891	-0.10089	9	11.27005	-2.27005	0.5	0.627338	-0.12734	9	11.86511	-2.86511
0.5	0.571437	-0.07144	9	10.60734	-1.60734	0.5	0.58216	-0.08216	9	10.8486	-1.8486
0.5	0.487631	0.012369	9	8.721686	0.278314	0.5	0.453523	0.046477	9	7.954264	1.045736
0.5	0.512127	-0.01213	9	9.272865	-0.27286	0.5	0.542091	-0.04209	9	9.947051	-0.94705
0.544444	0.595767	-0.05132	10	11.15476	-1.15476	0.544444	0.587552	-0.04311	10	10.96991	-0.96991
0.544444	0.553968	-0.00952	10	10.21427	-0.21427	0.544444	0.557872	-0.01343	10	10.30212	-0.30212
0.544444	0.541957	0.002487	10	9.944042	0.055958	0.544444	0.54965	-0.00521	10	10.11713	-0.11713
0.544444	0.618901	-0.07446	10	11.67526	-1.67526	0.544444	0.636096	-0.09165	10	12.06216	-2.06216
0.544444	0.627328	-0.08288	10	11.86488	-1.86488	0.544444	0.648409	-0.10396	10	12.3392	-2.3392
0.544444	0.456461	0.087984	10	8.02037	1.97963	0.544444	0.465071	0.079373	10	8.214105	1.785895
0.544444	0.466552	0.077892	10	8.247431	1.752569	0.544444	0.457793	0.086651	10	8.050347	1.949653
0.544444	0.598431	-0.05399	10	11.2147	-1.2147	0.544444	0.612853	-0.06841	10	11.53919	-1.53919
0.544444	0.571471	-0.02703	10	10.60809	-0.60809	0.544444	0.556475	-0.01203	10	10.27069	-0.27069
0.544444	0.633798	-0.08935	10	12.01045	-2.01045	0.544444	0.611742	-0.0673	10	11.5142	-1.5142

0.544444	0.623278	-0.07883	10	11.77376	-1.77376	0.544444	0.611675	-0.06723	10	11.51269	-1.51269
0.588889	0.650968	-0.06208	11	12.39678	-1.39678	0.588889	0.654289	-0.0654	11	12.47149	-1.47149
0.588889	0.634459	-0.04557	11	12.02533	-1.02533	0.588889	0.624107	-0.03522	11	11.79241	-0.79241
0.588889	0.629638	-0.04075	11	11.91686	-0.91686	0.588889	0.647308	-0.05842	11	12.31444	-1.31444
0.588889	0.705017	-0.11613	11	13.61289	-2.61289	0.588889	0.694349	-0.10546	11	13.37285	-2.37285
0.588889	0.639809	-0.05092	11	12.1457	-1.1457	0.588889	0.665245	-0.07636	11	12.71802	-1.71802
0.588889	0.721033	-0.13214	11	13.97323	-2.97323	0.588889	0.723771	-0.13488	11	14.03485	-3.03485
0.588889	0.595922	-0.00703	11	11.15826	-0.15826	0.588889	0.638663	-0.04977	11	12.11991	-1.11991
0.588889	0.624415	-0.03553	11	11.79933	-0.79933	0.588889	0.633244	-0.04436	11	11.998	-0.998
0.633333	0.688122	-0.05479	12	13.23276	-1.23276	0.633333	0.676039	-0.04271	12	12.96087	-0.96087
0.633333	0.675302	-0.04197	12	12.9443	-0.9443	0.633333	0.665002	-0.03167	12	12.71255	-0.71255
0.633333	0.699173	-0.06584	12	13.48139	-1.48139	0.633333	0.723146	-0.08981	12	14.02078	-2.02078
0.633333	0.684366	-0.05103	12	13.14825	-1.14825	0.633333	0.674208	-0.04087	12	12.91968	-0.91968
0.633333	0.633679	-0.00035	12	12.00777	-0.00777	0.633333	0.598614	0.034719	12	11.21882	0.781179
0.633333	0.71815	-0.08482	12	13.90837	-1.90837	0.633333	0.733728	-0.10039	12	14.25888	-2.25888
0.633333	0.645973	-0.01264	12	12.2844	-0.2844	0.633333	0.659332	-0.026	12	12.58498	-0.58498
0.633333	0.713715	-0.08038	12	13.80859	-1.80859	0.633333	0.748327	-0.11499	12	14.58737	-2.58737
0.633333	0.730492	-0.09716	12	14.18607	-2.18607	0.633333	0.711662	-0.07833	12	13.76239	-1.76239
0.633333	0.738803	-0.10547	12	14.37308	-2.37308	0.633333	0.718778	-0.08544	12	13.9225	-1.9225
0.633333	0.775643	-0.14231	12	15.20197	-3.20197	0.633333	0.78832	-0.15499	12	15.48719	-3.48719
0.633333	0.694537	-0.0612	12	13.37707	-1.37707	0.633333	0.697372	-0.06404	12	13.44087	-1.44087
0.633333	0.654974	-0.02164	12	12.48692	-0.48692	0.633333	0.654846	-0.02151	12	12.48403	-0.48403
0.677778	0.720806	-0.04303	13	13.96813	-0.96813	0.677778	0.706399	-0.02862	13	13.64398	-0.64398
0.677778	0.734824	-0.05705	13	14.28355	-1.28355	0.677778	0.722081	-0.0443	13	13.99682	-0.99682
0.677778	0.670179	0.007598	13	12.82903	0.170966	0.677778	0.654504	0.023274	13	12.47634	0.523661
0.677778	0.710441	-0.03266	13	13.73492	-0.73492	0.677778	0.701583	-0.0238	13	13.53561	-0.53561
0.677778	0.753488	-0.07571	13	14.70348	-1.70348	0.677778	0.724524	-0.04675	13	14.05178	-1.05178
0.677778	0.73946	-0.06168	13	14.38786	-1.38786	0.677778	0.732736	-0.05496	13	14.23656	-1.23656
0.677778	0.68625	-0.00847	13	13.19063	-0.19063	0.677778	0.6603	0.017478	13	12.60674	0.393256
0.677778	0.708137	-0.03036	13	13.68307	-0.68307	0.677778	0.655945	0.021833	13	12.50876	0.491242
0.677778	0.731759	-0.05398	13	14.21457	-1.21457	0.677778	0.717746	-0.03997	13	13.89928	-0.89928
0.677778	0.749635	-0.07186	13	14.6168	-1.6168	0.677778	0.735311	-0.05753	13	14.2945	-1.2945
0.722222	0.695498	0.026725	14	13.3987	0.601302	0.722222	0.697951	0.024271	14	13.4539	0.546097
0.722222	0.705544	0.016679	14	13.62473	0.375271	0.722222	0.739204	-0.01698	14	14.38209	-0.38209

0.722222	0.764601	-0.04238	14	14.95352	-0.95352	0.722222	0.778627	-0.0564	14	15.2691	-1.2691
0.722222	0.703282	0.01894	14	13.57385	0.426153	0.722222	0.750573	-0.02835	14	14.63789	-0.63789
0.722222	0.690281	0.031941	14	13.28133	0.718675	0.722222	0.670247	0.051975	14	12.83056	1.169437
0.722222	0.652539	0.069683	14	12.43213	1.567866	0.722222	0.681663	0.040559	14	13.08741	0.912587
0.722222	0.760857	-0.03863	14	14.86928	-0.86928	0.722222	0.764147	-0.04193	14	14.94332	-0.94332
0.722222	0.718841	0.003382	14	13.92391	0.076085	0.722222	0.730861	-0.00864	14	14.19438	-0.19438
0.722222	0.753209	-0.03099	14	14.6972	-0.6972	0.722222	0.741087	-0.01887	14	14.42446	-0.42446
0.722222	0.724093	-0.00187	14	14.04209	-0.04209	0.722222	0.732278	-0.01006	14	14.22625	-0.22625
0.722222	0.699648	0.022574	14	13.49209	0.507914	0.722222	0.695299	0.026923	14	13.39423	0.605767
0.722222	0.785771	-0.06355	14	15.42985	-1.42985	0.722222	0.781922	-0.0597	14	15.34325	-1.34325
0.766667	0.775306	-0.00864	15	15.19438	-0.19438	0.766667	0.781423	-0.01476	15	15.33202	-0.33202
0.766667	0.770097	-0.00343	15	15.07719	-0.07719	0.766667	0.763863	0.002804	15	14.93691	0.063094
0.766667	0.699794	0.066872	15	13.49537	1.504625	0.766667	0.711916	0.05475	15	13.76812	1.231879
0.766667	0.733795	0.032872	15	14.26038	0.73962	0.766667	0.758179	0.008488	15	14.80903	0.190971
0.766667	0.690702	0.075965	15	13.2908	1.709204	0.766667	0.635281	0.131385	15	12.04383	2.956167
0.766667	0.762453	0.004214	15	14.90519	0.094814	0.766667	0.770714	-0.00405	15	15.09107	-0.09107
0.766667	0.709332	0.057335	15	13.70997	1.290029	0.766667	0.733014	0.033653	15	14.24282	0.757182
0.811111	0.675023	0.136088	16	12.93802	3.061976	0.811111	0.674608	0.136503	16	12.92869	3.071312
0.811111	0.791055	0.020056	16	15.54873	0.45127	0.811111	0.779898	0.031213	16	15.2977	0.702301
0.811111	0.702278	0.108834	16	13.55125	2.448754	0.811111	0.699832	0.11128	16	13.49621	2.503791
0.811111	0.771042	0.040069	16	15.09844	0.901563	0.811111	0.789259	0.021852	16	15.50832	0.491679
0.811111	0.768822	0.042289	16	15.04849	0.951513	0.811111	0.763169	0.047942	16	14.92131	1.078687
0.811111	0.73492	0.076191	16	14.28571	1.714291	0.811111	0.700234	0.110877	16	13.50528	2.494724
0.811111	0.759866	0.051245	16	14.84699	1.153014	0.811111	0.738253	0.072858	16	14.36069	1.639311
0.811111	0.768394	0.042717	16	15.03887	0.96113	0.811111	0.758581	0.05253	16	14.81808	1.181924
0.811111	0.758427	0.052684	16	14.81462	1.185382	0.811111	0.771859	0.039253	16	15.11682	0.883182
0.855556	0.693997	0.161559	17	13.36493	3.635071	0.855556	0.66678	0.188775	17	12.75256	4.247441
0.855556	0.735337	0.120219	17	14.29507	2.704926	0.855556	0.736593	0.118963	17	14.32334	2.676665
0.855556	0.704995	0.15056	17	13.6124	3.387603	0.855556	0.709957	0.145598	17	13.72404	3.275957
0.855556	0.719926	0.135629	17	13.94834	3.051658	0.855556	0.750149	0.105407	17	14.62835	2.371648
0.855556	0.725594	0.129962	17	14.07586	2.924139	0.855556	0.692093	0.163462	17	13.3221	3.677897
0.855556	0.781936	0.07362	17	15.34355	1.656449	0.855556	0.775605	0.079951	17	15.20111	1.798887
0.9	0.6807	0.2193	18	13.06576	4.93424	0.9	0.692208	0.207792	18	13.32468	4.67532
0.9	0.672943	0.227057	18	12.89121	5.108787	0.9	0.635281	0.264719	18	12.04382	5.956181

0.9	0.709247	0.190753	18	13.70805	4.291952	0.9	0.720846	0.179154	18	13.96904	4.03096
0.9	0.689351	0.210649	18	13.2604	4.739598	0.9	0.707722	0.192278	18	13.67374	4.326263

APPENDIX P

NORMALIZED AND DENORMALIZED TEST DATASET OF TARGET FEATURE, OUTPUTS AND ERRORS FOR FEMALE HISPANIC POPULATION USING ANN APPROACH AND GA-ANN APPROACH AND THEIR DENORMALIZED TEST SET SQUARE ERRORS

ANN							GA-ANN						
Normalized			Denormalized				Normalized			Denormalized			
Target	Output	Error	Target	Output	Error	Error^2	Target	Output	Error	Target	Output	Error	Error^2
0.144444	0.160235	-0.01579	1	1.35528	-0.35528	0.1262236	0.144444	0.171211	-0.02677	1	1.602258	-0.60226	0.3627149
0.144444	0.185898	-0.04145	1	1.93271	-0.93271	0.8699482	0.144444	0.188988	-0.04454	1	2.00224	-1.00224	1.0044851
0.188889	0.217836	-0.02895	2	2.651318	-0.65132	0.4242148	0.188889	0.239164	-0.05027	2	3.131187	-1.13119	1.2795851
0.188889	0.208893	-0.02	2	2.450083	-0.45008	0.202575	0.188889	0.205634	-0.01675	2	2.376769	-0.37677	0.1419548
0.188889	0.190312	-0.00142	2	2.032009	-0.03201	0.0010246	0.188889	0.173359	0.01553	2	1.650577	0.349423	0.1220967
0.322222	0.314485	0.007737	5	4.825922	0.174078	0.0303032	0.322222	0.318016	0.004206	5	4.905369	0.094631	0.008955
0.366667	0.330118	0.036549	6	5.177644	0.822356	0.6762698	0.366667	0.337291	0.029376	6	5.339039	0.660961	0.4368701
0.366667	0.320472	0.046195	6	4.960624	1.039376	1.0803034	0.366667	0.320741	0.045926	6	4.966668	1.033332	1.0677757
0.366667	0.333974	0.032693	6	5.264418	0.735582	0.5410809	0.366667	0.351957	0.01471	6	5.669032	0.330968	0.1095401
0.366667	0.360853	0.005814	6	5.869192	0.130808	0.0171107	0.366667	0.355153	0.011514	6	5.740944	0.259056	0.0671099
0.411111	0.361824	0.049287	7	5.891033	1.108967	1.2298072	0.411111	0.334841	0.076271	7	5.283911	1.716089	2.9449606
0.411111	0.499053	-0.08794	7	8.978694	-1.97869	3.9152298	0.411111	0.412348	-0.00124	7	7.027829	-0.02783	0.0007745
0.411111	0.402584	0.008527	7	6.808137	0.191862	0.0368112	0.411111	0.43861	-0.0275	7	7.618726	-0.61873	0.3828213
0.411111	0.358334	0.052777	7	5.812521	1.187479	1.4101061	0.411111	0.404148	0.006963	7	6.843336	0.156664	0.0245435
0.411111	0.475403	-0.06429	7	8.446568	-1.44657	2.0925581	0.411111	0.48125	-0.07014	7	8.578115	-1.57811	2.4904463
0.455556	0.418337	0.037219	8	7.162575	0.837425	0.7012808	0.455556	0.392133	0.063422	8	6.572998	1.427002	2.0363357
0.455556	0.447337	0.008218	8	7.815088	0.184912	0.0341925	0.455556	0.47282	-0.01726	8	8.388447	-0.38845	0.1508907
0.5	0.538355	-0.03835	9	9.86298	-0.86298	0.7447344	0.5	0.519588	-0.01959	9	9.440739	-0.44074	0.194251
0.5	0.659258	-0.15926	9	12.58331	-3.58331	12.840116	0.5	0.611188	-0.11119	9	11.50172	-2.50172	6.2586081
0.5	0.617134	-0.11713	9	11.63552	-2.63552	6.9459483	0.5	0.613335	-0.11334	9	11.55004	-2.55004	6.5026916
0.5	0.643955	-0.14395	9	12.23898	-3.23898	10.491005	0.5	0.646051	-0.14605	9	12.28615	-3.28615	10.798797
0.5	0.446576	0.053424	9	7.797965	1.202035	1.4448877	0.5	0.504036	-0.00404	9	9.090821	-0.09082	0.0082484
0.544444	0.55614	-0.0117	10	10.26316	-0.26316	0.0692529	0.544444	0.580103	-0.03566	10	10.80231	-0.80231	0.6437025

0.544444	0.641034	-0.09659	10	12.17328	-2.17328	4.7231265	0.544444	0.678766	-0.13432	10	13.02223	-3.02223	9.1338448
0.544444	0.525876	0.018569	10	9.582207	0.417793	0.1745513	0.544444	0.538444	0.006	10	9.864999	0.135001	0.0182252
0.588889	0.510719	0.07817	11	9.241172	1.758828	3.0934744	0.588889	0.51797	0.070919	11	9.40432	1.59568	2.5461933
0.588889	0.677289	-0.0884	11	12.98901	-1.98901	3.9561596	0.588889	0.642245	-0.05336	11	12.20051	-1.20051	1.4412173
0.588889	0.676841	-0.08795	11	12.97893	-1.97893	3.9161718	0.588889	0.696297	-0.10741	11	13.41669	-2.41669	5.8403941
0.588889	0.7105	-0.12161	11	13.73625	-2.73625	7.4870672	0.588889	0.692444	-0.10355	11	13.32999	-2.32999	5.4288405
0.588889	0.66258	-0.07369	11	12.65805	-1.65805	2.7491175	0.588889	0.672237	-0.08335	11	12.87533	-1.87533	3.516855
0.588889	0.63075	-0.04186	11	11.94189	-0.94189	0.8871492	0.588889	0.649932	-0.06104	11	12.37348	-1.37348	1.8864476
0.588889	0.644503	-0.05561	11	12.25132	-1.25132	1.5658051	0.588889	0.655515	-0.06663	11	12.49909	-1.49909	2.2472766
0.633333	0.7633	-0.12997	12	14.92424	-2.92424	8.5511858	0.633333	0.775129	-0.1418	12	15.19039	-3.19039	10.178607
0.633333	0.633228	0.000105	12	11.99764	0.002363	5.586E-06	0.633333	0.58855	0.044783	12	10.99238	1.007624	1.0153071
0.677778	0.710951	-0.03317	13	13.74639	-0.74639	0.5570933	0.677778	0.67701	0.000768	13	12.98272	0.017276	0.0002985
0.677778	0.722019	-0.04424	13	13.99543	-0.99543	0.9908809	0.677778	0.700183	-0.02241	13	13.50413	-0.50413	0.254145
0.677778	0.692657	-0.01488	13	13.33478	-0.33478	0.1120801	0.677778	0.686474	-0.0087	13	13.19567	-0.19567	0.0382849
0.677778	0.72904	-0.05126	13	14.15341	-1.15341	1.3303506	0.677778	0.741731	-0.06395	13	14.43896	-1.43896	2.0706008
0.677778	0.735931	-0.05815	13	14.30844	-1.30844	1.7120282	0.677778	0.744738	-0.06696	13	14.5066	-1.5066	2.2698366
0.722222	0.7472	-0.02498	14	14.56199	-0.56199	0.3158322	0.722222	0.747734	-0.02551	14	14.57401	-0.57401	0.3294915
0.722222	0.768505	-0.04628	14	15.04137	-1.04137	1.0844461	0.722222	0.75182	-0.0296	14	14.66595	-0.66595	0.4434849
0.766667	0.668945	0.097722	15	12.80126	2.198745	4.8344794	0.766667	0.648801	0.117866	15	12.34801	2.651988	7.0330389
0.766667	0.777541	-0.01087	15	15.24468	-0.24468	0.0598674	0.766667	0.776177	-0.00951	15	15.21399	-0.21399	0.0457933
0.766667	0.743967	0.0227	15	14.48925	0.51075	0.2608654	0.766667	0.710237	0.05643	15	13.73032	1.269677	1.6120799
0.811111	0.754291	0.05682	16	14.72155	1.278452	1.6344391	0.811111	0.730238	0.080873	16	14.18035	1.819646	3.3111118
0.855556	0.708331	0.147225	17	13.68745	3.312554	10.973016	0.855556	0.696038	0.159518	17	13.41085	3.589153	12.882022
0.855556	0.744206	0.111349	17	14.49464	2.505358	6.2768201	0.855556	0.756936	0.098619	17	14.78106	2.218936	4.923677
0.855556	0.671879	0.183676	17	12.86728	4.132715	17.079336	0.855556	0.659169	0.196386	17	12.58131	4.418687	19.524793
0.855556	0.749239	0.106317	17	14.60787	2.392127	5.7222714	0.855556	0.736191	0.119364	17	14.31431	2.685692	7.2129436
0.9	0.749722	0.150278	18	14.61873	3.381266	11.432958	0.9	0.753526	0.146474	18	14.70434	3.295661	10.86138
0.9	0.779885	0.120115	18	15.29742	2.70258	7.3039394	0.9	0.777513	0.122487	18	15.24403	2.755965	7.5953434
0.9	0.705995	0.194005	18	13.63488	4.365124	19.054303	0.9	0.733892	0.166108	18	14.26258	3.737422	13.968326
0.9	0.773929	0.126071	18	15.1634	2.836596	8.0462773	0.9	0.786697	0.113303	18	15.45068	2.549322	6.4990433
0.9	0.71777	0.18223	18	13.89982	4.100182	16.811488	0.9	0.735001	0.164999	18	14.28753	3.712467	13.782411
0.9	0.670333	0.229667	18	12.83249	5.16751	26.703157	0.9	0.663309	0.236691	18	12.67444	5.325557	28.361559

APPENDIX Q

NORMALIZED AND DENORMALIZED TRAINING DATASET OF TARGET FEATURE, OUTPUTS AND ERRORS FOR MALE HISPANIC POPULATION USING ANN APPROACH AND GA-ANN APPROACH

ANN						GA-ANN					
Normalized			Denormalized			Normalized			Denormalized		
Target	Output	Error	Target	Output	Error	Target	Output	Error	Target	Output	Error
0.1	0.172571	-0.07257	0	1.632857	-1.63286	0.1	0.166519	-0.06652	0	1.496672	-1.49667
0.1	0.207328	-0.10733	0	2.414884	-2.41488	0.1	0.189582	-0.08958	0	2.015598	-2.0156
0.1	0.175667	-0.07567	0	1.702509	-1.70251	0.1	0.181696	-0.0817	0	1.838155	-1.83815
0.144444	0.254243	-0.1098	1	3.470469	-2.47047	0.144444	0.243028	-0.09858	1	3.218131	-2.21813
0.144444	0.186174	-0.04173	1	1.938922	-0.93892	0.144444	0.170479	-0.02603	1	1.585769	-0.58577
0.144444	0.215885	-0.07144	1	2.607419	-1.60742	0.144444	0.20733	-0.06289	1	2.414928	-1.41493
0.144444	0.162137	-0.01769	1	1.398087	-0.39809	0.144444	0.17621	-0.03177	1	1.714721	-0.71472
0.144444	0.196641	-0.0522	1	2.174425	-1.17443	0.144444	0.195107	-0.05066	1	2.139917	-1.13992
0.188889	0.207155	-0.01827	2	2.410987	-0.41099	0.188889	0.218731	-0.02984	2	2.671457	-0.67146
0.188889	0.21969	-0.0308	2	2.693025	-0.69303	0.188889	0.23199	-0.0431	2	2.969769	-0.96977
0.188889	0.195742	-0.00685	2	2.1542	-0.1542	0.188889	0.221564	-0.03268	2	2.73519	-0.73519
0.233333	0.262522	-0.02919	3	3.656738	-0.65674	0.233333	0.253572	-0.02024	3	3.455374	-0.45537
0.233333	0.208481	0.024852	3	2.440825	0.559175	0.233333	0.211307	0.022026	3	2.504407	0.495593
0.233333	0.175958	0.057376	3	1.709046	1.290954	0.233333	0.166172	0.067161	3	1.488871	1.511129
0.277778	0.303618	-0.02584	4	4.581398	-0.5814	0.277778	0.297836	-0.02006	4	4.451309	-0.45131
0.277778	0.260344	0.017434	4	3.607738	0.392262	0.277778	0.252396	0.025382	4	3.428909	0.571091
0.277778	0.300283	-0.02251	4	4.506367	-0.50637	0.277778	0.314114	-0.03634	4	4.817573	-0.81757
0.322222	0.320653	0.001569	5	4.964702	0.035298	0.322222	0.354719	-0.0325	5	5.73117	-0.73117
0.322222	0.381886	-0.05966	5	6.342426	-1.34243	0.322222	0.394602	-0.07238	5	6.628536	-1.62854
0.322222	0.360614	-0.03839	5	5.863814	-0.86381	0.322222	0.310325	0.011897	5	4.73231	0.26769

0.322222	0.366338	-0.04412	5	5.992612	-0.99261	0.322222	0.33592	-0.0137	5	5.308191	-0.30819
0.322222	0.248139	0.074084	5	3.333119	1.666881	0.322222	0.287523	0.0347	5	4.219259	0.780741
0.366667	0.330499	0.036168	6	5.186226	0.813774	0.366667	0.332674	0.033993	6	5.235156	0.764844
0.366667	0.430909	-0.06424	6	7.445459	-1.44546	0.366667	0.47508	-0.10841	6	8.439307	-2.43931
0.366667	0.400609	-0.03394	6	6.763714	-0.76371	0.366667	0.417345	-0.05068	6	7.140267	-1.14027
0.366667	0.36031	0.006356	6	5.856984	0.143016	0.366667	0.397307	-0.03064	6	6.689397	-0.6894
0.366667	0.436837	-0.07017	6	7.578844	-1.57884	0.366667	0.430617	-0.06395	6	7.438873	-1.43887
0.366667	0.395519	-0.02885	6	6.649169	-0.64917	0.366667	0.364779	0.001888	6	5.957523	0.042477
0.366667	0.399004	-0.03234	6	6.727586	-0.72759	0.366667	0.407628	-0.04096	6	6.92162	-0.92162
0.411111	0.475807	-0.0647	7	8.455651	-1.45565	0.411111	0.442613	-0.0315	7	7.708794	-0.70879
0.411111	0.352385	0.058726	7	5.67867	1.32133	0.411111	0.347416	0.063695	7	5.56686	1.43314
0.411111	0.442254	-0.03114	7	7.700722	-0.70072	0.411111	0.44309	-0.03198	7	7.719521	-0.71952
0.411111	0.491892	-0.08078	7	8.817571	-1.81757	0.411111	0.482505	-0.07139	7	8.606369	-1.60637
0.411111	0.48968	-0.07857	7	8.767806	-1.76781	0.411111	0.499551	-0.08844	7	8.989892	-1.98989
0.411111	0.56608	-0.15497	7	10.4868	-3.4868	0.411111	0.553425	-0.14231	7	10.20207	-3.20207
0.455556	0.418726	0.036829	8	7.171346	0.828654	0.455556	0.431625	0.023931	8	7.461555	0.538445
0.455556	0.513724	-0.05817	8	9.308789	-1.30879	0.455556	0.472117	-0.01656	8	8.372623	-0.37262
0.455556	0.503063	-0.04751	8	9.068929	-1.06893	0.455556	0.511397	-0.05584	8	9.256443	-1.25644
0.455556	0.445336	0.01022	8	7.770052	0.229948	0.455556	0.457168	-0.00161	8	8.036278	-0.03628
0.455556	0.457638	-0.00208	8	8.046853	-0.04685	0.455556	0.429479	0.026077	8	7.413273	0.586727
0.455556	0.546596	-0.09104	8	10.0484	-2.0484	0.455556	0.54986	-0.0943	8	10.12185	-2.12185
0.5	0.536275	-0.03627	9	9.816181	-0.81618	0.5	0.527892	-0.02789	9	9.627564	-0.62756
0.5	0.645299	-0.1453	9	12.26923	-3.26923	0.5	0.613822	-0.11382	9	11.56098	-2.56098
0.5	0.512461	-0.01246	9	9.280372	-0.28037	0.5	0.49497	0.00503	9	8.886825	0.113175
0.5	0.449227	0.050773	9	7.857611	1.142389	0.5	0.52493	-0.02493	9	9.560919	-0.56092
0.5	0.446992	0.053008	9	7.807316	1.192684	0.5	0.458223	0.041777	9	8.060014	0.939986
0.5	0.571867	-0.07187	9	10.61701	-1.61701	0.5	0.576743	-0.07674	9	10.72672	-1.72672
0.5	0.522621	-0.02262	9	9.508971	-0.50897	0.5	0.513448	-0.01345	9	9.302589	-0.30259
0.544444	0.525515	0.018929	10	9.574098	0.425902	0.544444	0.590817	-0.04637	10	11.04339	-1.04339

0.544444	0.440878	0.103567	10	7.669754	2.330246	0.544444	0.458818	0.085627	10	8.073402	1.926598
0.544444	0.621997	-0.07755	10	11.74492	-1.74492	0.544444	0.6876	-0.14316	10	13.22101	-3.22101
0.544444	0.534561	0.009883	10	9.777628	0.222372	0.544444	0.523619	0.020826	10	9.531418	0.468582
0.544444	0.420641	0.123804	10	7.214419	2.785581	0.544444	0.43003	0.114414	10	7.425675	2.574325
0.544444	0.548797	-0.00435	10	10.09794	-0.09794	0.544444	0.526863	0.017581	10	9.604424	0.395576
0.544444	0.63918	-0.09474	10	12.13156	-2.13156	0.544444	0.621112	-0.07667	10	11.72502	-1.72502
0.544444	0.578651	-0.03421	10	10.76965	-0.76965	0.544444	0.540208	0.004236	10	9.904683	0.095317
0.544444	0.643991	-0.09955	10	12.23979	-2.23979	0.544444	0.612106	-0.06766	10	11.5224	-1.5224
0.588889	0.614023	-0.02513	11	11.56553	-0.56553	0.588889	0.576067	0.012822	11	10.71151	0.288491
0.588889	0.587804	0.001085	11	10.97558	0.024418	0.588889	0.567007	0.021882	11	10.50766	0.492339
0.588889	0.605487	-0.0166	11	11.37346	-0.37346	0.588889	0.598919	-0.01003	11	11.22569	-0.22569
0.588889	0.528569	0.06032	11	9.642799	1.357201	0.588889	0.62242	-0.03353	11	11.75444	-0.75444
0.588889	0.700331	-0.11144	11	13.50744	-2.50744	0.588889	0.705795	-0.11691	11	13.63039	-2.63039
0.588889	0.618777	-0.02989	11	11.67248	-0.67248	0.588889	0.621765	-0.03288	11	11.7397	-0.7397
0.588889	0.302177	0.286712	11	4.548985	6.451015	0.588889	0.282876	0.306013	11	4.114708	6.885292
0.588889	0.295575	0.293314	11	4.40043	6.59957	0.588889	0.288837	0.300052	11	4.248823	6.751177
0.588889	0.74576	-0.15687	11	14.52959	-3.52959	0.588889	0.75224	-0.16335	11	14.6754	-3.6754
0.588889	0.613522	-0.02463	11	11.55424	-0.55424	0.588889	0.62318	-0.03429	11	11.77155	-0.77155
0.588889	0.523219	0.06567	11	9.522428	1.477572	0.588889	0.500172	0.088717	11	9.003865	1.996135
0.588889	0.618473	-0.02958	11	11.66563	-0.66563	0.588889	0.667352	-0.07846	11	12.76542	-1.76542
0.633333	0.648467	-0.01513	12	12.3405	-0.3405	0.633333	0.618815	0.014519	12	11.67333	0.326672
0.633333	0.654932	-0.0216	12	12.48596	-0.48596	0.633333	0.640328	-0.00699	12	12.15738	-0.15738
0.633333	0.703547	-0.07021	12	13.57981	-1.57981	0.633333	0.735365	-0.10203	12	14.29571	-2.29571
0.633333	0.674681	-0.04135	12	12.93033	-0.93033	0.633333	0.713418	-0.08008	12	13.80189	-1.80189
0.633333	0.67994	-0.04661	12	13.04864	-1.04864	0.633333	0.642078	-0.00874	12	12.19675	-0.19675
0.633333	0.673461	-0.04013	12	12.90286	-0.90286	0.633333	0.706479	-0.07315	12	13.64579	-1.64579
0.633333	0.582738	0.050596	12	10.8616	1.1384	0.633333	0.620006	0.013327	12	11.70014	0.299859
0.633333	0.701061	-0.06773	12	13.52387	-1.52387	0.633333	0.727709	-0.09438	12	14.12344	-2.12344
0.633333	0.749477	-0.11614	12	14.61324	-2.61324	0.633333	0.730584	-0.09725	12	14.18814	-2.18814

0.677778	0.693752	-0.01597	13	13.35943	-0.35943	0.677778	0.711081	-0.0333	13	13.74932	-0.74932
0.677778	0.613899	0.063878	13	11.56274	1.437264	0.677778	0.62726	0.050518	13	11.86335	1.136652
0.677778	0.720978	-0.0432	13	13.972	-0.972	0.677778	0.713451	-0.03567	13	13.80264	-0.80264
0.677778	0.694748	-0.01697	13	13.38183	-0.38183	0.677778	0.649463	0.028315	13	12.36291	0.637086
0.677778	0.679699	-0.00192	13	13.04323	-0.04323	0.677778	0.695068	-0.01729	13	13.38902	-0.38902
0.677778	0.680444	-0.00267	13	13.05999	-0.05999	0.677778	0.715178	-0.0374	13	13.84151	-0.84151
0.677778	0.706112	-0.02833	13	13.63752	-0.63752	0.677778	0.744311	-0.06653	13	14.497	-1.497
0.677778	0.752776	-0.075	13	14.68746	-1.68746	0.677778	0.764773	-0.08699	13	14.95739	-1.95739
0.722222	0.705072	0.017151	14	13.61411	0.385888	0.722222	0.725463	-0.00324	14	14.07293	-0.07293
0.722222	0.72569	-0.00347	14	14.07803	-0.07803	0.722222	0.714331	0.007891	14	13.82245	0.177553
0.722222	0.801988	-0.07977	14	15.79473	-1.79473	0.722222	0.773808	-0.05159	14	15.16069	-1.16069
0.722222	0.354683	0.367539	14	5.730364	8.269636	0.722222	0.312349	0.409873	14	4.777859	9.222141
0.722222	0.752835	-0.03061	14	14.68878	-0.68878	0.722222	0.758122	-0.0359	14	14.80774	-0.80774
0.722222	0.695112	0.02711	14	13.39003	0.609974	0.722222	0.723409	-0.00119	14	14.02671	-0.02671
0.722222	0.748063	-0.02584	14	14.58141	-0.58141	0.722222	0.745263	-0.02304	14	14.51841	-0.51841
0.722222	0.750667	-0.02845	14	14.64002	-0.64002	0.722222	0.740336	-0.01811	14	14.40756	-0.40756
0.722222	0.813307	-0.09108	14	16.0494	-2.0494	0.722222	0.804551	-0.08233	14	15.8524	-1.8524
0.722222	0.789849	-0.06763	14	15.5216	-1.5216	0.722222	0.742455	-0.02023	14	14.45524	-0.45524
0.722222	0.794395	-0.07217	14	15.62388	-1.62388	0.722222	0.806974	-0.08475	14	15.90692	-1.90692
0.766667	0.676968	0.089699	15	12.98177	2.01823	0.766667	0.719706	0.046961	15	13.94338	1.056616
0.766667	0.727742	0.038925	15	14.12419	0.875813	0.766667	0.745156	0.021511	15	14.51601	0.483994
0.766667	0.761329	0.005338	15	14.8799	0.120103	0.766667	0.745202	0.021465	15	14.51704	0.482959
0.766667	0.716616	0.05005	15	13.87387	1.126133	0.766667	0.739303	0.027363	15	14.38433	0.615672
0.766667	0.769363	-0.0027	15	15.06066	-0.06066	0.766667	0.769522	-0.00285	15	15.06424	-0.06424
0.766667	0.706076	0.060591	15	13.63671	1.363288	0.766667	0.690051	0.076616	15	13.27615	1.723849
0.766667	0.791042	-0.02438	15	15.54845	-0.54845	0.766667	0.777576	-0.01091	15	15.24546	-0.24546
0.766667	0.692865	0.073802	15	13.33946	1.660536	0.766667	0.701277	0.06539	15	13.52873	1.471266
0.811111	0.764902	0.046209	16	14.96029	1.039708	0.811111	0.766423	0.044688	16	14.99452	1.005477
0.811111	0.783664	0.027447	16	15.38245	0.617552	0.811111	0.775684	0.035427	16	15.20289	0.797114

0.811111	0.769367	0.041744	16	15.06076	0.939238	0.811111	0.749383	0.061728	16	14.61113	1.388872
0.811111	0.802824	0.008287	16	15.81355	0.186453	0.811111	0.799696	0.011415	16	15.74316	0.256839
0.811111	0.715967	0.095144	16	13.85925	2.140746	0.811111	0.702828	0.108283	16	13.56363	2.436373
0.811111	0.783583	0.027528	16	15.38063	0.619374	0.811111	0.768514	0.042597	16	15.04157	0.958432
0.811111	0.834138	-0.02303	16	16.51811	-0.51811	0.811111	0.811447	-0.00034	16	16.00756	-0.00756
0.811111	0.691439	0.119672	16	13.30738	2.692619	0.811111	0.666363	0.144748	16	12.74317	3.256827
0.855556	0.766086	0.08947	17	14.98693	2.013068	0.855556	0.74685	0.108706	17	14.55412	2.44588
0.855556	0.807282	0.048274	17	15.91384	1.086161	0.855556	0.803392	0.052163	17	15.82633	1.173669
0.855556	0.776523	0.079033	17	15.22177	1.778233	0.855556	0.769001	0.086555	17	15.05252	1.947484
0.855556	0.789478	0.066078	17	15.51325	1.486753	0.855556	0.769883	0.085672	17	15.07238	1.927621
0.855556	0.784492	0.071064	17	15.40106	1.59894	0.855556	0.760717	0.094839	17	14.86613	2.133869
0.855556	0.803659	0.051897	17	15.83232	1.167681	0.855556	0.778915	0.076641	17	15.27558	1.724415
0.9	0.75355	0.14645	18	14.70487	3.295134	0.9	0.764479	0.135521	18	14.95077	3.049228
0.9	0.823126	0.076874	18	16.27033	1.729673	0.9	0.785434	0.114566	18	15.42226	2.577741
0.9	0.795589	0.104411	18	15.65076	2.349243	0.9	0.78482	0.11518	18	15.40844	2.591559
0.9	0.782202	0.117798	18	15.34956	2.650444	0.9	0.766582	0.133418	18	14.9981	3.001898
0.9	0.777406	0.122594	18	15.24162	2.758375	0.9	0.777928	0.122072	18	15.25338	2.746617
0.9	0.73439	0.16561	18	14.27378	3.726219	0.9	0.757506	0.142494	18	14.79389	3.206105
0.9	0.791646	0.108354	18	15.56204	2.437956	0.9	0.771954	0.128046	18	15.11897	2.881027
0.9	0.792467	0.107533	18	15.58052	2.419484	0.9	0.769683	0.130317	18	15.06787	2.932126

APPENDIX R

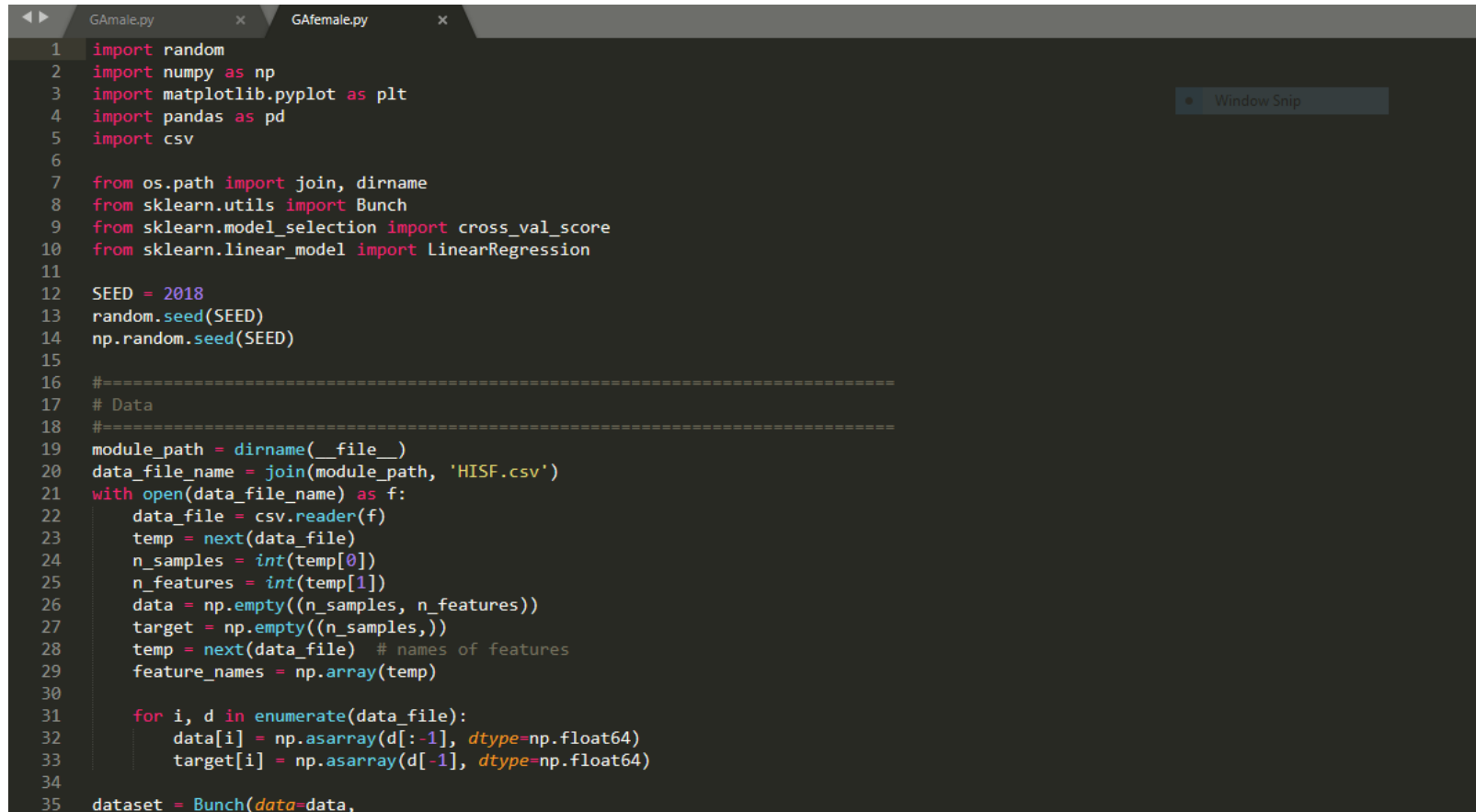
NORMALIZED AND DENORMALIZED TEST DATASET OF TARGET FEATURE, OUTPUTS AND ERRORS FOR MALE HISPANIC POPULATION USING ANN APPROACH AND GA-ANN APPROACH AND THEIR DENORMALIZED TEST SET SQUARE ERRORS

ANN							GA-ANN						
Normalized			Denormalized				Normalized			Denormalized			
Target	Output	Error	Target	Output	Error	Error^2	Target	Output	Error	Target	Output	Error	Error^2
0.1	0.201324	-0.10132	0	2.279791	-2.27979	5.1974449	0.1	0.183585	-0.08359	0	1.880664	-1.88066	3.5368987
0.188889	0.197235	-0.00835	2	2.187777	-0.18778	0.03526	0.188889	0.215019	-0.02613	2	2.587926	-0.58793	0.3456575
0.188889	0.233506	-0.04462	2	3.003875	-1.00387	1.0077643	0.188889	0.214525	-0.02564	2	2.576813	-0.57681	0.3327132
0.233333	0.279835	-0.0465	3	4.046295	-1.04629	1.0947323	0.233333	0.299638	-0.0663	3	4.491861	-1.49186	2.2256486
0.233333	0.211454	0.02188	3	2.507708	0.492292	0.242351	0.233333	0.219837	0.013496	3	2.696342	0.303658	0.0922084
0.277778	0.366258	-0.08848	4	5.990798	-1.9908	3.9632762	0.277778	0.321419	-0.04364	4	4.981924	-0.98192	0.9641748
0.277778	0.201166	0.076612	4	2.276225	1.723775	2.9714014	0.277778	0.208095	0.069683	4	2.432137	1.567863	2.4581941
0.322222	0.348731	-0.02651	5	5.596441	-0.59644	0.355742	0.322222	0.387335	-0.06511	5	6.465045	-1.46505	2.1463582
0.322222	0.431744	-0.10952	5	7.464246	-2.46425	6.0725074	0.322222	0.342664	-0.02044	5	5.459937	-0.45994	0.2115419
0.322222	0.332151	-0.00993	5	5.223394	-0.22339	0.049905	0.322222	0.369982	-0.04776	5	6.074604	-1.0746	1.154774
0.322222	0.347224	-0.025	5	5.56253	-0.56253	0.3164396	0.322222	0.373251	-0.05103	5	6.148144	-1.14814	1.3182336
0.366667	0.354582	0.012085	6	5.728094	0.271906	0.073933	0.366667	0.368918	-0.00225	6	6.050651	-0.05065	0.0025656
0.366667	0.359656	0.00701	6	5.842269	0.157731	0.024879	0.366667	0.394445	-0.02778	6	6.625014	-0.62501	0.3906426
0.411111	0.481509	-0.0704	7	8.583953	-1.58395	2.5089077	0.411111	0.450218	-0.03911	7	7.879906	-0.87991	0.7742341
0.411111	0.418609	-0.0075	7	7.168705	-0.16871	0.0284614	0.411111	0.382704	0.028408	7	6.36083	0.63917	0.4085383
0.411111	0.469562	-0.05845	7	8.315154	-1.31515	1.7296295	0.411111	0.45401	-0.0429	7	7.965232	-0.96523	0.9316724
0.411111	0.418849	-0.00774	7	7.174108	-0.17411	0.0303137	0.411111	0.412423	-0.00131	7	7.029521	-0.02952	0.0008715
0.455556	0.469473	-0.01392	8	8.313136	-0.31314	0.0980541	0.455556	0.504234	-0.04868	8	9.095267	-1.09527	1.1996099
0.455556	0.402567	0.052988	8	6.807768	1.192232	1.4214163	0.455556	0.491397	-0.03584	8	8.806426	-0.80643	0.6503225
0.455556	0.340921	0.114635	8	5.420717	2.579283	6.6527001	0.455556	0.415008	0.040548	8	7.087681	0.912319	0.8323259
0.455556	0.518221	-0.06267	8	9.409963	-1.40996	1.9879956	0.455556	0.493393	-0.03784	8	8.851337	-0.85134	0.724774
0.5	0.539811	-0.03981	9	9.895757	-0.89576	0.8023803	0.5	0.527366	-0.02737	9	9.615734	-0.61573	0.3791286
0.5	0.642738	-0.14274	9	12.21161	-3.21161	10.314416	0.5	0.576607	-0.07661	9	10.72366	-1.72366	2.9709889

0.5	0.624776	-0.12478	9	11.80746	-2.80746	7.8818342	0.5	0.608032	-0.10803	9	11.43072	-2.43072	5.9084008
0.544444	0.477199	0.067245	10	8.486977	1.513023	2.2892396	0.544444	0.475933	0.068512	10	8.458489	1.541511	2.3762555
0.544444	0.562167	-0.01772	10	10.39876	-0.39876	0.1590132	0.544444	0.575252	-0.03081	10	10.69318	-0.69318	0.480497
0.544444	0.61475	-0.07031	10	11.58187	-1.58187	2.5023213	0.544444	0.5905	-0.04606	10	11.03624	-1.03624	1.0737941
0.588889	0.535541	0.053348	11	9.799663	1.200337	1.4408096	0.588889	0.574432	0.014457	11	10.67472	0.325278	0.105806
0.588889	0.605067	-0.01618	11	11.36402	-0.36402	0.1325071	0.588889	0.629204	-0.04031	11	11.90709	-0.90709	0.8228051
0.633333	0.754195	-0.12086	12	14.71938	-2.71938	7.3950179	0.633333	0.751885	-0.11855	12	14.66741	-2.66741	7.1150809
0.633333	0.650181	-0.01685	12	12.37906	-0.37906	0.1436895	0.633333	0.627768	0.005566	12	11.87477	0.12523	0.0156827
0.633333	0.681152	-0.04782	12	13.07592	-1.07592	1.1576138	0.633333	0.671135	-0.0378	12	12.85053	-0.85053	0.7234027
0.633333	0.658568	-0.02523	12	12.56777	-0.56777	0.3223628	0.633333	0.678144	-0.04481	12	13.00823	-1.00823	1.0165321
0.633333	0.622084	0.01125	12	11.74689	0.253114	0.0640667	0.633333	0.624661	0.008672	12	11.80488	0.195123	0.0380731
0.633333	0.70065	-0.06732	12	13.51462	-1.51462	2.2940737	0.633333	0.70814	-0.07481	12	13.68316	-1.68316	2.8330133
0.677778	0.672703	0.005075	13	12.88582	0.114182	0.0130375	0.677778	0.668477	0.009301	13	12.79072	0.209278	0.0437974
0.677778	0.765351	-0.08757	13	14.9704	-1.9704	3.8824699	0.677778	0.772625	-0.09485	13	15.13407	-2.13407	4.5542406
0.677778	0.633847	0.043931	13	12.01156	0.988439	0.977012	0.677778	0.721206	-0.04343	13	13.97713	-0.97713	0.9547794
0.677778	0.726137	-0.04836	13	14.08809	-1.08809	1.1839369	0.677778	0.722753	-0.04498	13	14.01194	-1.01194	1.024021
0.677778	0.696694	-0.01892	13	13.42562	-0.42562	0.1811511	0.677778	0.636967	0.04081	13	12.08177	0.918231	0.8431488
0.677778	0.789076	-0.1113	13	15.50421	-2.50421	6.2710567	0.677778	0.729651	-0.05187	13	14.16714	-1.16714	1.3622203
0.677778	0.725624	-0.04785	13	14.07654	-1.07654	1.1589414	0.677778	0.696402	-0.01862	13	13.41905	-0.41905	0.1755997
0.722222	0.808195	-0.08597	14	15.93439	-1.93439	3.7418586	0.722222	0.77833	-0.05611	14	15.26242	-1.26242	1.5937016
0.722222	0.723004	-0.00078	14	14.01759	-0.01759	0.0003094	0.722222	0.738726	-0.0165	14	14.37134	-0.37134	0.1378948
0.722222	0.691355	0.030867	14	13.30549	0.694506	0.4823393	0.722222	0.697581	0.024642	14	13.44556	0.554436	0.3073996
0.766667	0.776064	-0.0094	15	15.21143	-0.21143	0.0447025	0.766667	0.764482	0.002185	15	14.95085	0.049152	0.0024159
0.766667	0.791099	-0.02443	15	15.54973	-0.54973	0.3022033	0.766667	0.784245	-0.01758	15	15.39551	-0.39551	0.1564279
0.811111	0.763783	0.047328	16	14.93511	1.064889	1.1339896	0.811111	0.763742	0.047369	16	14.93419	1.065813	1.1359574
0.811111	0.747164	0.063947	16	14.5612	1.4388	2.0701466	0.811111	0.740383	0.070728	16	14.40863	1.591374	2.5324713
0.855556	0.825701	0.029855	17	16.32826	0.671738	0.4512326	0.855556	0.787862	0.067693	17	15.4769	1.523102	2.3198399
0.855556	0.771538	0.084017	17	15.10961	1.89039	3.573573	0.855556	0.770061	0.085494	17	15.07638	1.923624	3.700331
0.855556	0.763251	0.092305	17	14.92314	2.076855	4.3133276	0.855556	0.779566	0.07599	17	15.29023	1.709767	2.9233016
0.855556	0.77824	0.077315	17	15.26041	1.739593	3.0261827	0.855556	0.780163	0.075393	17	15.30366	1.696342	2.8775757
0.9	0.755632	0.144368	18	14.75171	3.248289	10.551383	0.9	0.738611	0.161389	18	14.36874	3.631262	13.186061
0.9	0.714103	0.185897	18	13.81731	4.182687	17.494871	0.9	0.78023	0.11977	18	15.30516	2.694836	7.2621388

APPENDIX S

GA PYTHON SCRIPT FOR FEMALE FEATURE SELECTION



```
1 import random
2 import numpy as np
3 import matplotlib.pyplot as plt
4 import pandas as pd
5 import csv
6
7 from os.path import join, dirname
8 from sklearn.utils import Bunch
9 from sklearn.model_selection import cross_val_score
10 from sklearn.linear_model import LinearRegression
11
12 SEED = 2018
13 random.seed(SEED)
14 np.random.seed(SEED)
15
16 #=====
17 # Data
18 #=====
19 module_path = dirname(__file__)
20 data_file_name = join(module_path, 'HISF.csv')
21 with open(data_file_name) as f:
22     data_file = csv.reader(f)
23     temp = next(data_file)
24     n_samples = int(temp[0])
25     n_features = int(temp[1])
26     data = np.empty((n_samples, n_features))
27     target = np.empty((n_samples,))
28     temp = next(data_file) # names of features
29     feature_names = np.array(temp)
30
31     for i, d in enumerate(data_file):
32         data[i] = np.asarray(d[:-1], dtype=np.float64)
33         target[i] = np.asarray(d[-1], dtype=np.float64)
34
35 dataset = Bunch(data=data,
```

```

36         target=target,
37         # last column is target value
38         feature_names=feature_names[:-1],
39         filename=data_file_name)
40
41 X, y = dataset.data, dataset.target
42 features = dataset.feature_names
43 #features = ["Bone1" "Bone2" "Bone3" "Bone4" "Bone5" "Bone6" "Bone7" "Bone8" "Bone9" "Bone10" "Bone11" "Bone12" "Bone13" "Bone14" "Bone15"
44 #features = ["Bone16" "Bone17" "Bone18" "Bone19"]
45 #features = [Bone1, Bone2, Bone3, Bone4, Bone5, Bone6, Bone7, Bone8, Bone9, Bone10, Bone11, Bone12, Bone13, Bone14, Bone15, Bone16, Bone17,
46 #features = [Bone18, Bone19]
47
48 #=====
49 # CV MSE before feature selection
50 #=====
51 est = LinearRegression()
52 score = -1.0 * cross_val_score(est, X, y , cv=5, scoring="neg_mean_squared_error")
53 print("CV MSE before feature selection: {:.8f}".format(np.mean(score)))
54
55 #=====
56 # Class performing feature selection with genetic algorithm
57 #=====
58 class GeneticSelector():
59     def __init__(self, estimator, n_gen, size, n_best, n_rand,
60                 n_children, mutation_rate):
61         # Estimator
62         self.estimator = estimator
63         # Number of generations
64         self.n_gen = n_gen
65         # Number of chromosomes in population
66         self.size = size
67         # Number of best chromosomes to select

```



```

67     self.n_best = n_best
68     # Number of random chromosomes to select
69     self.n_rand = n_rand
70     # Number of children created during crossover
71     self.n_children = n_children
72     # Probability of chromosome mutation
73     self.mutation_rate = mutation_rate
74
75     if int((self.n_best + self.n_rand) / 2) * self.n_children != self.size:
76         raise ValueError("The population size is not stable.")
77
78     def initialize(self):
79         population = []
80         for i in range(self.size):
81             chromosome = np.ones(self.n_features, dtype=np.bool)
82             mask = np.random.rand(len(chromosome)) < 0.3
83             chromosome[mask] = False
84             population.append(chromosome)
85         return population
86
87     def fitness(self, population):
88         X, y = self.dataset
89         scores = []
90         for chromosome in population:
91             score = -1.0 * np.mean(cross_val_score(self.estimator, X[:,chromosome], y,
92                                                     cv=5,
93                                                     scoring="neg_mean_squared_error"))
94             scores.append(score)
95         scores, population = np.array(scores), np.array(population)
96         inds = np.argsort(scores)
97         return list(scores[inds]), list(population[inds,:])
98
99     def select(self, population_sorted):

```

```

100     population_next = []
101     for i in range(self.n_best):
102         population_next.append(population_sorted[i])
103     for i in range(self.n_rand):
104         population_next.append(random.choice(population_sorted))
105     random.shuffle(population_next)
106     return population_next
107
108     def crossover(self, population):
109         population_next = []
110         for i in range(int(len(population)/2)):
111             for j in range(self.n_children):
112                 chromosome1, chromosome2 = population[i], population[len(population)-1-i]
113                 child = chromosome1
114                 mask = np.random.rand(len(child)) > 0.5
115                 child[mask] = chromosome2[mask]
116                 population_next.append(child)
117         return population_next
118
119     def mutate(self, population):
120         population_next = []
121         for i in range(len(population)):
122             chromosome = population[i]
123             if random.random() < self.mutation_rate:
124                 mask = np.random.rand(len(chromosome)) < 0.05
125                 chromosome[mask] = False
126                 population_next.append(chromosome)
127         return population_next
128
129     def generate(self, population):
130         # Selection, crossover and mutation
131         scores_sorted, population_sorted = self.fitness(population)
132         population = self.select(population_sorted)

```

```

133     population = self.crossover(population)
134     population = self.mutate(population)
135     # History
136     self.chromosomes_best.append(population_sorted[0])
137     self.scores_best.append(scores_sorted[0])
138     self.scores_avg.append(np.mean(scores_sorted))
139
140     return population
141
142     def fit(self, X, y):
143
144         self.chromosomes_best = []
145         self.scores_best, self.scores_avg = [], []
146
147         self.dataset = X, y
148         self.n_features = X.shape[1]
149         self.population = []
150         population = self.initialize()
151         for i in range(self.n_gen):
152             population = self.generate(population)
153
154         self.population = population
155         return self
156
157     @property
158     def support_(self):
159         return self.chromosomes_best[-1]
160
161     def plot_scores(self):
162         plt.plot(self.scores_best, Label='Best')
163         plt.plot(self.scores_avg, Label='Average')
164         plt.legend()
165         plt.ylabel('Scores')

```

```

166         plt.xlabel('Generation')
167         plt.show()
168
169     sel = GeneticSelector(estimator=LinearRegression(),
170                           n_gen=200, size=200, n_best=40, n_rand=40,
171                           n_children=5, mutation_rate=0.05)
172     sel.fit(X, y)
173     sel.plot_scores()
174     score = -1.0 * cross_val_score(est, X[:,sel.support_], y, cv=5, scoring="neg_mean_squared_error")
175     print("CV MSE after feature selection: {:.8f}".format(np.mean(score)))
176     #print(sel.population)
177     np.savetxt('GApopulationFemale.csv', sel.population, fmt='%.2f', delimiter=',', header=" Bone1, Bone2, Bone3, Bone4, Bone5, Bone6, Bone7,
        Bone8, Bone9, Bone10, Bone11, Bone12, Bone13, Bone14, Bone15, Bone16, Bone17, Bone18, Bone19")

```

APPENDIX T

GA PYTHON SCRIPT FOR MALE FEATURE SELECTION

```
GAmale.py  x  GAFemale.py  x
1 import random
2 import numpy as np
3 import matplotlib.pyplot as plt
4 import pandas as pd
5 import csv
6
7 from os.path import join, dirname
8 from sklearn.utils import Bunch
9 from sklearn.model_selection import cross_val_score
10 from sklearn.linear_model import LinearRegression
11
12 SEED = 2018
13 random.seed(SEED)
14 np.random.seed(SEED)
15
16 #=====
17 # Data
18 #=====
19 module_path = dirname(__file__)
20 data_file_name = join(module_path, 'HISM.csv')
21 with open(data_file_name) as f:
22     data_file = csv.reader(f)
23     temp = next(data_file)
24     n_samples = int(temp[0])
25     n_features = int(temp[1])
26     data = np.empty((n_samples, n_features))
27     target = np.empty((n_samples,))
28     temp = next(data_file) # names of features
29     feature_names = np.array(temp)
30
31     for i, d in enumerate(data_file):
32         data[i] = np.asarray(d[:-1], dtype=np.float64)
33         target[i] = np.asarray(d[-1], dtype=np.float64)
34
35 dataset = Bunch(data=data,
```

```

36         target=target,
37         # last column is target value
38         feature_names=feature_names[:-1],
39         filename=data_file_name)
40
41 X, y = dataset.data, dataset.target
42 features = dataset.feature_names
43 #features = ["Bone1" "Bone2" "Bone3" "Bone4" "Bone5" "Bone6" "Bone7" "Bone8" "Bone9" "Bone10" "Bone11" "Bone12" "Bone13" "Bone14" "Bone15"
44 #features = ["Bone16" "Bone17" "Bone18" "Bone19"]
45 #features = [Bone1, Bone2, Bone3, Bone4, Bone5, Bone6, Bone7, Bone8, Bone9, Bone10, Bone11, Bone12, Bone13, Bone14, Bone15, Bone16, Bone17,
46 #features = [Bone18, Bone19]
47
48 #=====
49 # CV MSE before feature selection
50 #=====
51 est = LinearRegression()
52 score = -1.0 * cross_val_score(est, X, y , cv=5, scoring="neg_mean_squared_error")
53 print("CV MSE before feature selection: {:.8f}".format(np.mean(score)))
54
55 #=====
56 # Class performing feature selection with genetic algorithm
57 #=====
58 class GeneticSelector():
59     def __init__(self, estimator, n_gen, size, n_best, n_rand,
60                 n_children, mutation_rate):
61         # Estimator
62         self.estimator = estimator
63         # Number of generations
64         self.n_gen = n_gen
65         # Number of chromosomes in population
66         self.size = size
67         # Number of best chromosomes to select

```

```

67     self.n_best = n_best
68     # Number of random chromosomes to select
69     self.n_rand = n_rand
70     # Number of children created during crossover
71     self.n_children = n_children
72     # Probability of chromosome mutation
73     self.mutation_rate = mutation_rate
74
75     if int((self.n_best + self.n_rand) / 2) * self.n_children != self.size:
76         raise ValueError("The population size is not stable.")
77
78     def initialize(self):
79         population = []
80         for i in range(self.size):
81             chromosome = np.ones(self.n_features, dtype=np.bool)
82             mask = np.random.rand(len(chromosome)) < 0.3
83             chromosome[mask] = False
84             population.append(chromosome)
85         return population
86
87     def fitness(self, population):
88         X, y = self.dataset
89         scores = []
90         for chromosome in population:
91             score = -1.0 * np.mean(cross_val_score(self.estimator, X[:,chromosome], y,
92                                                     cv=5,
93                                                     scoring="neg_mean_squared_error"))
94             scores.append(score)
95         scores, population = np.array(scores), np.array(population)
96         inds = np.argsort(scores)
97         return list(scores[inds]), list(population[inds,:])
98
99     def select(self, population_sorted):

```

```

100     population_next = []
101     for i in range(self.n_best):
102         population_next.append(population_sorted[i])
103     for i in range(self.n_rand):
104         population_next.append(random.choice(population_sorted))
105     random.shuffle(population_next)
106     return population_next
107
108     def crossover(self, population):
109         population_next = []
110         for i in range(int(len(population)/2)):
111             for j in range(self.n_children):
112                 chromosome1, chromosome2 = population[i], population[len(population)-1-i]
113                 child = chromosome1
114                 mask = np.random.rand(len(child)) > 0.5
115                 child[mask] = chromosome2[mask]
116                 population_next.append(child)
117         return population_next
118
119     def mutate(self, population):
120         population_next = []
121         for i in range(len(population)):
122             chromosome = population[i]
123             if random.random() < self.mutation_rate:
124                 mask = np.random.rand(len(chromosome)) < 0.05
125                 chromosome[mask] = False
126             population_next.append(chromosome)
127         return population_next
128
129     def generate(self, population):
130         # Selection, crossover and mutation
131         scores_sorted, population_sorted = self.fitness(population)
132         population = self.select(population_sorted)

```



```

133     population = self.crossover(population)
134     population = self.mutate(population)
135     # History
136     self.chromosomes_best.append(population_sorted[0])
137     self.scores_best.append(scores_sorted[0])
138     self.scores_avg.append(np.mean(scores_sorted))
139
140     return population
141
142     def fit(self, X, y):
143
144         self.chromosomes_best = []
145         self.scores_best, self.scores_avg = [], []
146
147         self.dataset = X, y
148         self.n_features = X.shape[1]
149         self.population = []
150         population = self.initialize()
151         for i in range(self.n_gen):
152             population = self.generate(population)
153
154         self.population = population
155         return self
156
157     @property
158     def support_(self):
159         return self.chromosomes_best[-1]
160
161     def plot_scores(self):
162         plt.plot(self.scores_best, label='Best')
163         plt.plot(self.scores_avg, label='Average')
164         plt.legend()
165         plt.ylabel('Scores')

```

```

166         plt.xlabel('Generation')
167         plt.show()
168
169     sel = GeneticSelector(estimator=LinearRegression(),
170                           n_gen=200, size=200, n_best=40, n_rand=40,
171                           n_children=5, mutation_rate=0.05)
172     sel.fit(X, y)
173     sel.plot_scores()
174     score = -1.0 * cross_val_score(est, X[:,sel.support_], y, cv=5, scoring="neg_mean_squared_error")
175     print("CV MSE after feature selection: {:.8f}".format(np.mean(score)))
176     #print(sel.population)
177     np.savetxt('GApopulationMale.csv', sel.population, fmt='%.2f', delimiter=',', header=" Bone1, Bone2, Bone3, Bone4, Bone5, Bone6, Bone7,
        Bone8, Bone9, Bone10, Bone11, Bone12, Bone13, Bone14, Bone15, Bone16, Bone17, Bone18, Bone19")

```