# A review on prediction of diabetes type 2 by machine learning techniques

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Article History: Received: 10 November 2020; Revised 12 January 2021 Accepted: 27 January 2021; Published online: 5 April 2021

**Abstract:** Machine learning is considered to be one of the most promising tools when it comes to working with heterogeneous data. It provides a new dimension which enables one to extract relevant data and take decision for the effective functioning of the network, making use of network generated data. Every sphere of our life is now dependent on machine learning. It has flourished in every dimension. Making it versatile and ever demanding.

Department of healthcare contains very abundant and sensitive information which is needed to be carefully handled. Diabetes mellitus is increasing exponentially and is spreading like anything in the world. A reliable prediction system should be present for diagnosing diabetes. Variety of machine learning techniques find their use in the examination of data from variant perspectives and summarizing it into effective information. Usage of new patterns is done to elucidate these patterns in order to deliver relevant information for their users. By making use of techniques such as SVM, random forest, logistic regression, naïve bayes etc the prediction of diabetes can be done easily and accurately. In this study we will make use of different machine learning techniques and try to find accurate prediction regarding the same.

Keywords: Machine learning, diabetes type 2, supervised, unsupervised, reinforcement, training and algorithm

### 1. Introduction

Machinelearninghasthepotentialwhichenablesitto learn from previous data to generate futuristictrends in behavior. It has the capability to learn by the own. Machine learning can be applied onnumerous data making it very integral to thetelecommunication world today (Hang Lai et al2019) [1]. Machine learning methods detectlinearities/non linearitiesintherelationship

between dependent and independent variables (Geofrrey et al 2019) [2]. They can be used formaking predictions in case of continuous outcomes, known as regression type problems or can be used for making predictions in case of levels of categorical variable, which is known asclassification problems. It gives solution from the problems and learnshow to tack lewith the problem that may or may not be same by making

useoftrainingdatasetprovidedtothealgorithmearlier.

Diabetes is such a prolonged disease that canhappen when body cannot efficiently make use of the insulin it generates. As a result, diabetes affectsorgans which include heart diseases which could beheart stroke, high blood pressure and the rosclerosis, nerve damage that could lead tonumbness, gradually losing all sense of feelingespecially in the limbs, kidney failure is very common in diabetic patients, and hearing impairment is also seen in diabetic patients, the risk of Alzheimer's disease increases with type 2 diabetes.

Diabetescan becategorized into threetypes:-

- (a) ChildhoodorJuvenilediabetes
- (b) AdultorType2diabetes
- (c) Type3orGestationaldiabetes

Generally, type 1 diabetes occur because of the deficiency in insulin production and is commonly found in children. Diabetes type 2 is a chronic disease which affects how the human body metabolizes glucose. In case of diabetes type 2, the human body behaves in either of the 2 ways; firstly it resists the effect of insulin which is a hormoneresponsible for regulating the movement of sugarinthe cells. Secondly it doesn't produce ample insulin for the maintenance of normal glucose level.

Diabetes type 2 was known to be adult onsetdisease but nowadays much of the younger agegroupisbeingdiagnosed with the same, because of the rise in observing the result of the same because of the result of the result of the same because of the result of

available for the same but person can switch fromsedentary life style, follow balanced diet and canexercise well to manage the disease, as depicted infigure1. If this would not suffice then the personshould go for medications and insulin therapy. Theinsulin is secreted into the bloodstream by thepancreas. This insulin then circulates, enabling thesugartoenter thebodycells. Theamountofglucose in the bloodstream is lowered by theinsulin. Glucose i.e. sugar, is a major source of energy for cells that make up muscles and othertissues and it comes from food and liver. In case of lower glucose level the liver breaks down glycogeninto glucose in order to keep the glucose levelnormal. When it comes to type 2 diabetes, the sugarstarts to build up in the bloodstream instead of moving into the cells which lead to more release of insulinbybetacells inthepancreas, graduallythese cells become impaired and become incapable freleasing more insulin to fulfill the requirement body whereas in case of type 1 diabetes theimmune system by mistake destroys beta cellswhich leavethebody withlittleor no insulin.

Gestational diabetes is hyperglycemia whichhappens due to the change in hormones duringpregnancy.

Fot the past few decades we have seen that themachine learning discipline is assisting us to solvedifferent relevant biomedical problems. Themachinelearningtechniquesarefoundtooperateinboth real-life and scientific problems. In this study, we will be evaluating the performance of variousmachinelearningtechniquesfortheclassificationofpeople whether they are diabetic ornot.

## **1.2.Generalized** Architecture



### Figure1

## 2.Literature survey

Thissectionreviewsvariousresearch worksthatarerelatedtoourproposed work.

Arianna Dagliati et al [3] designed a machinelearningpredictionmodelfortype2diabeteswheredemographic data (i.e. age, gender, time todiagnosis), clinical data from the EHR (BMI,Hba1c,lipid

profile,smokinghabit),Administrative data (antihypertensive therapy) arecollected based on that predictive model formicrovascular complications in the population wasdesigned which focused on issues such asnephropathy, neuropathy and retinopathy. Themodel showed higher AUC values in case of SVMand Random Forest. The final model is based onlogistic regression with rebalanced classes whichsupportsnomograms.

D. Asir Antony Gnana Singh et al [4] designed amachinelearningmodelforpredictionofdiabetes. Three different type of supervised learning algorithm namely probabilistic based naïve bayes (NB), function based multilayer perceptron

(MLP), decision tree based random forests(RF) areused. Test methods such as 10 fold cross validation(FCV), makes use of percentage split (PS) with66% and training dataset(UTD). The preprocessing technique is used to increase the accuracy of the model. In case of pre-processing technique average accuracy for NB is increased as compared tomachinelearning algorithm.

K. Srinivas et al [5] developed data miningapplication techniques that can be used in case ofhealth care and prediction of heart attacks. In theresearch they made use of medical profiles such asblood pressure, age, blood sugar and sex and used this to predict the likeliness of getting kindneyproblemsandheartattack.

IdemudiaChristianUwaet el[6]designedamachinelearningpredictionmodelforpredictionof diabetes. On applying univariate selectionmethod with chi squared statistical test in case ofnon negative feature we obtain following attributeslikeplasma,bloodpressure,age,pedigreefunction,

B.M.I. Here the algorithms that were been applied arenaïve bayes, logistic regression, SVM, XG

Boost, KNN. The dataset were of 2 types one frompimaIndiandatasetandtheotherwasdr.schorlingdataset derived from hospital. We found that onboth the dataset, naïve bayes model showedconsistency and after naïve bayes logisticregression proved to be better wih accuracies of 83% and 81% respectively.

V. Ranjani et al [7] emphasized on the potentialuse of classification based data mining techniquesthat incluses aritifical neural network (ANN), rule-based methods, Naïve Bayes and decision treealgorithm to huge volume of data of health care. Intheresearch, medcalproblemshavebeen analysed and evaluated which include blood pressure and heart disease.

M.Durairaj et el [8] demonstrates a hybridprediction system consisting of Rough Set Theoryand Artificial Neural Network for depictingmedicaldata.Thisprocessof developmentofanew data mining technique and a software to helpcompetent answers in case of analysis of medicaldata is been explained. A hybrid tool is beenproposed that incorporates RST and ANN to makeefficient data analysis and indicative predictions.The experiments' on spermatological data set thatisbeenusedforthepredictingexcellenceofanimalsemen. The hybrid prediction system is beenapplied in case of pre-processing medical databaseand for the purpose of training the ANN for theprediction of production. The accuracy in case ofprediction is obtained in case of comparison that isbeen made between the observed and predictedcleavagerate.

S.M Hasan Mahmud et al [30] designed a machinelearningmodelfortheprediction ofdiabeteswhere

the comparison is been based on the performance valuation by 10-fold validation technique. Aframework is also been generated for diabetesprediction, monitoring and application (DPMA). Here the basic concept is that multiple machine learning classifiers are supposed to perform better than a single machine learning classifier.

Akm Ashiquzzaman, AbdulKawsar Tushar etal

designed a diabetes prediction model bymaking use of the application of the drop outmethod.Novelformofdeepneuralnetworkfortheprognosis of diabetes with increased accuracy isbeen discussed.

Ioannis Kavakiotis et al [32] designed machinelearning and data mining approaches that wereappliedonalltheaspectsofDMresearchand thatwere applied on biomarker identification and prediction diagnosis.

Muhammad Azeem Sarwar, Nasir Kamal et al [33]designed a model for the prediction of diabetes bymaking use of various machine learning learningtechniques where the data is divided into trainingdata and testing data. Enthought Canaopy methodand thus the result is been obtained. SVM andKNNhaveshown higher accuracy in themodel.

# 2.1.Literature review:-

S.No.	AUTHOR	DESCRIPTION/WORK	TITLE	METHODS/TOOLS	RESEARCHGAP
1.	mone Marini,Lucia Sacchi,GiuliaCogn i	-		LogisticRegression	Development ofpredictivemodel for theonset ofmicrovascularco mplicationsin case ofT2DM.
2.	Dr. E. Jebamalarleavline,	learningalgorithm is used for the the diagnosis and	usingmedicaldata	basedmultilayer perceptron(MLP), decision treebased randomforests(RF)	Thepreprocessingi ncreasedtheaccura cy of allthe models thatinclude naïvebayes , MLP,RF except the10 fold cross validationmethod.
3.	B. KavihtaRani, Dr.A. Govrdhan	ofsignificantpatternsisbeen established, on thebasis of calculated weight,the value	ofdata miningtechniques inhealthcare	DecisionTree	The role of textmining can beused inexpanded so towiden its rolein- case ofunstructureddata
4.	Uwa,NehikhareEfe hi	Data mining processesthat are been used in caseof medical diagnosis andthe usage of variousmachine learningtechniques for predictingdiabetes.	theperformance ofmachinelearning	regression,naïve bayes, supportvector machine,XGBoost,kNN	finetuningtechniq
5.	V.Ranjani	thanone data mining techniquefor diabetes	miningapplications inhealthcaresector:	Artificialneural network, ANNand HybridTechnique	Hybridtechniques when appliedfor variousdiseases canyield betteraccuracy.

r					
6.	K.Meena	ANNand RST is used as aalternativetotheconventiona	hybridpredictionsy stem usingrough	Theory(RST), ArtificialNeural Network(ANN)	Incorporationof biologicalinformat ion,systematiccom parison ofdifferentmachin elearningalgorith ms,hybridization ofrough sets andneural networkensembles tobuild predictorsfor improvingperform ance.
7.	,Md AltabHossin,Md. RazuAhmed	real time diabetesprediction, monitoringand application.	work fordiabetespredicti on.	foldvalidationtechnique along withnaïve bayes, ANN,Logistic Regression,Decision TreeRandom Forest andSVM	
8.	n,Abdul KawsarTushar,	dropoutmethod.	fitting inDiabetesPredicti	deepneural network withthe application ofdropoutmethod	Performance incase ofpredictivemodel s ofdiabetes canhave betterpredictionsc ores whichwill pave wayforbreakthrou gh inhealthprognostic ation.

9.	Olga Tsave,Athanasios, NicosMaglaveras,I oannisVlahavas,Io annaChouvarda	applications of machine learning, datamining techniques andtools with respect todiagnosis, prediction,genetic background andenvironment, and healthcareand managementisbeen done based ondifferentmachinelearningt	anddata miningmethods indiabetesresearch.	regression,SVM,ANN, NB, Linear DiscriminantAnalysis, KNN,fuzzy c- mean,Random forest,CART, MultifactorDimensional ityReduction.	With the adventof bio-technology, andwith the hugeamount of dataproduced, andwith theeverincreasinga mount ofEHRs thediagnosis, andtreatment ofdiseases can beenriched.
10.	MuhammadAzeem ,Sarwar,Nasir Kamal,Wajeeha Hamid,MunamAli Shah	machine learningalgorithmsareapplie	ofDiabetes UsingMachineLear	Bayes,KNN,SVM,LR,D T, andRandomForest	Theadvancementc an be made interms ofapplyingvarious techniques suchas big data,cloudcomputi ng withmachinelearni ngtools.

# 3 Machine Learning Algorithms:-

The significance of machine learning algorithmsdependsinthedevelopmentofmodelsthatisbasedontheexistingdataandconsequently,classification or prediction by making use of noveldata. Machine learning methods have been widelyusedin variousapplicationsindiversifieddomains

like system biology, genomics. Specificallyspeaking, supervised machine learning techniqueshave been finding immense importance in anumber of bioinformatics prediction techniques. The aim here is to showcase an overview of themachinelearningalgorithmsaswellasapplicationmethodsbasedonsame.



Figure 2: Types of machine learning algorithms:-

Machine learning techniques can be broadlycategorized as:-

Supervised learningUnsupervised learningReinforcementlearning

# 2..3. Supervised learning:

Supervised learning has the involvement of supervisor which works in the same way as ateacher in real life. It

is such a type of learning inwhichweteachortrainamachinebymaking useof data which has already been tagged with the correctanswer (PaulAkangah et el,2018) [9].

Further, we experiment the machine with new setsof data so that the supervised learning algorithmcananalyzethetrainingdataand cangiveacorrectoutcome on the basis of the previous labeled data(R.Sathya et al 2013) [10].

Supervised learning is classified into twocategories:

• **Regression:**IFtheoutputvariableincludesa real value, for example "dollars" or "weight" then we callitategression

problem. We see use of regression algorithmwhile dealing with decision trees, linearregression, logistic regression etc (GiovanniGrano et el, 2018) [11]

• **Classification** : IF the output variable includes a category, for example "Red" or "blue" or "disease" and "no disease" we callit a classification. We see use of classification algorithm while dealing withnaive bayes classifier, support vector machine, K- Nearest Neighbor (Kondi Srujan Kumarret el, 2019)[12]

## 3.1.1.Artificial Neural Network:-

This algorithm is conceptualized on basis ofbiological neurons. We can see that in case ofbiological learning process the process of learningisthoughttobebasedon minoradjustmentstothesynaptic connections between neurons whereas inANN the learning process is totally based on the interconnections between the processing elementswhich combineto formnetwork topology.

Basically, ANN consists of 3 layers i.e. input layer, hidden layer, and theoutputlayer. We see that

incase of ANN, the training of hidden layercontainingnetworkand makesuseofitsconnectedstructures for the purpose of pattern recognitionand classification. In case of bioinformaticsapplications of ANN, we employ different types of architectures with perceptron and multi layered perceptron being the simplest inthe category.

Radial basis function networks and Kohonen selforganizing mapsarealsofound useful.

ThemajorstepswhichareinvolvedinanANNalgorithm areasfollows:

- Byprocessingavailabledatatrainingandtestdatasetsare generated.
- Data is encoded into digital format bymakinguseofencodingsystems, such as binary systems.

• ANN architecture is designed anddevelopedbymakinguseof3layersforthepurpose of prediction.

• Ann is trained by making use of appropriate input data and parameters.

- ANN model is such selected which gives the validout put.
- Ann model is thus validated by using testdataset for the purpose of estimation of efficacy for prediction.

The biggest advantage we observe in case of ANNis its ability to analyze and process over largecomplex datasets, having non-linear relationships. This model includes more benefits like having theability to handle noisy data and the caliber of generalization. The limitation of the methodobserved is in the amount of time that would betakenincase of processing complex datasets. ANNhas extensively been used in case of generalization, sequence feature analysisetc.

### 3.1.2. Support Vector Machine

Support Vector Machine is a supervised learningmethod that is based on statistical learning theory. For linearly separable illustrations, SVM creates amaximum margin hyper-plane that separates thedata points into 2 different classes. The hyper-planeworks as a decision surface between two classes(Affsan Abbrar et al, 2018) [22]. In case of non-linearly separable data, firstly SVM changes datainto higher dimensional feature space and consequently makes use of a linear maximummarginhyper-plane. Thisleadsto the introduction

of computational intractability that requires atransformation to a higher dimensional space (An

T. Nguyen et al, 2018) [23]. SVM resolves this bydefining most appropriate kernel functions by thehelp of which the computations can be taken intoconsideration in the original space itself. The threepopular kernel functions that are used generally arelinear, polynomial and radial basis function (SandraVieira et al, 2019) [24]. In case of bioinformatics, we see many domain specific kernel functions suchasgraph

#### kernel,stringkernel(JesseH.Krijtheet

al, 2017)[25].Thisconcept canalsobe usedincase of multiclass classification. The two mostcommon multiclass classification methods that findtheirusehereareviz.,oneagainalland oneagainstone (Konstantinos Sechidis et al, 2017) [26]. Thesteps that are employed in SVM algorithm are given below:

• Feature vector is constructed in-order torepresent positive and negative dataset: this feature vector contains properties of the input data that could be amino acid, physio chemical properties etc.

• Appropriate kernel function is chosen soasto fitforthepredictiontaskby makinguseof classifiertraining.

The model is selected with bestperformancetomakepredictions.

• Theapplication of chosen model for doing predictions on the unknown input data set, the most robust classifier is SVM, it has the best generalization ability incase of unseen data in comparison to other methods.

SVM is the most commonly used machinelearning method that is used in case of computational biology and bioinformatics. It is also been used for secondary structureprediction, gene finding, fold recognitionas wellas binding site prediction.

Support vector machine is a distinguishingclassifier which is previously defined bysecludinghyperplanewhichmeans, on the given labeled training data, here supervised learning, the algorithm gives output in the form of a hyper-lane which will categories new examples. The hyper-lane is a linewhich divides a plane into two parts, incase of the two dimensional space where each of the classifier other states.



Figure3

Figure 2, in the above example diagram "b" shows that a line in this case separates the two different classes asdepicted in example "a". Hereweuse the equation of lineary=x.wemay also use the following y=mx+c.



### **Figure4**

Figure 3, in the above example we see different property of SVM where we are making use of  $z^2 = x^2 + y^2$ , here by making use of square values we can separate "c" as "d.

### 3.1.3.K-Nearest Neighbor

KNN classifier finds the k nearest examples in thereference set, and considering majority vote

from the classes of these kexamples to allocate a class to a query (P.Yasodhaet al 2014)[18]. Assignment of classes in decision boundaries are implicitly derived in case of KNN. Below are the relevant steps that are involved in the development of KNN classifier:

- Featuresetisconstructed and a distancemetric is used to compute distances between features.
- Numberofnearestneighborsisdeterminedforthetrainingset.

• Euclidian distances or any other distancemeasure such as Mahalanobis distance arecalculated between the query instance andthetrainingsamples.

• Distances are sorted and nearest neighborsare determined on the basis of the k-thminimum distance.

• Class label are predicted in case of new orunknown instance by making use of the classlabel of nearest neighbors.

The most significant advantage of KNN method is that it has higher efficiency on large datasets and robustness while processing noisy data (GopiBattineni et al, 2019) [19]. The drawback of KNN is thigh computation cost, which deduces its

speed. In case of bioinformatics, we observe thatKNN model is been employed successfully (Yun-leiCai et al, 2010) [20].

# 3.1.4Decision Tree

Decision tree are considered to be a branch testbased classifier. The construction of the sameinvolves the analysis of the set of trainingexamples, class labels are known for them. Newand unseen examples are classified by this information. A leaf node symbolizes a specific class and every branch represents a group of classes (Mikolas Janota et al, 2018) [21]. A test ona single attribute value is been represented by the decision node, with its one main branch and the subsequent classes are represented as possible outcomes (Sullivan hue et al, 2018) [29]. Themajor steps that are to be considered in case of decision tree algorithmis given below:-

• Training dataset is prepared in such an appropriate form in case of the classifier by the method of feature extraction from input data.

• Decisiontreeisconstructedbyputtingtheinstancesintrainingsetattheinitialnode.

• The instances are divided into twodistinguishable classes i.e. child nodesbased on their chosen testvalue.

• By the recursive application of the laststep it is checked that the fulfillment oftermination or pre- pruning condition ismet.

• The resultant tree is pruned with itsapplications for performing predictions.Decisiontreesaresimpleclassifiers and hence have better interpretability ascompared to other machine learning methods. They are widely used inbioinformatics for predicting genetic interactions and related applications.

# 3.1.5.Random Forests

Random Forests is a group of randomly created independent classifiers and decision trees (PaulAkangah et al, 2018) [9]. It generally depicts ubstantial performance improvisations over single tree classifiers such as C4.5, CART. Randomnessor Feasibility can be introduced in the RFalgorithm introduces:

1. .1.Bootstrapping, abootstrapset is created from the original training data set by making use of random sampling by doing replacement to generate each tree (Marcus Muller et al, 2018)[28].

1.2. Construction of bootstrapset is done by making use of original training dataset by thehelp of random sampling by the process of replacement in order of generate each tree.

2. Node Splitting: Here the selection of subset of attributes is carried out. On splitting a node, where there are M input attributes, then the number'm', where m<<M and is been specified in such a way that at eachnode, m attributes are randomly selected and the best split is considered on them. A value that is good of 'm' is by default selected by making use of various implementations, considering 'm' as sqrt (M) for the very purpose of classification. On the basis of the CART algorithm the classification tree is induced by making use of 'inbag' data. After that an out of bagdata, that is been formed after leaving out the in-bag samples from those of the original data is used in crossvalidation work. The steps involved in case of random for estal gorithm are given below:

• CART algorithm is been employed on data for the growth of random classification trees.

- Bootstrap data is beenused which is known asin-bag set that is used totraintheCARTalgorithm.
- On the basis of the bestcondition on a randomsubset of 'm' attributesnodesplittingisdone.

• By making use of majorityvote strategy in order todecideclassaffiliationincaseofeach OOBsample.

• Variable importance (VI)ranking,thatcanbeusedlater to retrain random forestby using a smaller subset of themostrelevant variables.

• Resistance to over fitting ofdata random forest and itsvariants are been applied tosolve a huge amount ofbioinformatics problemswhichincludesclassification of geneexpression, analysis of massspectroscopy data fordiabetes prediction, sequence annotation and prediction of diabetes 2 mellitus.

### **3.1.6.Ensemble Classifiers**

In case of ensemble classifiers, the individual decisions in case of asset of classifiers are joined with weighted or unweighted voting for the purpose of classification of new instances.

Ensemble classifiers are also called as multi-classifier systems. These classifiers are found to beefficient in prediction tasks because of the fact thatthey find use of a combined classifier and cancapture features that cannot even be captured bymaking use of any single model alone. These methods are been applied in differentbioinformatics problems because of their highprediction accuracy.

## **3.1.7.Unsupervised learning:**

Unsupervised learning is that type of training inmachine where we make use of information that isneither labeled nor is classified and so it lets thealgorithm to work on this information without anyprior guidance as in case of supervised learning(Nagdev Amruthnath et al, 2018) [13]. The task of the machine here is to group unsorted informationinto patterns or on the basis of differences and similarities without the prior training being done on the data (Memoona Khanametal, 2015)[14].

Unsupervisedlearningisclassified into two categories:

• Association:Dimensionalityreductionisthe other name of association rule learningproblem.Anassociationlearningproblemisone where one needs to find rules that couldbe applied to large data sets that may include for example people who wish to buy A and arealso intended to buyB.

• **Clustering**: A clustering problem is onewhere we want to find the inherentgroupingswithinthedata, which includes grouping various customers by their purchasing behavior (Oyelade et al 2010)[15].

Wefinduseofclusteringalgorithmwhiledealingwith K-means;meanshift,K-medoids.

# 3.1.8.Artificial Hidden Markov Models (HMM)

Hidden Markov Models have found their use invery popular machine learning approaches such asin case of bioinformatics. They are probabilistic models that are generally implied in timeseries and linear sequences. It can be used to describe the evolution of those events which are observable and these depend on internal factors, which themselves are not observable. Here we see that the observed

events arecalledassymbol andtheinvisible factors that are underlying the observations that are referred to as a state. An HMM comprises of several states, that are connected by means of transition probabilities, which leads to the formation of a Markov process. Every state herehas anobservable symbol that is been attached to it. An HMM comprises of visible process with observable events and a hidden process which includes internal states with their movement intandem. The goal here is to find the optimal pathfrom the states, which leads to maximization of theoccurrence of observed sequence of symbols. Therelevant steps that associated in the algorithm for the generation of HMM are given below:

• HMM architecture is been developed bymaking use of various states whichultimately represent the given set offeatures.

• Assignment is been done of the hiddenstates to the features and so is the construction of HMM model is been done.

• The HMM is thus trained using supervised technique or unsupervised technique inorder to let the model sufficiently fit theproblem that isunderstudy.

• Emission probabilities are derived that influence the distribution of observed symbols, which implies that the probability of a symbol being observed provided that HMM is in a specific state.

HMM is decoded for the prediction ofhidden statesfrom the data.

• The benefits associated with HMMs are the ease of their use, need of smaller datasets and precise comprehension of the process.

Among the major drawbacks associated withHMMs is their higher computational cost. HMMs found to be most effective in case of biologicalsequence analysis and so they are periodicallyapplied for multiple sequence alignments, genefinding,etc

## 3.1.9K-Means clustering

The k mean clusteringalgorithmprovides ageneralized methodto implement approximatesolution. The reason why k mean clusteringalgorithm is very popular is because of the ease and simplicity. Kmean can be considered to be a

gradient descent procedure, where the initiation in the algorithm is done at starting cluster centroids and it iteratively decreases the objective function. The convergence of the k meangenerally takes place at the local minimum. It basically performs the updation work unless the local minimum is found. The problem to find the global minimum is

NP- complete. The time complexity of the k-means clustering algorithm is O(nkl) where, therequired number of clusters is denoted by "k", thetotal number of objects in the dataset is denoted by "n" and the number of iterations is denoted by "I",  $k \le n, I \le n$ .



Figure5: Diagram depictingK-meansclustering

## **Reinforcement learning**

Reinforcement learning belongs to that area of Machine Learning where the actions are takenpurely to achieve maximize rewards in a specific situation. It can be used on different types of machines and even on software for finding the best path possible or behavior it is supposed to take in any specific situation (Jiachi Xie et al 15) [16]. It distinguishes itself from supervised learning in a

way that in case of supervised learning the trainingdata has the answer key with it and the model istrained with the correct answer by its own on theother hand, in case of reinforcement learning, answer key is not available but here we can see that the reinforcement agent decides what is to be donein order to perform the given task (Nicolas Bougieet al, 2019) [17]. In the absence of training data set, it is bound to learn from its own previous experiences.

Algorithm	Advantages	Drawbacks
ARTIFICIAL NEURALNETWORK	<ul> <li>Good performance in caseofnonlinearrelationshi ps.</li> <li>Capacity to handle noisydata.</li> </ul>	<ul> <li>Computationalburdenisg reater.</li> <li>Over fitting isaproblem.</li> </ul>
HIDDENMARKOVMODEL	<ul> <li>Precise comprehension ofbackground process.</li> <li>Easy to use and ispowerful.</li> </ul>	<ul> <li>Intensivecomputation</li> <li>Slower than othermethods.</li> <li>Pronetoover-fitting.</li> </ul>

Table1: Theadvantagesanddrawbacksrelated tovariousmachinelearning algorithmare:-

SUPPORTVECTORMACHINE	<ul> <li>Bestgeneralizationabilityi sprovided.</li> <li>Robustto noisy database.</li> <li>Susceptibilityislesstoo ver-fitting.</li> </ul>	<ul> <li>Computationisexpensivei n some cases such as incase of non- linearlyseparableproblem s.</li> </ul>
K-NEARESTNEIGHBOR	<ul> <li>Simpleand easy to learn.</li> <li>It is found efficient whenthetrainingdataislar ge.</li> </ul>	<ul> <li>Complexincomputation</li> <li>As the number of attributes gets increase d,</li> </ul>
	• Fasttraining.	performance becomesinconsistent.
DECISIONTREE	<ul> <li>Capability to handle bothcontinuous and discreteattributes.</li> <li>Interpretabilityisbetter.</li> <li>Resultsarebetterincaseof redundantattributes.</li> </ul>	<ul> <li>In the presence of largenumberofclasses,thed ataisprone toerrors.</li> <li>Sensitivitytowardssmall variationsindata.</li> </ul>
RANDOM FOREST	<ul> <li>Highspeedandaccuracy.</li> <li>Less prone towards over- fitting.</li> <li>Able to evaluate everyattributeforpredic tion.</li> </ul>	Tendency of over-fitting incaseof noisydata.
ENSEMBLECLASSIFIERS	<ul> <li>Incaseofprediction,g reater efficiency isobtained.</li> <li>Utilization ofdataismore.</li> </ul>	Computational complexityismore.

## 4. Machine Learning Advancements in diabetes prediction:-

Machine learning can be used in case of digitaldiagnosisofanydisease. Itcan detectpatternsofcertain diseases and help in providing a broaderperspective.

### 4.1.Diabots:

Itisfoundthatthischatbotiscapableofinteracting with patients seamlessly based on the symptoms. There are many generic text-to-text diabot i.e.diagnostic chatbot which makes use of NaturalLanguage Understanding (NLU) for the providing personalized prediction by making use of generalized health dataset and also on the basis of various symptoms soughtfrom the patient.

#### 4.2.Oncology:

Here the researchers are making use of deeplearning techniques for the purpose of training thealgorithm and to make it recognize carcinogenictissue (but at the same time it is taken intoconsiderationthatthebloodsugarlevelisnormal) at such a level that is comparable to even physicians.

## 4.3.Better Radiotherapy:

As the machine learning algorithms have thepotential to learn from the multitude of varioussamplesthatarebeenavailablein hand, it becomes highly effective to diagnose and find the variables if any. The example includes Google's DeepMindHealthwhich is assisting the healthcare

professional to distinguish between the healthy andunhealthy people. Here the advancement is beenmade in terms of diagnosing eye damage done byvariousdiseaseswhich includesdiabetestoo.

### 4.4.Outbreak Prediction:

Machine learning is used in monitoring and predicting epidemics around the globe. ANN can be used to collect information from different websites and predict information from dengueout break to severe chronic infectious diseases. This can also assist in knowing the world wide increase in the diabetes patients round the globe which ledus to the conclusion that India is the diabetic capital of the world.

# 4.5.Crowd sourced Data Collection:

Crowd sourcing has helped researchers and practitioners to get access to huge amount of information that are been uploaded by people basedon their consent. This helps in collected data that is been collected by the consent of the patients and is assisting in the research.

The various applications available for thepredication of diabetes includes Diabetik byUglyApps,DiabetesinCheckbyEverydayHealth,iCookbook Diabetic by Publications International,mySugr Junior by mySugr GmbH, HealthyOut byHealthyOutandmanymore.

## 5.Conclusion/ Future work:-

The applications of machine learning could beappliedforthediagnosisofvariousdiseases, theirsymptoms, their cause, their treatment. The suddendeaths occurring due to kidney failure, heart attack, strokes etc. accompanied with diabetes can be prevented through early treatment and diagnosis. In the study we saw various algorithms such as SVM, decision tree, KNN, naïve bayes, etc making theiruse in the prediction of incidence of diabetes. The classification techniques gived ifferent results when applied to different dataset. We found that various classification techniques are useful for different data sets. The variation in the model performance on be noticed for different datasets and the cause could be predicted accordingly.

Future study can be focused on acquiring newdataset that would lead to new insight andknowledge to improving the prediction of diabetesusing machine learning techniques. Based on theparameters like age, body mass index, obesitylevel, history of chronic disease, etc whenaccompanied by various machine learningtechniques will lead to better prediction levels. Thenew dimension which is extending is usage is deeplearning whichwhenassisted withmachinelearning can give tremendous results in terms ofpattern recognition andbetterpredicted values

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