

Dataset Link:

<https://www.kaggle.com/datasets/falgunipatel19/text-publication-classification>

For Biomedical text document classification, abstract and full papers(whose length less than or equal to 6 pages) available and used. This dataset focused on long research paper whose page size more than 6 pages. Dataset includes cancer documents to be classified into 3 categories like 'ThyroidCancer','ColonCancer','Lung_Cancer'. Total publications=7569. it has 3 class labels in dataset. number of samples in each categories: colon cancer=2579, lung cancer=2180, thyroid cancer=2810

```
In [1]: #Import necessary libraries
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.feature_extraction.text import TfidfVectorizer
from nltk.corpus import stopwords
from nltk.util import ngrams
from nltk.stem import WordNetLemmatizer
from nltk.sentiment.vader import SentimentIntensityAnalyzer
import re
from wordcloud import WordCloud
from textblob import TextBlob
from nltk import word_tokenize, sent_tokenize
import spacy
nlp = spacy.load('en_core_web_sm', disable=['ner'])
import string
from nltk.corpus import stopwords
from nltk.stem.porter import PorterStemmer
from sklearn.model_selection import train_test_split
from sklearn.linear_model import LogisticRegression
import warnings
warnings.filterwarnings('ignore')
```

```

2022-09-10 21:09:14.510426: I tensorflow/core/util/util.cc:169] oneDNN custom
operations are on. You may see slightly different numerical results due to flo
ating-point round-off errors from different computation orders. To turn them o
ff, set the environment variable `TF_ENABLE_ONEDNN_OPTS=0`.
2022-09-10 21:09:14.550292: W tensorflow/stream_executor/platform/default/dso_
loader.cc:64] Could not load dynamic library 'libcudart.so.11.0'; dlerror: lib
cudart.so.11.0: cannot open shared object file: No such file or directory
2022-09-10 21:09:14.550324: I tensorflow/stream_executor/cuda/cudart_stub.cc:2
9] Ignore above cudart dlerror if you do not have a GPU set up on your machin
e.
2022-09-10 21:09:16.405241: W tensorflow/stream_executor/platform/default/dso_
loader.cc:64] Could not load dynamic library 'libcuda.so.1'; dlerror: libcuda.
so.1: cannot open shared object file: No such file or directory
2022-09-10 21:09:16.405292: W tensorflow/stream_executor/cuda/cuda_driver.cc:2
69] failed call to cuInit: UNKNOWN ERROR (303)
2022-09-10 21:09:16.405322: I tensorflow/stream_executor/cuda/cuda_diagnostic
s.cc:156] kernel driver does not appear to be running on this host (vinod-Vost
ro-3400): /proc/driver/nvidia/version does not exist

```

```

In [2]: #Read the data using the pandas
data=pd.read_csv('/home/vinod/Downloads/alldata_1_for_kaggle.csv',encoding='la
data.head()

```

```

Out[2]:      Unnamed: 0      0      a
0      0  Thyroid_Cancer  Thyroid surgery in children in a single insti...
1      1  Thyroid_Cancer  " The adopted strategy was the same as that us...
2      2  Thyroid_Cancer  coronary arterybypass grafting thrombosis i~^b...
3      3  Thyroid_Cancer  Solitary plasmacytoma SP of the skull is an u...
4      4  Thyroid_Cancer  This study aimed to investigate serum matrix ...

```

```

In [3]: #data information in the dataset
data.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 7570 entries, 0 to 7569
Data columns (total 3 columns):
#   Column      Non-Null Count  Dtype
---  ---
0   Unnamed: 0  7570 non-null    int64
1   0           7570 non-null    object
2   a           7570 non-null    object
dtypes: int64(1), object(2)
memory usage: 177.5+ KB

```

```

In [4]: #Check the data shape of the dataset
data.shape

```

```

Out[4]: (7570, 3)

```

```

In [5]: #Check the null values in the dataset
data.isna().sum()

```

```
Out[5]: Unnamed: 0    0  
        0          0  
        a          0  
        dtype: int64
```

Data Preprocessing

```
In [6]: #Let's remove the unnecassary columns in the dataset  
data.drop(['Unnamed: 0'],axis=1,inplace=True)  
data.head().style.background_gradient(cmap='winter')
```

Out[6]:

0

0 Thyroid_Cancer

Thyroid surgery in children in a single institution from Osama Ibrahim Almosallama Ali A Alsobhib Saud AlShanafeyb From the aDepartment of Surgery College of Medicine Qass bDepartment of Surgery King Faisal Specialist Hospital and Research Center Riyadh Sa Specialist Hospital and Research Center Riyadh Saudi Arabia Correspondence Dr Osa College of Medicine Qassim University PO Box Buraidah Al Qassim Saudi Arabia osama_ Citation Almosallam OI Aseeri A Alhumaid A AlZahrani AS Alsobhi S AlShanafey S Thyro Ann Saudi Med Received January Accepted May Published August Copyright Copyright Ar access under the Creative Commons AttributionNonCommercialNoDerivatives International be accessed at <http://creativecommons/licenses/by-nc-nd/4.0/>Funding NoneBACKG scarceOBJECTIVE Analyze outcome data on thyroid surgery in a pediatric populationDf health care institutionPATIENTS AND METHODS We collected demographic and clinical dat surgery in the period to Descriptive data are presentedMAIN OUTCOME MEASURES complications length of stay and radioactive iodine treatment and recurrencesSAMP thyroidectomy procedures were females and the mean age at operation was years and wei type There was no history of radiation exposure Eightyone patients had fine needl histopathology in of cases Sixtysix patients had malignant cancer papillary of patients who h and had distant metastases to the lung Procedures included total thyroidectomy hemithyroid Twentythree patients developed hypocalcemia permanent and had unilateral recurrent followed up for a mean duration of months median months Of patients with thyroid cancer Malignancy is the commonest indication for thyroid surgery in children and FNA is highly di nerve injury are significant complications The recurrence rate in thyroid cancer is LIMITAT Noneoriginal ANN SAUDI MED JULYAUGUST WWWANNSAUDIMEDNET OcThyroid disease children compared to adults The prevalence of palpable thyroid nodules in children ranges fr is the most common endocrine malignancy in children accounting for of pediatric cancers in in adolescents aged year2 The most common indication for thyroid surgery in children var for malignant conditions is rising38 Data in children throughout the world are relatively scarce clinical data and outcome of thyroid surgery in a large series of children treated at a sing Research Center KFSHRC in RiyadhPATIENT AND METHODS With the approval of the medical records of all patients years old and younger who underwent a thyroid surgery I elected to include patients up to the year to ensure a reasonable followup period Patients operating room log for all procedures involving the thyroid gland for the specified age groupf outcomes were collected Specific data that were obtained included age at operation gender radiation exposure presence of multiple endocrine neoplasia type MEN thyroid function ultrasound presence of lymph nodes metastasis or distant metastasis fine needle a histopathology and length of followup Outcomes analyzed were postoperative complications transient or permanent recurrent laryngeal nerve paralysis wound infection and hematoma and recurrences Thyroid procedures in this series included hemithyroidectomy subtotal tota performed by either an endocrine adult surgeon or a pediatric surgeon No intraoperative procedures were performed by adult endocrine surgeons but lately a combined approach w endocrine surgeons collaborated in such cases procedures the normal range in or hypocalcemia was identified if it lasted for less than months while permanent hypocalc remained below normal range and the patient continued on calcium supplementation after history of MEN underwent genetic testing of the RET protooncogene to confirm the di thyroidectomy had a preoperative and postoperative vocal cords assessment at the Otolary and comparisons were conducted using the t test for continuous proportionsRESULTSBetween and patients underwent surgical procedures patients unde institution Eighty patients were females The mean age at operation was years median yea thyroidectomy was thyroid nodule which was present in of cases Table The mean SD size associated with MEN syndromes The final pathology in two patients with MEN syndrome s remaining patients had prophylactic procedures before development of MTC None of Eightyone patients FNA which correlated with the final histopathology in of cases There were Graves disease which did not require FNA The remaining cases underwent FNA at anotl institution or they came for completion thyroidectomy with documented pathology for malign hospitalThe most common diagnoses included papillary thyroid cancer and multinodular goit in patients IndicationNodulen MEN prophylaxisHyperthyroidismMultinodular goiterCompleti calcium levels below Data are number original PEDIATRIC THYRO WWWANNSAUDIMEDNET Ocnode Table Surgical procedures included total thyroidectom and subtotal thyroidectomy Neck dissection was performed in patients Operative cor common complication was hypocalcemia transient permanent and Table Thyroid pathology i tissueColloid noduleCystAdenomaThyroiditisGraves diseaseThyroid cancerPapillaryFol number Table Benign and malignant lesions in patientsBenignn37Malignantn66 P value / noduleHypocalcemiaRecurrent laryngeal nerve palsyBleedinghematomaWound infectionTrac stay daysMEN recurrent laryngeal nerve palsy transient permanent all were unilateral Tab node metastasis and patients had distant metastases to the lung None of the patients devel

tracheal injury Patients were followed up for a mean of months median range months radioa
with malignant lesions patients had recurrences were local recurrences and were local e
received radioactive iodine RAI before and after recurrence One case was low risk bef
recurrence One case had medullary thyroid cancer so did not receive RAI In the remaining fiv
patients received RAI before or only after a recurrence All local recurrences underwent rese
up There was no mortality in this study DISCUSSIONThe most common indication for thyroic
correlates with previously published reports in the pediatric population35 Children with thyroid
of developing thyroid cancer compared to adults910 The high incidence of malignancy in thi
should be carefully evaluatedFNA is a valuablemethod for preoperative evaluation of thyro
routine use of FNA in children including the need for sedation sampling errors and the limite
Many previous studies reported high sensitivity and specificity of FNA in evaluating thyroid
findingsOur data showed lymph node metastasis in of thyroid cancer cases which supp
frequently present with more extensive disease than adults Lymphnode involvement at diag
of adults with differentiated thyroid cancer1523 Because our hospital is the largest referra
cases this may explain the large number of lymph node and distant metastasis In this cohort
thyroidectomy in children is hypoparathyroidism with an incidence ranging between to which
SAUDI MED JULYAUGUST WWWANNSAUDIMEDNET 0ccorresponds with our results of wt
study found that total thyroidectomy central and bilateral neck dissection Graves disease and
after thyroid surgery3 In this cohort postoperative hypocalcemia was noted more in
significance Moreover there was no significant difference between benign and malignant
recurrent laryngeal nerve injury or overall complications a finding that was reported previousl
an inverse relationship between surgeon volume and complication rates2728 but similar data
found that highvolume endocrine surgeons have better outcomes and shorter lengths i
parathyroidectomy in children compared to pediatric surgeons general surgeons or otolar
concluded that a collaborative approach between pediatric and endocrine surgeons would h
to suggest that a combined approach with endocrine and pediatric surgeons in addition to p
of children with surgical thyroid disease given the low number of pediatric patients4 Ou
approaches given the late adoption of the combined approach The recurrence rate for thy
varied widely in reported studies ranging from to while it was in this cohort Only a few studi
node involvement multiple nodules male gender younger age histologic subtype and advance
recurrence17233033 In this study of patients with malignant lesions received RAI A
indications of postoperative RAI treatment in lowrisk patients the current recommendatio
RAI3436There are some limitations to this study The retrospective nature may affect the valid
cases in some categories did not enable us to compare groups and explore predictors relativ
adds to the scarce data on thyroid surgery in pediatric age group Malignancy is the common
FNA is highly diagnostic Hypocalcemia and recurrent laryngeal nerve injury are significant c
rare but recurrence is not uncommon and a significant number of patients with malignant ca
THYROID SURGERYANN SAUDI MED JULYAUGUST WWWANNSAUDIMEDNET 1
McLaren GD Nichaman MZ Iodine and goiter in children Pediatrics Ries LAG Melbert D Kray
et al SEER Cancer Statistics Review Bethesda National Cancer Institute Based on Novemb
PT Gaz RD Hodin RA Parangi S Randolph GW et al Pediatric thyroidectomy in a high
postoperative hypocalcemia J Pediatr Surg Aug5081316 Wood JH Partrick DA Barham HP B
thyroidectomy a collaborative surgical approach J Pediatr Surg May4658238 Scholz S Smi
Thyroid surgery at Childrens Hospital Boston a 35year singleinstitution experience J Pediat
Thyroid nodules and cancers in children Pediatr Endocrinol Rev Sep611423 Hameed F
adolescent thyroid cancer J Paediatr Child Health LugoVicente H Ortiz VN Irizarry I
management in the era of fine needle aspirationJ Pediatr Surg Mussa A De Andrea M Motta I
Malignancy in Children with Thyroid Nodules J Pediatr Oct167488692 Amirazodi E Propst E
thyroid FNA biopsy Outcomes and impact on management over years at a tertiary care ce
Cramer HM Chen S Wu HH Histologic and clinical followup of thyroid fineneedle aspirates in
Decoppi P Pierro A Brain C Hindmarsh P Butler G et al Thyroid Surgery in Children Clir
Kundel A Thompson GB Richards ML Qiu LX Cai Y Schwenk FW et al Pediatric Endocrine S
J Clin Endocrinol Metab February Jiang W Newbury RO Newfield RS Pediatric thyroid
institutional experience features and over a 10year period Int J Pediatr Endocrinol Burke JF S
Surgery at a Tertiary Medical Center Surg Res AlQahtani KH Tunio MA Al Asiri M Aljohan
treatment outcomes of differentiated thyroid cancer in Saudi children and adults J Otolaryngc
DJ Verrijn Stuart AA Lodewijk L Valk GD Van der Zee DC et al Postoperative Complica
Young Patients With Multiple Endocrine Neoplasia Type Medicine Baltimore 20159429e11C
Angelos P Reynolds M Total thyroidectomy for benign disease in the pediatric patientfeasi
PH Ko CY Yeh MW Surgeon volume as a predictor of outcomes in inpatient and outpatien
HM Tielsch JM Powe NR Gordon TA Udelsman R The importance of surgeon expe

thyroidectomy Ann Surg Tuggle CT Roman SA Wang TS Boudourakis L Thomas D Udels
operating on our children Surgery Dec144686977 Park S Jeong JS Ryu HR Lee C Park JH K
Children and Adolescents27Year Experience in the Yonsei University Health System J Ko
Kollars JP Moir CR Papillary thyroid carcinoma in children risk factors and complication
Sugino K Mimura T Nagahama M Kitagawa W Shibuya H et al Pediatric differentiated thy
disease free survival BMC Cancer D Danese Gardini A Farsetti A Sciacchitano S Andreoli M I
adolescents Eur J Pediatr Astl J Chovanec M Lukes P Kutra R Dvorakova M Vlcek P
adolescents years experience surgery of pediatric thyroid lymph node metastases carcin
Rangarajan V Nair N Nadkarni MS Pai PS Dcruz AK et al Pediatric thyroid cancer J Su
Matovic M Milovanovic Z et al Surgical management of welldifferentiated thyroid carcinoma i
of a single institution in Serbia Endocr J Scheumann GF Gimm O Wegener G Hundeshagen
management of locoregional in papillary thyroid cancer World J Surg Shi RL Qu N Yang SW
lymph node metastasis using a differentiated thyroid cancer risk model Onco Targets The
Ryan JJ Grant CS et al Papillary thyroid carcinoma in children and adults longterm fo
institution during three decades Surgery Collini P Mattavelli F Pellegrinelli A Barisella M Fe
thyroid gland of childhood and adolescence Morphologic subtypes biologic behavior and
cases treated at a single institution during a 30year period Am J Surg Pathol BorsonChazot
JL Predictive factors for recurrence from a series of children and adolescents with differen
HD Bauer AJ Isaza A MostoufiMoab S Kazahaya K Adzick NS Surgical management of pe
thyroidectomy at the Childrens Hospital of Philadelphia highvolume Pediatric Thyroid Center J
De Coppi P Thyroidectomy in Children InPediatric Surgery pp Springer Berlin Heidelberg F
Benvenega S et al Management Guidelines for Children with Thyroid Nodules and Differ
Association Guidelines Task Force on Pediatric Thyroid Cancer THYROID Volume Number
SAUDI MED

1 Thyroid_Cancer

" The adopted strategy was the same as that used in prior years [] and is based on four e
subsets The first query QPub_plain is based on a plaintext search in PubMed titles and s usi
relies on the PubMed indexing scheme using MeSH terms and results are m
QWoS_restricted is based on a plaintext search in WoS restricted to the two research areas
Services The fourth query QWoS_filtered is based on the same plaintext search used in Wo
Archeology Dance Zoology etc and the two research areas of the previous query It
nonPubMedindexed papers that are supposed to be caught by the two PubMed quer
citations was performed by the two section editors to select candidate best papers Followi
best papers were then individually reviewed and rated by both section editors the chief edit
reviewers from the international Medical Informatics community Based on the reviewe
committee then selected the best papers of the year in the decision support domainIM/
Thieme Verlag KG OcReview Results The literature search has been performed on Janu
distributed as follows for QPub_plain for QPub_indexed for QWoS_restricted and for QWo
PubMed and from WoS Compared to the previous year the global query retrieve
independently performed by both section editors based on the title and of papers not rejecte
two editors to achieve a final selection of candidate best papers After the external review of
three of them as best papers for [] Table They are discussed in the next section and s
AppendixDiscussion and OutlookIn the first paper Hendriks [] propose an approach to th
certainly builds on already existing approaches but which is systematically conducted in orde
guidelines They promote the formalism of clinical decision trees CDTs as they are both clir
and computerinterpretable thus suitable for implementation in datadriven CDSSs The disam
by the formal unequivocal specification of data items used as decision criteria using internat
and second by the representation of guideline knowledge as CDTs The method is applied to
were built involving a total of data items among which could not be linked to standard ter
certain criteria which could be subjective or had multiple definitions The resulting knowledg
application where it can be interactively browsed or automatically executed By modeling guid
in the sharing of encoded knowledgeIn the second paper KamiÅjaliÄ» [] tackled the i
processes used for managing chronic diseases and their execution in CDSSs They a
therapeutic management of chronic diseases like those known to increase the cardiovascu
strategy dosage adaptation and intolerance management To handle these different aspec
extended Timed Transition Diagram eTTD With eTTDs they illustrate the multilevel ar
contents of arterial hypertension management guidelines This detailed demonstration
management can be formalized to develop a CDSS could certainly be used in other medical c
conceptual and practical framework to help assess confidence in predictive tools GRASP f
method to look for evidence from the published literature and an analysis grid It standar

associated to a predictive tool and the grading of its level of proof Three phases of evaluation the tool to assess both its internal and external validity ii during the implementation to assess implementation to assess its effectiveness and safety In each phase the level of evidence qualitative summarizes the direction of evidence positive negative mixed This grid can be compared to the CONSORT statement for clinical trials However it gives a rigorous methodology for a CDSS extended to all kind of CDSSs It might be a useful tool to extend the evidencebased culture to the three best papers selected for the Decision Support section of the edition of the IMIA Yearbook literature review deserve to be cited Some of them deal with the personalization of decision support approach to develop personalized care plans that comply with clinical practice guidelines in complex situations Jafarpour [10] propose a solution to dynamically manage the conflicts that can arise when to introduce the use of health information technology involving multiple criteria decision support systems Interestingly other works promote the creation and sharing of operational knowledge based on CDSS to transform the textual STOPPSTART criteria into unambiguous definitions mapped to the EUCAST expert rules as an ontology and production rules to detect antimicrobial therapies and to build a knowledge base that can compete with commercial ones Replacing humans is another topic of interest to virtualize a doctor the automatic acquisition of data through sensors and speech recognition Rozenblum et al [11] propose a machine learning method to generate clinically valid alerts for CDSS CDSS is another key point Kannan [12] propose a method for a CDSS design to best meet a physician's needs Design alerts may also avoid rejection of CDSSs by caregivers Fernandes [13] created a decision support notifications delivered to healthcare professionals Amrose et al [14] tried to understand in real time the actions they triggered Finally it is always interesting to obtain varied evaluation results for CDSSs evaluated Watson for Oncology in thyroid carcinoma and reported a concordance rate with the physician the tool As evidenced by the number and the variety of works around decision support systems, the selection highlighted pragmatic works that promote the transparency and sharing of the IMIA Yearbook Table Best paper selection of 2013 for the IMIA Yearbook of Medical Informatics in the alphabetical order of the first authors surname Section Decision Support 10a7 Hendriks MP, Jansen J, MJC Strobbe LJA, Merkus JWS, Zonderland HM, Smorenburg CH, Jager A, Siesling S, Transformation of the National Breast Cancer Care Guidelines Into DataDriven Clinical Decision Support Systems KamiÅjaliÄtAtRia±otDtKerttStWelzertTtNemectZlatolastLtMultileveltmedicaltknowledge Engineering for chronic diseases Data Knowledge Engineering 10a7 Khalifa M, Magrabi F, Gallego J, Jafarpour B, evidencebased grading and assessment of predictive tools for clinical decision support BMC Med Inform Decis Mak 2013;13:10a7 support tools as well as the grading of their utility The ultimate goal of this work is to evaluate decision support themAcknowledgementWe would like to thank all the present and past editorial boards of the IMIA Yearbook and the IMIA Section Decision Support as well as the reviewers for their participation to the selection of the best papers for the IMIA Yearbook section We cannot end this synopsis without a meaningful thought for our colleague and friend who was very active in the IMIA Yearbook to tackle the tasks of a Decision Support section coeditor but passed away in last December 2013 Jankovic I, Chen JH Clinical Decision Support and Implications for the Clinician Burnouy O, Jankovic I, Contributions on Clinical Decision Support from the Literature Yearb Med Inform 2013;13:10a7 van der Slangen MJC, Strobbe LJA, Merkus JWS Transformation of the National Breast Cancer Care Guidelines Into DataDriven Clinical Decision Support Systems KamiÅjaliÄtAtRia±otDtKerttStWelzertTtNemectZlatolastLtMultileveltmedicaltknowledge Engineering for chronic diseases Data Knowledge Engineering 10a7 Khalifa M, Magrabi F, Gallego J, Jafarpour B, evidencebased grading and assessment of predictive tools for clinical decision support BMC Med Inform Decis Mak 2013;13:10a7 Sarigul B, Arvanitis TN, Lindman P, Chen R, A Collaborative Platform for Management of Chronic Diseases in Primary Care Plans Comput Struct Biotechnol J 2013;13:10a7 Jafarpour B, Raza Abidi S, Van Woensel W, Razafimanantsoa A, practice guidelines to provide decision support for comorbid conditions Artif Intell Med Ben Boussaid M, PARS a system combining semantic technologies with multiple criteria decision aiding for support of clinical decision Huibers CJA, Salleveld BTGM, de Groot DA, Boer MJ, van Campen JPCM, Davids CJ, Clinical Decision Support algorithms for software implementation A multidisciplinary consensus procedure Int J Med Inform 2013;13:10a7 Campos M, Palacios F, Impact of expert knowledge on the detection of patients at risk of hospitalization in decision support systems J Biomed Inform 2013;13:10a7 Mller L, Gangadharaiiah R, Klein SC, Perry J, Bernstein G, Clinical Decision Support for community driven diagnostic decision support system development BMC Med Inform Decis Mak 2013;13:10a7 Canbay A, Menrad K, Heider D, The virtual doctor An interactive clinicaldecisionsupport system for the prediction of diabetes Artif Intell Med Rozenblum R, RodriguezMonguio R, Volk LA, For the IMIA Yearbook Learning System to Identify and Prevent Medication Prescribing Errors A Clinical and Cost Analysis of a Decision Support System Kannan V, Basit MA, Bajaj P, Carrington AR, Donahue IB, Flahaven EL, User stories as ligaments for decision support development J Am Med Inform Assoc 2013;13:10a7 Fernandes CO, Miles S, Lucena CJP, Cowan J, Clinical Decision Support with Alarm Fatigue in Hospital Environments Because of Sensory Overload Algorithm 10a7 2013;13:10a7 20192111e15406 Amroze A, Field TS, Fouayzi H, Sundaresan D, Burns L, Garber L, et al, Use of a Decision Support System to Logs to Identify Physician Actions Following Noninterruptive Alert ing Descriptive Study J Am Med Inform Assoc 2013;13:10a7 Kim JM, Kim EH, Kim K, Pak K, Concordance in postsurgical radioactive iodine therapy recommendations for differentiated thyroid carcinoma Cancer Correspondence 2013;13:10a7 Lonard de Vinci rue Marcel Cachin Bobigny FranceEmail catherineduclosaphpfr IMIA Yearbook of Medical Informatics Considerations on Clinical Decision Support from the Literature 10cAppendix Content Summary

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Section of the IMIA YearbookHendriks MP Verbeek XAAM van Vegchel T van der Sangen I Smorenburg CH Jager A Siesling STransformation of the National Breast Cancer Guideline Cancer Inform May3114Since clinical practice guidelines are still narrative and described in la to model complex guidelines as datadriven clinical decision trees CDTs that could be still hum implementation in decision support systems The Dutch national breast cancer guidelin characterize the patient and the tumor and represent decisional criteria were encoded u coding systems related to breast cancer when feasible In total CDTs were necessary to cov all data items could be coded using existing classification and coding systems All CDTs repre guidelines could be transformed as systematically constructed modular datadriven CDTs tha decision support applicationKamiÅjaliÄ A Ria±o D Kert S Welzer T Nemec Zlatolas LMultile medical practice for chronic diseasesData Knowledge Engineering This research is focus medical processes involved in chronic diseases management which can be viewed as a proc An intuitive easy and effective mechanism for medical knowledge formalization is propos Transition Diagram eTTD This formalism allows for the consistent representation of three taken into account in the prescription and adaptation of longterm treatment therapy strategy be manually applied to build eTTDs from clinical practice guidelines eTTDs implementatio guidelines for the therapeutic management of arterial hypertension The obtained mode development of decision support systems involving medical proceduresKhalifa M M evidencebased grading and assessment of predictive tools for clinical decision supportBMC a clinical predictive tool in clinical practice should be guided by its correctly assessed effectiv a conceptual and practical framework to Grade and Assess Predictive tools GR evidencebased system to support their search for and selection of efficient predictive tools based on published evidence across three dimensions phase of evaluation level of evidence tool is based on the phase of evaluation that gets the hightest grade supported by the supports a positive This framework was successfully applied to five predictive tools GRA data base that documents the evidence of predictive tools IMIA Y

- 2 Thyroid_Cancer coronary arterybypass grafting thrombosis i-^brin i-^brinogen mutationIntroduction Intraoper harvesting is very rareCase Report We present a case of a 60yearold male patient with multiv nonST elevation acute coronary syndromeand type2 diabetes mellitus whombilateralintraoperative SV thrombosis occurred during graft harvesting Routinethromb cancer was excluded Compared with healthy controls we observed prolonged i-^brin c rei-^ected by endogenous thrombin potential Scanning electronmicroscopy of the thrombos layer on theclot surface with a solid mass of unusually compressed platelets and eryl i-^brinogen and factor F XIII polymorphismsand was found to be heterozygous for i-^br TConclusion i-^brinogen HaeIII and FXIII Val34Leu polymorphisms are rei-^ected inreduced might contribute to intraoperative SV thrombosis during vascular grafting procedures Car failure after bypass proceduresIntroductionCoronary artery bypass grafting CABG is a metho multivessel disease anddiabetes mellitus DM Although arterial grafts are preferredin se leftinternal thoracic artery LITA to bypass the left anteriordescending artery LAD and to p often chosen vascular graft the greatsaphenous vein SV offers decent durability and is easy t of caseswithin the i-^rst months and as many as may occludewithin i-^rst to weeks environment with disruption of bloodi-^low in vasa vasorum damage to the adventitia hypoxi focal endothelialdisruption2 Acute SV graft failure is usually a result of graftthron failuregrafttarget vessel disproportion etc may be caused byhypercoagulabilityreceivedMarc Thieme Verlag KGStuttgart · New York 0ce198Bilateral Saphenous Vein Thrombosis du male patient with multivessel coronary arterydisease who suffered from a nonST elevation a to admission a nonsmoker with type2 DM on metformin peptic ulcer diseaseand a history of CABG Just after the NSTEMI a left ventricle LV thrombus was seen on one echocar followup There was no deep venousthrombosis or bleeding diathesis history On admissi enoxaparin mg once daily Routine laboratory tests were withinnormal ranges °Table There apart from obesity body mass index kgm2 when the patient was admitted The lower extren veins nosigns or symptoms of venous insufi-^ciency and the pastmedical history was negativ venous insufi-^ciency or varicose veins Thepatient was operated on following the star surgery resident harvested theright SV using the technique The wall of the SV lookedgrc were tiedoff and clipped and a needle was placed at the distal end whilethe proximal end was Results of initial and followup laboratory testingVariableCoagulation testsRed blood 103µLPlatelet count 103µLPAPTT sPT sPT INRPPlatelet aggregation mmolL arachidonic acid µ gLAntithrombin III Ddimer µgLantiXa IUmLHomocysteine µmolLProtein C Protein S Fac cī-^brinogen 455G AFactor XIII 100G TLupus anticoagulant ratioLupus anticoagul IgMAnti2glycoprotein I IgG antibodyAnti2glycoprotein I IgM antibodyNormal rangesPreope

no mutationGG no mutation GPL MPL SGU SMUGG no mutationGG no mutationGA

SMUAbbreviations APTT activated partial thromboplastin time GPL IgG phospholipid unit I ratioMPL IgM phospholipid unit PT prothrombin time SGU standard IgG 2 glycoprotein unit S OcBilateral Saphenous Vein Thrombosis during CABG Mazur et al199to iush the vein wit normal saline mL while the distalend was closed with an atraumatic vascular clamp and vei the distal end aluminal thrombus was visible The left SV was then taken downu cardiacsurgeon with the same result Presence of a luminal thrombuswas conirmed upon s was administered and normal LITA outflow wasconirmed Concerns regarding safety of (suspected thrombotic issueand the approach was modifed The LITALAD anastomosis postoperative course was uneventful On postoperative day the patient received dual antiplate discharged on day with nosigns of thrombosis or myocardial ischemia Elective angiopla completethrombophilia screening was done °Table On the and12month followup the thrombophilia was suspected screening wasinitiated showing no abnormalities °Table (Positiveantibodies against neutrophil cytoplasm antigens pANCAand cANCA were excluded analyze i-brin phenotype using the previouslydescribed methodology34 Brieily plas hydrostatic pressure systemTubes containing i-brin clots formed from adding mmolLcal tocitrated plasma were connected through plastic tubing to abuffer reservoir M TrisHCl M Na measured within minutes A permeation coeficient K_s relecting poresize was calculated Q is the iflow rate in time t L is the length of a i-brin gel is the viscosity of liquid A is the cr in dyne cm^2 Lower K_s values indicated reduced permeability Fibrinogen was determined usi i-brinogenlevel was normal we identifed strongly decreased i-brin clotpermeability K_s from our previous report n ¼ K_s ¼ 9 cm^2 3 samples collected during late follow] up appointme controls n ¼ we observed prolonged clot lysistime CLT 06 vs 06 minutes and increased thrombinpotential ETP in the studied subject ETP ¼ 06 vs 06 nM 02 min respectively me with calibrated automated thrombography thrombinoscope BV Maastricht theNetherlands ac 96well plate iuorometer Ascent Reader Thermolabsystems OY Helsinki Finland equ Brieily microliters of plateletpoor plasma were diluted with μL of the reagent contain phosphatidylserinephosphatidylcholinephosphatidylethanolamine vesicle and μL c bovinealbumin and mmolL ZGlyGlyArg7amino4methylcoumarin Each plasma sample was c concentration of thrombin generated was used3Cryosectioned tissue sections were i-xed activity was quenchedwith H_2O_2 and unspecifc background was blocked with bc UnitedStates Primary adequate antibodies against i-brin or tissuefactor TF were applied b antibodies were followed by thecorresponding secondary antibodies conjugated with i Images were analyzed using Olympus BX microscope SVs immunostainingreve endothelium°Fig 1A and high TF °Fig 1B activity Within the thrombuswe found abunda suggesting the presence of proinflammatory monocyteswhich are known source of TF CD68 due to high unspecifc backgroundresulting from large amounts of i-brin The mi vessels °Fig 1C D Withinalmost every single vessel we found thrombi rich in bothprothromb clot phenotype relected by reduced K_s and prolonged CLT along with enhanced thrombi the immunostaining of the SVs prompted us to perform analysis ofwhole blood clot morpholo previously described6 After washing thethrombus was i-xed with glutaraldehyde pho: dehydrated goldcoated and photographed digitally with a JEOL JSM JEOL Tokyo Japan layer on the clot surface with a solid massof unusually compressed platelets and erythr veryhigh contractileforces during clot formation in a plateletdriven i-brinmediated mechani common i-brinogen and factor F XIII polymorphisms The patient was heterozygous for i-br TDiscussionA dramatic intraoperative SV thrombosis provoked by graftharvesting for CABG its cause remained unknown following thestandard thrombophilia screening The cases c period are very rareand as few as of grafts occlude within i-rst to weeks17TH Vol No 0ce2 CABG Mazur et alFig Representative images of SV graft immunostaining after massive thro green nucleistained blue using DAPI and scanning electron microscopic images E F of the

citrated blood obtained from the patient undergoing CABG. Box and arrow represent magnified pertinent stained fragments. See text. CABG, Coronary artery bypass grafting; SV, saphenous vein; intima, the media and the adventitia.⁸ The intima is built of the layer of endothelial cells or muscle cells and the adventitia forms the outer part.⁸ In a normal setting, the endothelium thrombosis⁹ and its focal disruption may predispose to vessel thrombosis.² SV manipulation integrity and elicits an inflammatory response with platelet adhesion and leukocyte recruitment, which is extremely rare in the operating room. SV dissection results in blood flow disruption, hypoxia, and vessel wall hypotension.¹⁰ Acute perioperative saphenous vein graft failure is a very uncommonly occurring event prior to graft placement. Surgical factors, like technical anastomotic failure, vessel and the graft may lead to thrombosis, but vessel injury and hypercoagulability are evident inflammatory processes in microscopy in our patient, but even if an inflammatory process is present, Saphenous Vein Thrombosis during CABG. Mazur et al.²⁰¹ preoperatively in our patients does not explain the dramatic intraoperative thrombosis. We hypothesized that increased thrombin phenotype were responsible for the clinical presentation. Conversion of fibrinogen to fibrin coagulation. It has been shown that fibrin clots with small pores between tight resistant.¹² Such clot phenotype has been evidenced in multiple thrombotic pathologies such as venous thromboembolism.⁴ The prothrombotic clot phenotype is affected by a tendency to form previously reported in patients with stent thrombosis.¹⁴ While routine thrombophilia screening in common hypercoagulable states,¹⁵ there are prothrombotic conditions that escape routine diagnosis and prothrombotic fibrin properties lead to the discovery of two mutations in thrombophilia screening, namely fibrinogen 455G/A and FXIII 100G/T. Elevated fibrinogen was associated with graft failure after CABG.^{11,16} Epidemiological studies have established that elevated fibrinogen is associated with cardiovascular diseases.¹⁷ A meta-analysis of individual records of participants from prospective studies revealed a 1 g/L increase in usual fibrinogen level for coronary heart disease was associated with a confidence interval [CI] as 95% CI. Risk of coronary disease progression was also linked to genetic polymorphisms of fibrinogen. The A allele of fibrinogen 455G/A was associated with more severe progression of coronary disease.¹⁸ Our colleagues in a meta-analysis of studies with patients found that A allele of the fibrinogen 455G/A was associated with coronary disease and also with ischemic stroke, odds ratio for stroke 1.4 [CI] for AA. ¹⁹ In a meta-analysis of atrial fibrillation, Hu and colleagues found that the A allele of fibrinogen 455G/A was associated with elevating the level of plasma fibrinogen.²⁰ On the other hand, in a meta-analysis of stroke, a polymorphism was shown to be associated with risk of myocardial infarction.²¹ FXIII is crucial for plasma concentration and reflects nonspecifically the extent of thrombosis as shown by Li et al.²² Interesting associations of FXIII Val34Leu polymorphism and thrombotic disorders were found in a meta-analysis of studies that FXIII Val34Leu polymorphism is associated with recurrent preoperative incidence of ischemic stroke was found for this polymorphism.²⁴ Apparently, when the stroke occurred, the severity of its outcome.²⁵ Furthermore, Kreutz and colleagues suggested that FXIII Leu34 allele is associated with recurrent MI and death in patients with angiographically established coronary artery disease.² In patients undergoing CABG, FXIII Leu34 allele is associated with decreased fibrinolysis.²⁷ Conclusion: Our extensive workup showed that fibrinogen HaeIII and FXIII Val34Leu polymorphisms are associated with decreased clot permeability and susceptibility to lysis. These mutations likely contributed to intraoperative graft failure. We needed to elucidate the role of these polymorphisms in early graft failure after bypass grafting. Funding: This study was funded by a grant from the Jagiellonian University Medical Research Fund. Interest: None declared. References: Bourassa MG. Fate of venous grafts: the past, the present, and the future. *Circulation*. 1995;92:1131-1136. Mazur P, Sokolowski G, Hubalska M, et al. Saphenous vein graft failure after CABG. *Ann Thorac Surg*. 2011;91:1131-1136. Undas A. Prothrombotic alterations in plasma fibrin clot properties in thyroid disorders and in patients with coronary artery disease. *Thromb Haemostasis*. 2004;83:1131-1136. Undas A, Zawilska K, Ciesla D, et al. Altered fibrin clot structure and function in patients with coronary artery disease and their relatives. *Blood*. 2005;106:1131-1136. Natorska J, Marek G, Hlawaty M, Sadowski J, Tracz W, Undas A. 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3 Thyroid_Cancer

Solitary plasmacytoma SP of the skull is an uncommon clinical entity that is charac

monoclonal plasma cells This case report describes a 50 year old male that presented with

occiput The diagnosis of SP was based on the pathological results and imaging examinatio

skull reconstruction and lower trapezius myocutaneous flap LTMF transplantation under gener

extended to the subcutaneous and the subdural space through the dura mater with skull defe

large areas of scalp and subcutaneous tissue which resulted in a large postopera

transplantation All of the tumour was removed and the transplanted flap grew well Followup

on the right frontal lobe The patient received six cycles of the PAD chemotherapy regimen be

the lesion was significantly reduced This case demonstrates that LTMF is an alternative ap

soft tissue defects caused by the excision of a large malignant tumour of the occipital regi

neoplastic recurrence Keywords Solitary plasmacytoma lower trapezius myocutaneous flap

July accepted March 1 Department of Neurosurgery Hunan Cancer Hospital and the Affiliated

Central South University Changsha Hunan Province China 2 Department of Head and Neck S

Cancer Hospital of Xiangya School of Medicine Central South University Changsha Hunan P

and Zheng Wen He Department of Neurosurgery Hunan Cancer Hospital and the Affiliated

Central South University Tongzipo Road Yuelu District Changsha Hu

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work without further permission provided the original work is attributed as specified on the SA

access at sage 0 Introduction Solitary plasmacytoma SP is the pathological manifestation of

an SP that originates in bone tissue is called a solitary plasmacytoma of bone SPB 1 Bone de

the most common sites are the pelvis spine femur humerus and ribs 2 An SPB of the skull is

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rarely mentioned in the literature³⁴ Complete tumour removal is the first and best approach for the body³ This current case report describes a rare case of SPB of the occipital bone with successful radical resection and reconstruction of the occipital bone. *Journal of International Medical Research*

trapezius myocutaneous flap LTMF transplantation Case report A 50-year-old male patient from the Department of Neurosurgery, Hunan Cancer Hospital and the Affiliated Cancer Hospital of Xiangya University, Changsha, Hunan Province, China, presented with a headache and an exophytic mass on the occipital bone. Computed tomography (CT) showed a large mass with homogeneous enhancement on the occipital bone, and the neurological examination was normal. Preoperative imaging examinations showed a preoperative appearance of the mass on CT scan, c) preoperative enhanced magnetic resonance imaging (MRI) scan, d) preoperative digital subtraction angiography (DSA). The colour version of this figure is available at <http://jims.sagepub.com>. The figure revealed a solitary osteolytic lesion involving the whole entire of the occipital bone. The extracerebral expansile osseous lesion (20 mm) mass was mostly isointense on T2-weighted images. Enhanced Figures 1c and 1d Digital subtraction angiography showed hypervascularity that was supplied from the occipital artery. In order to decrease bleeding, a blood vessel was embolized during DSA homogeneously. The patient underwent occipital craniotomy and transplantation under general anaesthesia. The tumour was capsulized and extended to the dura mater with skull defects. Grossly, the tumour had a fishmeat-like appearance, mixed with haemorrhage, and had a rich blood supply. Despite embolization of the main blood supply artery during the operation, the tumour mass underwent extended resection, including the marginal flap window and a 20 cm scalp defect. The skull defect was reconstructed using titanium mesh and the trapezius and the skin island (20 cm) and the supplying vessels of the transverse cervical artery. The trapezius and the skin island (20 cm) was excised and its muscle pedicle dissected up to the occipital bone. The examinations a) the tumour was fishmeat soft tan in appearance, b) the trapezius and transverse cervical artery and the dorsal scapular artery marked out on the skin, c) the skin island flap was set into the defect with a well-perfused distal end, e) the stitches were removed. Postoperative enhanced MRI scan (sagittal view), g) postoperative enhanced MRI scan (axial view), h) postoperative enhanced MRI scan (coronal view). <http://jims.sagepub.com>. 0) rotation point at the medial superior edge of the scapula. c) The LTMF was transposed through the neck posterior subcutaneous tunnel. d) Two weeks after the operation, the transplanted trapezius was undisturbed. Figure 2a shows the removed tumour. Figures 2f and 2g show the MRI completely indicating the presence of atypical plasma cells with typical eccentric round nuclei staining strongly for epithelial membrane antigen (melanin A), CD38, λ CD138. Immunohistochemical *Journal of International Medical Research*

Lambda λ glial fibrillary acidic protein (GFAP), S100, CD68, λ thyroid transcription factor 1 (TTF1), Vim, CD138, radiotherapy for financial reasons. After a follow-up period of around 6 months, he was symptomatic for the 5-month follow-up visit. MRI revealed no residual recurrence but an aggressive mass lesion. Chemotherapy (PAD regimen: bortezomib, pegylated liposomal doxorubicin, idarubicin, melphalan, thalidomide, and prednisone) was administered. Representative photomicrographs of the tumour: a) haematoxylin and eosin (H&E) staining, b) immunohistochemical staining for CD138 showing strong positivity in the tumour cells, c) CD38 showing strong positivity in the tumour cells, d) the positive expression of Ki67. <http://jims.sagepub.com>. Scale bar: mm. 0 cm. Wang et al. Figure 3 shows the magnetic resonance imaging (MRI) scans at the follow-up visit showing no recurrence in situ but an aggressive mass lesion with enhancement. After consecutive cycles of chemotherapy showing no recurrence in situ and the tumour was reduced. Department of Haematology, Hunan Cancer Hospital and the Affiliated Cancer Hospital of Xiangya University, Changsha, Hunan Province, China. After six consecutive cycles of chemotherapy, the tumour was significantly reduced. Figures 4c and 4d show the postoperative review after 6 months, showing no tumour recurrence. This was a case report; the Institutional Review Board of Hunan Cancer Hospital waived the need for informed consent for publication that was approved by the Institutional Review Board. Anonymized Discussion Huge intra and extracranial SPs of the occipital bone are very rare and characterized by the presence of a solitary lytic lesion due to monoclonal plasma cells without account for all SP cases and they occur primarily in red marrow-containing bones.⁶ Radiation therapy remains the only treatment option for SPs. The recommendations from a European expert panel: a total fractionated dose of 40 Gy should be employed.⁶ In this current case, it was unfortunate that the patient refused radiation therapy after the operation and a new mass was found on the right frontal lobe. After six cycles of chemotherapy, the tumour was reduced, which suggests that chemotherapy has a positive impact on the growth of recurrent tumours. The management of SPB in the skull is complex and can easily lead to misdiagnosis. Enhanced CT scan and MRI are credible means by which to diagnose SPB and they could provide more information. The MRI examination allowed for the identification of the location, size, and shape of the tumour and the surrounding structures. In our opinion, preoperative DSA is necessary for the identification of

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embolization During the tumour recurrence current operation the tumour was found to involve the probability of the involved scalp underwent an extended resection LTMF was used to facilitate available muscle compartments transferred on a reliable vascular pedicle to the dorsal sup LTMF include well vascularized tissue ease of harvest and the provision of a large flap local The main blood supply to the LTMF originates from the transverse cervical artery and the dorsal the problem of insufficient blood supply caused by titanium plate implantation In addition subcutaneous cavity created by the huge tumour resection preventing occipital scalp hydrops approach for the repair of scalp and subcutaneous soft tissue defects caused by excision of contributions LW studied the case collected the references and wrote the paper ZH designed the paper XP analysed the data NR served as the first chief during surgery and wrote the final manuscript Declaration of conflicting interest The authors declare that there are no supported by grants from the Scientific Research Project of Hunan Provincial Health Commission Science Foundation of China No 2019JJ40182 and the Sailing Programme of Hunan Province ID Lei Wang References Sabattini E Bacci F Sagrarnoso C et al WHO classification of tumours an overview Pathologica Gee ED and Sadovsky R Multiple myeloma recognition and management Ghehit, 15a K Let al Neurosurgical rare disease solitary O Wang et al plasmacytoma of the Morphol Embryol Chang MY Shih LY Dunn P et al Solitary plasmacytoma of bone J Fore extramedullary plasmacytoma Hematology Am Soc Hematol Educ Program Caers J P response assessment in solitary plasmacytoma updated recommendations from a European Zouhair A Tsang RW et al Prognostic factors in solitary plasmacytoma of the bone a multic Lieboss RH Ha CS Cox JD et al Solitary bone plasmacytoma outcome and prognostic factors Phys Mohos G Vass G Kemeny L et al Extended lower trapezius myocutaneous flap to cover new application J Plast Surg Hand Surg U 15 gurlu K Ozcelik D Hu et al Extended and neck reconstruction as a salvage procedure Plast Reconstr Surg Baek SM Bille myocutaneous flap Ann Plast Surg Netterville JL and Wood DE The lower trapezius flap

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This study aimed to investigate serum matrix metalloproteinase MMP2 and MMP9 levels in PTC Methods Fortyone patients with PTC undergoing ultrasound guided radiofrequency ablation MMP2 and MMP9 levels were determined by enzyme linked immunosorbent assay before and after surgery were evaluated by logistic regression analysis Results Serum MMP2 and MMP9 levels compared with controls and decreased significantly after surgery According to receiver operating characteristic curve values for preoperative serum MMP2 and MMP9 levels were 0.85 and 0.82 respectively There was no contrast enhancement within or at the lesion edge in The volume reduction at months followup was 45.2% diameter and number were influencing factors for PTC Age and lesion diameter and number and morphology were protective factors Conclusion Serum MMP2 and MMP9 levels before and treatment of PTC by RFA Preoperative serum MMP2 and MMP9 levels combined with ultrasound guided radiofrequency ablation can predict the prognosis Department of Ultrasound Beilun Peoples Hospital of Ningbo Beilun Branch of the First Affiliated Hospital of Ningbo Zhejiang China These authors contributed equally to this study Corresponding author: Dr. Qiang Ding, Department of Ultrasound, Beilun Peoples Hospital of Ningbo Beilun Branch of the First Affiliated Hospital of Zhejiang University, 315000, China Email: qiangding02@sohu.com Creative Commons Non Commercial CC BY-NC This is distributed under the Attribution Non Commercial License creative commons licenses/by-nc/4.0 which permits non commercial work without further permission provided the original work is attributed as specified on the SAGE Publishing website access at sagepub.com/journalPermissions/permissions.html Journal of International Medical Research Keywords Papillary thyroid carcinoma PTC is a common thyroid malignancy accounting for about 60% of systemic malignant thyroid tumors with high incidence and low mortality 2 clinical Recent improvements in high frequency ultrasound guided puncture techniques have led to an apparent increase in the incidence of thyroid microtumors in the general population is about 10% compared with 1% in the past which has thus greatly improved disease diagnosis diagnosis of thyroid microtumors by ultrasound is usually high and the consequent reduction in thyroid function can seriously affect the increasing detection rate of thyroid tumors and the pursuit of minimally invasive treatment

gradually applied in the clinic RFA uses local hyperthermia to cause tissue necrosis The t and most lesions can be completely eliminated by RFA 89 RFA has thus become a novel loc benign and malignant thyroid tumors currently depend on the clinical manifestation clinical manifestations are mostly derived from involvement subjective empirical analysis while a i with less satisfactory specificity It is therefore necessary to identify appropriate predictive collagenases matrix metalloproteinase MMP2 and MMP9 can degrade ty important tumor angiogenesis and tumor cell invasion and metastasis 10 MMP2 and MMP9 ex thyroid cancer tissue 11 however these studies mostly examined pathological tissues after inva serum levels of MMP2 and MMP9 have been less well considered In this study we detected se PTC before and after ultrasound-guided RFA We also determined the therapeutic effects of RFA relevant prognostic factors Materials and methods Study subjects Patients who underwent to October were included in this study The inclusion criteria were as follows patients dia cytology no history of neck surgery and patients requiring minimally invasive treatment for a es OcPan et al with anxiety The exclusion criteria were as follows benign lesions confirmed by surgery and severe coagulopathy Peripheral venous blood samples were obtained from the i the operation and serum levels of MMP2 and MMP9 were determined Additional subjects RFA were included as a control group Prior written informed consent was obtained from all pa review board of our hospital Preoperative preparation calcium i-cation The number size nature (nodular blood i-cow distribution of the tumors were assessed before the operation After skin d lidocaine solution A total of mL Sonovue Bracco Milan Italy was injected via the elbow vei evaluated by contrast-enhanced ultrasound CEUS of the ablation targeted lesions using a Myla Shenzhen Guangdong China According to the location of the thyroid nodules the thyroid and c and esophageal space and posterior thyroid space recurrent laryngeal nerve were separate the intraoperative conditions to form a liquid separation zone to protect these structures ultrasound guidance the tip of the RFA needle rated power W output frequency kHz was was performed using an Olympus Celon Power RFA System Germany in mobile mode 12 followi were subjected to multipointed and multifaceted ablation until the thyroid tissue layer with the echogenerated by heat accumulation The whole process was carried out under contin produced in the ablation zone during the ablation treatment The position of the electrode lesion size After ensuring that there was no residual enhancement in the ablation zone the abl completed After the operation an ice compress was applied for h to avoid skin burns Serum blood was collected from the elbow vein under fasting conditions before and after control group were collected after ultrasound contrast examination The blood samples were subjected to centrifugation at 02 g for minutes The serum was harvested and serum lev enzymelinked immunosorbent ELISA kits Boster Bioengineering Wuhan Hubei China assay Fol the operation the ablation range was evaluated by CEUS If residual tissues were detected detection was performed at and months after surgery to determine the nodule sizes and volum according to the following formula volume reduction rate $\frac{1}{4}$ preoperative volume foll blood i-cow changes in the ablation zone were also observed and analyzed The effi-cacy i treating tumors 13 disappearance of nodules indicated by complete disappearance of blood i complete cure nodule by 15 indicated volumemarked ei improvement reduced Clinicopathological features Information on ultrasound-based clinicopath of the lesion were obtained features version Statistical analysis Data were expressed as mea performed using IBM SPSS Statistics for Windows IBM Corp Armonk NY USA Com v2 tests Potentially related factors were analyzed by univariate or multivariate logistic regre: MMP2 and MMP9 levels were evaluated by receiver operating characteristic ROC signii-cant Results Patients Journal of International Medical Research men mean age 06 year before and after treatment The characteristics of the ultrasound images in the included subj and MMP9 were measured before and after treatment Serum levels of MMP2 and MMP9 compared with the control subjects P Serum levels of MMP2 and MMP9 had declined at surgery but the difference was not signii-cant However serum levels of MMP2 and MMP9 ha and months all P Table These results suggest that changes in serum MMP2 and MMP9 lev signii-cant implications for the therapy of PTC ROC curve analysis of preoperative serum MMP and MMP9 levels were used as potential diagnostic indicators In the patients with PTC the was used as the diagnostic results and the gold standard classification criteria were us obtained accordingly The area under curve AUC values for serum MMP and MMP9 lev suggest that serum levels of MMP2 and MMP9 could contribute to the disease dia enrolled including women and men mean age 06 years range to 65 years The c benign thyroid nodules including women and Evaluation of RFA efficacy We also evaluated t ablation showed hypoenhancement in nodules isoenhancement in nodules and slight hyper thyroid ultrasound images PTC patients Normal control Le cm Calcification Microcalcification Coarse calcification Morphology Regular Irregular A

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showing no obvious enhancement in the lesion with lowerperfusion performance c Inthe 2
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alDiscussionitis difi~cultPTC is a type of thyroid tumor with a highincidence14 wl
Mostthyroidtumor cases are currently diagnosed by hiscytological detectiontopathologic
malignant papillaryhyperplastic nodules and it is therefore difi~cult to diagnose PT
prognostic molecular markers for PTC17 The relationshipbetween MMPs and tumors
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alternatingelectromagnetic wave only circulates in theeffective region between the two
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was thus relatively greaterWe analyzed the serum levels of MMP2and MMP9 in PTC patie
enzymeswere signii~cantly higher in patients withPTC compared with patients with ber
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MMP2 and MMP9 were secretedby the tumor The lesions disappearedafter PTC ablation t
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summary the results of this studys showed that RFA could shrink or eliminatethyroid lesions t
effective methodforserumlevels of MMP2 and MMP9 before RFAcould provide a val
serological indexes combined with relevant risk factors mayalso help to predict the prognos

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 focused ultrasound HIFU ablation of benign thyroid nodules: a retrospective analysis. Int J Hyp
 Can malignant thyroid nodules be distinguished from benign thyroid nodules in children and ad
 pe

```
In [7]: #Rename the column names
data.rename({'0': 'Label', 'a': 'Text'}, axis=1, inplace=True)
```

EDA Process

```
In [8]: #Check the data columns
data.columns
```

```
Out[8]: Index(['Label', 'Text'], dtype='object')
```

```
In [9]: data[data.Label=='Lung_Cancer'].Text[539]
```

Out[9]: "We lack biomarkers for identifying aggressive primary tumor subsets that give rise to metastases and impact early cancer detection and treatment. Many solid tumors are known to accumulate hyaluronan (HA) a glycosaminoglycan which is also produced by the tumor cells themselves. We report a quantitative approach for uncovering breast cancer heterogeneity using fluorescent HA to detect differential binding patterns to CD44 and RHAMM/HMMR receptors. This approach permits identification of tumor-cell subsets that bind high levels of HA and may be applicable to other ligands/receptors and disease models. Despite representing the invasive/metastatic subset of parental tumors unexpectedly the high HA-binding subset was slow-growing and is thus likely to be a source of dormancy and relapse. Tumor heterogeneity confounds cancer diagnosis and the outcome of therapy necessitating analysis of tumor cell subsets within the tumor mass. Elevated expression of hyaluronan (HA) and HA receptors receptor for HA-mediated motility (RHAMM)/HA-mediated motility receptor and cluster designation 44 (CD44) in breast tumors correlates with poor outcome. We hypothesized that a probe for detecting HA \times HA receptor interactions may reveal breast cancer (BCa) cell heterogeneity relevant to tumor progression. A fluorescent HA (F-HA) probe containing a mixture of polymer sizes typical of tumor microenvironments (10 \times 93480 kDa) multiplexed profiling and flow cytometry were used to monitor HA binding to BCa cell lines of different molecular subtypes. Formulae were developed to quantify binding heterogeneity and to measure invasion in vivo. Two subsets exhibiting differential binding (HA \uparrow /low vs. HA \uparrow high) were isolated and characterized for morphology growth and invasion in culture and as xenografts in vivo. F-HA \times binding amounts and degree of heterogeneity varied with BCa subtype were highest in the malignant basal-like cell lines and decreased upon reversion to a nonmalignant phenotype. Binding amounts correlated with CD44 and RHAMM displayed but binding heterogeneity appeared to arise from a differential ability of HA receptor-positive subpopulations to interact with F-HA. HA \uparrow high subpopulations exhibited significantly higher local invasion and lung micrometastases but unexpectedly lower proliferation than either unsorted parental cells or the HA \uparrow /low subpopulation. Querying F-HA binding to aggressive tumor cells reveals a previously undetected form of heterogeneity that predicts invasive/metastatic behavior and that may aid both early identification of cancer patients susceptible to metastasis and detection/therapy of invasive BCa subpopulations. tumor cell heterogeneity hyaluronan binding heterogeneity index PLoS One one 1932-6203 Public Library of Science San Francisco USA 24454921 38932 58 PONE-D-13-41069 .0085702 Research Biology Biochemistry Bioenergetics Energy-Producing Processes Metabolism Carbohydrate Metabolism Metabolic Pathways Oxygen Metabolism Protein Metabolism Cofactors Drug Discovery Enzymes Genetics Gene Expression Medicine Drugs and Devices Drug Research and Development Drug Discovery Hematology Hematologic Cancers and Related Disorders Leukemias Acute Lymphoblastic Leukemia Nutrition Obstetrics and Gynecology Breast Cancer Oncology Cancers and Neoplasms Hematologic Cancers and Related Disorders Leukemias Breast Tumors Oncology Agents Metabolic Effects of Acute Thiamine Depletion Are Reversed by Rapamycin in Breast and Leukemia Cells Thiamine Depletion and Metabolism in Cancer Cells Liu Shuqian 1 Miriyala Sumitra 2 Keaton Mignon A. 3 Jordan Craig T. 4 Wiedl Christina 5 Clair Daret K. St. 2 Moscow Jeffrey A. 1 * 1 Department of Pediatrics University of Kentucky College of Medicine Lexington Kentucky United States of America 2 Graduate Center for Toxicology University of Kentucky College of Medicine Lexington Kentucky United States of America 3 Metabolon Inc Durham North Carolina United States of America 4 Division of Hematology Hematologic Malignancies and Stem Cell Transplantation University of Colorado Denver Colorado United States of America 5 Department of Pediatrics Virginia Commonwealth University Richmond Virginia United States of America Ahmad Aamir Editor Wayne State University School of Medicine United States of America * E-mail: jmoscow@uky.edu Competing Interests: One of the authors of this paper Mignon A. Keaton was employed by Metabolon Inc. during the data acquisition and analysis phases of the study. Dr. Keaton is no longer employed by Metabolon. Her employment history does not alter the authors' adherence to all the PLOS ONE policies on sharing data and materials. Conceived and designed the

experiments: MAK CTJ DKS JAM. Performed the experiments: SL SM MAK CTJ CW. Analyzed the data: MAK DKS JAM. Contributed reagents/materials/analysis tools: SM MAK CTJ CW DKS JAM. Wrote the paper: MAK CTJ JAM. 2014 15 1 2014 9 1 e85702 8 10 2013 5 12 2013 2014 Liu et al This is an open-access distributed under the terms of the Creative Commons Attribution License which permits unrestricted use distribution and reproduction in any medium provided the original author and source are credited. Thiamine-dependent enzymes (TDEs) control metabolic pathways that are frequently altered in cancer and therefore present cancer-relevant targets. We have previously shown that the recombinant enzyme thiaminase cleaves and depletes intracellular thiamine has growth inhibitory activity against leukemia and breast cancer cell lines and that its growth inhibitory effects were reversed in leukemia cell lines by rapamycin. Now we first show further evidence of thiaminase therapeutic potential by demonstrating its activity against breast and leukemia xenografts and against a primary leukemia xenograft. We therefore further explored the metabolic effects of thiaminase in combination with rapamycin in leukemia and breast cell lines. Thiaminase decreased oxygen consumption rate and increased extracellular acidification rate consistent with the inhibitory effect of acute thiamine depletion on the activity of the TDEs pyruvate dehydrogenase and 2-oxoglutarate dehydrogenase complexes; these effects were reversed by rapamycin. Metabolomic studies demonstrated intracellular thiamine depletion and the presence of the thiazole cleavage product in thiaminase-treated cells providing validation of the experimental procedures. Accumulation of ribose and ribulose in both cell lines support the thiaminase-mediated suppression of the TDE transketolase. Interestingly thiaminase suppression of another TDE branched chain amino ketoacid dehydrogenase (BCKDH) showed very different patterns in the two cell lines: in RS4 leukemia cells it led to an increase in BCKDH substrates and in MCF-7 breast cancer cells it led to a decrease in BCKDH products. Immunoblot analyses showed corresponding differences in expression of BCKDH pathway enzymes and partial protection of thiaminase growth inhibition by gabapentin indicated that BCKDH inhibition may be a mechanism of thiaminase-mediated toxicity. Surprisingly most of thiaminase-mediated metabolomic effects were also reversed by rapamycin. Thus these studies demonstrate that acute intracellular thiamine depletion by recombinant thiaminase results in metabolic changes in thiamine-dependent metabolism and demonstrate a previously unrecognized role of mTOR signaling in the regulation of thiamine-dependent metabolism. No current external funding sources for this study.

Introduction Thiamine (vitamin B1) is a cofactor for enzymes involved in critical metabolic processes involving energy production biomass generation and amino acid catabolism. Despite the requirement for this vitamin in these central processes the role of thiamine and thiamine-dependent enzymes (TDEs) in cancer development and treatment has received little attention although a recent review has summarized the potential importance of TDEs in cancer metabolism [1]. Unlike antifolates which have a well-established role in cancer therapy a analogous small molecule thiamine antagonists are relatively inert leading to a that TDE pathways could not be important as anticancer targets. However the limitations of small molecule TDE inhibitors should not be confused with the potential role of TDEs as anticancer therapeutic targets. Antifolates can be effective because intracellular folates only transiently associate with enzymes during the catalytic process allowing for inhibition of enzyme activity by molecules designed to bind more tightly than the intracellular substrates. In contrast intracellular thiamine activated by phosphorylation remains tightly bound to enzyme complexes during the catalytic cycle leaving little opportunity for inhibitors to displace it once the complex has assembled. This inherent pharmacologic challenge could disguise the potential of targeting TDEs for cancer therapy. We have previously shown down-regulation of thiamine transporter gene expression in tumors compared to normal tissues [2] [3] and more recently have shown that a low thiamine diet delays onset of mammary tumors in MMTV(her2) mice [4] an effect that is abrogated by a high fat diet. These observations have led to our hypothesis that TDE pathways are altered as part of the overall changes in energy metabolism that occurs in cancer cells and that these changes c

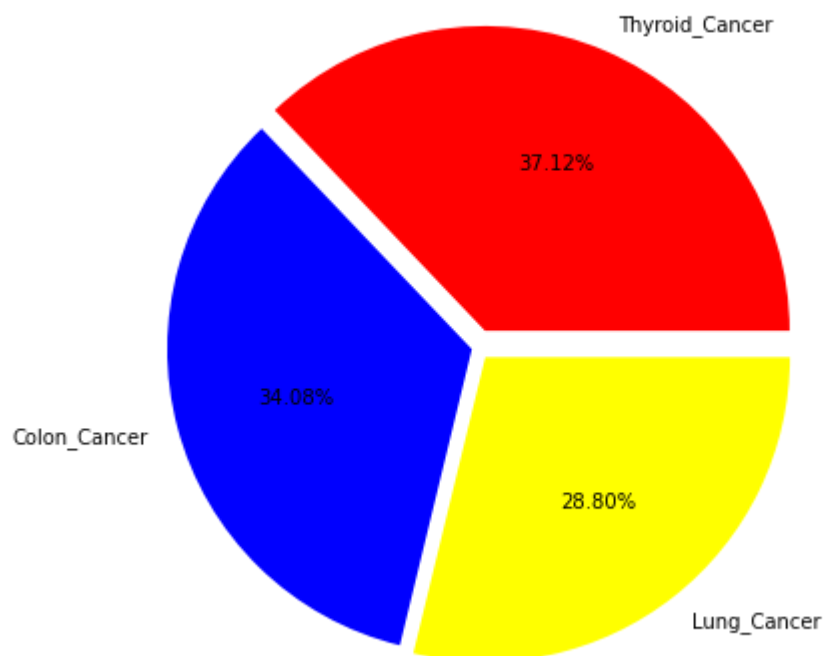
ould produce metabolic vulnerabilities that could be exploited by therapies aimed at TDE activities. To take a novel path in the exploration of TDEs in cancer we have studied the cytotoxic activity of the bacterial enzyme thiaminase which cleaves thiamine into its pyrimidine and thiazole moieties [5]. Thiaminase overcomes the limitations of small molecule TDE "

```
In [10]: #Let's count the label values in the dataset
label=data['Label'].value_counts()
label
```

```
Out[10]: Thyroid_Cancer    2810
Colon_Cancer    2580
Lung_Cancer    2180
Name: Label, dtype: int64
```

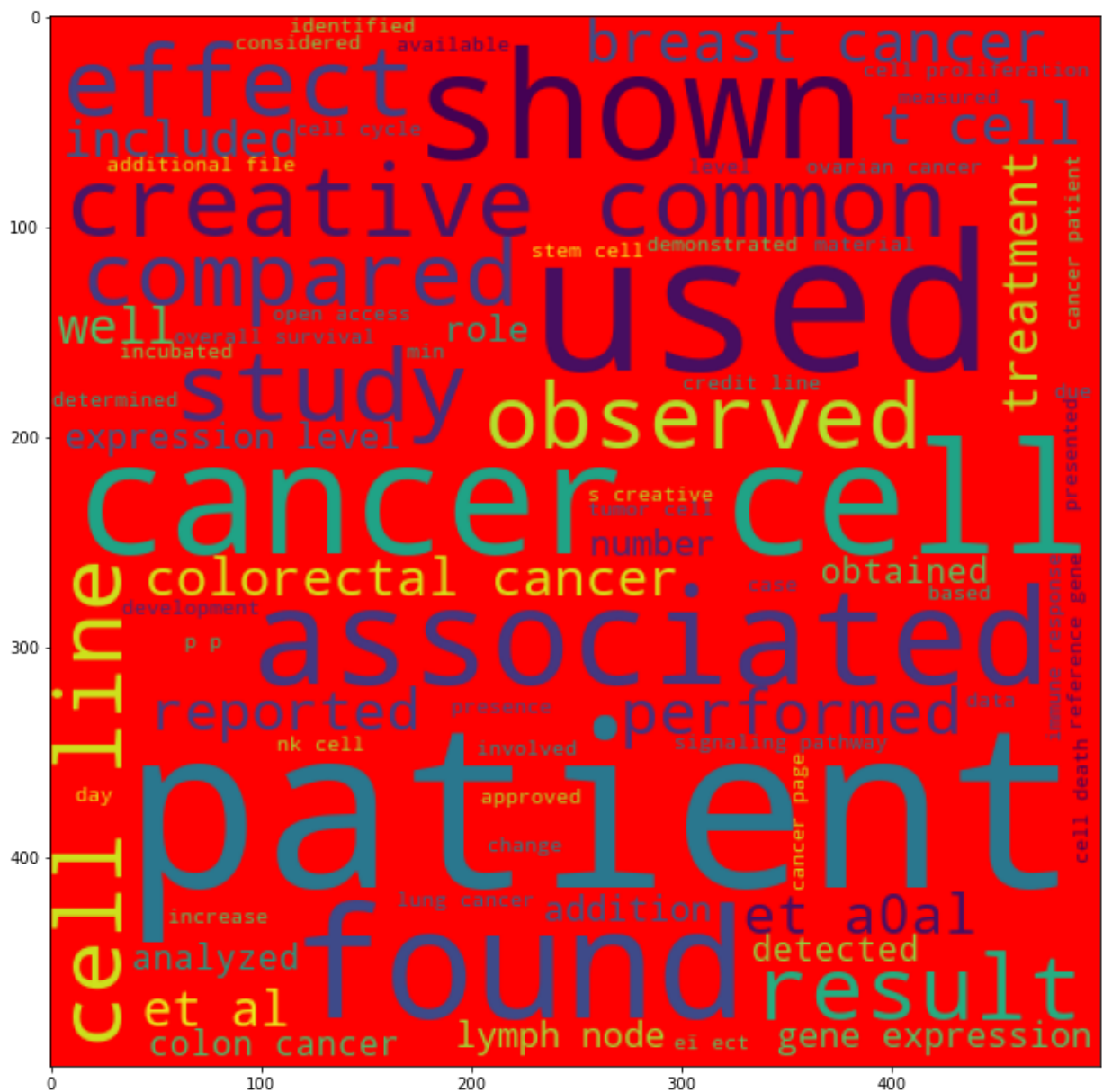
```
In [11]: #Let's visualize the above information on the dataset
plt.figure(figsize=(19,7))
plt.pie(label,
        labels=['Thyroid_Cancer','Colon_Cancer','Lung_Cancer'],
        colors=['Red','blue','yellow'],
        autopct='%1.2f%%',explode=[0.06,0.02,0.04])
plt.title("The label percentage in the dataset",fontsize=32)
plt.show()
```

The label percentage in the dataset



```
In [12]: #Visualize the WordCloud in the Thyroid_Cancer from the dataset
plt.figure(figsize=(12,15))
wc=WordCloud(height=500,width=500,min_font_size=10,background_color='blue')
w_c=wc.generate(data[data['Label']=='Thyroid_Cancer']['Text'].str.cat(sep=" "))
plt.imshow(w_c)
```

```
Out[12]: <matplotlib.image.AxesImage at 0x7fc925a8b5e0>
```

```
In [14]: #Visualize the WordCloud in the Colon_Cancer from the dataset
plt.figure(figsize=(12,15))
wc=WordCloud(height=500,width=500,min_font_size=10,background_color='green')
wc_2=wc.generate(data[data['Label']=='Lung_Cancer']['Text'].str.cat(sep=" "))
plt.imshow(wc_2)
```

```
Out[14]: <matplotlib.image.AxesImage at 0x7fc919dd5fd0>
```



```
In [16]: #Define the function to clean the text
def clean_text(text):
    pattern = r'^a-zA-Z\s]'
    text=re.sub(pattern,'',text)
    return text
#Apply to the function to the dataset
data['Text']=data['Text'].apply(clean_text)
```

Remove the stopwords

```
In [17]: #Create function to the remove the stopwords
names = ['Colon_Cancer', 'Lung_Cancer', 'Thyroid_Cancer']
def clean_stop(text):
    stop_words = stopwords.words('english')
    for name in names:
        stop_words.append(name)
    return " ".join([w.lower() for w in text.split() if w.lower() not in stop_
#And finally apply the above function to the dataset
data['Text']=data['Text'].apply(clean_stop)
```

Tokenization

```
In [18]: #Define the tokenize function
def tokenize(d):
    return word_tokenize(d)
data['Text']=data['Text'].apply(tokenize)
```

```
In [19]: data.head()
```

```
Out[19]:
```

	Label	Text
0	Thyroid_Cancer	[thyroid, surgery, children, single, instituti...
1	Thyroid_Cancer	[adopted, strategy, used, prior, years, based,...
2	Thyroid_Cancer	[coronary, arterybypass, grafting, thrombosis,...
3	Thyroid_Cancer	[solitary, plasmacytoma, sp, skull, uncommon, ...
4	Thyroid_Cancer	[study, aimed, investigate, serum, matrix, met...

Remove the special characters

```
In [20]: #removing special character
def remove_special_char(list):
    y=[]
    for string in list:
        if string.isalnum():
            y.append(string)
    return y
data['Text']=data['Text'].apply(lambda x: remove_special_char(x))
```


In [21]: `data.head()`

Out[21]:

	Label	Text
0	Thyroid_Cancer	[thyroid, surgery, children, single, instituti...
1	Thyroid_Cancer	[adopted, strategy, used, prior, years, based,...
2	Thyroid_Cancer	[coronary, arterybypass, grafting, thrombosis,...
3	Thyroid_Cancer	[solitary, plasmacytoma, sp, skull, uncommon, ...
4	Thyroid_Cancer	[study, aimed, investigate, serum, matrix, met...

Stemming process

In [22]:

```
#Create a function to the stemming processs
ps=PorterStemmer()
def stemming(list):
    #Create a empty list
    y=[]
    #Create a for loop for text in list
    for text in list:
        #Then i finallay append to the empty list
        y.append(ps.stem(text))
    #then return to the empty list
    return y
data['Text']=data['Text'].apply(lambda x:stemming(x))
#join the words
data['Text']=data['Text'].apply(lambda x:" ".join(x))
```

In [23]: `data.head()`

Out[23]:

	Label	Text
0	Thyroid_Cancer	thyroid surgeri children singl institut osama ...
1	Thyroid_Cancer	adopt strategi use prior year base four exclus...
2	Thyroid_Cancer	coronari arterybypass graft thrombosi brin bri...
3	Thyroid_Cancer	solitari plasmacytoma sp skull uncommon clinic...
4	Thyroid_Cancer	studi aim investig serum matrix metalloprotein...

Modeling

In [24]:

```
#Divided the data into two variables
X=data['Text']
y=data['Label']
```

In [25]:

```
#install the TfidfVectorizer
vector=TfidfVectorizer()
#Fit the X data to the TfidfVectorizer
vector.fit(X)
```

```
#And transform the
X=vector.transform(X)
```

```
In [26]: #And Divided the data into traing and testing and finally split the data and
X_train,X_test,y_train,y_test=train_test_split(X,y,test_size=0.25,random_state
```

LogisticRegression

```
In [27]: #Install the logisticregression model
logistic=LogisticRegression()
#And fit the model to the train data
logistic.fit(X_train,y_train)
```

```
Out[27]: LogisticRegression()
```

```
In [28]: #Prediction of the logisticregression algorithm
logistic_pred=logistic.predict(X_test)
logistic_pred
```

```
Out[28]: array(['Colon_Cancer', 'Colon_Cancer', 'Colon_Cancer', ..., 'Lung_Cancer',
                'Thyroid_Cancer', 'Colon_Cancer'], dtype=object)
```

```
In [29]: from sklearn.metrics import classification_report,accuracy_score,confusion_mat
```

```
In [30]: #Check the test score and train score to the logisticregression algorithm
print(f'The Test_accuracy: {logistic.score(X_test,y_test)*100:.2f}')
#Train score for the data
print(f'The Train_accuracy: {logistic.score(X_train,y_train)*100:.2f}')
#Check the accuracy_score to the model
print(f'The Accuracy_score: {accuracy_score(y_test,logistic_pred)*100:.2f}')
```

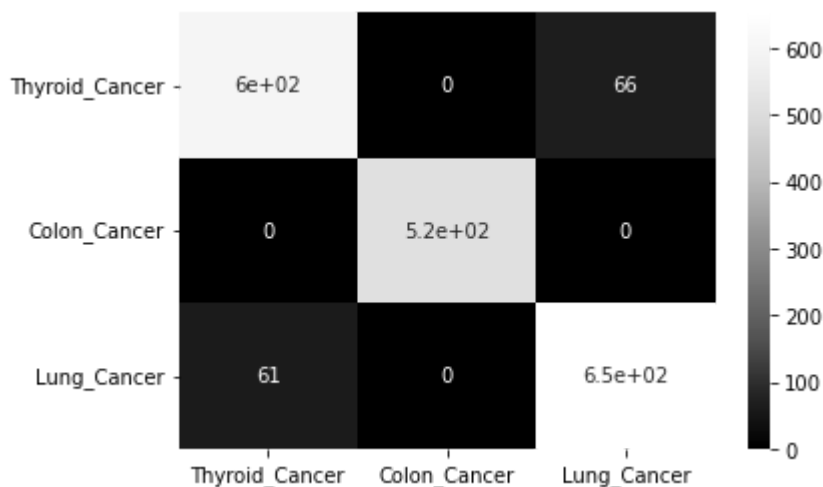
```
The Test_accuracy: 93.29
The Train_accuracy: 95.90
The Accuracy_score: 93.29
```

Classification_report and Confusion_matrix

```
In [31]: #Classification report
print(classification_report(y_test,logistic_pred))
#confusion matrix
cn=confusion_matrix(y_test,logistic_pred)
sns.heatmap(cn,annot=True,cmap='Greys_r',xticklabels=['Thyroid_Cancer','Colon_
```

	precision	recall	f1-score	support
Colon_Cancer	0.91	0.90	0.90	664
Lung_Cancer	1.00	1.00	1.00	515
Thyroid_Cancer	0.91	0.91	0.91	714
accuracy			0.93	1893
macro avg	0.94	0.94	0.94	1893
weighted avg	0.93	0.93	0.93	1893

```
Out[31]: <AxesSubplot:>
```



DecisionTreeClassifier

```
In [32]: #Import the DecisionTreeClassifier algorithm
from sklearn.tree import DecisionTreeClassifier
#install the DecisionTreeClassifier model
tree=DecisionTreeClassifier()
#Fit the train data to the model
tree.fit(X_train,y_train)
```

```
Out[32]: DecisionTreeClassifier()
```

```
In [33]: #Prediction of the DecisionTreeClassifier algorithm
tree_pred=tree.predict(X_test)
tree_pred
```

```
Out[33]: array(['Colon_Cancer', 'Colon_Cancer', 'Thyroid_Cancer', ...,
                'Lung_Cancer', 'Thyroid_Cancer', 'Colon_Cancer'], dtype=object)
```

```
In [34]: #Check the test score and train score to the DecisionTreeClassifier algorithm
print(f'The Test_accuracy: {tree.score(X_test,y_test)*100:.2f}')
```

```
#Train score for the data
print(f'The Train_accuracy: {tree.score(X_train,y_train)*100:.2f}')
```

```
#Check the accuracy_score to the model
print(f'The Accuracy_score: {accuracy_score(y_test,tree_pred)*100:.2f}')
```

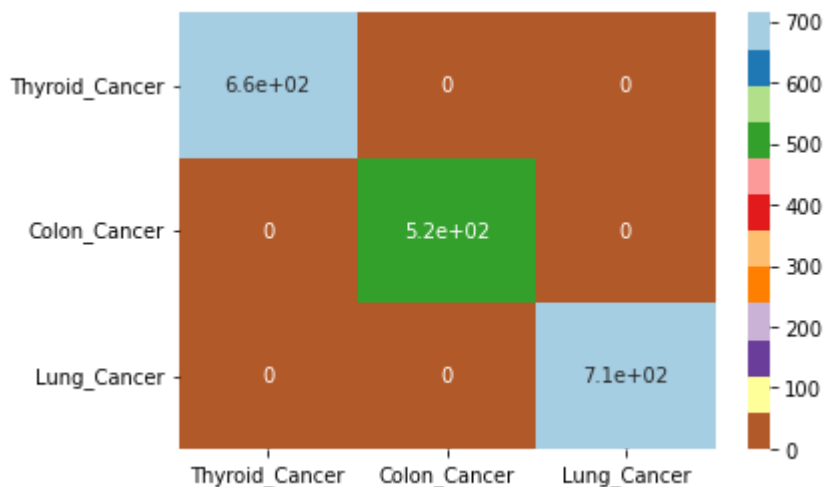
```
The Test_accuracy: 100.00
The Train_accuracy: 100.00
The Accuracy_score: 100.00
```

Classification_report and Confusion_matrix

```
In [35]: #Classification report
print(classification_report(y_test,tree_pred))
#confusion matrix
cn=confusion_matrix(y_test,tree_pred)
sns.heatmap(cn,annot=True,cmap='Paired_r',xticklabels=['Thyroid_Cancer','Colon
```

	precision	recall	f1-score	support
Colon_Cancer	1.00	1.00	1.00	664
Lung_Cancer	1.00	1.00	1.00	515
Thyroid_Cancer	1.00	1.00	1.00	714
accuracy			1.00	1893
macro avg	1.00	1.00	1.00	1893
weighted avg	1.00	1.00	1.00	1893

Out[35]: <AxesSubplot:>



RandomForestClassifier

```
In [36]: #Import the RandomForestClassifier algorithm
from sklearn.ensemble import RandomForestClassifier
#install the DecisionTreeClassifier model
random=RandomForestClassifier()
#Fit the train data to the model
random.fit(X_train,y_train)
```

Out[36]: RandomForestClassifier()

```
In [37]: #Prediction of the RandomForestClassifier algorithm
random_pred=random.predict(X_test)
random_pred
```

Out[37]: array(['Colon_Cancer', 'Colon_Cancer', 'Thyroid_Cancer', ...,
'Lung_Cancer', 'Thyroid_Cancer', 'Colon_Cancer'], dtype=object)

```
In [38]: #Check the test score and train score to the RandomForestClassifier algorithm
print(f'The Test_accuracy: {random.score(X_test,y_test)*100:.2f}')
```

```
#Train score for the data
print(f'The Train_accuracy: {random.score(X_train,y_train)*100:.2f}')
```

```
#Check the accuracy_score to the model
print(f'The Accuracy_score: {accuracy_score(y_test,random_pred)*100:.2f}')
```

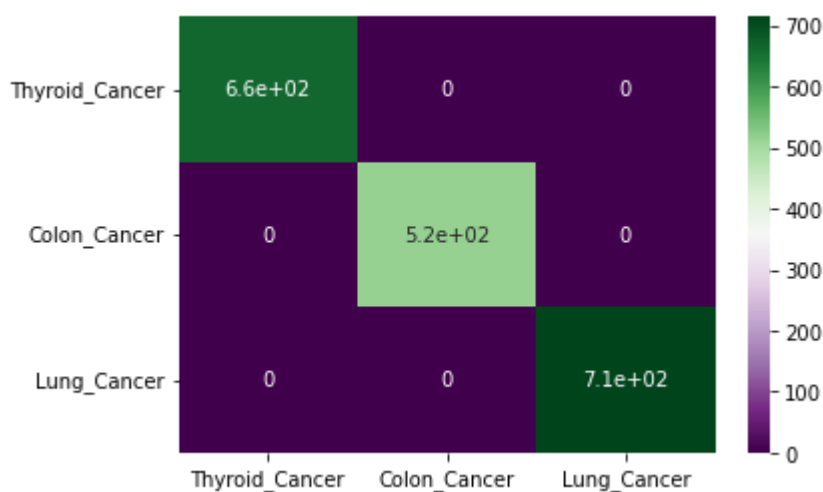
The Test_accuracy: 100.00
The Train_accuracy: 100.00
The Accuracy_score: 100.00

Classification_report and Confusion_matrix

```
In [39]: #Classification report
print(classification_report(y_test,random_pred))
#confusion_matrix
cn=confusion_matrix(y_test,random_pred)
sns.heatmap(cn,annot=True,cmap='PRGn',xticklabels=['Thyroid_Cancer','Colon_Can
```

	precision	recall	f1-score	support
Colon_Cancer	1.00	1.00	1.00	664
Lung_Cancer	1.00	1.00	1.00	515
Thyroid_Cancer	1.00	1.00	1.00	714
accuracy			1.00	1893
macro avg	1.00	1.00	1.00	1893
weighted avg	1.00	1.00	1.00	1893

Out[39]: <AxesSubplot:>



MultinomialNB

```
In [40]: #Import the MultinomialNB algorithm to train the our model
from sklearn.naive_bayes import MultinomialNB
#install the model
multinomial=MultinomialNB()
#fit the train data to our model
multinomial.fit(X_train,y_train)
```

Out[40]: MultinomialNB()

```
In [41]: #Prediction to the test data MultinomialNB
multinomial_pred=multinomial.predict(X_test)
multinomial_pred
```

Out[41]: array(['Colon_Cancer', 'Colon_Cancer', 'Thyroid_Cancer', ...,
'Lung_Cancer', 'Thyroid_Cancer', 'Colon_Cancer'], dtype='<U14')

```
In [42]: #Check the test score and train score to the MultinomialNB algorithm
print(f'The Test_accuracy: {multinomial.score(X_test,y_test)*100:.2f}')
```

```
#Train score for the data
print(f'The Train_accuracy: {multinomial.score(X_train,y_train)*100:.2f}')
#Check the accuracy_score to the model
print(f'The Accuracy_score: {accuracy_score(y_test,multinomial_pred)*100:.2f}')

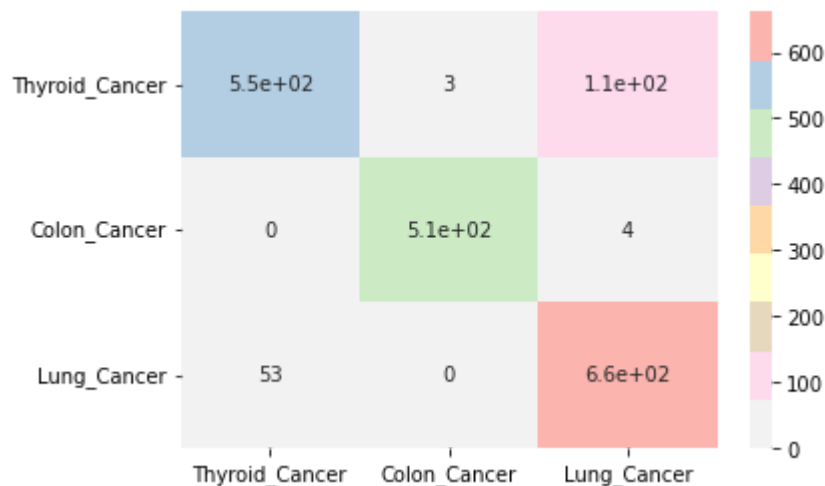
The Test_accuracy: 90.86
The Train_accuracy: 93.62
The Accuracy_score: 90.86
```

Classification_report and Confusion_matix

```
In [43]: #Classification report
print(classification_report(y_test,multinomial_pred))
#confusion_matrix
cn=confusion_matrix(y_test,multinomial_pred)
sns.heatmap(cn,annot=True,cmap='Pastell_r',xticklabels=['Thyroid_Cancer','Colo
```

	precision	recall	f1-score	support
Colon_Cancer	0.91	0.83	0.87	664
Lung_Cancer	0.99	0.99	0.99	515
Thyroid_Cancer	0.85	0.93	0.89	714
accuracy			0.91	1893
macro avg	0.92	0.91	0.92	1893
weighted avg	0.91	0.91	0.91	1893

Out[43]: <AxesSubplot:>



XGBClassifier

```
In [44]: #Import the XGBClassifier model and install the model
from xgboost import XGBClassifier
#install the XGBClassifier
xgb=XGBClassifier()
#And finally fit the data to train data
xgb.fit(X_train,y_train)
```

```
Out[44]: XGBClassifier(base_score=0.5, booster=None, colsample_bylevel=1,
                colsample_bynode=1, colsample_bytree=1, gamma=0, gpu_id=-1,
                importance_type='gain', interaction_constraints=None,
                learning_rate=0.300000012, max_delta_step=0, max_depth=6,
                min_child_weight=1, missing=nan, monotone_constraints=None,
                n_estimators=100, n_jobs=0, num_parallel_tree=1,
                objective='multi:softprob', random_state=0, reg_alpha=0,
                reg_lambda=1, scale_pos_weight=None, subsample=1,
                tree_method=None, validate_parameters=False, verbosity=None)
```

```
In [45]: #Prediction to the test data XGBClassifier
xgb_pred=xgb.predict(X_test)
xgb_pred
```

```
Out[45]: array(['Colon_Cancer', 'Colon_Cancer', 'Thyroid_Cancer', ...,
                'Lung_Cancer', 'Thyroid_Cancer', 'Colon_Cancer'], dtype=object)
```

```
In [46]: #Check the test score and train score to the XGBClassifier algorithm
print(f'The Test_accuracy: {xgb.score(X_test,y_test)*100:.2f}')
#Train score for the data
print(f'The Train_accuracy: {xgb.score(X_train,y_train)*100:.2f}')
#Check the accuracy_score to the model
print(f'The Accuracy_score: {accuracy_score(y_test,xgb_pred)*100:.2f}')

```

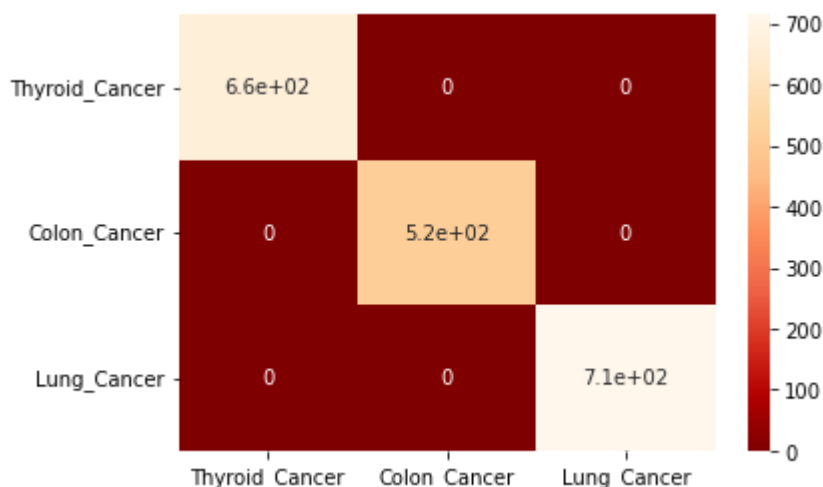
```
The Test_accuracy: 100.00
The Train_accuracy: 100.00
The Accuracy_score: 100.00
```

Classification_report and Confusion_matrix

```
In [47]: #Classification report
print(classification_report(y_test,xgb_pred))
#confusion matrix
cn=confusion_matrix(y_test,xgb_pred)
sns.heatmap(cn,annot=True,cmap='OrRd_r',xticklabels=['Thyroid_Cancer','Colon_C
```

	precision	recall	f1-score	support
Colon_Cancer	1.00	1.00	1.00	664
Lung_Cancer	1.00	1.00	1.00	515
Thyroid_Cancer	1.00	1.00	1.00	714
accuracy			1.00	1893
macro avg	1.00	1.00	1.00	1893
weighted avg	1.00	1.00	1.00	1893

```
Out[47]: <AxesSubplot:>
```



Deep Learning models

```
In [48]: #Import the necessary librairys
from tensorflow.keras.utils import to_categorical
from gensim.models import Word2Vec
from gensim.models.keyedvectors import KeyedVectors
import time
from keras.layers import Dense, Input, Flatten, LSTM, Bidirectional, Embedding,
from keras.layers import Conv1D, MaxPooling1D, Embedding
from keras.models import Sequential, load_model
from keras import losses
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.models import Model
from keras.utils import pad_sequences
from keras.utils.vis_utils import plot_model
from keras.callbacks import EarlyStopping
from keras.preprocessing.text import Tokenizer
```

```
In [61]: #Create the dummies values to the dataset and divided the data train test split
y = pd.get_dummies(data.Label)
X_trn, X_tst, y_trn, y_tst = train_test_split(X, y, test_size=0.2, random_state=42)
X_trn, X_val, y_trn, y_val = train_test_split(X_trn, y_trn, test_size=0.3, random_state=42)
```

```
In [50]: #Creat the max_word and max_len variables
max_words = 5000
max_len = 300
#Then create a function for the padding the text data
def tokenize_pad_sequences(text):
    # Text tokenization
    tokenizer = Tokenizer(num_words=max_words, lower=True, split=' ')
    tokenizer.fit_on_texts(text)
    # Transforms text to a sequence of integers
    X = tokenizer.texts_to_sequences(text)
    # Pad sequences to the same length
    X = pad_sequences(X, padding='post', maxlen=max_len)
    # return sequences
    return X, tokenizer

print('Before Tokenization & Padding \n', data['Text'][0], '\n')
```



```
X, tokenizer = tokenize_pad_sequences(data['Text'])  
print('After Tokenization & Padding \n', X[0])
```

Before Tokenization & Padding

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After Tokenization & Padding

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```

```
In [51]: #Vocab_size
vocab_size = 5000
#Embeddig_size
embedding_size = 32
epochs=50
#Let's install the Sequential model
model= Sequential()
#then add the Embedding to the model with the vocab_size and input_length
model.add(Embedding(vocab_size, embedding_size, input_length=max_len))
#Add the convlution 1d to the model with filters 32 and padding, relu activati
model.add(Conv1D(filters=32, kernel_size=3, padding='same', activation='relu'))
#Add another macpooling layer to the model
model.add(MaxPooling1D(pool_size=2))
#Add the LSTM to the model
model.add(Bidirectional(LSTM(32)))
#Add the dropout function to the model
model.add(Dropout(0.4))
#And finally add to the dense layer to the model
model.add(Dense(3, activation='sigmoid'))

plot_model(model, show_shapes = True)
```

2022-09-10 21:17:07.968023: I tensorflow/core/platform/cpu_feature_guard.cc:193] This TensorFlow binary is optimized with oneAPI Deep Neural Network Library (oneDNN) to use the following CPU instructions in performance-critical operations: AVX2 AVX512F AVX512_VNNI FMA
To enable them in other operations, rebuild TensorFlow with the appropriate compiler flags.

You must install pydot (`pip install pydot`) and install graphviz (see instructions at <https://graphviz.gitlab.io/download/>) for plot_model/model_to_dot to work.

```
In [52]: #Let's compile the model the categorical_crossentropy loss function and adam optimizer
model.compile(loss='categorical_crossentropy', optimizer='adam', metrics=['accuracy'])
print(model.summary())
```

Model: "sequential"

Layer (type)	Output Shape	Param #
embedding (Embedding)	(None, 300, 32)	160000
conv1d (Conv1D)	(None, 300, 32)	3104
max_pooling1d (MaxPooling1D)	(None, 150, 32)	0
bidirectional (Bidirectional)	(None, 64)	16640
dropout (Dropout)	(None, 64)	0
dense (Dense)	(None, 3)	195

```
=====  
Total params: 179,939  
Trainable params: 179,939  
Non-trainable params: 0
```

None

```
In [62]: #Create a earlystopping function
es = EarlyStopping(monitor = 'val_loss', patience=5)
batch_size = 64
#And fit the model to the train and validation data
history = model.fit(X_trn, y_trn,
                    validation_data=(X_vld, y_vld),
                    batch_size=batch_size, epochs=epochs, verbose=1,
                    callbacks = [es])
```

Epoch 1/50
67/67 [=====] - 8s 74ms/step - loss: 0.9791 - accuracy: 0.4713 - val_loss: 0.6943 - val_accuracy: 0.6247

Epoch 2/50
67/67 [=====] - 4s 67ms/step - loss: 0.4512 - accuracy: 0.8351 - val_loss: 0.2462 - val_accuracy: 0.9290

Epoch 3/50
67/67 [=====] - 4s 65ms/step - loss: 0.1701 - accuracy: 0.9566 - val_loss: 0.1000 - val_accuracy: 0.9719

Epoch 4/50
67/67 [=====] - 4s 65ms/step - loss: 0.1443 - accuracy: 0.9533 - val_loss: 0.1154 - val_accuracy: 0.9565

Epoch 5/50
67/67 [=====] - 4s 65ms/step - loss: 0.0598 - accuracy: 0.9866 - val_loss: 0.0546 - val_accuracy: 0.9868

Epoch 6/50
67/67 [=====] - 4s 67ms/step - loss: 0.0452 - accuracy: 0.9882 - val_loss: 0.0473 - val_accuracy: 0.9840

Epoch 7/50
67/67 [=====] - 4s 65ms/step - loss: 0.0316 - accuracy: 0.9925 - val_loss: 0.0397 - val_accuracy: 0.9884

Epoch 8/50
67/67 [=====] - 4s 65ms/step - loss: 0.0411 - accuracy: 0.9917 - val_loss: 0.0413 - val_accuracy: 0.9890

Epoch 9/50
67/67 [=====] - 5s 68ms/step - loss: 0.0381 - accuracy: 0.9913 - val_loss: 0.0395 - val_accuracy: 0.9890

Epoch 10/50
67/67 [=====] - 4s 66ms/step - loss: 0.0298 - accuracy: 0.9913 - val_loss: 0.0315 - val_accuracy: 0.9917

Epoch 11/50
67/67 [=====] - 4s 66ms/step - loss: 0.0215 - accuracy: 0.9932 - val_loss: 0.0360 - val_accuracy: 0.9917

Epoch 12/50
67/67 [=====] - 4s 66ms/step - loss: 0.0239 - accuracy: 0.9939 - val_loss: 0.0372 - val_accuracy: 0.9906

Epoch 13/50
67/67 [=====] - 5s 68ms/step - loss: 0.0251 - accuracy: 0.9925 - val_loss: 0.0284 - val_accuracy: 0.9923

Epoch 14/50
67/67 [=====] - 4s 66ms/step - loss: 0.0164 - accuracy: 0.9936 - val_loss: 0.0315 - val_accuracy: 0.9912

Epoch 15/50
67/67 [=====] - 4s 66ms/step - loss: 0.0131 - accuracy: 0.9965 - val_loss: 0.0311 - val_accuracy: 0.9917

Epoch 16/50
67/67 [=====] - 5s 70ms/step - loss: 0.0158 - accuracy: 0.9943 - val_loss: 0.0264 - val_accuracy: 0.9934

Epoch 17/50
67/67 [=====] - 5s 70ms/step - loss: 0.0136 - accuracy: 0.9960 - val_loss: 0.0268 - val_accuracy: 0.9923

Epoch 18/50
67/67 [=====] - 4s 67ms/step - loss: 0.0121 - accuracy: 0.9950 - val_loss: 0.0272 - val_accuracy: 0.9945

Epoch 19/50
67/67 [=====] - 4s 67ms/step - loss: 0.0100 - accuracy: 0.9965 - val_loss: 0.0270 - val_accuracy: 0.9945

Epoch 20/50
67/67 [=====] - 5s 72ms/step - loss: 0.0103 - accuracy: 0.9960 - val_loss: 0.0273 - val_accuracy: 0.9945

Epoch 21/50

67/67 [=====] - 5s 68ms/step - loss: 0.0106 - accuracy: 0.9955 - val_loss: 0.0267 - val_accuracy: 0.9950

In [63]: `loss, accuracy = model.evaluate(X_tst, y_tst, verbose=0)`

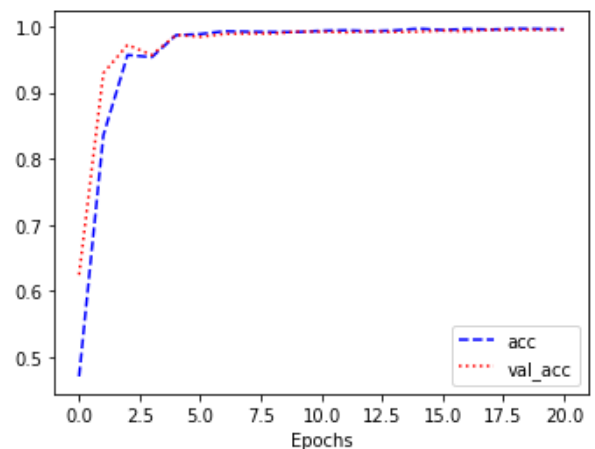
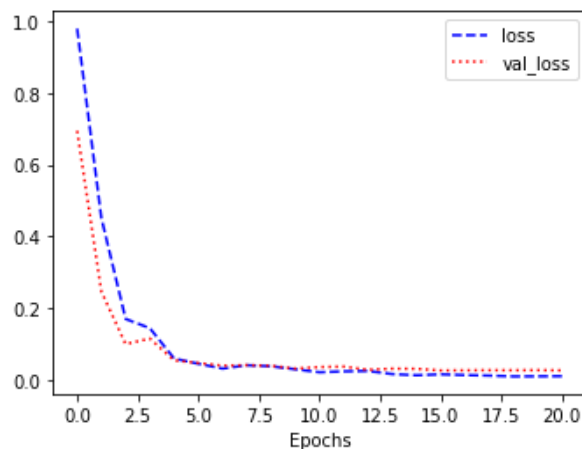
```
# Print metrics
print('Accuracy : {:.4f}'.format(accuracy))
```

Accuracy : 0.9941

In [64]: `#To visualize the the accuracy score to the model using the matplotlib`

```
plt.figure(figsize=(12, 4))
#Create a subplots to the model
plt.subplot(1, 2, 1)
#plot the loss of the mode history
plt.plot(history.history['loss'], 'b--', label = 'loss')
#plot the val_loss of the history
plt.plot(history.history['val_loss'], 'r:', label = 'val_loss')
#On x-axis Epochs
plt.xlabel('Epochs')
plt.legend()
#Create a subplots to the model
plt.subplot(1, 2, 2)
#plot the accuracy of the mode history
plt.plot(history.history['accuracy'], 'b--', label = 'acc')
#plot the val_accuracy of the mode history
plt.plot(history.history['val_accuracy'], 'r:', label = 'val_acc')
#On x-axis Epochs
plt.xlabel('Epochs')
plt.legend()

plt.show()
```



Test The model predict the good result

```
In [65]: text='the endothelium is crucial for vein integrity andprevention of thrombosi
text=[text]
text_int=vector.transform(text)
prediction=logistic.predict(text_int)
f" Biomedical text document classification is {prediction[0]}"
```



```
Out[65]: ' Biomedical text document classification is Thyroid_Cancer'
```

```
In [68]: text='bacteroid fragili fragili produc biofilm colonis intestin tract caus ser
text=[text]
text_int=vector.transform(text)
prediction=logistic.predict(text_int)
f" Biomedical text document classification is {prediction[0]}"
```

```
Out[68]: ' Biomedical text document classification is Colon_Cancer'
```

```
In [67]: text='lack biotmark identifi aggress primari tumor subset give rise metastas im
text=[text]
text_int=vector.transform(text)
prediction=random.predict(text_int)
f" Biomedical text document classification is {prediction[0]}"
```

```
Out[67]: ' Biomedical text document classification is Lung_Cancer'
```

The LogisticRegression and DecisionTreeClassifier, RandomForestClassifier give the best result to the model.

CONCLUSION

About the data

In the data we use predict the Biomedical text document classification is whether it's Thyroid_Cancer, Lung_Cancer, Colon_Cancer based on the performed basic EDA, text preprocessing, build different models, such as LogisticRegression, DecisionTreeClassification, RandomForestClassification, XGBoostClassifier. For the above model only all algorithms have good accuracy score compare to the other model. After that we run deep learning model to the dataset. And create the Sequential model to the data and fit the data to the model in this model we use conv1d and several input layers used and finally we use 50 epochs to the model we get the 99% accuracy_score to the deep learning model.

```
In [ ]:
```