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## APPENDIX I

### Swami Rama Himalayan University

Swami Ram Nagar, P.O. Jolly Grant, Dehra dun 248016 (INDIA) Phone: 91-135-2471111, Extn. 328, Fax 910135-24711122

"Ethics Committee"
(Reg. No. ECR/483/Inst/UK/2013/RR-16)

SRHU/HIMS/ETHICS/2020//93

Dated: 28.02.2020

"Ethics Committee"
(Registration No.: ECR/483/Inst/UK/ 2013/RR-16, Dt. 23.8.2017)

The Ethics committee in its meeting held on 21.12.2019 approved the Research Project entitled:

Vitamin D Receptor (VDR) gene polymorphisms and steroid receptor status in breast cancer patients.

Submitted by Principal investigator, Ashok Kumar Dogra, Ph.D scholar, Under the guidance of Dr. Archana Prakash, Professor, Deptt. of Biochemistry, HIMS, Swami Rama Himalayan University.

Dt: 28.02.2020

Dr. D.C. Dhasmana Member Secretary, Ethics Committee

## **APPENDIX II**

## **Thesis Protocol, Dated:**

Ethics committee approval letter no

receptor status among Breast Cancer patients

## INFORMED CONSENT DOCUMENT

(To be administered before recruiting/ screening of the study subject/patient. A signed copy of it must be given back to the subject/patient.)

Part I:<u>Subject/Patient Information document</u>
(To be written in non technical language understandable to a layman;Strike

dated

off which ever point is not applicable)

Title of the study/protocol: Vitamin D Receptor (VDR) gene polymorphisms and steroid receptor status in Breast Cancer patients

• •	o. o. c o o.
CTRI/UHID Registration no, (if applicable)	••••••
Principal Investigator with qualifications: Dr. Archana Prakash, l Dept. of Biochemistry.	Professor,
Name & address of institution: SRHU, Jolly Grant, Dehradun	
Contact No:	
Subject's /Patient's Name	
Introduction: matched case-control study.	
Purpose: The study is to assess VDR gene polymorphisms	and steroid

Methods: Blood Sampling (~5 ml) for DNA extraction and vit D level. PCR-RFLP Methodology for identify VDR gene polymorphisms and assess the Estrogen, progesterone and Her2 receptors from medical records Risk involved: *None*.

Potential benefits: The information collated from this research work can contribute to improve health and healthcare of patients and population affected with breast cancer or having high prognosis for breast cancer occurrence.

Reasonable alternatives/ possible variant treatment available: None.

Subject's responsibility: To provide the relevant information being asked

Compensation: No

Confidentiality: Yes

Voluntary participation: Yes

Financial cost of participation involved: None

Contact person: Dr. Archana Prakash, Professor, Dept. of Biochemistry, Dr. Sanjay Gupta, Prof. and Head, Dept. of Biosciences. Dr. Meenu Gupta, Professor, Dept. of Radiotherapy & Ethics committee (Member secretary) for further information on any query at any time in an event of a problem.

Patient /legal representative initials	
Date	

(Thank you for taking time to read this document .If you decide to take part in this study, you will be given a copy of this information document and signed consent form to keep with you)

Principal Investigator' Name: Dr. Archana Prakash, Professor, Dept. of Biochemistry.

Name of the Institute: SRHU, Jolly Grant, Dehradun.

## Part II: Informed consent form

Name of the study/trial: The study is to assess VDR gene polymorphisms and steroid receptor status among Breast Cancer patients.

of Dr. Archana Prakash, Professor, Dept. of Biochemistry and the Co-Supervisor-Dr. Sanjay Gupta, Prof. and Head, Dept. of Biosciences. Dr. Meenu Gupta, Professor, Dept. of Radiotherapy.

Study code:	
Patient Name	
Date of birth	age

- 1. I confirm that I have read and understood the patient information sheet dated...... for the above study on .......(drugs /procedures etc).... and had the opportunity to ask questions which were answered to my satisfaction
- 2. I have been well informed about the potential anticipated risks, discomfort and side effects associated with ....(the trial drugs/procedures etc).... and what I will be expected to do?
- 3. I understand that my participation is voluntary and I am free to withdraw from the study at any time without giving any reason, without affecting my future medical care or legal rights. I shall inform the principal investigator in this regard for any precaution/ medical care required to follow.
- 4. I understand that the principal investigator, others workers on the principal investigator's behalf and the ethics committee HIHT University will not need my permission to look at my health record both in the respect of the current study and any further research that may be conducted in relation to it, even if I withdraw from the study/ trial. I agree to this access. However, I understand that my identity will not be revealed in any information released to third parties or published.
- 5. I agree not restrict the use of any data or results that arise from this study provided such a use is only for scientific purpose.

- 6. I am aware that investigator will inform, whenever the situation arises, about any new finding that develop anywhere in the world, related to my treatment which may affect any decision to continue participation in the study.
- 7. I had have time to make my decision whether or not to take part in this study/trial. I agree to take part in the above study; I have received a signed and dated copy of this consent form for my records.

Patient/ Legally Acceptable Representative(LAR) Name	Patient's LAR's Signature	Date and Time
Investigator/ Designee Name	Investigator/ Designee's Name	Date and Time
-		
Witness Name	Witness's Signature	Date and Time

## APPENDIX III

## **Patient Information Form - Breast Cancer**

1.	Name of the Participa	ting Centre		
2.	Registration ID/UHID	of Registration		
	2.4. Case Registered Others	as Out Patient (OP)	In Patient (IP)	
3.	Date of First			
	Diagnosis	1 1 1 6	4: 12	
	(Date of first attendar	ice to any hospital fo	or this diseases)	
4.	Patient's			
5.	Name of the Spouse/I	Father/Mother/Careta	ıker.	
	Name	Mobile No.	Name	Mobile No.
5.	Age		7.	
	Sex			
	Menopausal tus			_
10.	Details of Socioecono	mic status, Family ir	ncome, Occupation, etc.	
11.	Co-Morbid Condition	s		
	/Hypertension/Isc (Specify)	chaemic Heart Diseas	nma/Hepatitis/HBsAg +ve/AIDS/HIV se/Allergic Conditions	'+ve
12.	Method of Diagnosis			
j	i. Clinical only	ii. Microscopic	iii. X-Ray/Imaging Techniques	iv. Others

Microscopic (If ii above) above)		X-Ray/Imaging Techniques(If iii abo	e) Others(If iv.	
b) c)	Histology of Primary Histology of Metastasis Cytology of Primary Cytology of Metastasis	<ul><li>b) Isotopes</li><li>c) Angiography</li></ul>	a) Surgery or Autopsy Without Histology b) Biochemical/Immuno logical tests. Specify test(s)	
	hological Diagnosis (With cological Diagnosis)	omplete description of Primary site of tu	umour and	
	rimary/Secondary site of Turogy			
13.2 M	[orphology			
14. De	etails of Stage			
14.1 St	aging System Followed:			
a)	TNM staging b) Others (Specify)		_	
14.2 St	aging Done at			
	Reporting Institution b)	Previous Institution c) Others		
15. Cli	inical Stage – UICC			
15.1 TI	NM with Description			
I.	Tumour size (in cms)	×		
II.	Axillary Lymph Node(s):  If Present, Number	1) Not Present 2) Present Size (in cms) of largest not	de	
	Whether Matted a) No	b) Yes		
	Whether Fixed a) No	b) Yes		
III.	Supra-Clavicular Node(s): If Present, Number	1) Not Present 2) Present Size (in cms) of largest not	de	

		Wheth Wheth			a) No a) No	)		Yes ) Yes					
IV		If Yes, Ulcer Infiltra Satelli Others (specif	ntion te nod	lule		ot Prese	nt	o) Yes Present					
	T T2	TX ,	Т0 Т3	Tis	Tis(l	OCIS)	Tis(LCI	S) Tis(	(Paget)	T1	T1a	T1b	T1c
	T4	T4a	Т	4b	T4c	T4d	Un	known					
	N Un	NX known	N0	N1	N2	N2a	N2b	N3	N3a	N3b	N3	3c	
	M	MX	<b>M</b> 0		M1 (e	.g. PUI	<u>(</u> )	Unkr	nown				
V]	[.	Stage	Group	oing (	Tick (v	) as app	propriate)						
I 16.		IA nvestiga	IB tions		IIA aging	IIB	IIIA	IIIB	IIIC	IV		Unkn	own
		Mamn Bone Secify)_	Scan			5. Oth	Thest X-ra		3.	Ultasou	ınd-Ab	domen o	& Pelvis
	*If	Mamm	ograp	hy is	done		'Normal	' 'Al	onormal'	'Su	spiciou	ıs/Incon	clusive'
	•	ecify an dings	•										
17.	De	tails of	Cance	er dire	ected tr	eatmen	t (CDT) (	Tick (√)	as appro	priate)			
17.	1 Т	reatme	nt giv	en pri	ior to re	egistrati	ion at repo	orting ins	stitution	(RI)			
	a.	No			b. Ye	es	c. Uı	nknown					
If Y	Yes,												

17.2 Type of I	Prior Tre	atment gi	ven		
1.Surgery treatment		No	Unknown	If Yes, Date of completion o	f
2.Radiotherapy treatment		No	Unknown	If Yes, Date of completion of	
3.Chemotherap treatment	•	No	Unknown	If Yes, Date of completion of	•
4.Others(Special treatment	•	No	Unknown	If Yes, Date of completion of	f
(Including Harr therapy)					_
18. Treatment a	t reporti	ng institu	tion (RI)		
18.1 Intention t	o treat				
1) Curativ	e/Radica	al	2) Palliative	3) No Treatment 4) Unk	nown
If Palliative Ye			2) 1 411141111	e, 10 11 <b>0</b>	
1) Palliati Surgery		1ly 2)	Palliative RT	C+CT 3) Palliative CT only	4) Palliative
5) Pain & (specify)	• •		· ·		
6) Others (specify)					
18.2 Type of C	ancer D	irected T	reatment Plann	ed at Reporting Institution:	
1) Surgery		Yes	No	Unknown	
2) Chemothera	ару	Yes	No	Unknown	
3) Radiotheraj	ру	Yes	No	Unknown	
4) Others (specify)					
19. Performano	ce Status	(WHO)	Before Treatme	ent	

1) Able to carry out all normal activity without restriction.

2) Restricted in physically strenuous activity by	ut ambulatory and	able to car	ry out light w	ork.
3) Ambulatory and capable of all self-care but than 50% of waking hours.	unable to carry out	any work;	up and about	more
4) Capable of only limited self-care; confined to	o bed or chair more	e than 50%	of waking ho	ours.
5) Completely disabled; cannot carry on any se	elf-care; totally cor	ifined to be	ed or chair.	
6) Unknown				
20. Surgery				
1) Surgery not planned 2) Yes, done as plant	anned 3) Surger	y planned	but not taken	
4) Others (specify)				
20.1 If Surgery done, Type of Surgical Procedu	are (specify)			
21. Surgical Histopathology Findings				
21.1pT size (cms)	_×		<del></del>	
21.2 Tumour origin: Sin	gle	Multice	entric	
21.3 Modified Richardson Bloom Score		_		
21.4 Extensive Intraductal Component(EIC)	Not Applicable	Positive	Negative	Unknown
21.5 Lymphatic/Vasculuar Invasion	Not Applicable	Positive	Negative	Unknown
21.6 Nipple/Skin Involvement	Not Applicable	Positive	Negative	Unknown
21.7 Oestrogen Receptor Status	Not Applicable	Positive	Negative	Unknown
21.8 Progesterone Receptor Status	Not Applicable	Positive	Negative	Unknown
21.9 Androgen Receptor Status	Not Applicable	Positive	Negative	Unknown
21.10 C-erb – B2/HER – 2	Not Applicable	Positive	Negative	Unknown

21.11 Number of Axillary nodes removed

Number showing Tumour

21.12 Pathological Stage

21.13 R Classification

pT pTX pT0 pTis pTis(DCIS) pTis(LCIS) pTis(Paget) pT1 pT1a pT1c pT1b pT4b Unknown pT2 pT3 pT4 pT4a pT4c pT4d pN1mi pΝ pNX pN0 pN1 pN1a pN1b pN1c pN2 pN2a pN2b pN3 pN3a pN3b pN3c Unknown Unknown pMpMXpM0 pM1 (specify)\_\_\_\_

R0

R1

R2

Unknown

RX

## APPENDIX IV

## **Patient Information Form - Controls**

7.	7. Name of the Participating Centre		
8.	8. Registration ID/UHID	IP)	
9.	9		
	(Date of first attendance to hospital for Routine Checkup/ San	nple collection	)
10.	10. Patient's Name		
	11. Name of the Spouse/Father/Mother/Caretaker.		
	Name Mobile No. Na	me	Mobile No.
12.	12. Age Sex	7.	
8.	8. Menopausal Status		
9.	9. Address		
10.	10. Details of Socioeconomic status, Family income, Occupation,	etc.	
11.	11. Co-Morbid Conditions		
	Tuberculosis/Diabetes/Bronchial Asthma/Hepatitis/HBsA/Hypertension/Ischemic Heart Disease/Allergic Condition (Specify) Others (Specify)	ns	
12.	12. Biochemical/Immuno-logical tests.		
Spe	Specify test(s)		

## **APPENDIX V**

## LIST OF PUBLICATIONS

- Dogra AK, Prakash A, Gupta S, Gupta M. An effective and rapid method of DNA extraction protocol from samples of human blood. Asian J Biol Life Sci [Internet].
   2023;12(1):187–91. Available from: http://dx.doi.org/10.5530/ ajbls.2023.12.25
   Impact factor:1.36
- Dogra AK, Prakash A, Gupta S, Gupta M, Bhat SA. Genetic variations of vitamin
   D receptor gene and steroid receptors status in breast cancer risk: An updated
   review. Adv Biomark Sci Technol [Internet]. 2022;4:1–11. Available from:
   http://dx.doi.org/10.1016/j.abst.2022.01.001 Impact factor:5.1
- 3. Ashok Kumar Dogra, Pranav Prakash, Sanjay Gupta, Meenu Gupta, Archana Prakash, Haamid Bashir, Role of Vitamin D Receptor Gene Polymorphism with Steroid Receptors in Breast Cancer: an Update. (2022).Int. J. Life Sci. Pharma Res.12(3), 1-6 http://dx.doi.org/10.22376/ijpbs/lpr.2022.12.2.L1-6 Impact factor:7.1
- 4. Dogra AK. Prakash A, Gupta S, Gupta M, Vitamin D and Vitamin D Receptor FokI, ApaI and BsmI gene polymorphisms and their relation with the risk of breast carcinoma: A case-control Study. J Clin Diagn Res. 2024; Available from:http://dx.doi.org/10.7860/jcdr/2024/69296.19241 Impact factor:1.14

## **APPENDIX VI**

## LIST OF PAPER/POSTER PRESENTATION

- Paper presentation at International Conference on Health and Medicine (ICHM- 21)
  held in Chandigarh, India on 12th December 2021, entitled "Polymorphisms of the
  Vitamin D Receptor Gene and Their Relationship with Steroid Receptor Status An
  Update"
- Paper presentation at International Conference on Biology, Applied Science & Medicine 28-Jan-2022, entitled "The Association of Vitamin D Receptor Gene Polymorphisms in Women With Breast Cancer"
- 3. Presented a poster in the CME & Workshop on "Recent Advances in Breast Cancer Management" at Cancer Research Institute, HIMS, Dehradun.

## **APPENDIX VII**

## **Awards/Certificates**





International Conference on Biology, Applied Science & Medicine 28-Jan-2022 | Webinar

## CERTIFICATE OF PARTICIPATION

This is to certify that

#### Mr. ASHOK KUMAR DOGRA

PHD SCHOLAR, SWAMI RAMA HIMALAYAN UNIVERSITY, DEHRADUN, INDIA.

#### Has Presented a Paper titled on

## "THE ASSOCIATION OF VITAMIN D RECEPTOR GENE POLYMORPHISMS IN WOMEN WITH BREAST CANCER"

In the International Conference on Biology, Applied Science & Medicine held on 28-Jan-2022

Organized by VDGOOD Professional Association.

Mr.M.Dinesh
Association Director









# CME and Workshop Recent Advances in Breast Cancer Management

Organized by

Department of Surgical Oncology, Cancer Research Institute, Himalayan Institute of Medical Sciences, SRHU
Asian Society Of Mastology (ASOMA)
American Society of Surgeons – India Chapter

## Certificate

This is to certify that

Ashok Kumar Dogra

has participated as Delgate and Poster Presentation in this "Recent Advances in Breast Cancer Management" at Cancer Research Institute, HIMS, Dehradun.

Prof. Sunil Saini Organizing Chairman Prof. Chintamani President-ASOMA

Dr. Anshika Arora Organizing Secretary

Conference accredited from Uttarakhand Medical Council for 3 Hours (U.K.M.C./2022-23/1479)





E-certificate No.: JUN22C05942





## NATIONAL ELIGIBILITY TEST FOR ASSISTANT PROFESSOR

NTA Ref. No.: 221610228410	Roll No.: UK0102430100
Certified that ASHOK KUMAR DOGRA	TI as
Son/Daughter of GEETA DEVI	
and LAKSHMAN DASS	has qualified
in June 2022 Joint CSIR-UGC Test for	eligibility for Assistant Professor in the subjec
Life Sciences	

As per information provided by the candidate, he/she had completed/appeared or was pursuing his/her Master's degree or equivalent examination in the concerned/related subject at the time of applying for Joint CSIR-UGC Test.

The date of eligibility for Assistant Professor is the date of declaration of Joint CSIR-UGC Test result, i.e., 28-10-2022, or the date of completion of Master's degree or equivalent examination with required percentage of marks within two years from the date of declaration of Joint CSIR-UGC Test result, i.e. by 27-10-2024, whichever is later.

This is an electronic certificate only, its authenticity and category in which the candidate had appeared should be verified from National Testing Agency (NTA) by the institution/appointing authority. This electronic certificate can also be verified by scanning the QK Code.

The validity of this electronic certificate is forever.

Date of issue: 29.11.2022

Series Director NIT

Senior Director, NTA

Note: NTA has issued the electronic certificate on the basis of information provided by the candidate in his/her online Application Form. The appointing authority should verify the original records/certificates of the candidate while considering him/her for appointment, as the NTA will not be liable for any false information provided by the candidate. The NTA is only responsible for the result which can be verified from the repository available in the website of NTA (csimet.nta.nic.in). The candidate must fulfil the minimum eligibility conditions as laid down in the notification for Joint CSIR-UGC Test.

## An Effective and Rapid Method of DNA Extraction Protocol from Samples of Human Blood

Submission Date: 19-02-2023; Revision Date: 13-03-2023; Accepted Date: 12-04-2023.

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## ABSTRACT

Introduction: Several different protocols ranging from a variety of manual and automated DNA extraction protocols, are available to extract nucleic acids from whole blood samples, one of the primary sources of DNA. These methods have one or more limitations in terms of low yield, Quality issues, cost, and time efficacy, utilization of toxic organic solvents, and others as well. This study aims to develop an effective protocol for extracting DNA from 500 µl of human blood. Materials and Methods: The extraction procedure was standardized using 500 microliters of fresh human blood samples. The disruption and cell lysis done by Lysis Buffers R (RBC) and N (Nucleic acid) contain detergents and salts, followed by the removal of proteins and other contaminants and recovery of DNA. The DNA samples were investigated for quality and quantity by measuring their absorbance at 260 and 280 nm, respectively (A260/A280). Results: DNA was checked by Gel docking on 0.8% Agarose gels. According to our protocol, we yielded 19 to 25 µg DNA, respectively, from 500 µL of fresh blood. Conclusion: Furthermore, our protocol yields bulk amounts of DNA while avoiding toxic organic solvents like Phenol. Consequently, downstream applications can be performed with the DNA because its quality has not been affected.

Keywords: Buffers, DNA extraction, Human blood.

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#### INTRODUCTION

In molecular biology, DNA, RNA, and proteins are needed for human health studies. DNA of high quantity and quality is crucial for successful downstream applications. [1-3] Furthermore, the rapid and cost-effective DNA extraction method would make it much more research-friendly. [4]

Blood DNA isolation protocols have been published in several publications. Albarino *et al.* Parzer *et al.* Robbins *et al.* Rudbeck *et al.* Sambrook *et al.* Wang *et al.* Albarino et al. However, some of

these methods require significant amounts of blood samples, making them unsuitable for low volume DNA extraction procedures. Lahiri and Nurnberger et al.<sup>[1:3]</sup> and Miller et al.<sup>[1:3]</sup> There were some protocols that used enzymes and organic solvents to get high-quality, PCR-inhibitor-free DNA, while others incorporated salting-out procedures to increase the yield of DNA. Castella et al.<sup>[1:4]</sup> and Cattaneo et al.<sup>[1:5]</sup> Therefore, the costs and time involved in some protocols are high by Nasiri et al.<sup>[1:6]</sup> Besides, in some cases, the quality of the DNA has been compromised. El Bali et al.<sup>[1:7]</sup> Chacon-Cortes et al.<sup>[1:8]</sup> and Santos et al.<sup>[1:9]</sup>

As a result, to meet the requirement for rapid, low volume and cost-effective genomic DNA extraction, our objective was to developed a protocol that would reflect the wide range of scientific interest pertaining to this field for extracting pure DNA from fresh human blood without costly enzymes and toxic organic solvents.

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#### **MATERIALS AND METHODS**

#### **Blood Collection**

An EDTA-containing vacutainer tube containing blood samples was collected from 30 healthy individuals randomly selected from areas nearby the Swami Rama Himalayan University's campus situated at Jolly Grant, Dehradun, Uttarakhand, India. In order to participate in this study, the volunteers provided consent prior to participation, and no prevalent diseases were present in the volunteers. As a part of the research work, ethics committee approval was obtained prior to the study. The DNA extraction process was performed on fresh blood after 1 hr after collection. While handling the blood samples, appropriate precautions was taken to prevent biohazards. Troutman et al.<sup>[20]</sup>

#### **Chemicals and Reagents**

The chemicals used in this method are standard chemicals found at major suppliers, Himedia Chemicals, for this study.

#### **Reagents Preparation**

Thereagents were prepared using different concentrations as RBC lysis buffer denoted as Lysis buffer R (10X) containing NH<sub>4</sub>Cl (1.54 M), NaHCO<sub>3</sub> (0.14 M) and 0.5 M EDTA (pH-8) dissolved in 100 mL distilled water and pH adjusted to 8. The solution was converted to the 1X working solution. Another lysis buffer, nucleic acid lysis buffer denoted as Lysis buffer N containing Tris-Cl (50 mM), MgCl<sub>2</sub> (50 mM), EDTA (2 mM), NaCl (0.5M), 1% Triton X-100 and 2% of 2- mercaptoethanol dissolved in 25 mL of autoclaved distilled water and pH was adjusted to 8. This protocol also involves 10% SDS followed by high salt concentration preparation of 6M NaCl. Further, reagents preparation involves chemicals like TAE buffer, Chloroform: Isoamyl alcohol (24:1), Isopropanol and 70% Ethanol.

#### Methodology for DNA extraction

The standardization of the DNA extraction method was done for the fresh blood samples.

Step 1. A whole Blood sample of only 500  $\mu L$  from the vacutainer was centrifuged at 2500 rpm for three minutes at 4°C and the serum was carefully aspirated.

Step 2. Approximately 1mL of Lysis buffer R (1X) was added to the pellet, mixed gently with periodic inversions, and was kept for 5 min at room temperature.

Step 3. Centrifugation of the cell's mixture was done for 5 min at 5000 rpm and carefully supernatant was discarded. Repeated the steps 2 and 3 for the pellet becomes white.

Step 4. Approximately, 500µL of Lysis Buffer N was added to the white pellet obtained from the previous steps, was mixed gently with a wide bore pipette, followed by adding 50µL of 10% SDS. The mixture was then incubated at 55-60°C for 30 min.

**Step 5.** At the end of incubation, 200µL of NaCl was added, vortexed vigorously, and centrifugation was done at 8000 rpm for 5 min.

**Step 6.** The Supernatant was then taken in a clean Eppendorf tube and added an equal volume of Chloroform: Isoamyl alcohol (24:1). The mixture was mixed well by gentle inversions and centrifuged at 12,000 rpm for 1 min.

Step 7. The Aqueous phase (DNA present) was removed carefully without disturbing the base layer and transferred into a clean Eppendorf tube containing an equal volume of chilled Isopropanol. Then the tube was vigorously shaken for few seconds resulted in fine white threads that appeared in the solution.

**Step 8.** The floating precipitate was transferred into a clean Eppendorf tube, washed with chilled 70% ethanol to remove any salts trapped with DNA, and was centrifuged at 13,000 rpm for 3 min.

**Step 9.** The tube was drained and evaporated to remove the ethanol completely. The pellet was then allowed to dried at 37°C.

Step 10. Finally, added sufficient (1X) TAE solution and dissolved the precipitate by light fingertip vibration. A solution of DNA was then stored at -20°C for further use.

#### **RESULTS**

A total of thirty human whole blood samples were used for DNA extraction and samples prepared manually showed no failures. Our optimized method was accelerated and economized by using red cell and nucleic acid lysis instead of separation of buffy coats and replacement of proteinase K. Our method observed human whole blood samples with A260/A280 absorbance ratios, a consistent range of 1.8. This effectively indicates that the samples were pure and successfully deproteinized. (Figure 1) It was also assumed that RNA was not found in the extracted DNA



Figure 1: DNA extraction was demonstrated by electrophoresis on agarose gel containing 0.8 percent agarose.

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Table 1: Ratios of Optical Density (OD) and the amount of DNA that can be extracted from each 500 µL Human blood DNA.

Coding sample	DNA quantity in ug/500ul blood	OD260/OD280 ratio
H1	23.00	1.78-0.02
H2	24.12	1.82-0.04
H3	19.11	1.72-0.05
H4	25.19	1.81-0.02
H5	22.12	1.85-0.03
H6	25.16	1.89-0.04
H7	22.14	1.87-0.07
H8	23.18	1.86-0.02

samples. In accordance with our protocol, we yielded an average of  $25\,\mu g$  of genomic DNA per  $500\,\mu L$  of fresh blood. The mean concentrations were calculated and tabulated using the experiment results (Table 1). There were varying intensities of bands in DNA samples run on 0.8% agarose gels. Despite this, all DNA bands were prominent and homogeneous, with minimal lane smearing. It appears that none of the DNA has degraded, despite being exposed to several chemical

At this point, DNA can be quantified by spectrophotometer and diluted to a working concentration, or you can simply use 1-2  $\mu$ L per for the PCR reaction by Huberman *et al.*<sup>[21]</sup>

#### DISCUSSION

Genomic DNA extracted from whole blood samples is being used for diagnostic and genotype studies, thereby facilitating personalized medicine. DNA extraction protocols are designed according to the chemical activity of the reagents on various organelles of cells by Push et al. [23] and Tan et al. [23] The RBC Lysis Buffer contains NH<sub>4</sub>Cl, NaHCO<sub>3</sub>, and EDTA. NH<sub>4</sub>Cl produces an increase in osmotic pressure within RBCs until they burst due to water influx. Despite this, it has a negligible effect on other cellular contents of the blood, especially on leucocytes. EDTA and NaHCO, act as buffer components, and NaHCO, increases the RBCs' swelling rate by Thoms et al.[24] A higher concentration of Tris buffer was used for the DNA extraction buffer, which maintains the buffer's pH at a steady level. Following this, EDTA was added to the extraction buffer to bind calcium and divalent magnesium cations that maintain the membrane integrity. Triton X-100 is a non-ionic surfactant to lyse cells and maintains DNA integrity, but higher concentrations lead to cell death by Yee et al.[25] NaCl forms ionic bonds with the phosphates in DNA that neutralize the negative charges, otherwise causing DNA molecules to repel one another and keep the DNA in solution. Anionic detergents like SDS can solubilize lipids and proteins by assisting in breaking down the membranes and nuclear envelopes, exposing the DNA-containing chromosomes. 2-mercaptoethanol is a powerful reducing agent used in DNA extraction buffer through disulfide bond breaking. It linearizes proteins, causing the molecules' denaturation and removal during the centrifugation by Koley et al.[26] The next step involves the addition of chloroform and isoamyl alcohol, which help to bind and precipitate proteins and lipids of cell membranes. DNA was generated in an aqueous phase, and lipids and proteins were formed in a non-aqueous phase. The hydration shell is formed by water molecules surrounding DNA at this stage. The DNA can therefore be separated from the remaining soluble components by centrifugation by adding Isopropanol, which acts as a dehydrating agent and disrupts the hydration shell.

Thomas et al.<sup>[27]</sup> published an earlier DNA extraction protocol using CTAB, but the buffer compositions and the sequences of steps differed considerably from our experiment. However, our present method is unique because it is simpler, faster, and more robust than many other methods for separating DNA from human blood samples. Additionally, there are no toxic reagents used in this protocol., so extractions are safe. Ness et al.<sup>[28]</sup> The average DNA yield was within the normal range using our protocol, and it takes roughly 2-3 hr for the protocol to be successfully completed.

#### CONCLUSION

This protocol could prove to be efficient in obtaining considerable quantities of genomic DNA from fresh human blood samples. The versatility of this universal method can be extended from fresh to frozen samples. Additionally, eliminating time-consuming steps like enzymatic incubation of Proteinase K and RNAase treatment and the absence of toxic organic solvents such as Phenol permitted an efficient and time-saving protocol in a way that could be used for advanced molecular biological techniques. Further, laboratories with limited funds may benefit from it by pursuing basic molecular biological research.

## ACKNOWLEDGEMENT

The Authors like to thank all the volunteers who provided their consent to participate in our study and

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the medical staff of Swami Rama Himalayan University, Dehradun, who assisted us in collecting samples.

#### **CONFLICT OF INTEREST**

The authors declare that there is no conflict of interest.

#### **ABBREVIATIONS**

µl: Microlitre; nm: Nanogram; µg: Microgram; A260/A280: Absorbance; OD: Optical density; EDTA: Ethylenediamine tetraacetic acid; NaCl: Sodium Chloride; NaHCO<sub>3</sub>: Sodium bicarbonate; NH<sub>4</sub>Cl: Ammonium chloride; SDS: Sodium Dodecyl Sulfate; TAE: Tris-acetate-EDTA.

#### SUMMARY

A successful application of this method has been made in freshly isolated human whole blood samples. Genomic DNA isolated by this method has an average quantity of 25 µg, and according to the measurements, this DNA had a quality of 1.7 to 1.8. Further agarose gel electrophoresis was performed to ensure that the DNA obtained was of high quality, without RNA and protein contamination (Figure 1). In this method, however, only a few chemicals were used in addition to ethanol (tris-HCl, EDTA, NaCl, and SDS), and they are readily available in every routine laboratory worldwide.

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Contents lists available at ScienceDirect



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journal homepage: www.keaipublishing.com/ABST

## Genetic variations of vitamin D receptor gene and steroid receptors status in breast cancer risk: An updated review

Ashok Kumar Dogra $^{a,\,^*},$  Archana Prakash $^a,$  Sanjay Gupta $^b,$  Meenu Gupta $^c,$  Showkat Ahmad Bhat $^d$ 

#### ARTICLEINFO

# Keywords: Breast cancer Steroid receptors Vitamin D Vitamin D receptor (VDR) gene polymorphism

#### ABSTRACT

Breast cancer, the most predominant type of cancer reported in females, is a heterogeneous disease classified into various subcategories depending on the presence of hormone receptors. Epidemiological studies show a strong correlation between reduced 1.25 dihydroxy vitamin D3 (1,25(OH)2D3) levels, the active component of vitamin D, and increased breast cancer risk in diverse populations. In a ligand-dependent manner, vitamin D receptor (VDR) transcriptionally modulates its target genes belonging to cell proliferation, differentiation, and apoptosis pathways, thus imparting protective function against cancer growth and progression. The coding and regulatory regions of the VDR gene contain several polymorphisms (BsmI, Fok1, Taq1, Apa1, Cdx2, poly (A), etc.) that modulate its transcription, translation, and mRNA stability. Despite this, research in this area has not yet led to many conclusions. In this review, we analyzed in a systematic way that the association of VDR allelic variants with breast cancer risk among patients from various populations. This analysis has revealed that Fok1, Bsm1, Apa1were to some extent associated with breast cancer risk, Taq1 shows no association, and Cdx2, poly(A), Tru91 gene polymorphisms may be susceptible for breast cancer development. We have highlighted the new insights of the current understanding of molecular mechanisms of the VDR gene polymorphisms related to breast cancer risk and also examined the interaction between VDR polymorphisms and steroid hormone (estrogen, progesterone, and androgen) receptors and their modifying effects on breast cancer risk, cancer severity, progression rate, and disease outcome. Therefore, with a lack of studies and inconsistent results, we recommend that further studies focus on genetic variations of the VDR gene that should be integrated with the assessment of steroid hormone receptor status

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#### 1. Introduction

Cancer is a multifaceted disease characterized by the unregulated proliferation of cells and is the primary cause of mortality globally, resulting in 9.6 million deaths. Despite tremendous progress in cancer therapy, heterogeneity of the disease and development of resistance to anti-cancer drugs continue to pose challenges for effective treatment. Breast cancer commonly arises from the uncontrolled proliferation of the epithelial cells present within the ducts or lobules of the mammary gland, leading to deregulated cellular signaling. It is the most commonly diagnosed malignancy noted among women in nearly 154 countries and results in ~630, 000 deaths annually. Breast cancer contributes to 29% of the reported cancers in the United States and is a significant cause of cancer-related mortality among women. On the recent years, statistical analyses have documented a steady rise in breast cancer diagnosis among the Indian female population, wherein 25.8 females against every 100,000 females are at risk of developing cancer, and 12.7 female deaths against every 100,000 females reported in India. In addition to age, the well-known risk factors of breast cancer include genetic (BRCA1 and BRCA2 mutations), environmental, and reproductive factors (early menarche and late menopause). Also, modifiable risk factors such as alcohol consumption and a high-fat diet contribute to the disease incidence. Some of the risk factors that account for the differences between patients are nulliparity or first pregnancy after 30, combined estrogen and progesterone use, breastfeeding, drinking alcohol, obesity, and insufficient exercise.

Vitamin D and Vitamin D receptor (VDR) has gained increasing relevance during the past two decades and are well-known for their roles in calcium homeostasis and metabolism. <sup>10</sup> Apart from regulation in calcium homeostasis, they promote cell differentiation and inhibit the proliferation of specific cells exhibiting cancer prevention properties. <sup>11</sup> Growing evidence demonstrates the inverse correlation between vitamin D and the development of breast cancer. <sup>12</sup> Several evidence-based studies have shown that calcitriol, and active form of vitamin D, plays a crucial role in inhibiting cell proliferation and angiogenesis and inducing cell differentiation and cell death in breast cancer through the intervention of VDR. <sup>13–15</sup> We have illustrated new insights into the current understanding of molecular mechanisms concerning breast cancer risk.

The polymorphisms of the VDR gene highly contribute to the risk of breast cancer, as per several studies conducted by researchers to identify its association. <sup>16-18</sup> Studies concerning the association of VDR gene polymorphisms and steroid receptors and their modifying effects on breast cancer risk are limited. <sup>19-21</sup> The purpose of this study is to provide a comprehensive review of the genetic variations of the VDR gene that may elucidate the association of VDR polymorphism with steroid hormone receptor status in breast cancer.

#### 2. Search strategy

We conducted a systematic analysis of published literature using the National Library of Medicine (NLM) of PubMed database

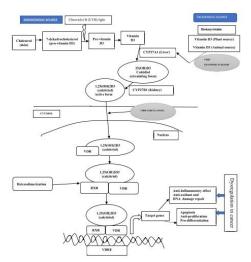


Fig. 1. Diagrammatic pathway representation of the endogenous vitamin D synthesis and downstream signaling. The active hormone 1,25(OH)2D3 is synthesized from precursor cholesterol mediated by ultraviolet-B (UVB) light. Vitamin D receptor (VDR) is activated by its ligand 1,25(OH)2D3 and it later forms a heterodimeric complex with retinoid X receptor (RXR). The 100-kDa sized complex identifies the vitamin D response element (VDRE) sequences in the 5' end of its target genes in the nucleus. Vitamin D signaling transcriptionally modulates genes functioning in cellular processes such as cell proliferation, apoptosis, and angiogenesis etc., which are cellular pathways pivotal for cancer development. The key enzymes and intermediate metabolites involved in the pathway are indicated.

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search from the year 2010–2020. In the search, keyword combinations were used as 'vitamin D' or 'vitamin D receptor gene polymorphisms' and 'Breast cancer' also, keywords: 'VDR gene polymorphisms' and 'Steroid receptors' concerning breast cancer. Our study included articles investigating VDR gene polymorphisms and steroid hormone receptor status related to breast cancer risk and incidence. The animal studies and studies focused on breast cancer in men or adolescents were excluded from this study.

#### 3. Vitamin D and vitamin D receptor (VDR) concerning breast cancer

Vitamin D is a lipophilic secosteroid that is endogenously synthesized from cholesterol in a multi-step process under the skin or can be obtained through dietary sources (exogenously). Via photolytic reaction, ultraviolet B (UVB) light mediates the formation of

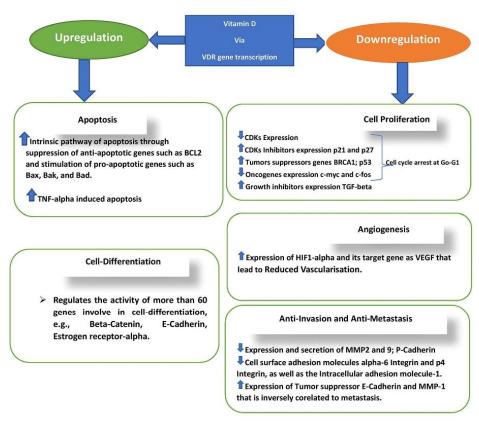


Fig. 2. Diagrammatic representation of Vitamin Ds effects on Breast cancer. 1,25 (OH)2D3-VDR complex activates the transcription of CDK inhibitors like p21 and p27 and represses cell cycle promoting factors including cyclin D1, cyclin A1, and cyclin E1. These transcriptional changes induce G1 cell cycle arrest, promote cell cycle exit, and limit the expansion of the cell cycle. Vitamin D signaling suppresses the expression of c-myc and c-fos oncogenes and upregulates tumor suppressor BRCA1, both of which regulate cell proliferation. Furthermore, 1,25(OH)2D3 hormone exerts its control on the apoptosis process by stimulating pro-apoptotic factors (BAX, BAK, BAD) and by downregulating anti-apoptotic factors (BCL-2, BCL-XL). In addition to controlling cellular proliferation, 1,25(OH)2D3 represses the hypoxia-inducible factor (HIF)-1 pathway and inhibits angiogenesis. Vitamin D signaling suppresses the HIF-1 and its target pro-survival genes such as Glut-1, ET-1, and vascular endothelial growth factor (VEGF). Vitamin D's anti-invasion or anti-metastasis properties are attained by inhibiting the expression of matrix metalloproteinase (MMP), urokinase-type plasminogen activator (uPA), and P-cadherin promotes metastasis, accompanied by stimulation of cell adhesion molecules such as E-cadherin and MMP inhibitor 1. Vitamin D, in addition to restricting cell proliferation, also promotes cellular differentiation that facilitates normal development. This is achieved through inhibition of epidermal growth factor receptor (EGFR) and insulin-like growth factor 1 (IG-1) that suppress downstream mitogen-activated protein kinase (MAPK) and extracellular signal-regulated kinase (ERK) 1 and 2 pathways essential for cell proliferation. These data imply that VDR is necessary and sufficient to carry out the anticarcinogenic functions mediated by vitamin D, specifically in breast malignancies, yet, the mechanism of such functions is still unknown. (Up arrows indicates stimulation and down arrows indicate repression in the boxes).

vitamin D3 (cholecalciferol) from 7-dehydrocholesterol via pre-vitamin D3 intermediate-generation step. <sup>22,23</sup> Further, 1, 25-dihydroxy vitamin D3 (1,25(OH)2D3; calcitriol), the biologically relevant active molecule, is derived by the action of two key CYP450-mediated hydroxylase enzymes - CYP27A1 and CYP27B1. <sup>24</sup> The intermediate metabolite 25-hydroxyvitamin D3 (25(OH)D3) is synthesized from vitamin D3 via the first hydroxylation step mediated by CYP27A1 in the liver. <sup>25</sup> The levels of the stable and most circulating 25(OH)D3 metabolite are commonly measured as an indicator of the vitamin D levels of an individual. CYP27B1 mediates the second hydroxylation reaction and catalyzes the conversion of circulating 25(OH)D3 to the potent steroid hormone 1,25(OH)2D3 in the kidney. <sup>26</sup> After entering the circulation, calcitriol binds to vitamin D binding protein (VDBP). It is administered to target tissues via interaction of its high-affinity receptor, vitamin D receptor, and it heterodimerizes with auxiliary proteins such as retinoid X receptors (RXR) to carry out diverse biologic functions of vitamin D<sup>27</sup> (Fig. 1).

The VDR protein belongs to the nuclear receptor superfamily, and its gene comprises at least 11 exons that stretch over 60 kb of DNA region on long arm of chromosome 12 (12q 12–14).<sup>26</sup> However, exon 1 constitutes a non-coding region, and exon 2–9 codes for the VDR protein. 29 Multiple studies have shown that vitamin D regulates vital cellular pathways that prevent cancer, such as apoptosis, differentiation, cell proliferation, and promote the inhibition of angiogenesis, metastasis, invasion, and inflammation. 30-32 Therefore, Vitamin Ds anti-cancer properties suggest that as an agent that could be used in the apeutics. Moreover, VDRs can be found in a large number of tumors, suggesting the receptor affects the aetiology of cancer. In a normal mammary gland, VDR expression is observed, and epithelial cells of mammary glands possess the same enzyme system as the kidneys; hence, vitamin D is essential for normal mammary gland development.33 Therefore, the effects of vitamin D on breast cancer are biologically reasonable. We have illustrated updated information on molecular mechanisms of the Vitamin Ds effects of breast cancer (Fig. 2), and in relation to this figure, further studies pertain to a complete understanding of the mechanism of action concerning breast cancer in the vitamin D research is required.<sup>34</sup> The literature has provided much information concerning vitamin D preventive function in breast cancer.<sup>35–37</sup> The correlation between circulating 25(OH)D levels and breast cancer risk has been highlighted in case-control studies, showing that higher levels of 25(OH)D may reduce breast cancer risk. 30,39 In addition, a significant reduction in 25(OH) levels were observed in women diagnosed with highly aggressive triple-negative and basal-like breast cancer. 40 Accordingly, these studies suggest that there is an increased risk of breast cancer with the deficiency of vitamin D and is associated with more aggressive cancers. On the other hand, different studies showed no link between the plasma levels of 25(OH)D or supplementation with the risk of breast cancer. 41,42 cording to these contradictory findings, vitamin D might only exert its effects on distinct breast cancer subtypes with different designs of tumors, which may reflect the disease's heterogeneity. As a result, further research is needed to fully understand this issue.

# 4. Findings in VDR gene polymorphisms and breast cancer

# 4.1. Polymorphic sites in the human VDR gene

Multiple research groups have identified more than 200 variations in the DNA sequences of the VDR gene, known as polymorphisms, including restriction fragment length polymorphisms (RFLPs) and a variable number of tandem repeats (VNTRs).  $^{43}$  Numerous population-based studies demonstrate a link between breast cancer and VDR gene polymorphisms as major polymorphisms in the coding region are Fok1 and Taq1 in exons 2 and 9, respectively, whereas Cdx2 polymorphism in exon 1, Bsm1, Apa1 in intron 8, and poly (A) repeats in the 3' UTR are found in the regulatory regions. Also, some of the rarer polymorphisms include Tru91 in intron 8 and A-1012G near the transcription start-site in exon  $1.^{44-47}$  The Fok1 and Cdx2 are near the 5' UTR region, and the rest of all polymorphisms are near the 3' UTR region. Allele frequencies differ widely over populations for polymorphisms in the 3'-UTR region, suggesting that these polymorphisms were linked to linkage disequilibrium (LD). (Fig. 3).

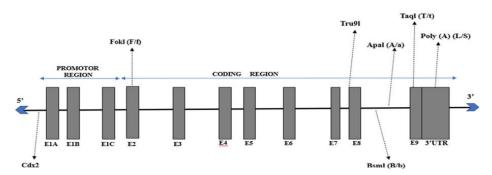


Fig. 3. Schematic of the human Vitamin D receptor (VDR) gene structure marked with the location and alleles of the different VDR gene polymorphisms. E1 to E9 indicate the exons of the VDR gene.

# 4.2. Fok1 polymorphism and breast cancer

In exon 2, Fok1 polymorphism consists of T to C mutation (T/C), leading to an alternate upstream start codon (ACG to ATG) that creates a differently sized VDR protein and has shown differences in the activity of the two isoforms of the transcription factor. The polymorphism results in F (absence of Fok1 restriction site) and f (presence of Fok1 restriction site) alleles resulted in three genotypes FF, Ff, and ff. Numerous case-control studies on different ethnic populations have investigated the relationship between VDR gene polymorphism of Fok1 with breast cancer, summarized in Table 1. Only one study found a significant inverse relationship between Fok1ff genotype and breast cancer (OR = 0.71; 95% CI: 0.57-0.88). This study examines a large sample size with estimates age-adjusted and shows a marked deviation from Hardy-Weinberg equilibrium in this polymorphism.

In contrast, six studies \$^{1,50-54}\$ significantly positive associations were reported between Fok1 ff genotype and risk for breast cancer. In the other nine studies, \$^{43,46,55-61}\$ no significant associations were found. One study (Amadori et al.) was conducted on a mixed population, i.e., African and Caucasian, but the effect was only seen in Africans, and a relatively limited sample size (53/50) may have influenced the results. Most of the studies have been done in African and Caucasian populations and have shown inconclusive results. As far the populations and ethnicities of the Indian subcontinent are pretty diverse, there is limited information available about the variations found in the Fok1 VDR gene. A recent study in India shows that the Fok1 ff allele shows a strong association with breast cancer. S A meta-analysis of 34 studies including 26,372 cases and 32,883 controls reaffirmed the risk posed by Fok1 in taking with other gene variants like Bsml, Apal, and poly (A) VDR variants are susceptible to develop breast cancer. There is still a need for further studies to be conducted in more ethnic groups in order to determine its functional polymorphism effects related to the risk of breast cancer.

# 4.3. Bsm1 polymorphism and breast cancer

The Bsm1 polymorphism (A/G) assigned the genotypes BB, Bb, bb considered silent polymorphism (SNPs) that do not alter the amino acid sequence as in Fok1, and it may affect gene expression via modulation of mRNA stability. <sup>16</sup> Several research studies have been conducted, but some conclusions vary by ethnic group and author, as shown in Table 2. Most of the studies have found no association between Bsm1 and breast cancer risk. <sup>43,49–51,56,59,6,54</sup> Few studies in this regard show Bsm1 bb and Bb genotype associations. <sup>46,57,50,60,61</sup> A study by shabbazi in Iranians shows significantly increased breast cancer risk between Bsm1 bb and BB genotypes (OR 1.74, CI 1.06–2.87 and OR 2.08, CI 1.31–3.29) in the BRCA1/2 mutation of the non-carriers subgroup. In an Egyptian

Table 1
Fok1 polymorphism and Breast cancer.

Author's Studies (Year)	Study Population	Type of study	Polymorphism studied	Number of Cases/ Controls	Study Findings
Anderson et al (2011)	Caucasian	PCCS	Fok1, Bsm1, Apa1, Taq1, Cdx2	1777/1839	The Fok1 ff genotype is inversely related to breast cancer risk.
Engel et al (2012)	Caucasian	NCCS	27 SNPs including Fok1	270/554	No association
Rollison et al (2012)	Hispanic and Non- Hispanic	PCCS	Fok1, Bsm1, Poly(A)	Hispanic-1527/791 Non- Hispanic-1599/922	No association
Fuhrman et al (2013)	Caucasian	PCCS	6 SNPs in CYP24A1 including Fok1, Bsm1	484/845	No association
Mishra et al (2013)	African-American and Hispanic	CCS	Fokl, Bsml, Apal, Taq1	232/349	The Fok1 f allele shows an increased association with breast cancer risk.
Shahbazi et al (2013)	Iranians	CCS	Fokl, Bsml	140/156	No association
Akilzhanova et al (2014)	Other	CCS	Fok1, Bsm1	315/604	The Fok1 ff genotype is related to increase breast cancer risk.
Abd-Elsalam et al (2015)	African	CCS	Fokl, Bsml, Apal, Taq1	130/100	No association
Nemenqani et al (2015)	Asian	CCS	Fok1, Taq1	95/100	The Fok1 ff genotype is related to breast cancer risk.
Rashid et al (2015)	Other	CCS	Fokl, Bsml	463/1012	No association
Deshasaux et al (2016)	Caucasian	NCCS	Fok1, Bsm1	233/466	No association
Amadori et al (2017)	African and Caucasian	PCCS	Fok1, A1012G, Cdx2	53/50	The Fok1 ff genotype is related to increase breast cancer risk. (African)
Shahabi et al (2018)	Other	CCS	Fok1, Bsm1	203/214	No association
Shaker et al (2019)	African	CCS	Fok1, Bsm1	115/120	No association
Raza et al (2019)	Indian	CCS	Fok1	125/125	The Fok1 ff genotype is related to increase breast cancer risk.
Ahmed et al (2019)	African	CCS	Fok1, Apa1, Taq1	392/193	The Fok1 ff genotype is related to breast cancer risk

PCCS: population-based case-control study, CCS: case-control study, NCCS: nested case-control study.

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Table 2

Author's Studies (Year)	Study Population	Type of study	Polymorphism studied	Number of Cases/ Controls	Study Findings
Anderson et al (2011)	Caucasian	PCCS	Bsm1, Fok1, Apa1, Taq1, Cdx2	1777/1839	No association
Rollison et al (2012)	Hispanic and Non- Hispanic	PCCS	Bsm1, Fok1, Poly(A)	Hispanic-1527/791 Non-Hispanic-1599/ 922	No association
Fuhrman et al (2013)	Caucasian	PCCS	6 SNPs in CYP24A1 including Bsm1,Fok1	484/845	No association
Mishra et al (2013)	African-American and Hispanic	CCS	Bsm1, Fok1, Apa1, Taq1	232/349	No association
Shahbazi et al (2013)	Iranians	CCS	Bsm1, Fok1	140/156	BsmI bb and Bb genotype is significantly associated with an increased risk of breast cancer.
Akilzhanova et al (2014)	Other	CCS	Bsm1, Fok1	315/604	No association
Abd-Elsalam et al (2015)	African	CCS	Bsm1, Fok1, Apa1, Taq1	130/100	BsmI bb genotype is significantly associated with an increased risk of breast cancer.
Guo et al (2015)	Chinese	CCS	Bsm1, Apa1, Taq1	219/391	No association
Rashid et al (2015)	Other	CCS	Bsm1, Fok1	463/1012	BsmI bb genotype is associated with an increased risk of breast cancer.
Deshasaux et al (2016)	Caucasian	NCCS	Bsm1, Fok1	233/466	No association
El-Shorbagy et al (2017)	African	CCS	Bsm1, Apa1, Taq1	100/50	No association
Shahabi et al (2018)	Other	CCS	Bsm1, Fok1	203/214	BsmI bb and Bb genotype is significantly associated with an increased risk of breast cancer.
Shaker et al (2019)	African	CCS	Bsm1, Fok1	146/130	BsmI bb and Bb genotype is significantly associated with an increased risk of breast cancer.

 $\hbox{PCCS: population-based case-control study, CCS: case-control study, NCCS: nested case-control study.}$ 

study of 130 cases and 100 controls, it was found that the B allele was shown to be protective for risk of breast cancer with significant risk reduction by 44 and 60% those with genotypes Bb and BB in comparison to bb genotype.  $^{46}$  Studies conducted in Kazakhstan,  $^{51}$  Pakistan,  $^{50}$  and Iran  $^{61}$  also found that the BB genotype significantly reduced breast cancer risk by 32, 33, and 47%, respectively. In short, on average, it appears there is no association between Bsm1 and breast cancer risk. Some recent studies have found significant differences among estimates even in groups of similar ethnicities, which should be investigated further and maybe through subgroup analysis in the future.

# 4.4. Taq1 polymorphism and breast cancer

The Taq1 is the ATT nucleotide substitution for ATC in exon nine, resulting in a synonym change in codon 252 (isoleucine) and its polymorphism assigned genotypes as TT, Tt, and tt.  $^{16}$  An analysis of a case-control study (PCCS) has found a significant association

Table 3
Taq1 polymorphism and Breast cancer.

Author's Studies (Year)	Study Population	Type of study	Polymorphism studied	Number of Cases/ Controls	Study Findings
Anderson et al (2011)	Caucasian	PCCS	Taq1, Fok1, Bsm1, Apa1, Cdx2	1777/1839	No association
Engel et al (2012)	Caucasian	NCCS	27 SNPs including Taq1	270/554	No association
Mishra et al (2013)	African-American and Hispanic	CCS	Taq1, Apa1, Fok1, Bsm1	232/349	No association.
Abd-Elsalam et al (2015)	African	CCS	Taq1, Fok1, Bsm1, Apa1	130/100	No association.
Nemenqani et al (2015)	Asian	CCS	Taq1, Fok1	95/100	No association.
Guo et al (2015)	Chinese	CCS	Taq1, Bsm1, Apa1	219/391	No association.
Atoum et al (2017)	Other	CCS	Taq1	122/100	No association.
Ahmed et al (2019)	African	CCS	Taq1, Fok1, Apa1	392/193	No association
Matini et al (2020)	Other	CCS	Taq1, Apa1	150/150	The tt genotype is significantly associated with increased breast cancer risk.

PCCS: population-based case-control study, CCS: case-control study, NCCS: nested case-control study.

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among premenopausal women for VDR Taq1 Tt (OR = 1.33; 95% CI: 1.01-1.74) and no association for vitamin D or calcium intake. <sup>49</sup> A nested case-control study (NCCS) review haplotypes analysis defines different haplotypes blocks according to their study population and showed that block B, which includes Taq1, was associated with a decreased risk of breast cancer (OR = 0.5). <sup>55</sup> Another study in African-Americans and Hispanics observed the Taq1 is in the strongest linkage disequilibrium with Apa1 in Hispanics than in African-Americans. <sup>50</sup> The Jordanian population found that Taq1 tt, TT, Tt genotypes and 25(OH)D levels are significantly associated. <sup>65</sup> Only one study found a significant association of Taq1 tt genotype with the breast cancer risk. <sup>66</sup> In conclusion, Taq1 VDR polymorphism has not been associated with breast cancer risk in many studies. Additional studies with a larger sample size in conjunction with different environmental and genetic factors are required to obtain more accurate findings. An analysis of the association of the Taq1 polymorphism with Breast cancer is shown in Table 3.

# 4.5. Apa1 polymorphism and breast cancer

The Apa1 vitamin D receptor gene assigned genotypes are AA, Aa, aa and contain a C to A base substitution in Intron 8 considered silent polymorphism (SNPs) that do not alter the amino acid sequence as in Fok1, to influence mRNA stability. A large population-based case-control study shows higher risk of breast cancer is associated with the AA genotype. The Apa1 is in linkage disequilibrium and is associated with 50% reduced breast cancer risk in a nested case-control study. A genotype has increased by 1.9 times greater than that of the population as a whole, which is statistically significant in an increased breast cancer risk. Also present were Aa and as genotypes, although at a lower frequency. And the other hand, Mishra et al. show no association with Apa1 and breast cancer risk but explore a significant association between VDR Apa1 as genotype and poorly differentiated tumors (p = 0.04). Also, the frequency of the aa allele is more common in Hispanics/Latinas in younger age groups than African-Americans indicates that VDR-Apa1 plays a role in advancing cancer in younger breast cancer patients.

Similarly, Guo et al. have found no association in the Chinese population  $^{63}$ , and Ahmed et al. have found no association in the African population.  $^{54}$  Abd-Elsalam et al. observed a significantly increased risk for breast cancer in Apa1 aa genotype (OR = 2.2).  $^{46}$  A recent study in Iran shows that the Apa1 aa allele appears to be associated with an increased risk of breast cancer, and Taq1 variations did not significantly correlate with tumor size or lymph node metastasis based on a stratified analysis.  $^{66}$  Accordingly, although the Apa1 allele association has not been studied extensively across races, most studies have noted it is associated with breast cancer. An analysis of the association of the Apa1 polymorphism with Breast cancer is shown in Table 4.

# 4.6. Poly (A), Cdx2, Tru91 polymorphism, and breast cancer

The Poly (A) polymorphism yields two alleles – S (short) allele containing less than 18 A nucleotides and L (long) allele containing more than 168 A nucleotides. Differing number of adenine (A) repeats in the 3' UTR, deviant from the ordinarily occurring poly (A) repeats in this loci, generated the poly (A) polymorphism that can potentially influence the post-transcriptional regulation of the VDR gene. The Cdx2 polymorphism referred to G to A sequence variation in the VDR promoter and suggested modulation of promoter activity. The Tru91 polymorphism in the VDR gene is a G (U allele) to A (u allele) is thought to modify gene expression via its effects on the affinity of enhancer regions. A population-based case-control study found that reduction in risk of breast cancer was associated with the Cdx2 AG genotype (OR = 0.83), and a higher risk of cancer is associated with the Cdx2 AA genotype (OR = 1.36). Another population-based case-control study for Poy(A) genotype is not associated with increased breast cancer risk but with dietary intake of vitamin D; women with LS or SS Poly (A) genotypes are at greater risk of breast cancer with higher vitamin D intake (OR = 1.41). The study by Yao et al. revealed that the Cdx2 G allele is marginally associated with an increased risk of breast cancer. A case-control study in southern Pakistani females for Cdx2 polymorphism observed that breast cancer risk was not significantly associated with VDR Cdx2 polymorphism, but the risk was associated with the GG genotype (OR = 1.8). Another case-control study

Table 4
Apa1 polymorphism and Breast cancer.

Author's Studies (Year)	Study Population	Type of study	Polymorphism studied	Number of Cases/ Controls	Study Findings
Anderson et al (2011)	Caucasian	PCCS	Apa1, Fok1, Bsm1, Taq1, Cdx2	1777/1839	The AA genotype shows an increased risk for breast cancer.
Engel et al (2012)	Caucasian	NCCS	27 SNPs including Apa1	270/554	No association
Dalessandri et al (2012)	Caucasian	Pilot-CCS	Apa1	164/174	The AA genotype shows a significantly increased risk for breast cancer.
Mishra et al (2013)	African-American and Hispanic	CCS	Apa1, Taq1, Fok1, Bsm1	232/349	No association
Abd-Elsalam et al (2015)	African	CCS	Apa1, Taq1, Fok1, Bsm1	130/100	The an genotype shows a significantly increased risk for breast cancer.
Guo et al (2015)	Chinese	CCS	Apa1, Taq1, Bsm1	219/391	No association
Ahmed et al (2019)	African	CCS	Apa1, Fok1, Taq1	392/193	No association
Matini et al (2020)	Other	CCS	Apa1, Taq1	150/150	The aa genotype is significantly associated with increased breast cancer risk.

PCCS: population-based case-control study, CCS: case-control study, NCCS: nested case-control study.

in Iranian females assessed an increased risk of breast cancer in women carrying the Poly(A) LS and LL alleles (OR = 1.8), and women with a low 25(OH)D level and the L carrier genotype are at greater risk. Sa Amadori et al. found that the Cdx2 AA genotype was linked to breast cancer susceptibility in Africans only. Sa In the VDR Tru9I polymorphism study, premenopausal breast cancer was not significantly associated.

In contrast, breast cancer risk was associated with the 'uu' genotype (OR = 1.1) in a case-control study. <sup>47</sup> In conclusion, poly(A) seems controversial, but the variant L allele apparently increases breast cancer risk, although further verification is undoubtedly warranted. Also, further studies are required for Cdx2, Tru91 VDR polymorphisms as the data are still limited. An analysis of the association of the Poly(A), Cdx2, Tru91 polymorphism with Breast cancer is shown in Table 5.

# 5. Steroid hormone receptor status and breast cancer

Like VDR, hormone receptors ER, PR, and AR (Estrogen, Progesterone, and Androgen Receptors) are part of the nuclear receptor superfamily. Binding to the respective ligands (estrogen, progesterone, and androgen) induces the dimerization of the receptors and translocation to the nucleus. The hormone-receptor complex transcriptionally modulates the expression of its downstream target genes by identifying the DNA sequences corresponding to hormone response elements (HREs) in the promoter region of these genes.

Two forms of ER, encoded by different genes –  $Er\alpha$ ,  $ER\beta$ , are of interest in the context of breast cancer. Depending on the presence of  $ER\alpha$ , breast cancers categorized as ER-positive (ER+) or ER-negative (ER+) and ER-cancers can be positive for PR. Therefore, ER+ cancers positively respond to hormone therapy and are less aggressive with better survival outcome. Notably, studies have demonstrated that  $ER\alpha$  transcriptionally regulates the expression of PR in the mammary gland. Accordingly, it becomes clear that mammary epithelial cells depend on estrogen and/or progesterone for proliferation, which significantly contributes to different subtypes of breast cancer. The Androgen receptor (AR) is strongly linked to prostate cancer pathology, but evaluation of AR expression in primary breast cancer samples showed reduced levels of AR in triple-negative cancers compared with the levels in AR-/HER2-or HER2+ cancers. However, increased AR expression was noted in less aggressive forms (AR) of tumors. Furthermore, AR expression was positively associated with the co-expression of several luminal genes, steroid hormone genes, and HER2 pathway members. In triple-negative breast cancer, the presence of AR with the concurrent expression of luminal genes is also characterized as the luminal-androgen receptor (AR) subtype. We have the formula of AR with the concurrent expression of luminal genes is also characterized as the luminal-androgen receptor (AR) subtype. We have the formula of AR with the concurrent expression of luminal genes is also characterized as the luminal-androgen receptor (AR) subtype. We have the formula of AR and AR in the formula of AR in t

A study conducted in India recorded 49.2% of patients to be ER+, and 50.8% to be ER-including 24.8% of triple-negative tumor cases  $^{7.0}$  However, ER-breast cancer patients were double in proportion (50.8%) in comparison to ER+ patients (23.4%), as per another independent report. Further, 15% of patients were diagnosed with triple-negative tumors. The study also revealed that the ER negativity rates increased by 63.5% among young women (less than 50 years), and this trend diminished with the rise in age. However, in the United States, the majority of the cases were ER+, and the distribution of the patients was found to be 67.2% ER+ PR+, 19.0% ER+ PR-, 1.6% ER- PR+, and ER- PR-, 12.2%. $^{79}$  Further, individuals suffering from ER+ PR+ breast cancer had 30%-60% lower mortality risk from the disease. Individuals showing ER positivity within the range of 1%-9% are more likely to receive chemotherapy as they are considered to have lower ER expression than patients that show more than 10% ER positive rates and are treated with endocrine therapy. $^{50}$  Although ERa, PR, and HER2 are the commonly studied prognosis biomarkers, current efforts have been intensified to investigate the role of other less-studied receptors such as ER $\beta$  and AR. $^{51}$ 

# 6. VDR polymorphisms and hormone receptor status in breast cancer

The steroid hormone receptor status is known to influence the relationship between vitamin D and breast cancer risk. Reduced levels of 25(OH)D show association with increased risk of advanced forms of breast cancers, and to a lesser extent for ER+ PR+ breast cancers with a higher prognosis. Treatment with 1,25(OH) $_2$ D $_3$  halts explicitly the cell cycle progression in ER expressing breast cancer cells. Multiple observational studies have highlighted the pronounced reduction of 25 (OH)D metabolite levels in triple-negative breast cancers, unlike other molecular subtypes, demonstrating that vitamin D is negatively correlated with this subtype. Supporting this finding, increased circulating 25 (OH)D concentrations exerted maximum protective effect in triple-negative breast cancer patients.

Epidemiological studies that evaluate the relationship between VDR polymorphism with different molecular subtypes of breast cancer will provide more detailed insights into the heterogeneity of this malignancy. VDR polymorphism alone was not earlier considered a breast cancer prognostic marker owing to lack of significant association of VDR with various factors or markers of breast cancer checked such as tumor type, lymph node status, hormone receptors, Ki-67 expression, p53 levels, etc. <sup>96</sup> However, another study reported a negative association between VDR expression and factors including hormone receptor status, Ki-67, triple-negative status, tumor size etc. <sup>97</sup> Due to the conflicting observations, further conclusive studies are warranted in this aspect. It is proposed that interactions of VDR with the steroid receptor status can possibly modify the breast cancer risk and further our understanding of the disease pathogenesis and the resistance to therapies noted in patients. <sup>98</sup>

Due to insufficient power and low sample size, minimal studies have considered the hormone receptor's status when assessing the link between VDR polymorphism and breast cancer. A study demonstrated a significant interaction of FokI polymorphisms (carrying at least one copy of 'f' allele) on ER+ cancer development in Saudi women patients.  $^{21}$  A study assessing the effects of common VDR variations Taq1, Apa1, and Bsm1 on breast cancer susceptibility among Egyptian women showed a significant correlation for Apa1 and Taq1 variants and the absence of a link between the VDR genotypes and ER and PR expression levels in the tumor tissues.  $^{64}$  In summary, these studies highlight the contribution of VDR variant alleles in response to treatments, specifically targeting hormone receptors such as estrogen modulators in breast cancer patients.

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Table 5 Poly(A), Cdx2, Tru91 polymorphism, and Breast cancer.

Author's Studies (Year)	Study Population	Type of study	Polymorphism studied	Number of Cases/ Controls	Study Findings
Anderson et al (2011)	Caucasian	PCCS	Cdx2, Fok1, Bsm1, Apa1, Taq1	1777/1839	The Cdx2 AA genotype is related to increase breast cancer risk.
Rollison et al. (2012)	Hispanic and Non- Hispanic	PCCS	Poly(A), Fok1, Bsm1	Hispanic-1527/791 Non-Hispanic-1599/922	No association
Yao et al. (2012)	African- American, and Caucasian	PCCS	Cdx2	928/843	The Cdx2 G allele is related to a lower risk of breast cancer in African-Americans.
Iqbal et al (2015)	Other	CCS	Cdx2	103/161	The Cdx2 GG genotype is related to breast cancer susceptibility.
Colagar et al (2015)	Other	CCS	Poly(A)	134/127	The Long Poly (A) L allele is significantly associated with an increased risk for breast cancer.
Amadori et al. (2017)	African and Caucasian	PCCS	Cdx2, Fok1, A1012G	53/50	The Cdx2 AA genotype is related to breast cancer susceptibility.
Iqbal et al (2018)	Other	CCS	Tru91	228/503	The Tru91 uu genotype is related to increased breast cancer risk.

PCCS: population-based case-control study, CCS: case-control study, NCCS: nested case-control study,

### 7. Conclusions

Breast cancer is caused by the complex interplay of genetic, epigenetic, environmental, and reproductive factors and is the most predominant malignancy among women. In the past two decades, a number of large population-based studies have been conducted to determine whether variants in the VDR gene polymorphism are associated with breast cancer risk. However, studies have yielded conflicting results. On the basis of the data collected from different studies conducted on different ethnicities, we cannot make a definitive statement regarding the role of VDR polymorphisms in breast cancer and the interaction of steroid hormone receptor status. However, our illustrated overview of vitamin D-related effects on breast cancer that will provide new insights. Nevertheless, because vitamin D via VDR acts in a multitude of ways on cancer cell biology, including the cell cycle regulation, apoptosis, invasion, and metastasis, as well as angiogenesis, it has brought together a wide-ranging impact on tumor development under its domain. Wellpowered epidemiological studies with a large sample size accompanied by molecular investigations are warranted in the future to understand the relationship between VDR variants, hormone receptor status, and 1,25(OH)2D3 concentrations, and the underlying molecular mechanisms that contribute to the pathophysiology of breast cancer and would enable the development of new prevention strategies.

# Author's contribution

All the authors have contributed equally. All the authors read and approved the final manuscript.

# Declaration of competing interest

Authors declare that they have no conflict of interests.

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Review Article

Biochemistry for better diagnosis and therapy



# Role of Vitamin D Receptor Gene Polymorphism with Steroid Receptors in Breast Cancer: an Update

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Abstract: This review touches on new insights into the possible role of Vitamin D receptors genetic polymorphisms, and steroid receptors in the patients diagnosed with breast cancer. This is the most common cancer in women; further, it creates diverse illnesses among the diseased patients, and the prognosis is linked with the different subtypes present in the hormone receptors. Most of the review studies focus on the epidemiology of the disease. However, fewer studies are done on the genes polymorphism prognosis, which plays a more significant role in the prognosis and diagnosis of breast cancer, so by proper screening at the genetic level as early, can predict the disease in females which will help the clinicians in better management of the disease. Hence the burden of disease and its ill effects can be reduced in the patient care system. Thus the Vitamin D levels in the serum and the vitamin D receptors (VDR) transcriptionally controls its target genes in the cell proliferation, differentiation, and death pathways in a ligand-dependent manner, providing protection against cancer growth and progression. Also, there are strong relationships between VDR polymorphisms and steroid hormone (estrogen, progesterone, and androgen) receptors, which will help in the prognosis and diagnosis of breast cancer disease. We recommend early screening of these receptors using advanced molecular biology techniques like Real-time Polymerase chain reaction, Chemiluminescence, Western blotting which will help to detect the genotyping of these genes at the earlier stages and are non-invasive, patient-friendly, reliable, and accurate. Vitamin D receptor gene polymorphism and steroid receptors themselves can act as early predictive biomarkers for many studies that are to be warranted further in the different ethnic populations with large sample sizes.

Keywords- Breast cancer, steroid receptors, prognosis, Vitamin D Receptor (VDR), polymorphisms.

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# I. INTRODUCTION

Breast cancer is the most frequent cancer in women, accounting for more than one out of every ten new cancer diagnoses each year. It is also the second most common cancer among females 1,2. Despite remarkable advancements in cancer therapy, the heterogeneity of breast cancer and the development of anticancer drug resistance remain key hurdles for effective treatment 3. The majority of breast cancers are caused by the uncontrolled multiplication of epithelial cells in the mammary gland's ducts and lobules 4. Having a first-degree relative with the disease, extremely thick breasts, past benign breast biopsy results, being on oral contraceptives, and having the first child at 30 years or older are all risk factors for breast cancer 5. Furthermore, modifiable risk factors such as alcohol consumption and a high-fat diet contribute to the disease incidence 6. The vitamin D receptor (VDR) is a mediator for the cellular effects of vitamin D and interacts with other cell signalling pathways that influences the cancer development. The biologically most active form of vitamin D is 1,25(OH)2D, which mainly exerts its anti-proliferative effects by binding to the vitamin D receptor (VDR) and acting in the complex as a transcriptional factor for a variety of genes, including those involved in cell differentiation and cell growth 7,8. Higher vitamin D levels are thought to protect against a variety of malignancies, including breast cancer, through genomic effects controlled by the vitamin D receptor (VDR) 9. There is mounting evidence that vitamin D is linked to a lower risk of breast cancer 10. Apart from calcium homeostasis, vitamin D and its receptor (VDR) promote cell differentiation and suppress the proliferation of cells that have cancer-preventive capabilities 11. The human VDR gene, which is located on chromosome 12q13-14, has about 470 single-nucleotide polymorphisms (SNPs). Fok1 (rs2228570), Bsm1 (rs1544410), Poly A (rs17878969), Apal (rs7975232), and Taql (rs7975232) are the most investigated SNPs 12. Breast cancer risk is controlled by the prognosis of VDR gene polymorphism and steroid receptor status 13,14. The polymorphism of the vitamin D receptor (VDR) is based on single nucleotide polymorphisms that might interfere with the activity of vitamin D which affects the prognosis of tumor <sup>15</sup>. This review touches on the role of Vitamin D receptor gene polymorphisms and steroid receptors status in prognosis and diagnosis of breast cancer as early biomarkers that can help clinicians to manage the disease better. Previously, many literature reviews on breast cancers touched on the epidemiology and prevalence, but no such study has been directly found which has been done on the plausible role of vitamin D receptor gene polymorphisms and on the steroid receptors status, if any are less and being done on single genes. For that, a comprehensive literature review has been done systematically to assemble all the latest material in all ethnic populations and prepare the same for the better management and prognosis of the disease. The aim and objective of this review are to offer a complete assessment of the role of VDR polymorphism and steroid receptor status in terms of modifying impacts on breast cancer risk, severity, progression rate, and disease prognosis, which will help the clinicians in improving the health index of patients.

# 2. METHODOLOGY

Using the National Library of Medicine's (NLM) PubMed, Medline and Scopus database search of previous years, we conducted an updated comprehensive evaluation of published research. Keywords including 'VDR gene polymorphisms,' 'Steroid receptors status,' and 'breast cancer' were utilized in

the search. Articles on the role of VDR gene polymorphisms and steroid receptor status in relation to breast cancer risk and incidence were included in our analysis. Animal studies and research on breast cancer in men or teenagers were omitted from our investigation. The quality of the evidence for each part of the statement was rated as high (randomised controlled trial (RCT) evidence - level 1), moderate (intervention short of RCT or large observational studies level 2), or low (case series, case reports, expert opinion level 3). The clinical significance and weight of opinion favouring each statement were used to determine whether it was strong (S - recommendation) or weak (W - suggestion). Strong suggestions are clinically relevant best practices that will be applied to the majority of patients in the majority of circumstances. In contrast, weak statements should be examined by the clinician and will be applicable best practices only to specific patients or in certain circumstances.

# 3. Vitamin D receptors and breast cancer prognosis

Vitamin D interact with VDR in the breast epithelium in the same cell or in nearby cells in order to keep the cell differentiated and quiescent 16. By studying case-control samples from breast cancer samples, Hemida et al., showed that the expression of VDR in breast cancer tissues was elevated, and an association exists with the expression of estrogen receptor alpha 17. Heublein et al., and Huss et al., conducted retrospective studies which indicated that low VDR expression is a poor prognostic indicator for breast cancer 18,19. Currently, the number of tumor node metastasis (TNM) is the most significant factor in determining how breast cancer patients will respond to treatment. However, individual differences reduces the accuracy of prediction. It is therefore imperative that breast cancer patients have a reliable biological indicator that can help predict their prognosis 20. Despite the fact that vitamin D levels and VDR expression are associated with breast cancer prognosis <sup>21,22</sup>, it seems that the expression of VDR is more reliable for considering the prognosis of breast cancer since vitamin D levels changes that are highly dependent on dietary exposure to sunlight. Furthermore, VDR protein expression has prognostic significance, but it has also been reported that expression of VDR mRNA has prognostic value in breast cancer <sup>23,24</sup>. Further studies are needed to elucidate the roles of expression and polymorphism of VDR mRNA in breast cancer.

# 4. Steroid receptor status concerning Breast cancer

Estrogen, Progesterone, and Androgen receptors (ER, PR, and AR), like VDR, are members of the nuclear receptor superfamily  $^{25}$ .

# 4.1 Estrogen Receptor

The estrogen Receptor (ER) is widely acknowledged as a significant predictor of the start of breast cancer in women. According to a survey done by the Department of Surgical Oncology and the Division of Molecular Medicine in India, 50.8 percent of patients were found to be ER-negative, whereas 49.2 percent were found to be ER-positive <sup>26</sup>. Surveillance, Epidemiology, and End Results (SEER) studies, on the other hand, showed that ER is negative in more than double the proportion of patients with breast cancer. It was discovered that 50.8 percent of breast cancer patients had negative ER, compared to 23.4 percent who had positive ER. The SEER reports also revealed that 24.8 percent of patients had Triple-

negative (TPN) tumors, compared to 15 percent of patients with TPN tumors, and that 76.4 percent of patients had a median age of 53 years. The study found that ER-negative increased among young women, with 63.5 percent of those under 50 years old, and dropped as women's ages grew 27. In the primary tumor stage, ER staining was found in 74.2 percent of Whites, 13.7 percent of Blacks, 8.5 percent of Hispanics, 3.0 percent of Asians, and 0.6 percent of Others, respectively. It was discovered that patients with ER-positive rates ranging from 1% to 9% are more likely to get chemotherapy than patients with ER-positive rates greater than 10% who are treated with endocrine treatment. The presence of a positive estrogen receptor in women indicates an increased risk of breast and ovarian cancer. As a result, positive estrogen receptor cells boost surrounding cell proliferation, release paracrine hormones, and promote epithelial cell development<sup>28</sup>. Negative ER cells, on the other hand, aid mammary gland growth and restore the proliferation process. Positive estrogen receptor cells are identified in 20% of sporadic breast cancer cases, while negative estrogen receptor cells are seen in 70% to 80% of instances. When the size of the tumor grows large enough to reduce the estrogen effect on the patients, the ER-positive cells undergo metamorphosis and eventually perish. As a result, the negative estrogen receptor cell, like the positive estrogen receptor cell, undergoes neoplastic change as it grows older. As the tumor increases in size, positive estrogen receptors become differentiated, while negative estrogen receptors become stimulated by mitogenic signaling 29.

# 4.2 Progesterone Receptor

Progesterone receptors are members of the nuclear subfamily 3 (NR3C3) and belong to the protein group C. Positive progesterone receptors are malignant cells in the breast that are extremely sensitive to progesterone and have receptors that help hormones proliferate. Hormone therapy is used to inhibit estrogen receptors in those who have positive progesterone receptor cancer. Patients with negative progesterone receptor cancer, on the other hand, are not suggested to receive hormone therapy because it was ineffective for them 30. When looking at the positive and bad aspects of progesterone receptors in India, it was discovered that they were present in 49.8% of patients suffering from breast cancer. There are 204 cases of lobular carcinoma and 50 cases of invasive lobular carcinoma. Invasive ductal carcinoma was found in 817 Indian women, and breast cancer was found in 16 of every 1087 Indian women. The Indians had a more significant percentage of progesterone receptors, which exhibited negative breast cancer proportions 31. According to a survey conducted by the National Institutes of Health in the United States, 67.2 percent of women with breast cancer in the United States have positive progesterone receptors as well as positive estrogen receptors, while 19.0 percent of women with breast cancer have negative progesterone receptors and negative estrogen receptors. It was also shown that 12.2 percent of breast cancer patients had negative progesterone-receptors with positive estrogen-receptors, whereas 1.6 percent had positive progesterone-receptors with negative estrogen-receptors. It was also discovered that women with breast cancer who had positive progesterone-receptor and positive estrogen-receptor symptoms have a 30%-60% decreased likelihood of dying from the

# 4.3 Androgen Receptor

The nuclear receptor known as the androgen receptor is strongly linked to the development of the prostate. It's a

breast cancer expression that also recognizes luminal genes and the luminal-androgen receptor (LAR) subtype. The expression of numerous luminal genes is linked to positive androgen receptors, which are also linked to the HER2 pathway. Furthermore, when compared to positive androgen receptors, androgen receptors were negatively related with grade I/II vs III malignancies in high proportions <sup>33,34</sup>.

# 5. VDR polymorphism and Steroid receptor status concerning Breast Cancer prognosis.

The steroid hormone receptor status has an impact on the link between vitamin D and breast cancer risk. Reduced 25(OH)D has been linked to an increased incidence of advanced breast cancer and, to a lesser extent, ER+ PR+ breast cancer, which has a better prognosis 35. Due to its lack of relationship with numerous determinants or markers of breast cancer, such as tumour type, lymph node status, hormone receptors, Ki-67 expression, and p53 levels, the VDR polymorphism was previously not considered a predictive factor for breast cancer  $^{36}$ . Hormone receptor status, Ki-67, triple-negative status, and tumour size are all linked to lower VDR expression, according to a recent study37. Because of the lack of conclusive research in this area, more research is needed. VDR expression varies greatly between different types of breast cancers, with studies reporting almost 90% expression in ER+ tumours compared to only 27% in basal/triple-negative tumours, implying an inverse association between VDR expression and cancer severity 38. Abbas et al., found a link between the Taql VDR polymorphism (containing at least one copy of the t allele) and ER-positive postmenopausal breast cancer <sup>39</sup>. A recent investigation found that Fokl polymorphisms have a similar favorable effect on the development of ER+ cancer in Saudi women patients 1 Premenopausal Chinese women with both the aa genotype of the Apal polymorphism and the ER haplotype experienced a delayed onset of menarche. This finding is substantial as the age of menarche is a significant risk factor for breast cancer 40. The steroid hormone receptor status is known to influence the relationship between vitamin D and breast cancer risk Various processes such as frequency identification of FFLL and FfLL genotypes against Fokl and poly(adenylate) grouping, comparing tumor grade, lymph node involvement, and estrogen receptor (ER) status among cancer patients, and VDR genotype are some of the methods that aid in determining the levels, stages, and progression of cancer in patients. For example, adjusted odds ratios (OR) for age at sampling, HRT use, and menopausal state at diagnosis were observed to be 1.12 (0.62-2.04) in the identification of total tumor grade in tumor grade I, FFLL or FfLL. As a result, it can be concluded that the use of Fokl, poly (adenylate) grouping, and other techniques aided in gaining a thorough understanding of the tumor grade present in breast cancer patients 42. VDR polymorphisms, which include the bb genotype, are directly linked to the spread of breast cancer. The bb genotype, which was derived from VDR Bsml polymorphisms, was found to have four times the chance of developing metastases as the BB genotype 43. Furthermore, the TT genotype derived from the VDR Tagl polymorphism is strongly linked to an elevated risk of 1.8 lymph node metastases. Females that have more of the haplotype baTL have a higher chance of acquiring metastatic illness, especially in Caucasian female communities 44. As a result, VDR polymorphisms are connected to the development of breast cancer risks in individuals.

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# Heterogeneity in VDR expression concerning Steroid receptors status

The effects of vitamin D endocrine signaling on distinct cell types in mammary tissues are poorly understood. Breast cancer is a diverse illness that develops from a variety of mammary epithelial cell types <sup>45</sup>. The inner luminal layer and outer basal layer of myoepithelial cells make up the mammary epithelium. A recent genome-wide transcriptome research in human tissues found that CYP24AI is expressed differently in luminal progenitor cells, suggesting that the vitamin D pathway may play a role in mammary cell lineage development 46. In normal human breast tissues, major steroid receptors VDR, ER, and AR revealed differential expression between luminal and basal cell types. VDR expression, which is commonly coupled with ER and/or AR, was observed during particular stages of luminal cell differentiation and suppression during other stages, according to Santagata et al. 47. In some cases of breast cancer, variations in VDR expression may lead to unresponsiveness or resistance to vitamin D supplementation. Additionally, epigenetic alterations in VDR and CYP24A1 have been linked to vitamin D resistance 48. Individuals with malignancies that are positive for ER, VDR, and AR all have a better prognosis 49. These findings highlight the need of assessing VDR polymorphisms and steroid receptor status in breast cancer samples at the same time 50,51.

# 7. CONCLUSIONS

VDR gene polymorphism is a highly effective indicator for predicting and assessing the beginning and progression of

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breast cancer when taken collectively. Further research on the link between VDR genetic polymorphisms and breast cancer incidence in patients who are classified based on their steroid receptor (ER, PR, and AR) status is required. This will allow researchers to explore the relationship between VDR and the molecular subtypes of breast cancer, which will aid in the development of more tailored therapies for patients. In this review article more has been added related to breast cancer. prognosis, and diagnosis which will help the clinicians to establish new treatment protocols in systematically managing the diseases. Further studies are needed for eradicating the diseases by targeting the gene at the protein level in the human body system, in order to improve the health index of our woman gender. Therapeutic approaches are needed for best treatment protocols and will curtain fewer surgical procedures.

# 8. AUTHOR'S CONTRIBUTION STATEMENT

Ashok Kumar Dogra, conceptualized, prepared the original draft, reviewed, edit the draft, and designed the study; Dr. Pranav Prakash curated data; Dr. Sanjay Gupta, discussed methodology; Dr. Meenu Gupta, review the draft; Dr. Archana Prakash, analyzed and revised the draft; Haamid Bashir, provided valuable inputs towards designing the manuscript. All authors approved, read, and approved the final version of the manuscript.

# 9. CONFLICT OF INTEREST

Conflict of interest declared none

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# Vitamin D and Vitamin D Receptor Fokl, Apal, and Bsml Gene Polymorphisms and their Relation with the Risk of Breast Carcinoma: A Case-control Study

Biochemistry Section

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# **ABSTRACT**

Introduction: Breast cancer stands as the leading cause of mortality among women in developing nations. The potential role of Vitamin D in mitigating the incidence of breast cancer is thought to stem from its ability to impede cell proliferation by interacting with the Vitamin D Receptor (VDR). The VDR gene is responsible for encoding the VDR, which plays a pivotal role in mediating the effects of vitamin D.

**Aim:** To analyse vitamin D levels and the association of VDR Fokl, Apal, and Bsml genotypic distribution frequency with the risk of breast cancer.

Materials and Methods: The case-control study included 220 samples, including 110 breast cancer patients and 110 agematched control women aged 30-70 years. The Polymerase Chain Reaction-Restriction Fragment Length Polymorphism (PCR-RFLP) genotyping was performed using Deoxyribonucleic acid (DNA) extracted from blood, and the circulating levels of 25-hydroxyvitamin D by case/control were estimated by chemilluminescence immunoassay.

Results: The 3' VDR polymorphism Bsml sequence showed minimal association with breast cancer risk. The bb genotype

had a significantly lower odds ratio of 0.056 (p-value <0.05). Conversely, the BB and Bb genotypes exhibited no statistically significant associations with odds ratios of 1.76 (95% CI: 0.36-8.54; p-value >0.05) and 1.30 (95% CI: 0.27-6.25; p-value >0.05), respectively. Isolated analysis of the Fokl variant revealed a significant association with increased breast cancer risk, with odds ratios of 5.49 (FF) and 6.00 (Ff), both demonstrating statistical significance (p-value <0.05), and a Chi-square value of 0.006. Additionally, the p-value for serum Vitamin D levels was found to be highly significant at p-value <0.001, indicating that the levels were significantly lower in individuals newly diagnosed with breast cancer compared to those in the healthy control group.

Conclusion: The study found a significant link between breast cancer susceptibility and VDR (Fokl) polymorphism FF and Ff genotypes, with minimal impact observed for (Bsml) polymorphism bb genotype. This implies that certain genetic variations, especially in the Fokl polymorphism of the VDR gene, are associated with an elevated risk of breast cancer.

**Keywords:** Apal single nucleotide polymorphisms, Breast cancer, Bsml single nucleotide polymorphisms, Fold single nucleotide polymorphisms

# INTRODUCTION

Every year, breast cancer leads the world in morbidity and mortality rates for women [1]. In India, this disease accounts for 30.1% of all female cancer cases (Globocan 2020; http://globocan.iarc.fr/). The Indian subcontinent has seen an increase in cancer incidence, mortality, and morbidity [2-5]. The increasing mortality is likely due to ineffective screening strategies, advanced-stage diagnosis, and inadequate medical facilities. Recent studies have indicated that vitamin D may play a role in the development of breast cancer. In several studies, there is evidence that low levels of 25hydroxyvitamin D {25(OH)D} increases the risk of breast cancer and have also shown an association between dietary intakes, dietary supplements, and skin production of vitamin D and breast cancer risk [6-8]. Aside from the classical role of vitamin D in calcium and phosphorus homeostasis, calcitriol exerts anticancer properties through transcriptional and/or non genetic mechanisms [9]. The active vitamin D3 (1 $\alpha$ ,25-dihydroxy vitamin D3) exerts its biological effects via the VDR. It belongs to the family of nuclear receptors and is a ligand-dependent transcription factor [10]. Cell cycle arrest, senescence, differentiation, and apoptosis are induced by the Vitamin D-ligand VDR in a variety of tumour types [11].

There are several polymorphisms in the coding and non coding regions of VDR on 12q13.11. Numerous Single Nucleotide

Polymorphisms (SNPs) have been identified in and around exons 2-9 as well as in the 3' UTR region of the VDR gene [12]. The most commonly studied SNPs are those containing Restriction Fragment Length polymorphisms (RFLPs) rs1544410, rs2228570, and rs7975232, determined by restriction endonucleases Bsml, Fokl, and Apal [13-15]. The VDR contains the Bsml SNP (A/G) in intron eight near the 3' end, and its effect on VDR protein expression and activity is unclear. However, in Caucasians, Chinese, and Japanese Americans, it is in strong linkage disequilibrium with a polyadenosine microsatellite repeat, which may affect mRNA stability or translation activity [16]. A Fokl site present in the 5' promoter region substitution results in thymine (T) to cytosine (C) that changes the first of two possible translation initiation sites, resulting in different-sized VDR proteins. An f allele is three amino acids longer than an F allele and transcriptionally less active [17]. An Apal SNP (C/A) variable site is located in intron 8 of the VDR gene. VDR polymorphisms may alter expression and function in breast cells, thereby modulating breast cancer risk [18].

Several studies performed on Caucasian populations have given inconsistent results regarding Fokl, Bsml, and Apal SNPs and breast cancer risk [19-22]. There have been very few studies on Asian populations. An association was found between the Fokl SNP and Japanese-American women in the Hawaii-Los Angeles

Multiethnic Cohort (MEC), but not in large Chinese studies or small Iranian studies [23-25]. A Bsml SNP was associated with Iranian women and Japanese-American women from the Multiethnic Cohort (MEC) study, but not with two Chinese studies [26]. The Apal SNP showed mixed results in different populations and was primarily studied in African Americans, Caucasians, and Chinese [27-29]. In the present study, three polymorphisms, one from the 5' region (Fokl) and two from the 3' region of the VDR gene (Apal and Bsml), and vitamin D, were investigated to assess the association with the risk of breast cancer. The study aims to offer insights into genetic factors impacting breast cancer susceptibility for personalised risk assessment and prevention.

# **MATERIALS AND METHODS**

The present study was a hospital-based age-matched case-control study conducted at the Cancer Research Institute, a tertiary care centre located at Jolly Grant, Dehradun, Uttarakhand, India during the period from the year 2020 to 2023. This study included 110 freshly diagnosed breast cancer patients and age-matched 110 healthy controls aged 30 to 70 years. The institute's ethics committee approved the study, which was conducted in accordance with all the provisions of the Declaration of Helsinki (Letter No. SRHU/HIMS/ETHICS/2020/193). A written informed consent was obtained from all study participants.

Inclusion and Exclusion criteria: The inclusion and exclusion criteria were primarily used to select patients. The inclusion criteria included breast cancer patients selected based on histopathological confirmation, both pre- and postmenopausal women, and excluding those on hormonal therapy, with other cancers, recent Vitamin D supplementation, or pregnant or lactating. Age-matched healthy female volunteers were included as controls.

Sample size estimation: The sample size was estimated using the n-Master software for a matched case-control study (1:1) matching. Assuming that the proportion of exposed controls is 50% and the level of significance is 5% with a power of 90% to detect a two-fold increase in risk. The minimum number of required discordant pairs is 110.

Data collection: A comprehensive proforma was used to capture demographics, co-morbidities, family history, and anthropometric information, as well as pertinent clinical information from our online hospital database. The tumour morphology was classified according to criteria; Elston and Ellis used architectural aspects, nuclear differentiation levels, and mitotic index based on the 8<sup>th</sup> edition of the TNM staging system for breast cancer developed by the American Joint Committee on Cancer (AJCC) [30]. The quality of genomic DNA was assessed through agarose gel electrophoresis, and VDR gene polymorphisms were genotyped using PCR-RFLP analysis.

The Vitamin D levels in serum were determined for all freshly diagnosed cases and controls using the Chemiluminescent Immunoassay (CIA) method by trained laboratory technicians. Serum 25(OH)D levels were classified based on our institution's laboratory reference standards as sufficient/normal (75-250 nmol/L), insufficient (50-<75 nmol/L), and deficient (<50 nmol/L) [31]. Results were expressed in nmol/L.

**Genotyping analysis:** Fold, Bsml, and Apal genotyping utilised PCR-RFLP analysis, employing agarose gels for DNA quality confirmation. The Fold polymorphism was detected using the following primers:

Forward: 5'GAT GCC AGC TGG CCC TGG CAC TG 3' and Reverse: 5'ATG GAA ACA CCT TGC TTC TCC CTC 3', yielding a 272 bp fragment spanning the Fokl site (Raza S et al., 2019) [32]. The Bsml polymorphism was detected using the following primers: Forward: 5'CAACAAGACTACCAGTACCGC GTCAGTGA3' and Reverse:

5'AACCAGCGGAAGAGGTCAAG GGG 3', generating an 825 bp fragment surrounding the Bsml site (Raza S et al., 2017) [33]. The Apal-RFLP was detected by the following primers: Forward: 5' CAG AGC ATG GAC AGG GAG CAA G 3' and Reverse: 5' CGG CAG CGG ATG TAC GTC TGC AG 3', yielding a 352 bp fragment spanning the Apal site (El-Shorbagy HM et al., 2017) [29]. The following conditions were used for the PCR: initial denaturation at 94°C for three minutes, followed by 34 cycles of cyclic denaturation at 94°C for one minute, annealing 50 seconds at 71°C for Fokl, 71°C for Apal, and 58°C for Bsml, then extension at 72°C for one minute and one final cycle of final extension at 72°C for eight minutes, and final hold at 4°C. After PCR, the amplified PCR products were digested according to the manufacturer's instructions with Fokl, Bsml (New England Biolabs, USA), and Apal (Promega). In 2% agarose, fragments were stained with ethidium bromide to determine whether the enzyme recognition site was present (lowercase) or absent (uppercase). The genotypes for VDR-Fokl (FF, Ff, ff), VDR-Bsml (BB, Bb, bb), and VDR-Apal (AA, Aa, aa) polymorphisms were assigned. Randomly selected samples of three genotypes including Homozygous dominant, recessive, and heterozygous were confirmed by SNP sequencing, and the results were 100% concordant.

# STATISTICAL ANALYSIS

Statistical analysis, including odds ratios and Chi-square tests, were used to evaluate associations between specific VDR gene polymorphisms and breast carcinoma risk. Data entered into Microsoft Excel 2010 were analysed using statistical software version Statistical Package for Social Sciences (SPSS) 20.0. Normality was assessed by the Kolmogorov-Smirnov test. An Independent t-test was used for two groups, and Analysis of Variance (ANOVA) for more than two groups to compare mean differences. The deviation from Hardy-Weinberg Equilibrium (HWE) was tested for polymorphisms by examining the differences between genotype frequencies observed and those expected, utilising the  $\chi^2$  test. Descriptive statistics and graphical representations were used to enhance the result interpretation. It is considered statistically significant when the p-value <0.05. and statistically insignificant if the p-value >0.05.

# RESULTS

In this hospital-based case-control study, 110 patients and 110 healthy controls were compared. Among the study participants, demographic characteristics and risk factors were analysed, with predominantly 37 (33.6%) cases and 40 (36.4%) controls falling within the 40-49 age range. Urban residency accounted for 70% of cases and 86.4% of controls, while rural and Semiurban areas had lower frequencies. Premenopausal status was balanced, with 58.2% of cases and 61.8% of controls. Normal BMI was observed in 58.2% of cases and a higher percentage in controls (85.5%). Notably, 12.7% of cases had a positive first-degree family history, contrasting with the absence of such history in controls [Table/Fig-1]. Based on TNM staging, tumour morphology among 110 breast cancer patients showed Grade-I tumours with one case having vitamin D <50 nmol/L, two cases with 50 to <75 nmol/L, and

Characteristics		Cases (%)	Controls (%)	
	30-39	31 (28.2)	34 (30.9)	
A	40-49	37 (33.6)	40 (36.4)	
Age (years)	50-59	28 (25.5)	27 (24.5)	
	60-69	13 (11.8)	9 (8.2)	
	≥70	1 (0.9)	0	
	Rural	22 (20)	6 (5.5)	
Areas	Urban	77 (70)	95 (86.4)	
	Semiurban	11 (10)	9 (8.2)	
	Premenopausal	64 (58.2)	68 (61.8)	
Menopausal	Postmenopausal	46 (41.8)	42 (38.2)	

	Normal	64 (58.2)	94 (85.5)	
	Underweight	5 (4.5)	9 (8.2)	
BMI category	Overweight	37 (33.6)	7 (6.4)	
	Obese Class-I	4 (3.6)	0	
First degree family history	Yes	14 (12.7)	0	

[Table/Fig-1]: Demographic characteristics of breast cancer patients and their matched controls.

none with 75-250 nmol/L (p-value=0.433). For Grade-II tumours, 51 (67.1%) had <50 nmol/L, 20 (26.3%) had 50 to <75 nmol/L, and 5 (6.6%) had 75-250 nmol/L (p-value 0.433). Grade-III tumours had 21 (67.7%) with <50 nmol/L, 8 (25.8%) with 50 to <75 nmol/L, and 2 (6.5%) with 75-250 nmol/L (p-value 0.433). No significant associations between tumour grades and vitamin D levels were found, contributing to understanding the tumour grade-vitamin D status relationship in this patient population.

The genotypes and allele frequencies of Fokl, Apal, and Bsml were illustrated in [Table/Fig-2]. In both cases and controls, the HWE of 0.15 indicates equilibrium. In accordance with HWE, genotypic data were found to be reliable, indicating that selection or genetic drift has little influence on the genotypic distribution.

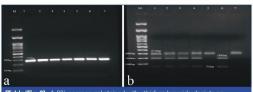
The examination of the Apal (rs 7975232) VDR polymorphisms revealed an amplification product with a size of 352 bp [Table/Fig-4a]. On agarose gels, 352 bp bands were genotyped as AA homozygotes. The homozygote (AA) produces 216 bp and 135 bp fragments, while heterozygotes (AA) display three fragments, 352 bp, 216 bp, and 135 bp [Table/Fig-4b]. 40.9% were heterozygous AA, 49.1% were homozygous AA, and 10% were homozygous aa with respect to the Apa1 polymorphism among 110 cases and 110 controls. The Apal genotypes showed no association with breast cancer risk.

The analysis of Bsml (rs1544410) polymorphisms showed an amplified product size of 825bp [Table/Fig-5a]. Two or three fragments showing the Bsm1 restriction site (b) indicated intact amplification (B) reveals the absence. The undigested bands of 825 bp indicated a homozygous BB genotype. bb homozygotes produced two fragments (650 bp and 175 bp), and Bb heterozygotes produced three fragments (825 bp, 650 bp, and 175 bp) on agarose gel [Table/Fig-5b]. Bsm1 polymorphism cases comprised 53.6% heterozygous Bb, 42.7% homozygous BB, and 3.7% homozygous bb. Additionally, 47.3%, 50%, and 2.7% of the control groups were genotyped.

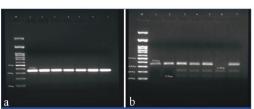
Genotypes		Cases	Allele probabilities (p-value HWE)	Control	Allele probabilities (p-value HWE*)	OR (95% CI)	p-value*	χ² value
	FF	60		64		5.49 (1.72-17.64)	0.004	
Genotypes (Fokl)	Ff	32	F=0.69, f=0.30 (0.15)	42	t=0.22 (0.15)	6.00 (1.83-19.67)	0.003	0.006
(i OKI)	ff	18		4		2	0.181	
	AA	54	A=0.69, a=0.30 (0.15)	62	A=0.75, a=0.24 (0.15)	2.87 (0.92-8.97)	0.069	
Genotypes (Apal)	Aa	45		43		2.31 (0.72-7.37)	0.157	0.241
( spear)	aa	11	u=0.00 (0.10)	5		=	0.518	
000 72	BB	47	Will makes	55		1.76 (0.36-8.54)	0.482	0.546
Genotypes (Bsml)	Bb	59	B=0.69, b=0.30 (0.15)	52	B=0.73, b=0.26 (0.15)	1.30 (0.27-6.25)	0.743	
(Dorrii)	bb	4	5-0.00 (0.10)	3	5-0.20 (0.10)	0.056	0.009	
Total		110		110				

[Table/Fig-2]: An association of the genotypes of breast cancer patients and controls was made in the study "Hardy Weinberg equilibrium." 74.0 05. Significant

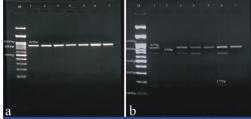
An analysis of FokI VDR polymorphisms yielded the amplification product with a size of 272bp [Table/Fig-3a]. An amplification product without FokI restriction site (f), while present in two or three fragments, indicates FokI restriction site (f). A non digested, single 272bp band genotype FF as homozygous, while Homozygotes (ff) showed two fragments of 198 and 72 bp, and heterozygotes (Fi) showed three fragments of 272, 198, and 72 bp [Table/Fig-3b]. The distribution of polymorphism in VDR FokI showed that 29.1% constituted the Heterozygous Ff, 54.5% were Homozygous FF, and 16.4% presented as homozygous ff cases, whereas the corresponding Control group genotype frequencies were 58.2%, 38.2%, and 3.6%, respectively. There was a significant association between the FokI genotypes FF and Ff and breast cancer risk. With a 95% Confidence Interval (CI), the odds ratios were 5.49 (1.72, 17.64) and 6.00 (1.83, 19.67) with p-values of 0.004 and 0.003.



[Table/Fig-3]: A 2% agarose gel stained with ethidium bromide depicts an amplification product of 272bp was obtained for Fok1. Ladder 100bp is Lane M. Lanes 1-7 show 272bp Fok1 amplified PGR products (a), F (Tallele) is in upper band. C (C allele) is in lower bands. 100bps ladder on lane M. Lanes, 1-3, represer H heterozygotes, Lanes 4-6, represents FF homozygotes and Lane 7 represent if homozygotes (b).



[Table/Fig-4]: A two percent agarcse gel stained with ethidium bromide revealed an amplification product of size 352 bp for Apal polymorphism. Lane M shows a 100 bp ladder and lanes 2-7 show a 352 bp product (a); The upper band indicates a (C allele), lower band indicates a (A allele), 100bp ladder in lane M. There are Aa homozygotes in lanes 1.2 and Aa heterovondes in I ares 3.4 5.7 and as homozogotes in lanes 1.6 and Aa heterovondes in I ares 3.4 5.7 and as homozogotes in ares 6.16 in the control of the contr



[Table/Fig-5]: A two percent agarcse gel stained with ethicium bromide revealed an amplification product of size 825 bp for Bsml polymorphism. Lane M shows a 100 bp ladder and lanes 2-7 shows 825 bp product (a). The upper band indicates B (A allele) lower band indicates b (G allele). 100 bp ladder in lane M. There are BB homozygotes in Lanes 5,7 and Bb heteroxygotes in Lanes 1,3,4,6 and bb homozygote in Lane 2 (b).

A comparative analysis of vitamin D levels in breast cancer patients and the control group indicated mean values and standard deviations. Breast cancer patients had a mean vitamin D level of  $43.54\pm19.58\ \text{nmol/L}$ , while the control group had a higher mean of  $89.89\pm26.13\ \text{nmol/L}$ . The independent t-test showed a highly significant p-value of <0.001, signifying a substantial difference.

These findings highlight potential biomarkers for breast cancer, suggesting that low levels of vitamin D are related to increased breast cancer risk and lead to implications for diagnosis and prognosis in this patient population.

Moreover, a comparative analysis of genotypes and vitamin D levels in breast cancer patients and the control group, focusing on specific genotypes (FF, Ff, ff for Fokl; AA, Aa, aa for Apal; BB, Bb, bb for Bsml) was conducted. Mean vitamin D levels with standard deviations were assessed for both cases and controls. It shows significance in uncovering potential associations between distinct genetic variations and vitamin D levels. Significant differences in mean Vitamin D levels were observed among both cases (p-value=0.006) and controls (p-value=0.001) for different genotypes of Bsml. The data suggested varying degrees of association between genotypes and vitamin D levels [Table/Fig-6].

		Cases	Control	
Genot	ypes	Mea	n±SD	
	FF	42.98±21.32	46.33±26.08	
E-IV	Ff	55.58±19.04	72.96±25.59	
Fokl	ff	81.77±16.40	114.71±18.66	
	p-value*	0.756	0.067	
	AA	43.66±17.37	48.90±19.14	
A1	Aa	65.39±21.56	67.90±27.14	
Apal	aa	89.54±19.43	102.67±21.42	
	p-value*	0.316	0.001	
	BB	44.76±22.41	87.70±24.24	
Desert	Bb	56.71±15.52	88.02±22.23	
Bsml	bb	72.23±16.79	162.63±6.05	
	p-value*	0.006	0.001	

[Table/Fig-6]: Comparison of genotypes and Vitamin D levels (nmol/L) among Breast cancer patients and control groups.

# DISCUSSION

Breast cancer prevails as the predominant cancer in women globally and in India, with an age-adjusted prevalence of 25.8 cases per 100,000 women and a fatality rate of 12.7 per 100,000 women [34]. While developed regions still show higher occurrence rates, emerging countries, including India, face increased death rates from breast cancer [35]. Age is a significant risk factor, and present study indicates a notable prevalence of 33.6% in women aged 40-49. Tumour size and lymph node involvement are pivotal prognostic factors, with a common occurrence of lymph node infiltration at diagnosis (30-50% of cases). Metropolitan regions, age 50-59, and premenopausal status were prominent in these cases.

Present study revealed a significant association between the Fokl genotype and the occurrence of breast cancer within the study population. Specifically, the FF genotype showed a substantial increase in breast cancer risk (OR: 5.49, p-value: 0.004), while the Ff genotype was associated with a significant risk reduction (OR: 6.00, p-value: 0.003). Conversely, the ff genotype was less frequent in the control group compared to the case group. These findings are consistent with Mishra DK et al., study on African American and Hispanic populations, as well as Chakraborty M et al., study within the Indian population, highlighting the elevated risk associated with Fok1 FF and Ff genotypes in breast cancer susceptibility [36,37].

In this study, analysis of the Apal genotype distribution revealed a lower frequency of AA genotypes in cases (OR: 2.87, p-value: 0.069), with Aa genotypes showing a slightly higher frequency (OR: 2.31, p-value: 0.157). However, no significant association with breast cancer risk was observed for Apal genotypes. This aligns with the ongoing study by Ahmed JH et al., on the African population, where no significant association was found between the Apa1 polymorphism and the condition under study [28].

Examining Bsml genotypes in present study, it was found that there were no significant associations with breast cancer risk. The Apal genotypes (Aa and aa) revealed notable differences in vitamin D levels between cases and controls (p-values: 0.316 and 0.001). suggesting a potential relationship with breast cancer. These findings were in line with Reimers LL et al., population-based casecontrol study conducted on Long Island, New York, emphasising the influence of vitamin D-related gene polymorphisms on breast cancer susceptibility [38]. However, Bsml genotypes in present study showed no significant association with vitamin D levels. The p-value for serum vitamin D levels was highly significant at 0.001, signifying a substantial decrease in levels among individuals newly diagnosed with breast cancer compared to those in the healthy control group within our studied population. In a study by Ingles SA et al., African-American women with LS and LL poly(A) variations demonstrated a 50% lower risk of breast cancer than those with the SS genotype, particularly in the presence of the FF (Fokl) mutation [39]. Whitfield KG et al., study on human fibroblast cell lines highlighted the statistical significance of VDR activity when both Fokl and poly(A) genotypes were considered together [40]. Present study, aligning with previous research, conducted a comparative analysis of Fokl, Apal, and Bsml genotypes along with vitamin D levels (<50, 50 to <75, and 75 to 250 nmol/L) in breast cancer patients and controls [41,42]. Notably, statistical comparison between the groups for vitamin D levels for those carrying Apal genotypes (Aa and aa) showed significant differences in vitamin D concentrations in cases and controls, and Bsml genotypes (Bb and bb) showed significant differences in vitamin D concentrations in cases and controls, suggesting that vitamin D status may be influenced by these genotypes.

# Limitation(s)

As with any research study, the present study has both strengths and limitations. For the studied analysis, only a few factors were taken into account, and quite a few factors were unmatched. Moreover, only a few VDR polymorphisms are considered for the 5' and 3' ends of this gene. Despite these limitations, this study significantly advanced the understanding of VDR polymorphisms (Fokl, Bsml, and Apal genotypes) and breast cancer risk.

# CONCLUSION(S)

This study identifies an association between VDR(Fokl) polymorphism FF and Ff genotypes and minimal impact for (Bsml) polymorphism bb genotype in breast cancer susceptibility. These findings could be useful in predicting breast cancer risk or whether a woman who has breast cancer will develop metastases. Highly significant serum vitamin D levels between breast cancer and control groups highlight the significant influence of VDR polymorphisms, particularly Fokl, stressing the need for comprehensive studies across diverse ethnic populations to understand VDR gene variations' impact on breast cancer development thoroughly. Moreover, consideration of prognostic risk factors is needed for therapeutic applications in the context of breast cancer. Vitamin D's potential preventive role in breast cancer, achievable through safe and affordable supplementation, emphasises its modifiability. The documented link between vitamin D deficiency and increased breast cancer risk

underscores its public health significance, necessitating larger-scale investigations. VDR abundance in breast cancer tissues suggests potential treatment targets. Research on VDR Fokl polymorphism gains importance, considering its potential moderation by family history.

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# PLAGIARISM CHECKING METHODS: [Jain H et al.]

Plagiarism X-checker: Jan 01, 2024
 Manual Googling: Feb 02, 2024
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72	Ranjit Kaur	3163121	Cancer Research Institute, SRHU, Del Radiation Oncology	5/19/2021	12/6/2021	Haridwar, Uttarakhand	2	7018599451
73	Anandi Chauhan	3183089	Cancer Research Institute, SRHU, Del Radiation Oncology	9/14/2021	12/14/2021	Near OBC bank Balawala, Dehradu		7251021233
74	Neelu Sharma Rihana	3228660	Cancer Research Institute, SRHU, Del Radiation Oncology	11/27/2021 12/15/2021	12/14/2021	Defence Colony, Uttarakhand	2	9719263398
75 76	Rihana Santosh Devi	3229520 3225261	Cancer Research Institute, SRHU, Del Radiation Oncology Cancer Research Institute, SRHU, Del Radiation Oncology	12/15/2021 12/7/2021	12/16/2021 2/3/2022	Saharanpur, U.P. Gaon-Thana Mohalla, Dehradun, I	-	7906920930 7895662402
77	Akila Khan	3229757	Cancer Research Institute, SRHU, Del Radiation Oncology	10/9/2021	2/17/2022	Village- Sirsal, Saharanpur, U.P.	1	9997777745
78	Mamta Darrhani Dovi	3246346	Cancer Research Institute, SRHU, Del Radiation Oncology	1/11/2022	2/17/2022	Chandrabani, Dehradun, Uttarakh		8755958325
79 80	Darshani Devi Geeta Devi	3234837 3247576	Cancer Research Institute, SRHU, Del Radiation Oncology Cancer Research Institute, SRHU, Del Radiation Oncology	12/22/2021 1/28/2022	2/18/2022 3/15/2022	Dandi Motharowala, Uttarakhand New Nahar Colony, Bahadrabad, H		7425902032 9719314839
81	Israna	3263025	Cancer Research Institute, SRHU, Del Medical Oncology	3/5/2022	3/17/2022	Dehradun, Uttarakhand	2	8923440311
82	Savita Sharma	3307335	Cancer Research Institute, SRHU, Del Radiation Oncology	5/16/2022	5/20/2022	Muzaffarnagar, U.P.	2	8057032846
83 84	Swati Sharma Rina	3302697 3307155	Cancer Research Institute, SRHU, Del Surgical Oncology Cancer Research Institute, SRHU, Del Radiation Oncology	5/21/2022 5/14/2022	5/21/2022 5/21/2022	Haridwar, Uttarakhand Moradabad, U.P.	2	9997005385 7300768877
85	Tarannum Bee	3307155	Cancer Research Institute, SRHU, Del Radiation Oncology  Cancer Research Institute, SRHU, Del Radiation Oncology	5/21/2022	5/21/2022	US Nagar, Dehradun	2	9756610755
86	Saroj	3317214	Cancer Research Institute, SRHU, Del Radiation Oncology	3/11/2022	6/3/2022	Bijnor, U.P.	2	8006584159
87	Shanti Ray	3321223	Cancer Research Institute, SRHU, Del Medical Oncology	5/10/2022	6/10/2022	US Nagar, Dehradun	2	8126138372
88 89	Meena Bisht Tabassum	3321300 3321396	Cancer Research Institute, SRHU, Del Medical Oncology Cancer Research Institute, SRHU, Del Radiation Oncology	6/1/2022 5/17/2022	6/10/2022 6/10/2022	Lane 7, E-Block, Dehradun Mohalla Neel Khudan, Haridwar	2	9410317791 7037850320
90	Laxmi Devi	3324685	Cancer Research Institute, SRHU, Del Radiation Oncology  Cancer Research Institute, SRHU, Del Radiation Oncology	6/10/2022	6/16/2022	Village, Sambhal, U.P.	1	8941080715
91	Urmila Rawat	3296137	Cancer Research Institute, SRHU, Del Radiation Oncology	5/4/2022	6/17/2022	Village Dungri, Tehri-Garhwal	3	9520606317
92	Nardei	3360721	Cancer Research Institute, SRHU, Del Radiation Oncology	8/20/2022	8/30/2022	Uttarkashi, Uttarakhand	2	9084366689
93 94	Rihana Parveen Arti Devi	3360354 3362592	Cancer Research Institute, SRHU, Del Radiation Oncology Cancer Research Institute, SRHU, Del Radiation Oncology	8/18/2022 7/12/2022	9/29/2022 11/22/2022	Sharanpur U.P. 104 Chhabra Ward No. 2, Dehradu	2	9058106263 9719019316
95	Suman Chaudhary	3404129	Cancer Research Institute, SRHU, Del Radiation Oncology	11/28/2022	11/28/2022	Sahabpur, Bijnor, U.P.	2	9847590342
96	Kusumlata	3377974	Cancer Research Institute, SRHU, Del Radiation Oncology	9/27/2022	11/30/2022	Shekhpur, Bijnor, U.P.	2	9837589045
97	Sureshna Devi	3357571	Cancer Research Institute, SRHU, Del Radiation Oncology	8/20/2022	11/30/2022	96- Subhash Nagar, Dehradun, Utt		7906610175
98 99	Urmila Negi Neha Nautiyal	3389695 3403883	Cancer Research Institute, SRHU, Del Medical Oncology Cancer Research Institute, SRHU, Del Radiation Oncology	10/28/2022 6/25/2022	12/2/2022 12/5/2022	Village Tuntowala, Mehuwala, Utta Raipur, Dehradun, Uttarkhand	: 1 2	7535908143 7895531917
100	Poonam	3385899	Cancer Research Institute, SRHU, Del Radiation Oncology	9/15/2022	12/7/2022	Gram Majra, Malakpur, Haridwar,		7409663675
101	Laxmi Devi	3400862	Cancer Research Institute, SRHU, Del Radiation Oncology	11/28/2022	12/16/2022	Village Chaudel, Pauri, Uttarakhar	13	9548769730
102 103	Santosh Devi Sushma Kalra	3394775 3400711	Cancer Research Institute, SRHU, Del Radiation Oncology Cancer Research Institute, SRHU, Del Radiation Oncology	11/9/2022 8/31/2022	12/23/2022 12/24/2022	Village Asmauli, U.P. Teachers Colony, Udham Singh Na	1	8923375043 9368466587
103	Bharti Chand	3383213	Cancer Research Institute, SRHU, Del Radiation Oncology Cancer Research Institute, SRHU, Del Radiation Oncology	8/31/2022 8/31/2022	12/24/2022 12/27/2022	Shiv Mandir, Saspur, Dehradun, U		7906076547
105	Indu Vaish	3408003	Cancer Research Institute, SRHU, Del Radiation Oncology	12/8/2022	1/3/2023	Prem Nagar, Dehradun, Uttarakha	2	8864823232
106	Shivani Aniana Davi	3408114	Cancer Research Institute, SRHU, Del Radiation Oncology	12/6/2022	1/3/2023	Nathuwala, Dehradun, Uttarakhar		9897383045
107 108	Anjana Devi Neema Bora	3410629 3420610	Cancer Research Institute, SRHU, Del Surgical Oncology Cancer Research Institute, SRHU, Del Radiation Oncology	12/31/2022 1/10/2023	1/20/2023 1/20/2023	Aamwala Nalapani, Tapowan, Deh 208- Siv Nagar, Dehradun, Uttarak		9761852539 7989415106
109	Farjana	3406601	Cancer Research Institute, SRHU, Del Radiation Oncology	11/19/2022	1/24/2023	Mawakot, Pauri, Uttarakahand	3	7055245983
110	Savitri Devi	3422664	Cancer Research Institute, SRHU, Del Radiation Oncology	1/13/2023	1/25/2023	Dehradun, Uttarakhand	2	9084925659

Age 46		Menopausal status( Pre=1, post=2) 1		First Degree Family history (Yes=1, None=2) 2	History of any other Cancer None	Weight(kg) 52	Height(cm) 159	BMI(kg/m²) 20.6	BMI Category (Normal=1, Underweight=2, Overweight=3, Obese class I=4) 1
50 47		2	50 47	1 2	None	54 70	154 162	22.8 26.7	1 3
39		1	39	1	None None	50	157	20.3	1
51 46		2	51 46	1 2	None None	54 65	157 157	21.9 26.4	1 3
54	Female	2	54	2	None	58	152	25.1	3
54 51		2 2	54 51	1 2	None None	62 53	157 162	25.2 20.2	3 1
42		1	42	2	None	40	152	17.3	2
63 67		2 2	63 67	1 1	None None	38 66	142 157	18.8 26.8	1 3
45 43		1	45 43	2 2	None None	49 61	164 159	18.2 24.1	2 1
39	Female	1	39	1	None	63	157	25.6	3
57 48		2	57 48	2 2	None None	65 51	144 152	31.3 22.1	4
51	Female	2	51	2	None	66	160	25.8	3
43 59		1 2	43 59	2 2	None None	80 64	162 157	30.5 26	4 3
44 50		1 2	44 50	2 2	None None	65 60	161 154	25.1 25.3	3 3
35	Female	1	35	2	None	55	158	22	1
38 40		1	38 40	2 2	None None	62 67	158 160	24.8 26.2	1 3
30	Female	1	30	2	None	64	164	23.8	1
57 30		2 1	57 30	1 2	None None	53 58	155 165	22.1 21.3	1 1
34 68		1 2	34 68	2 1	None None	56 52	157 168	22.7 18.4	1 2
37	Female	1	37	2	None	62	155	25.8	3
38 42		1	38 42	2 2	None None	60 65	158 158	24 26	1 3
34		1	34	2	None	53	156	21.8	1
65 44		2	65 44	2 2	None None	51 63	155 166	21.2 22.9	1 1
33 44		1 1	33 41	2 2	None None	69 59	150 161	30.7 22.8	4 1
56	Female	2	56	2	None	64	158	25.6	3
37 52		1 2	37 52	2 2	None None	72 56	160 165	28.1 20.6	3 1
30	Female	1	30	2	None	51	159	20.2	1
44 55		1 2	44 55	2 2	None None	58 60	156 165	23.8 22	1 1
47 46		1	47 46	1 2	None None	50 57	164 153	18.6 24.3	1 3
51	Female	2	51	2	None	64	166	23.2	1
37 52		1 2	37 52	2 2	None None	59 53	153 165	25.2 19.5	3 1
62	Female	2	62	2	None	50	162	19.1	1
47 36		2 1	47 36	1 2	None None	49 68	162 162	18.7 25.6	1 3
56 52		2 2	56 52	2 2	None None	70 61	157 160	28.4 23.8	3 1
58	Female	2	58	2	None	66	165	24.2	1
45 33		1	45 33	2 2	None None	53 58	152 156	22.9 23.8	1
31		1 2	31	2 2	None	62	154	26.1	3
55 37		1	55 37	2	None None	79 62	179 164	24.7 23.1	1 1
65 42		2 1	65 42	2 2	None None	58 60	144 161	28 23.1	3 1
67	Female	2	67	2	None	60	168	21.3	1
41 30		1	41 30	2 2	None None	58 58	157 162	23.5 22.1	1 1
49 33		2	49 33	1 2	None None	67 65	163 165	25.2 23.9	3 1
57	Female	2	57	2	None	50	163	18.8	1
40 43	Female Female		40 43	1 2	None None	55 81	164 160	20.4 31.6	1 4
52	Female	2	52	2	None	58	165	21.3	1
55 46	Female Female			1 2	None None	65 60	162 154	24.8 25.3	1 3
43 45	Female Female		43 45	2 2	None None	66 68	161 162	25.5 25.9	3 3
36	Female	1	36	2	None	54	164	20.1	1
62 45	Female Female		62 45	2 2	None None	60 59	154 154	25.3 24.9	3 1
51 70	Female Female		51 70	2 2	None None	62 50	156 163	25.5 18.8	3 1
38	Female			2	None	41	150	18.2	2
47 38	Female Female		47 38	2 2	None None	55 65	154 155	23.2 27.1	1 3
32	Female	1	32	2	None	60	152	26	3
38 45	Female Female			2 2	None None	68 57	154 165	28.7 20.9	3 1
32	Female		32	2	None	55	155	22.9	1
57 41	Female Female	1	41	2 2	None None	58 68	152 158	25.1 27.2	3 3
48 52	Female Female			2 2	None None	50 68	159 166	19.8 24.7	1 1
60	Female	2	60	2	None	52	165	19.1	1
40 40	Female Female		40 40	2 2	None None	59 58	151 150	25.9 25.8	3 3
62	Female	2	62	2 2	None	59	155 158	24.6	1 3
61 51	Female Female	2	51	2	None None	65 48	149	26 21.6	1
32 33	Female Female	1	32 33	2 2	None None	58 53	156 153	23.8 22.6	1 1
39	Female	1	39	2	None	58	159	22.9	1
68 48	Female Female		68 48	2 2	None None	58 55	154 150	24.5 24.4	1 1
50	Female	2	50	2	None	40	150	17.8	2
42 44	Female Female		44	2 2	None None	47 61	152 156	20.3 25.1	1 3
41 54	Female Female		41 54	2 2	None None	59 67	157 168	23.9 23.1	1 1
36	Female	1	36	2	None	60	149	27	3
32 69	Female Female			2 2	None None	61 50	159 155	24.1 20.8	1 1

-Morbid conditions(None=1, Type-2 Diabetes=2, Hypertension=3, Hypothyroidism=4)	Method of Diagnosis	Pathological Diagnosis		Benign/Malignant		
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	2	Malignant Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant	pT2N3aMo	IIIC
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical, Microsopic, Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	1 2	Malignant Malignant		IIA IIB
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant	pT2N3aMo	IIIC
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	2	Malignant Malignant	pT3N1aMo pT1NoMx	IIIA I
	Clinical: Microsopic: Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant	pT1cN1aMo	IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	2	Malignant Malignant		II B II B
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIC
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant	pT2N2aMx	IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	2 2	Malignant Malignant		III A II A
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		II B
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant	pT1NoMx	1
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant	PT3NoMx	IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIB
	Clinical; Microsopic; Imaging techniques Clinical: Microsopic: Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	2 2	Malignant Malignant		IIIA IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		III A
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		II A
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant	pT2N3aMx	IIIC
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIA
	Clinical: Microsopic: Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	1	Malignant Malignant		II A
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IV
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant	pT2NoMo	IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIC
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	2	Malignant Malignant		II B
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIC
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		II B
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1 2	Malignant	pT1NoMx	IIA
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	1	Malignant Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant	pT2NoMo	IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIA
	Clinical: Microsopic: Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIA IIA
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	1 2	Malignant Malignant		IIA IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant	pT3N2Mo	IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	2 2	Malignant Malignant		IIIA
	Clinical; Microsopic; Imaging techniques  Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant	pT1cNoMo	I
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical: Microsopic: Imaging techniques	Infiltrating Ductal Carcinoma	2 2	Malignant		IIIC
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	2	Malignant Malignant		IIIA
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIA
	Clinical: Microsopic: Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant	pT1NoMx	1
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma Infiltrating Ductal Carcinoma	1 2	Malignant Malignant	pT1cNoMx cT4bN2Mo	IIIB
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		III A
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		IIB
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		IIIA
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant	pT2N1aMx	II B
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	2	Malignant		II A
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		II B
	Clinical: Microsopic: Imaging techniques	Infiltrating Ductal Carcinoma Ductal Carcinoma in situ	2	Malignant		III C
	Clinical; Microsopic; Imaging techniques Clinical; Microsopic; Imaging techniques	Ductal Carcinoma in situ Infiltrating Ductal Carcinoma	1	Malignant Malignant		NA II B
			-		IVIU	•
	Clinical; Microsopic; Imaging techniques	Infiltrating Ductal Carcinoma	1	Malignant		II A

Pathological Primary Tumor Size(cm)	Tumor size (upto 2cm=1, >2 and upto 5cm=2, >5cm=3)	Histologic Sub type(TumorGrade)	Tumor Grade (Well differentiated=1, Moderately differentiated=2, Poorly differentiated=3)
3.4×2.4×2.4 cm	2	Grade II	2
8.2×7.0×4.4 cm 3.0×2.5×2.3 cm	3 2	Grade III Grade II	3 2
5.9×4.6×3.0 cm	3	Grade II	2
3.5×3.0×0.5 cm	2	Grade II	2
6.5×3.5×3.0 cm	3	GradeII	2
4.0×3.2×3.0 cm	2	Grade II	2
4.0×3.0×3.0 cm 4.4×3.5×2.6 cm	2	Grade II Grade II	2 2
3.4×3.2×1.4 cm	2	Grade II	2
4.0×2.0×1.8 cm	2	Grade II	2
4.0×3.5×2.2 cm	2	Grade III	3
3.0×1.5×0.8 cm	2	Grade II	2
2.8×2.4×2.2cm	2	Grade III	3
4.8×1.8×2.0 cm 5.1×4.2×4.0 cm	3	Grade II Grade II	2 2
1.2×1.0×0.5 cm	1	Gradel	1
3.1×2.0 cm	2	Grade III	3
4.5×3.5×3.0 cm	2	Grade III	3
1.2×1.0×0.8 cm	1	Grade II	2
5.8×4.0×3.5cm 4.4×2.8×2.5 cm	3 2	Grade III	2 3
5.5×5.0×5.0cm	3	Grade II	2
3.8×3.1×2.2cm	2	Grade II	2
4.8×2.0×2.0cm	2	Grade II	2
3.5×3.0×2.5cm	2	Grade II	2
3.5×3.0×3.0 cm	2	Grade II	2
2.3×1.2×0.6 cm 2.2×1.8×1.2cm	2 2	Grade III Grade III	3
5.5×4.5×3.5 cm	3	Grade II	2
4.1×3.5×2.8 cm	2	Grade III	3
2.1×1.8×1.0 cm	2	Grade II	2
5.5×3.5×3.0cm	3	Grade II	2
1.5×1.0×1.0 cm 4.0×3.0×2.8 cm	1 2	Grade II	2 2
4.0×3.0×2.8 cm 5.1×5.0×1.8cm	3	Grade III	3
2.2×1.0×0.8 cm	2	Grade III	3
3.1×2.3×1.8cm	2	Grade II	2
5.6×3.0×2.0 cm	3	Grade II	2
6.5×3.5×4.8 cm	3	Grade III	3
2.0×1.5×1.1 cm	1	Grade III	3
4.0×3.8×2.6 cm 2.2×1.8×1.2cm	2 2	Grade III Grade II	3 2
2.8×2.7×1.8cm	2	Grade II	2
4.5×3.3×7.7 cm	2	Grade II	2
2.1×1.4×1.0cm	2	Grade II	2
4.0×3.5×3.0cm	2	Grade II	2
3.8×3.2×2.4 cm 4.5×4.0×1.5 cm	2 2	Grade II Grade II	2 2
5.3×2.0×5.3 cm	3	Grade II	2
6.0×5.0×4.0 cm	3	Grade III	3
2.2×1.2×1.2 cm	2	Grade II	2
2.9×2.0×1.2 cm	2	Grade III	3
4.0×3.6×2.8 cm	2	Grade III	3
3.4×2.3×2.0cm 4.2×2.2×1.2 cm	2 2	Grade II Grade II	2 2
2.8×2.5×1.8 cm	2	Grade II	2
3.8×2.0×1.5cm	2	Grade II	2
3.0×2.5×1.5cm	2	Grade II	2
4.8×4.0×3.0cm	2	Grade III	3
3.4×3.0×2.0 cm	2	Grade II	2
3.0×2.0×1.5 cm 1.7×1.0×0.8cm	2	Grade II	2 2
2.0×1.4×1.2 cm	1	Grade II Grade II	2
2.8×1.8×1.5cm	2	Grade III	3
5.5×3.0×2.8 cm	3	Grade II	2
2.8×1.8×1.6 cm	2	Grade II	2
3.5×3.2×3.5 cm 1.7×1.2×1.1 cm	2	Grade III Grade II	3 2
3.5×2.9	2	Grade III	3
2.2×1.8×1.3cm	2	Grade II	2
1.8×1.5×0.8 cm	1	Grade II	2
3.7×2.8×1.7 cm	2	Grade II	2
3.0×2.5×2.1cm	2	Grade II	2
2.7×2.3cm 3.4×2.9×1.5 cm	2 2	Grade III	2 3
4.5×3.5×2.5 cm	2	Grade II	2
1.9×1.8×1.6 cm	1	Grade II	2
4.0×2.5×2.0 cm	2	Grade II	2
2.5×1.5×1.3 cm 2.5×2.5×1.0 cm	2 2	Grade III	2
2.5×2.5×1.0 cm 1.9×1.6×1.4 cm	1	Grade III Grade II	3 2
2.6×2.4×2.2 cm	2	Grade II	2
1.9×1.8×1.6 cm	1	Grade II	2
1.5×1.2×0.8 cm	1	Grade II	2
6.4×5.2×2.6 cm 1.2×1.0×0.8 cm	3 1	Grade II	2 2
1.2×1.0×0.8 cm 3.8×3.2×3.2 cm	2	Grade III	3
1.5×1.8×0.2 cm	1	Grade III Grade I	1
4.0×4.0×3.0 cm	2	Grade II	2
2.4×2.0×1.8 cm	2	Grade II	2
5.1×5.3cm	3	Grade II	2
0.8×0.5×0.3 cm 6.0×4.0×3.0 cm	1 3	Grade II Grade II	2 2
5.3×3.0×2.2 cm	3	Grade III	3
2.5×2.5×2.0 cm	2	Grade II	2
1.8×1.6×1.3 cm	1	Grade II	1
1.3×1.2×1.0cm	1	Grade II	2
5.3×5.0×2.8 cm	3	Grade II	2
3.0×1.8×1.5 cm 2.9×2.3×1.8 cm	2 2	Grade III Grade II	3 2
2.9×2.3×1.8 cm 6.5×5.0×2.5 cm	3	Grade III	3
3.8×3.5×2.0 cm	2	Grade III	3
4.0×2.5×1.5 cm	2	Grade II	2
3.1×3.0×2.7 cm	2	Grade II	2
7.0×6.0×2.6 cm NA	3 NA	Grade III NA	3
NA 2.4×1.5×1.9 cm	NA 2	NA Grade II	2
3.0×2.0×2.5 cm	2	Grade III	3
2.0×1.0×0.2 cm	1	Grade II	2

Lymph node status (Presents-1, Absents-2, Unknown)   No. of lymph nodes involved /showing turns of Axillary lymph nodes removed   Modifical fichiardson Bloom Score   Total MRB Score   Residual Turns   Aspenditures   No. of lymph nodes involved /showing turns of Axillary lymph nodes removed   Modifical fichiardson Bloom Score   Total MRB Score   Residual Turns   Aspenditures   No. of land   No. of land	sitive=1, Negative=2)
2	
1	
5         6         2-3-2-7         7         Ro         1           2         0         17         2-2+2-6         6         Ro         1           1         4         18         2-2+2-2-6         6         Ro         1           2         0         19         2-3+3-7         7         Ro         1           1         2         25         2-2+2-6         6         Ro         1           1         1         12         3-2+2-7         7         Ro         1           1         1         12         2-2+2-6         6         Ro         1           1         1         1         12         3-2+2-6         6         Ro         1           1         1         1         12         3-2+2-6         6         Ro         1           2         0         0         10         3-3+2-8         8         Ro         1           2         0         0         10         3-3+2-7         7         Unknown         1           1         1         3         12         3-2+2-7         7         Unknown         1           1         <	
2	
1	
1         2         25         2+2+2=6         6         Ro         1           1         1         1         12         3+2+2=7         7         Ro         1           1         9         12         2+2+2=6         6         Ro         1           1         1         12         3+2+3=8         8         Ro         1           2         0         10         3+3+2=8         8         Ro         1           2         0         10         3+3+2=8         8         Ro         1           2         0         10         3+3+2=8         8         Ro         1           1         3         12         3+2+2=7         7         Unknown         1           1         3         12         3+2+2=7         7         Unknown         1           1         8         15         3+3+2=8         8         Ro         1           2         0         18         1+2+2=5         5         Ro         2           1         8         15         3+3+2=8         8         Ro         1           2         1         4         2+2+2=6	
1     1     12     3+2+2=6     6     Ro     1       1     9     12     2+2+2=6     6     Ro     1       1     1     12     3+2+3=8     8     Ro     1       1     20     23     2+2+2=6     6     Ro     1       2     0     10     3+3+2=8     8     Ro     1       1     3     12     3+2+2=7     7     Unknown     1       1     3     15     3+3+2=8     8     Ro     2       1     2     0     15     3+3+2=8     8     Ro     1       1     2     4     2+2+2=6     6     Ro     1       1     2     17     2+2+2=6     6     Ro     1       1     2     17     2+2+2=6     6     Ro     1       1     4     19     3+2+3=8     8     Ro     1       1     1     1     1     1 <td< td=""><td></td></td<>	
1       9       12       2+2+2=6       6       Ro       1         1       1       12       3+2+3=8       8       Ro       1         1       20       23       2+2+2=6       6       Ro       1         2       0       10       3+3+2=8       8       Ro       1         2       NA       NA       NA       2+2+2=6       6       Unknown       1         1       3       12       3+2+2=7       7       Unknown       1         2       0       18       1+2+2=5       5       Ro       2         1       8       15       3+3+2=8       8       Ro       1         2       0       10       3+3+2=8       8       Ro       1         1       2       4       2+2+2=6       6       Ro       1         1       2       4       1       2+2+2=6       6       Ro       1         1       2       1       1       3+2+2=8       8       Ro       1         1       1       1       1       3+2+2=8       8       Ro       1         1       1       1	
1 1 2 3+2+3=8 8 Ro 1 1 20 20 23 2+2+2=6 6 Ro 1 2 0 0 10 3+3+2=8 8 Ro 1 2 1 NA NA NA 2+2+2=6 6 Unknown 1 3 3 12 3+2+2=7 7 Unknown 1 2 1 3+2+2=7 7 Ro 1 1 8 15 3+3+2=8 8 Ro 1 1 12 3+2+2=7 7 Ro 1 1 8 8 15 3+3+2=8 8 Ro 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	
2 0 10 10 3+3+2=8 8 Ro 1 2 NA NA NA 2+2+2=6 6 Unknown 1 3 3 12 3+2+2=7 7 Unknown 1 2 0 18 1+2+2=5 5 Ro 2 1 10 3+3+2=8 8 RO 1 2 0 18 15 3+3+2=8 8 RO 1 2 0 0 10 3+3+2=7 7 RO 1 1 2 0 0 10 3+3+2=7 7 RO 1 1 2 0 10 3+3+2=7 7 RO 1 1 2 1 2 17 2+2+2=6 6 RO 1 1 1 2 1 19 3+2+3=8 8 RO 1 1 19 3+2+3=8 8 RO 1 1 1 1 1 15 22 17 2+2+2=6 6 RO 1 1 1 1 15 22 17 2+2+2=6 6 RO 1 1 1 1 15 22 17 2+2+2=6 6 RO 1 1 1 1 15 22 17 2+2+2=6 6 RO 1 1 1 1 15 22 17 2+2+2=6 6 RO 1 1 1 1 15 22 2+2+2=6 6 RO 1 1 1 1 1 15 22 2+2+2+3=7 7 RO 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	
2         NA         NA         2+2+2=6         6         Unknown         1           1         3         12         3+2+2=7         7         Unknown         1           2         0         18         1+2+2=5         5         Ro         2           1         8         15         3+3+2=8         8         Ro         1           2         0         0         10         3+3+2=7         7         Ro         1           1         2         4         2+2+2=6         6         Ro         1           1         2         17         2+2+2=6         6         Ro         1           1         4         19         3+2+3=8         8         Ro         1           2         1         17         2+2+2=6         6         Ro         1           1         4         19         3+2+3=8         8         Ro         1           1         NA         NA         2+2+2=6         6         Ro         1           1         1         1         2+2+3=7         7         Ro         1           1         2         2         2+2+3=7	
1	
1 8 15 3+3+2=8 8 Ro 1 2 0 10 10 3+3+2=7 7 Ro 1 1 2 4 4 2+2+2=6 6 Ro 1 1 2 17 2+2+2=6 6 Ro 1 1 4 19 3+2+3=8 8 Ro 1 1 1 4 19 3+2+3=8 8 Ro 1 1 1 1 1 15 2 2 17 Ro 1 1 1 15 2 2 17 Ro 1 1 1 15 2 2 2 2 2 2+2+3=7 7 Ro 1 1 1 1 15 2 2 2 2 2+2+3=7 7 Ro 1 1 1 1 2 1 1 2 2 7 7 2+2+2=6 6 Ro 1 1 1 1 1 1 3+3+3=9 9 Ro 1 1 1 1 1 1 1 3+3+3=9 9 Ro 1 1 1 1 1 1 1 3+3+3=9 9 Ro 1 1 1 1 1 1 3+3+3=8 8 Ro 1 1 1 1 1 3 3 15 3+2+2=7 7 Ro 1	
2 0 10 10 3+3+2=7 7 Ro 1 1 2 4 4 2+2+2=6 6 Ro 1 1 2 17 2+2+2=6 6 Ro 1 1 2 17 2+2+2=6 6 Ro 1 1 2 19 3+2+3=8 8 Ro 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	
1 2 4 2-2-2-6 6 Ro 1 1 2 17 2-2-2-6 6 Ro 1 1 2 17 2-2-2-6 6 Ro 1 1 4 19 3-2-3-8 8 Ro 1 2 NA NA NA 2-2-2-6 6 Ro 1 1 NA NA NA 2-2-2-6 6 Ro 1 1 15 22 2-2-2-6 7 Ro 1 1 1 2 2 7 2-2-2-6 6 Ro 1 1 1 3 2 2 7 2-2-2-6 6 Ro 1 1 1 4 2 3 2-2-2-3-7 7 Ro 1 1 1 3 4 9 2-2-2-3-7 7 Ro 1 1 1 3 3 15 3-3-2-2-7 7 Ro 1	
1 4 19 3+2+3=8 8 Ro 1 2 NA NA NA 2+2+2=6 6 Ro 1 1 NA NA 2+2+3=7 7 NA 1 1 15 22 2+2+3=7 7 Ro 1 1 2 7 2+2+2=6 6 Ro 1 1 1 2 15 22 2+2+3=7 7 Ro 1 1 2 1 4 23 2+2+2=6 6 Ro 1 1 4 9 2+2+3=7 7 Ro 1 1 1 3+3+3=9 9 Ro 1 1 4 9 2+3+3=8 8 Ro 1 1 3 3 15 3+2+2=7 7 Ro 1	
2 NA NA NA 2+2+2=6 6 Ro 1 1 NA NA NA 2+2+3=7 7 NA 1 1 15 22 2+2+3=7 7 Ro 1 1 2 2 7 2+2+3=7 7 Ro 1 1 4 2 23 2+2+3=7 7 Ro 1 1 1 1 11 3+3+3=9 9 Ro 1 1 4 9 2+3+3=8 8 RO 1 1 3 15 3+2+2=7 7 Ro 1	
1     NA     NA     2+2+3=7     7     NA     1       1     15     22     2+2+3=7     7     Ro     1       1     2     7     2+2+2=6     6     Ro     1       1     4     23     2+2+3=7     7     Ro     1       1     1     11     3+3+3=9     9     Ro     1       1     4     9     2+3+3=8     8     Ro     1       1     3     15     3+2+2=7     7     Ro     1	
1     2     7     2+2+2=6     6     Ro     1       1     4     23     2+2+3=7     7     Ro     1       1     1     11     3+3+3=9     9     Ro     1       1     4     9     2+3+3=8     8     Ro     1       1     3     15     3+2+2=7     7     Ro     1	
1 4 23 2+2+3=7 7 Ro 1 1 11 3+3+3=9 9 Ro 1 1 4 9 2+3+3=8 8 Ro 1 1 3 15 3+2+2=7 7 Ro 1	
1 1 11 3+3+3+9 9 Ro 1 1 4 9 2+3+3=8 8 RO 1 1 3 15 3+2+2=7 7 Ro 1	
1 3 15 3+2+2=7 7 Ro 1	
2 0 27 2+2+2=6 6 Ro 1	
2 0 10 2+2+2=6 6 Ro 1	
2 0 14 2+2+2=6 6 Ro 2	
2 0 18 2+2+2=6 6 Ro 1 2 0 18 3+3+3=9 9 Ro 1	
2 0 16 313137 9 NO 1 1 7 26 313139 9 Ro 1	
1 NA NA 2+2+2=6 6 Unknown 1	
1 4 15 2+2+2=6 6 Ro 1 2 0 19 3+3+3=9 9 Ro 1	
2 0 15 3-3-3-9 9 Ro 1	
1 5 26 3+3+3=9 9 Ro 1	
2 0 10 2+2+3=7 7 Ro 1 1 23 29 2+2+2=6 6 Ro 1	
1 23 29 2+2+2=6 6 Ro 1 2 0 9 2+2+2=6 6 Ro 1	
2 0 10 2+2+2=6 6 Ro 1	
2 0 33 2+2+3=7 7 Ro 1	
2 0 13 2+2+2=6 6 Ro 2 1 4 22 2+3+1=6 6 Ro 1	
1 NA NA 2-2-2-2-6 6 Unknown 1	
2 0 27 3+2+3=8 8 Ro 1	
2 0 11 3+2+2=7 7 Ro 1 1 31 31 2+3+3=8 8 Ro 1	
1 1 15 243438 8 R0 1	
2 NA NA 2+3+2=7 7 Unknown 1	
1 5 21 342-2-7 7 Ro 2	
1 1 15 2+2+2=6 6 Ro 1 1 4 35 2+2+3=7 7 Ro 1	
1 16 28 3+2+2=7 7 Ro 1	
2 0 15 3+3+2=8 8 Ro 1	
1 9 16 2+2+3=7 7 Ro 1 1 NA NA 3+2+2=7 7 NA 2	
2 0 10 2+2+2=6 6 Ro 2	
1 NA NA 2+3+2=7 7 NA 1	
2 0 21 3+3+3=9 9 Ro 1 1 7 7 3+3+1=7 7 Ro 1	
1 1 13 224226 6 RO 1	
1 7 23 3+2+3=8 8 Ro 1	
2 0 25 2+2+2=6 6 Ro 1 1 NA NA 2+3+3=8 8 Unknown 1	
1 NA NA 27272-6 6 UINIOWII 1 2 NA NA 22422-6 6 UINIOWII 1	
1 1 23 3+2+2=7 7 Ro 1	
2 0 2 3+2+2=7 7 Ro 1 2 0 13 3+2+2=7 7 Ro 2	
2 0 13 3*2*2=7 / KU 2 1 NA NA 32*2=7 7 Unknown 1	
2 0 6 3+3+3=9 9 Ro 1	
1 4 15 2+2+2=6 6 Ro 1 2 0 10 2+2+2=6 6 Ro 1	
1 7 18 2+2+2=6 6 Ro 1	
1 3 11 2+2+2=6 6 Ro 1	
1 10 14 3+3+2=8 8 Ro 1 2 NA NA 2+2+2=6 6 NA 2	
2 NA NA 2+2+2=6 6 NA 2 1 12 13 3+2+2=7 7 RO 1	
2 NA NA 2+2+2=6 6 NA 2	
2 0 19 2+2+2=6 6 Ro 1	
1 5 NA 2+2+2=6 6 NA 1 1 4 21 3+2+2=7 7 RO 1	
1 6 9 3+2+2=8 8 Ro 1	
2 NA NA 1+2+2=5 5 NA 2	
1 4 NA 2+2+2=6 6 NA 1 1 2 12 2+2+2=6 6 RO 1	
1 NA NA 2+2+2=6 6 Unknown 1	
2 0 4 2+3+3=7 7 Ro 1 1 7 7 2+2+2=6 6 Ro 1	
1 7 7 7 2+2+2=6 6 Ro 1 1 NA NA 3+3+3=9 9 NA 1	
2 0 10 3+2+2=7 7 Ro 1	
2 0 10 3+2+2=7 7 Ro 1	
2 0 8 2+2+2=6 6 Ro 1 1 NA NA 2+2+2=6 6 NA 1	
1 6 15 3+3+3=9 9 Ro 1	
1 NA NA 2+2+2=6 6 NA 1	
1 3 13 3+3+2=8 8 Ro 1 1 1 23 3+3+3=9 9 Ro 1	
2 0 10 2+2+2=6 6 Ro 1	
1 NA NA 3+2+2=7 7 NA 1	
1 16 23 2+3+3=8 8 Ro 1 2 NA NA NA NA RO 2	
1 NA NA 2+2+2=6 6 NA 2	
2 NA NA 3+2+2=8 8 NA 1	
2 2+2+2=6 6 NA 1	

Nipple/Skin Involment(Positive=1, Negative=2) 2	ER Status(Positive=1, Negative=2) F 2		Her-2 Status(Positive=1, Negative=2) 2	ki-67 expression(%) 70	Molecular Phenotypes Triple Negative	Vitamin D level(nmol/L) 25.1
	2 1		2	60	Luminal B	14.4
	1 1		2	40 60	Her2- Luminal B	69.9
	1 1		1 2	NA	Her2+ Luminal B Her2- Luminal B	42.8 76.1
2	2 2	2	2	80	Triple Negative	48.2
	2 2		2 2	30 55	Triple Negative Triple Negative	39.3 39.6
2	1 :	Į.	1	60	Her2+ Luminal B	35.6
	2 2		1	50 80	Her2 High Her2 High	50.3 51.5
	1 2		2	60	Her2- Luminal B	63.6
	1 :		2	50	Her2- Luminal B	10.7
<del>-</del>	1 1		2 1	30 60	Her2- Luminal B Luminal B	33.6 29.2
2	1 :	l	1	70	Her2+ Luminal B	50.4
	2 2		2 2	40 80	Triple Negative Triple Negative	66.38 31.8
	2		2	80	Triple Negative	28.7
	1 1		2	60	Her2- Luminal B	49.32
	2 2		2 2	60 60	Triple Negative Her2- Luminal B	23.21 56.9
	1 1		2	70	Her2- Luminal B	48.2
	2 1		1	35 80	Her2 High Her2+ Luminal B	29.6 53.12
	2		1	70	Her2 High	41.2
	1 1		2 2	40	Her2- Luminal B	24.8
	2 2		2	70 65	Triple Negative Triple Negative	18.8 21.3
2	2		1	40	Her2 High	53.2
	2 1		2 2	70 40	Her2- Luminal B Her2- Luminal B	32.6 42.2
2	1 1		2	30	Her2- Luminal B	22.6
	1		2	60	Her2- Luminal B	36.1
	1 1		2	60 50	Her2- Luminal B Her2 High	38.5 39.1
2	1 :	<u>l</u>	2	40	Her2- Luminal B	58.4
	2 2		1 2	60 10	Her2 High Luminal A	39.1 55
2	2	2	2	50	Triple Negative	21.32
	2 2		2	80	Triple Negative	23.2
	2 1		2 2	NA	Her2- Luminal B Triple Negative	38.4 24.3
	1 1		2	80	Her2- Luminal B	64.09
	1 1		1 2	70 25	Her2+ Luminal B Her2- Luminal B	57.34 36.42
	1		2	30	Her2+ Luminal B	48.34
	1		2	70	Her2- Luminal B	53.3
	1 1		1 2	NA	Her2+ Luminal B Her2- Luminal B	7.5 12.8
1	2		2	70	Triple Negative	17.6
	1 1		2 2	70 60	Her2- Luminal B Her2- Luminal B	22.2 39.71
	1		1	60	Her2+ Luminal B	26.1
	2		1	60	Her2 High	34.67
	2 2		1 2	20 70	Her2 High Triple Negative	49 32.3
2	1 :	l	1	80	Her2+ Luminal B	54.12
	1 1		2 1	40 60	Her2 High Her2+ Luminal B	44.31 51.23
	1		2	80	Her2- Luminal B	45.18
	1		2	50	Her2- Luminal B	50.4
<del>-</del>	1 1		2 2	10 50	Luminal A Her2- Luminal B	74.1 51.2
2	2		2	67	Triple Negative	23.6
	1 1		2 2	60 20	Her2- Luminal B Her2- Luminal B	39.8 59.9
1	1 :	l	2	15	Her2- Luminal B	133.4
	1 1		1 2	NA 10	Her2- Luminal B Her2- Luminal B	31.1 43.1
	2		1	20	Her2 High	40.23
	1 1		2	70	Her2- Luminal B	25.5
	1 1		2 1	40 NA	Her2- Luminal B Her2 High	93.45 38.12
2	2	2	2	90	Triple Negative	21.1
	2 2		2 2	80 25	Triple Negative Her2- Luminal B	27.43 44.36
2	1 :	l	2	40	Her2- Luminal B	29.34
	1 1		2	30 87	Her2- Luminal B Her2- Luminal B	55.1
	1 2		2 2	87 50	Her2- Luminal B	23.12 51.2
2	1 :	<u>l</u>	2	10	Luminal A	56.44
	1 2		1 2	70 10	Her2+ Luminal B Luminal A	17 59.31
2	1 2	2	2	80	Her2- Luminal B	70.1
	2 2		1 2	40 50	Her2 High Her2- Luminal B	30.7 42.7
	2		2	80	Triple Negative	89.5
	1		2	NA	Her2- Luminal B	65.3
	1 1		2 1	NA 60	Her2- Luminal B Her2 High	54.9 28.3
2	1 :	Į.	2	NA	Her2- Luminal B	48.37
	1 1		2 2	40 30	Her2- Luminal B Her2- Luminal B	40.5 31.46
1	1 :	Į.	2	30	Her2- Luminal B	48.78
	1 1		2	70	Her2- Luminal B	40.3
	2 2		1 2	50 60	Her2 High Her2- Luminal B	23.89 46.32
1	2 2	2	2	NA	Triple Negative	37.7
	1 1		2 1	60 50	Her2- Luminal B Her2 High	62.2 83.69
2	2 2	2	1	50	Her2 High	22.04
	2		1	70 80	Her2- Luminal B	52.4 65.27
	2 2		2 2	80 60	Triple Negative Her2- Luminal B	65.27 81.37
2	1 :	<u>l</u>	2	80	Her2- Luminal B	26.45
	1 1		2 1	8 30	Luminal A Her2+ Luminal B	48.54 44.7
2	1 :	<u>l</u>	2	50	Her2- Luminal B	39.46
2	1 1	l	1	75	Her2+ Luminal B	82.43

Vitamin D level nmol/L (<50 as Deficient=1, 50 to <75 as Insufficient=2, 75-250 as Normal=3, >250 as Intoxication=4)		DNA A280		Genotypes Fokl FF=1, Ff=2, ff=3	Genotypes Apal AA=1, Aa=2, aa=3	Genotypes Bsml BB=1, Bb=2
1 1	0.079 0.141	0.044	1.79 1.76	3	1	2 2
2	0.091	0.05	1.82	1	1	2
1 3	0.062 0.109	0.034 0.064	1.82 1.7	3 2	2	2 1
1	0.109	0.064	1.7	3	1 1	2 2
1	0.14 0.151	0.079 0.086	1.77 1.75	3	1	1
1 2	0.121 0.112	0.071 0.065	1.7 1.72		1 2	2 1
2	0.102	0.06	1.7		1	2
2 1	0.132 0.132	0.073 0.075	1.8 1.76		1 2	2 2
1	0.177	0.098	1.8	1	2	1
1 2	0.126 0.143	0.074 0.089	1.7 1.76		1	1 2
2 1	0.127 0.119	0.07 0.069	1.81 1.72	3 2	2 3	2 1
1	0.11	0.064	1.71	1	3	2
1	0.186 0.134	0.101 0.074	1.84 1.81	1	1 2	2
2	0.111	0.064	1.73	3	1	1
1	0.161 0.131	0.089 0.072	1.8 1.81	3	2	2
2	0.164 0.176	0.091 0.097	1.8 1.81		2	1 2
1	0.108	0.06	1.8	1	2	2
1	0.139 0.175	0.077 0.097	1.8 1.8		3	1 2
2	0.169	0.094	1.79	3	1	1
1	0.125 0.103	0.069 0.057	1.81 1.8	2	3 2	1 2
1 1	0.174 0.102	0.096 0.057	1.81 1.78		1 1	1 2
1	0.144	0.08	1.8		2	2
1 2	0.175 0.138	0.097 0.076	1.8 1.81		1 3	2
1	0.179	0.099	1.8	3	2	1
2 1	0.159 0.121	0.087 0.066	1.82 1.83	2	1 3	2
1	0.15	0.083	1.8	3	2	2
1	0.141 0.143	0.078 0.08	1.8 1.79	3	3	2 2
2 2	0.119 0.133	0.066 0.073	1.8 1.82	2	2	1 2
1	0.124	0.068	1.82	3	1	2
1 2	0.121 0.155	0.067 0.086	1.8 1.8	1 2	1	1
1	0.128	0.071	1.8	1	1	1
1	0.115 0.117	0.067 0.068	1.71 1.72		2	1 2
1	0.126	0.072	1.75	1	2	2
1	0.101 0.154	0.055 0.09	1.83 1.71	2	3	2 1
1	0.181 0.152	0.101 0.089	1.79 1.7	1	2 2	1 2
1	0.154	0.085	1.81	1	1	1
2 1	0.168 0.165	0.093 0.091	1.8 1.81		2 2	2
2	0.031	0.018	1.72		2	2
1 2	0.109 0.127	0.061 0.07	1.78 1.81		1	1 2
2 2	0.182 0.131	0.101 0.072	1.8 1.81	3 2	2 2	1 2
1	0.173	0.096	1.8	2	1	1
1 2	0.146 0.103	0.081 0.057	1.8 1.8	1 2	2 2	2 1
3	0.141	0.078	1.8	1	2	1
1	0.122 0.127	0.067 0.073	1.82 1.73	1	1 2	2 2
1	0.126 0.12	0.07 0.069	1.8 1.73	2	2 2	1
3	0.131	0.072	1.81	2	2	2
1 1	0.129 0.111	0.071 0.062	1.81 1.79		1 2	1
1	0.145	0.081	1.79	2	1	1
1 1	0.147 0.101	0.081 0.058	1.81 1.74	1	2 1	1 2
2 1	0.124 0.109	0.069 0.06	1.79 1.81		2	1 2
2	0.117	0.065	1.8	1	2	2
2 1	0.132 0.145	0.073 0.082	1.8 1.79		1 2	3 2
2	0.149	0.082	1.81	1	1	3
2 1	0.148 0.147	0.083 0.081	1.78 1.81		1 2	1 2
1 3	0.121 0.124	0.068 0.069	1.77 1.79		1 3	1 3
2	0.102	0.056	1.8	2	2	1
2 1	0.128 0.123	0.071 0.068	1.8 1.8		2	1
1	0.191	0.106	1.8	2	2	2
1	0.021 0.098	0.012 0.054	1.75 1.81	1	1	2
1 1	0.122 0.119	0.067 0.066	1.82 1.8		2 1	2 1
1	0.124	0.068	1.82	2	1	2
1	0.181 0.132	0.1 0.073	1.8 1.8		2	2
2	0.126	0.07	1.8	1	1	2
3 1	0.101 0.085	0.056 0.047	1.8 1.8	2	1 3	3 1
2 2	0.13 0.125	0.072 0.069	1.8 1.81		2 2	2 1
3	0.133	0.074	1.79	2	1	1
1 1	0.135 0.099	0.075 0.055	1.8 1.8		2	2
1	0.146	0.081	1.8	1	2	2
1 3	0.103 0.125	0.057 0.069	1.8 1.81			2 1

S.No.	Name Anupa	UHID 2981750	Name of Participating Centre Swami Rama Himalayan University, Dehradun, Uttarakhand	Name of the Department Dermatology	Date of first attendance to hospital for Routine Checkup/ Sample collection 6/3/2020	Address Athoorwala, Dehradun
2	Payal Chauhan	2840224	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-I	6/3/2020	Jolly grant, Dehradun
3	Savita Pande	2407251	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-I	7/22/2020	Pauri, Garhwal, Uttarakhand
4	Sunita	3001398	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-I	7/23/2020	Bhaniyawala, Dehradun
5 6	Anita Dabral Rani	3007900 3008725	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-I Ortho Unit-I	8/10/2020 8/13/2020	Rishikesh, Uttarakhand Bijnor, U.P.
7	Bahdi Devi	3009920	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	8/17/2020	Village Sunargaon, Uttarakhand
8	Tabasum	3371134	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	9/10/2020	Haridwar, Uttarakhand
9 10	Mamta Aswal Prakash Devi	2693764 3024273	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS HIMS	9/17/2020 9/21/2020	4/1, Teg Bhadur Road, Dehradun Bapu gram, Rishikesh, Uttarakhand
11	Raj Kumari	3030282	Swami Rama Himalayan University, Dehradun, Uttarakhand	Ortho Unit-II	10/6/2020	Bijnor, U.P.
12	Vibha Ghildiyal	3032080	Swami Rama Himalayan University, Dehradun, Uttarakhand	Dermatology	10/10/2020	Staff, Jolly grant, Dehradun
13 14	Neha Sharma Sushila	3044254 3047617	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	11/13/2020 11/23/2020	Bijnor, U.P.
15	Haira	3059574	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Dermatology Ortho Unit-II	12/22/2020	Peeth Bazar, Haridwar, Uttarakhand Biinor, U.P.
16	Mohini Sethi	2318370	Swami Rama Himalayan University, Dehradun, Uttarakhand	Ortho Unit-I	1/1/2021	Staff, Jolly grant, Dehradun
17	Vandana Kukreti	3065767	Swami Rama Himalayan University, Dehradun, Uttarakhand	Ortho Unit-II	1/6/2021	Staff, Jolly grant, Dehradun
18 19	Kamlesh Prem Lata	3062434 3066932	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-II Medicine Unit-III	1/7/2021 1/9/2021	Village Satpura, U.P. Sai Dham, Phase 4, Haridwar, Uttarakand
20	Amita	3067923	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	1/11/2021	228, Salawala, Dehradun
21	Sunita	3068184	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	1/12/2021	Staff, Jolly grant, Dehradun
22 23	Manju Rastogi Santosh Devi	3071701 3079723	Swami Rama Himalayan University, Dehradun, Uttarakhand swami Rama Himalayan University, Dehradun, Uttarakhand	General Medicine Unit-III Medicine Unit-I	1/21/2021 2/8/2021	Kila Afjalgarh, Bijnor, U.P. Amroha, U.P.
24	Alma Panwar	3080918	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-II	2/11/2021	Village Chidderwala, Uttarakhand
25	Meenu Pradhan	3082546	Swami Rama Himalayan University, Dehradun, Uttarakhand	General Medicine Unit-III	2/13/2021	Solani puram, Roorkee, Uttarakhand
26 27	Kamla Sharma Adesh Devi	3083879 3084158	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III Medicine Unit-I	2/17/2021 2/18/2021	Bhaniyawala, Dehradun Surjannagar, Moradabad, U.P.
28	Manju Panday	3084311	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	2/18/2021	Awas vikas Modal, Haridwar, Uttarakhand
29	Sushma Devi	3077931	Swami Rama Himalayan University, Dehradun, Uttarakhand	Ortho Unit-II	2/26/2021	Rani Pokhri, Dehradun
30	Jahana Contra Tanana	3096776	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III General Medicine Unit-III	3/13/2021	Village Mebla, U.P.
31 32	Gunjan Tomar Raija Devi	2915835 3110128	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-II	3/24/2021 4/5/2021	Jolly grant, Dehradun Dehradun
33	Meenu Majumdar	3313819	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	4/10/2021	Bijnor, U.P.
34	Priyanka Gupta	2819744	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	5/24/2021	Dehradun
35 36	Munni Devi Neeru goyal	3130745 3137308	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Dermatology Medicine Unit-I	6/1/2021 6/17/2021	Jolly grant, Dehradun Haridwar, Uttarakhand
37	Reena	3141515	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	6/26/2021	Village Sherpur, Uttarakhand
38	Vinita Devi	3142924	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-I	6/30/2021	Bhaniyawala, Dehradun
39 40	Anita Rani Savita Devi	3071908 3154522	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III HIMS	7/17/2021 7/23/2021	Haridwar, Uttarakhand Dehradun
41	Sarojni Kukreti	3154924	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	7/24/2021	Pauri, Garhwal, Uttarakhand
42	Rajni	3160572	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	8/4/2021	Saharanpur, U.P.
43 44	Kavita Bisht	3160977	Swami Rama Himalayan University, Dehradun, Uttarakhand	E.N.T	8/4/2021	Tehri Garhwal, Uttarakhand
44	Sarla Kiran Rawat	3161642 3156226	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS Dermatology	8/6/2021 8/7/2021	Saharanpur, U.P. Doiwala, Dehradun
46	Amrit Kaur	3164327	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	8/11/2021	Kheri Doiwala, Dehradun
47	Usha Devi	3173356	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	8/27/2021	Moradabad, U.P.
48 49	Namita Nautiyal Babita Rani	3084192 3187724	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	E.N.T Ortho Unit-III	8/31/2021 9/22/2021	Shaheed Dwar, Athoorwala, Dehradun Haridwar, Uttarakhand
50	Baljeet Kaur	3198598	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	10/12/2021	Doiwala, Dehradun
51	Urmila Devi	3200366	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	10/16/2021	Sivalik nagar, Haridwar, Uttarakhand
52 53	Chanchal Rani Anisa	3200509 3204201	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Ortho Unit-III Medicine Unit-III	10/16/2021 10/23/2021	83-Shivalik nagar, Haridwar, Uttarakhand Village Sambhal, U.P.
54	Munni Devi	3214524	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	11/13/2021	Tehri Garhwal, Uttarakhand
55	Mirdula Rani	3215148	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-I	11/15/2021	Bijnor, U.P.
56 57	Shanti Devi Geetanjali	3223017 3228007	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-I Medicine Unit-I	12/2/2021 12/13/2021	Dehradun Chamoli, Uttarakahand
58	Anita Kapurwan	3097471	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Ortho Unit- II	1/4/2022	Rani Pokhri, Dehradun
59	Sundra Devi	3244960	Swami Rama Himalayan University, Dehradun, Uttarakhand	Ortho Unit- III	1/19/2022	Bhaniyawala, Dehradun
60	Renu gaur	3258468	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	2/25/2022	H.No. 14-Jeewani mai wali gali, Rishikesh, Utt
61 62	Sunita Devi Charanjeet Kaur	3256103 3262763	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Ortho Unit-II Medicine Unit-III	3/4/2022 3/5/2022	62-Shiv nagar, Dehradun U S nagar, Uttarakhand
63	Savita	3265129	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	3/9/2022	Bijnor, U.P.
64 65	Deepa Devi Manju	3271294 3274617	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS Medicine Unit-II	3/21/2022 3/25/2022	Rishikesh, Uttarakhand 322, Gurukul Kangari, Haridwar, Uttarakhand
66	Lalita Naval	2639625	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	3/28/2022	Dehradun
67	Raksha Chauhan	3283227	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-II	4/8/2022	Bijnor, U.P.
68	Sunita Rani	3284753	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	4/11/2022	Bijnor, U.P.
69 70	Jonny Devi Rima Gupta	3306496 3320277	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III Medicine Unit-III	5/18/2022 6/8/2022	Saharanpur, U.P. Moradabad, U.P.
71	Abha Gupta	3318958	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	6/9/2022	Dehradun
72	Anusuiya Bartwal	3323904	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	6/15/2022	Athoorwala, Dehradun
73 74	Savita Singh Urmila Devi	3327137 3352464	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS Medicine Unit-I	6/20/2022 8/1/2022	Haridwar, Uttarakhand Vinayal hills, Shastradhara road, Dehradun
75	Seema Tyagi	3358196	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	8/13/2022	Bijnor, U.P.
76	Rambatauri	3358731	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	8/17/2022	Dehradun
77 78	Nirdesh Sushma Devi	3362728 3364622	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III Medicine Unit-III	8/24/2022 8/27/2022	Bijnor, U.P. Saharanpur, U.P.
79	Vinita Kandwal	3367562	Swami Rama Himalayan University, Dehradun, Uttarakhand	E.N.T	9/2/2022	Amit gram gumaniwala, Dehradun
80	Shabana Khan	3368228	Swami Rama Himalayan University, Dehradun, Uttarakhand	Ortho Unit-I	9/5/2022	Haridwar, Uttarakhand
81 82	Asha Kunj Sunita Sharma	2242429 2665780	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-I Medicine Unit-I	9/17/2022 9/26/2022	Staff, Jolly grant, Dehradun Staff, Jolly grant, Dehradun
83	Rajeshwari Devi	3378874	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	9/29/2022	Pauri, Garhwal, Uttarakhand
84	Yukta Rajput	3378874	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	10/10/2022	Bijnor, U.P.
85 86	Saraswati Anuradha	3255718 2723712	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS HIMS	10/13/2022 10/15/2022	U S nagar, Uttarakhand Bijnor, U.P.
87	Nazneen Bano	3388635	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	10/22/2022	Roorkee, Uttarakhand
88	Alka Devi	3158771	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	10/28/2022	Saharanpur, U.P.
89 90	Gauri Manju Kaushal	3392386 3096906	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS Medicine Unit-II	11/1/2022 11/11/2022	WNo. 13, Ranipokhri, Dehradun Vikas nagar, Dehradun
91	Shivani Sharma	3396723	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Ortho Unit-II	11/11/2022	Haridwar, Uttarakhand
92	Sutlesh	3396862	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	11/14/2022	WNo. 5, Ranjhawala, Dehradun
93 94	Pooja Sunita Devi	3398920 3400332	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS Medicine Unit-III	11/16/2022 11/19/2022	Haridwar, Uttarakhand Tehri Garhwal, Uttarakhand
94 95	Rupa Devi	3397731	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	11/19/2022	Haridwar, Uttarakhand
96	Kamlesh	3410905	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	11/23/2022	U S nagar, Uttarakhand
97 98	Usha Devi	3404810	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	11/30/2022 12/1/2022	Athoorwala, Dehradun
98 99	Ritika agarwal Ganeshi Devi	2763389 3407259	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-I HIMS	12/1/2022 12/6/2022	Staff, Jolly grant, Dehradun Pauri, Garhwal, Uttarakhand
100	Geeta rana	3257903	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	12/8/2022	Staff, Jolly grant, Dehradun
101	Anjali arora	3414951	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-I	12/26/2022	U S nagar, Uttarakhand
102 103	Manju Kothari Anita Bisht	3415459 2916705	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-II Medicine Unit-III	12/27/2022 1/7/2023	Doiwala, Dehradun Bhaniyawala, Dehradun
104	Sulochana	3420749	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-I	1/9/2023	Tehri Garhwal, Uttarakhand
105	Sheetal	3421600	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	1/11/2023	U S nagar, Uttarakhand
106 107	Snehlata sharma Ashiya	3307451 3430666	Swami Rama Himalayan University, Dehradun, Uttarakhand Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	1/27/2023 2/4/2023	Staff, Jolly grant, Dehradun Haridwar, Uttarakhand
108	Monika Biswas	3431731	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS	2/7/2023	Staff, Jolly grant, Dehradun
109	Surama Devi	3434736	Swami Rama Himalayan University, Dehradun, Uttarakhand	HIMS Medicine Unit III	2/14/2023 8/24/2023	Tehri Garhwal, Uttarakhand
110	Parvati Devi	3362964	Swami Rama Himalayan University, Dehradun, Uttarakhand	Medicine Unit-III	الما المالية	Athoorwala, Dehradun

2 14/11 2/2 111 2							214/1 / 21	
Rural=1/Urban=2/Semi-Urban=3 2	Mobile no. 9411142880	Age 38	Sex Female	Menopausal status ( Pre=1, post=2) 1	Weight (kg) 56	Height (cm) 159	BMI(kg/m²) 22.2	BMI category (Normal=1, Underweight=2, Overweight=3, Obese class I=4) 1
2	7017960496	30	Female	1	51	160	19.9	1
3 2	9557877664 7414598764	47 55	Female Female	1 2	53 50	161 154	20.4 21.1	1
2	8126202021	48	Female	2	54	154	22.8	1
2	6397129708	32	Female	1	49	161	18.9	1
1 2	7302717417 7906942233	50 30	Female Female	2	53 49	167 162	19 18.7	1
2	9410314098	53	Female	2	58	152	25.1	3
2	958249226	65	Female	2	52	159	20.6	1
2 2	9368263627 9411156293	40 57	Female Female	1 2	51 58	161 165	19.7 21.3	1
2	8941014997	36	Female	1	60	163	22.6	1
2	9897750170	44	Female	1	51	165	18.7	1
2 2	8057306401 9438567623	46 66	Female Female	1 2	61 64	161 158	23.2 25.6	1 3
2	7351116177	41	Female	1	48	149	21.6	1
1 2	7253050830	41	Female	1	48	166	17.4	2
2	9958383086 9456590746	55 36	Female Female	2	39 55	150 160	17.3 21.5	2
2	8218539943	49	Female	2	57	165	20.9	1
2 2	9410600001 8532870716	57 51	Female Female	2 2	62 51	159 155	24.5 21.2	1
1	8859449223	60	Female	2	49	162	18.7	1
2 2	7895298039	59 46	Female	2	65 51	159 161	25.7 19.7	3
2	9719298959 7017556475	45	Female Female	1	57	165	20.9	1
2	9897425471	42	Female	1	60	154	25.3	3
2	9412985064 7417248559	51 35	Female Female	2	50 59	162 165	19.1 21.7	1
2	9412158901	40	Female	1	56	162	21.3	1
2	7248121568	45	Female	1	52	161	20.1	1
2 2	8449955170 6261883112	55 40	Female Female	2	48 53	162 160	18.3 20.7	2
2	9534493059	62	Female	2	45	158	18	2
2	9837326188 8865894326	42 32	Female Female	1	59 54	153 164	25.2 20.1	3 1
2	7415639874	36	Female	1	47	159	18.6	1
2	9760325992	50	Female	2	54	165	19.8	1
2	9411771547 9410744733	46 51	Female Female	1 2	57 49	163 152	21.5 21.2	1
2	9719678085	33	Female	1	47	159	18.6	1
3	8968876612	30	Female	1	54	162	20.6	1
2 2	8859018204 7895483129	57 37	Female Female	2	54 51	163 160	20.3 19.9	1
2	9837154452	32	Female	1	57	163	21.5	1
2 2	9649479679 8449891263	55 35	Female Female	2	57 49	163 158	21.5 19.6	1 1
2	830754546	45	Female	1	52	160	20.3	1
2	7983488117	34	Female	1	55	163	20.7	1
2 2	7705042267 9911046969	40 60	Female Female	1 2	48 48	159 162	19 18.3	1 2
1	6398692576	44	Female	1	61	165	22.4	1
3	9647862391	40	Female	1	56	159	22.2	1
2 2	9410263125 8057688048	45 35	Female Female	1	62 56	164 166	23.1 20.3	1
3	8979566361	43	Female	1	55	159	21.8	1
2 2	9756612755 9084083818	40 65	Female Female	1 2	52 61	163 155	19.6 25.4	1 3
2	9867563522	49	Female	2	51	167	18.3	2
2	877434947	46	Female	1	57	164	21.2	1
2 2	8006000254 9760905160	43 38	Female Female	1	58 58	159 164	22.9 21.6	1
2	8861899450	38	Female	1	49	161	18.9	1
2 2	7456961833	45	Female	1	58	158	23.2	1
2	9410170728 8057923429	42 63	Female Female	1 2	61 54	164 161	22.7 20.8	1
2	9568464890	41	Female	1	56	163	21.1	1
2 2	9761422683 9058298310	32 49	Female Female	1 2	57 56	162 149	21.7 25.6	1 3
2	9897256189	50	Female	2	51	157	20.7	1
2	7078287118	32	Female	1	47	155	19.6	1
2 2	9045775183 7070834349	46 58	Female Female	1 2	42 62	161 164	16.2 23.1	2 1
2	9675063599	38	Female	1	56	163	21.1	1
2 2	9528178050 9759012549	39 41	Female Female	1	48 58	160 159	18.7 22.9	1
2	9837013422	48	Female	2	57	165	20.9	1
2 2	9997952238 9997211116	34 33	Female	1	49 54	160 159	19.1 21.4	1
2	8650820711	50	Female Female	1	56	164	20.8	1
2	8650384890	36	Female	1	58	159	22.9	1
3 2	9675656504 9634810273	56 31	Female Female	2	49 56	160 164	19.1 20.8	1
2	7300740113	30	Female	1	58	160	22.7	1
2 2	9927560126 9319540500	33 48	Female	1 2	60 57	164 161	22.3 22	1
2	9756692464	48 51	Female Female	2	63	162	24	1
2	6397698163	49	Female	2	58	159	22.9	1
2 2	8273719044 9690310698	51 30	Female Female	2	63 55	164 162	23.4 21	1
2	9837520465	62	Female	2	51	163	19.2	1
2	9927931446	35	Female	1	48	162	18.3	2
3 2	9557934352 9149105759	30 33	Female Female	1	54 57	164 161	20.1 22	1
2	8534849970	46	Female	1	59	165	21.7	1
2 2	8527375553 9411714685	66 37	Female Female	2	61 45	162 159	23.2 17.8	1 2
3	8126138832	56	Female	2	45 59	162	22.5	1
2	9411642829	36	Female	1	48	158	19.2	1
2 2	7302041166 7895242122	45 43	Female Female	1	53 56	165 164	19.5 20.8	1
2	9418751796	52	Female	2	49	161	18.9	1
3 2	9557681443 7017195433	50 52	Female Female	2 2	59 54	164 161	21.9 20.8	1
2	7017195433 8449741172	46	Female	1	59	164	20.8	1
2	9012647754	54	Female	2	48	161	18.5	1
2 2	8979151623 8979423039	49 53	Female Female	2 2	56 61	160 164	21.9 22.7	1
2	9675318190	54	Female	1	61	164	22.7	1

History of any cancer	Co-Morbid conditions	Vitamin D level(nmol/L)	Vitamin D level nmol/L (<50 as Deficient=1, 50 to <75 as Insufficient=2, 75-250 as Normal=3, >250 as Intoxication=4)	DNA A260	DNA A280	Ratio A260/A280
None	None	114.7	3	0.162	0.09	1.8
None None	None None	68 107.7	2 3	0.171 0.191	0.096 0.11	1.78 1.73
None	None	99	3	0.108	0.06	1.8
None None	None None	156 51	3 2	0.147 0.139	0.081 0.076	1.81 1.82
None	None	79.2	3	0.142	0.081	1.75
None	None	89.32	3	0.122	0.069	1.76
None None	None None	77.5 102.7	3 3	0.117 0.119	0.068 0.07	1.72 1.7
None	None	167.7	3	0.181	0.101	1.79
None None	None None	125.2 99.7	3 3	0.184 0.166	0.102 0.093	1.8 1.78
None	None	100.5	3	0.129	0.072	1.79
None	None	71.5	2	0.125	0.069	1.81
None None	None None	27.5 47.2	1	0.112 0.132	0.065 0.075	1.72 1.76
None	None	164.2	3	0.179	0.101	1.77
None None	None None	43.2 68.5	1 2	0.141 0.178	0.081	1.74 1.76
None	None	114	3	0.178	0.101	1.8
None	None	112.2	3	0.129	0.072	1.79
None None	None None	68.2 46.5	2 1	0.152 0.119	0.084	1.8 1.72
None	None	103	3	0.182	0.101	1.8
None None	None None	48.9 53.2	1	0.127 0.152	0.071 0.084	1.78 1.8
None	None	63.6	2	0.132	0.082	1.79
None	None	51.2	2	0.115	0.067	1.71
None None	None None	50.8 76.2	2 3	0.121 0.159	0.071 0.088	1.7 1.8
None	None	59.4	2	0.135	0.075	1.8
None None	None None	101 88.5	3 3	0.159 0.142	0.089 0.079	1.78 1.79
None	None	94.1	3	0.142	0.079	1.79
None	None	72.4	2	0.156	0.087	1.79
None None	None None	71.4 59.3	2 2	0.151 0.163	0.083	1.81 1.77
None	None	121.7	3	0.153	0.09	1.7
None	None	101 90.9	3 3	0.155	0.086	1.8
None None	None None	90.9 98.2	3	0.161 0.108	0.089	1.8 1.8
None	None	94.5	3	0.131	0.075	1.74
None None	None None	89.2 101.4	3 3	0.105 0.145	0.061 0.08	1.72 1.81
None	None	39.2	1	0.136	0.08	1.7
None	None	106.3	3	0.154	0.09	1.71
None None	None None	89.2 89.1	3 3	0.166 0.141	0.092 0.081	1.8 1.74
None	None	88.7	3	0.106	0.061	1.73
None None	None None	66.7 106.2	2 3	0.119 0.178	0.066 0.101	1.8 1.76
None	None	66.5	2	0.101	0.057	1.77
None	None	115	3	0.175	0.102	1.71
None None	None None	152 71.5	3 2	0.184 0.111	0.104 0.061	1.76 1.81
None	None	78.7	3	0.181	0.101	1.79
None None	None None	87.8 108.7	3 3	0.191 0.177	0.106 0.102	1.81 1.73
None	None	79.6	3	0.17	0.1	1.7
None	None	106	3	0.169	0.094	1.79
None None	None None	75.8 71.3	3 2	0.173 0.115	0.096 0.064	1.8 1.79
None	None	81.3	3	0.11	0.061	1.8
None None	None None	99.6 95	3 3	0.115 0.119	0.063	1.82 1.8
None	None	89	3	0.138	0.077	1.79
None	None	62.4	2	0.125	0.071	1.76
None None	None None	77 89.78	3 3	0.129 0.129	0.071 0.072	1.81 1.79
None	None	77.46	3	0.158	0.089	1.77
None None	None None	78.42 81.22	3 3	0.181 0.183	0.104 0.107	1.74 1.71
None	None	85.3	3	0.141	0.078	1.8
None None	None	77.65 80.41	3 3	0.163 0.159	0.092	1.77
None None	None None	80.41 84.7	3	0.159 0.116	0.091 0.068	1.74 1.7
None	None	82.39	3	0.108	0.06	1.8
None None	None None	100.56 91.01	3 3	0.117 0.133	0.068 0.078	1.72 1.7
None	None	75.59	3	0.141	0.081	1.74
None None	None None	77.46 76.67	3 3	0.155 0.166	0.091 0.092	1.7 1.8
None	None	109.22	3	0.166	0.092	1.8
None	None	110.96	3	0.111	0.065	1.7
None None	None None	83.26 91.94	3 3	0.151 0.102	0.087 0.057	1.73 1.78
None	None	155.5	3	0.109	0.06	1.81
None	None	80.82	3	0.177	0.102	1.73
None None	None None	87.19 84.96	3 3	0.171 0.164	0.1 0.091	1.71 1.8
None	None	80.37	3	0.116	0.068	1.7
None None	None None	85.99 108.67	3 3	0.161 0.119	0.089 0.07	1.8 1.7
None	None	96.99	3	0.183	0.105	1.74
None	None	102.43	3	0.192	0.109	1.76
None None	None None	130.47 101.42	3 3	0.101 0.133	0.056 0.078	1.8 1.7
None	None	77.77	3	0.141	0.078	1.8
None None	None None	125.76 115.74	3 3	0.125 0.117	0.069 0.065	1.81 1.8
None	None	103.73	3	0.116	0.064	1.81
None	None	81.45	3	0.137	0.08	1.71
None None	None None	148.14 148.4	3 3	0.139 0.147	0.081	1.71 1.79
None	None	99.61	3	0.191	0.109	1.75
None None	None None	75.33 75.37	3 3	0.196 0.181	0.109 0.105	1.79 1.72
None	None	83.6	3	0.187	0.106	1.72
None	None	91.2	3	0.144	0.08	1.8

Genotypes Fokl FF=1, Ff=2, ff=3	Genotypes Apal AA=1, Aa=2, aa=3	Genotypes Bsml BB=1, Bb=2, bb=3
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