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LIVESTOCK DISEASE FOREWARNING METHODOLOGY

Powered by Artificial Intelligence & Machine Learning

ICAR – NATIONAL INSTITUTE OF
VETERINARY EPIDEMIOLOGY AND
DISEASE INFORMATICS
(ICAR – NIVEDI)



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**JUNE
2024**

**NOW LINKED WITH FARMER REGISTRATION
AND UNIFIED BENEFICIARY INFORMATION
SYSTEM (FRUITS)**



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Prepared By: Dr. K. P. Suresh
Dr. Rajeswari Shome
Dr. D. Hemadri
Dr. S. S. Patil
Dr. P. Krishnamoorthy
Dr. S. J. Siju
Dr. Gulati BR (Overall supervisor)

Front Page Design by: Krupakar K. R

Technical support: Sushma R, Navnath M Kamble, Krupakar K R, Chetan A.J.

Disclaimer

The forewarnings are based on the retrospective disease data available in the NADRES database. Hence, for those states wherein data is limited/less, the forewarning may not be realistic. Further the forewarning will not take into consideration the control measures that are *in situ*.

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Director
ICAR- NIVEDI

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1. ABOUT THE METHODOLOGY...

Livestock sector plays a crucial role in the rural economy of India as around 20.5 million people depend upon livestock for their livelihood. Even though the investment in the livestock sector is meagre, tremendous achievements have been observed in the sector during the last decade. As it is an important component in poverty alleviation programmes, continuous emphasis is being laid on this sector for enhancing the quality of the primary and secondary products in the international market, which in turn demands improved animal health. Therefore, livestock development programmes cannot succeed unless a well-organized animal health service is built up and in place for safeguarding the livestock against economically important diseases.

India has made a noteworthy success in the eradication of Rinderpest (RP), CBPP, AHS and Dourine. However, there are several other infectious and non-infectious diseases prevailing in the country causing huge annual economic loss. Prevention, control and eradication of the animal diseases need a thorough understanding of the epidemiology as well their economic impact.

National Institute of Veterinary Epidemiology and Disease Informatics (NIVEDI) has the mandate to carry out research activities in the area of veterinary epidemiology and disease informatics. With the eradication of RP successfully, India has not only proved its ability to face the challenges but also to succeed, despite various limitations. Similar efforts are needed to control and eradicate diseases like FMD, PPR, Brucellosis, CSF, HS etc., which cause huge economic loss annually to the livestock industry. To this end, ICAR-NIVEDI has identified 15 priority livestock diseases, based on the past incidence patterns and has built a strong database of these diseases. The database, which is the backbone of the National Animal Disease Referral Expert System (NADRESv2), is used for providing monthly livestock disease forewarning, which is compiled in this monthly bulletin to alert the animal husbandry departments, both at the National/state level, to take appropriate control measures. We hope users/stakeholders find this bulletin useful in their quest to control livestock diseases. This forewarning bulletin will assist the field Veterinarians in adopting appropriate preventive and control measures, thereby reducing the occurrence of livestock disease outbreaks.

2. SUMMARY OF THE FOREWARNING METHODOLOGY....

The association between infectious diseases and the climate was known from ancient times. Hippocrates observed in the 5th century that epidemics were associated with natural phenomena rather than divinities or demons. In modern times, our increasing capabilities to detect and predict climate variations joined with growing evidence for global climate change, have powered interest in understanding the impacts of climate on animal health, particularly the emergence and transmission of infectious disease agents. Simple reasoning suggests that climate can affect infectious disease patterns because the pathogens (viruses, bacteria, and parasites) and their vectors are sensitive to temperature, moisture, and other ambient environmental conditions.

India being an agriculture-based country, the livestock sector plays a vital role in contributing to the economy. A robust reporting and forewarning system enable the concerned authorities in disease preparedness and awareness of the risk associated with livestock diseases. Therefore, the economic loss due to morbidity and mortality of the animals is reduced thereby helps to increase the productivity in terms of egg, meat, and dairy products. National Animal Disease Referral Expert System database is a weather-based forewarning system enabled with an artificial intelligence system developed by ICAR-National Institute of Veterinary Epidemiology & Disease Informatics Bengaluru, Karnataka state, India that forecast potential threats from pathogens two months in advance to provide the stakeholders with sufficient timeline for awareness and preparedness to act. Artificial Intelligence (AI) and Machine Learning (ML) models use the programmed algorithms that receive and analyse input data to predict output (Infectious risk prediction) values within an acceptable range. As new data fed into these algorithms, they learn and optimize their operations to improve performance, developing intelligence over time.

The livestock disease forewarning for June 2024 revealed Jharkhand (185), Uttar Pradesh (99), Assam (80) and Karnataka (70) as the leading states anticipated to experience significant outbreaks of livestock diseases.

In terms of projected disease occurrences, Jharkhand is poised to encounter a higher number of anticipated Babesiosis, Fasciolosis, Trypanosomosis, Theileriosis and PPR outbreaks. Moreover, the simultaneous presence of FMD and HS is likely to be observed in Assam, Himachal Pradesh, Jharkhand, Karnataka, Kerala, Meghalaya, Odisha, Rajasthan and West Bengal. Among the different diseases in livestock, the predicted outbreaks are expected to be high for Babesiosis (95) and Theileriosis (88).

The primary obstacles facing an efficient disease control program include inadequate comprehension of the intricate disease dynamics, the extensive host range of pathogens, and the broadening pathogen niches attributed to climate change. Moreover, India has observed the emergence and resurgence of numerous infectious agents over the past decade, with a majority of these ailments having zoonotic attributes. This underscores the pressing need to enhance the nation's monitoring and surveillance systems. To effectively tackle significant livestock diseases within the country, a strategic approach involves implementing existing control measures in high-risk regions, coupled with routine disease surveillance and monitoring. This combination can contribute to a well-structured control program capable of addressing major livestock diseases

Table S1. Summary of Statewise Livestock Disease forewarning for June- 2024

Sl. No	State Name	ASF	Anthrax	Babesiosis	BQ	BT	CSF	ET	Fasciolosis	FMD	HS	LSD	PPR	S&G Pox	Theileriosis	Trypanosomosis	Total number of districts predicted for risk of disease
1	Andaman & Nicobar	0	0	1	0	0	0	0	3	2	0	0	0	0	0	0	6
2	Andhra Pradesh	0	4	0	1	0	0	0	0	0	2	0	3	2	0	0	12
3	Arunachal Pradesh	0	0	0	0	0	0	0	3	2	0	7	0	0	0	0	12
4	Assam	4	1	10	13	0	9	6	3	1	6	10	10	4	3	0	80
5	Bihar	0	0	1	0	0	0	0	0	0	0	1	1	0	0	1	4
6	Chhattisgarh	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	1
7	Goa	0	0	0	0	0	0	0	0	0	1	1	0	0	1	0	3
8	Gujarat	0	0	0	0	0	0	0	0	0	2	6	1	2	0	0	11
9	Haryana	0	0	0	0	0	2	2	0	0	0	0	2	0	1	0	7
10	Himachal Pradesh	0	0	0	0	0	0	0	0	2	1	0	3	2	0	0	8
11	Jammu & Kashmir	0	0	0	0	0	0	0	0	0	0	0	0	4	0	0	4
12	Jharkhand	0	4	25	12	0	13	9	25	17	6	0	20	5	24	25	185
13	Karnataka	0	6	1	12	0	0	14	0	12	5	7	9	3	1	0	70
14	Kerala	3	2	14	0	0	2	1	0	14	7	5	2	0	13	0	63
15	Madhya Pradesh	0	0	1	3	0	0	0	0	0	3	0	2	0	0	0	9
16	Maharashtra	0	1	0	0	0	0	0	0	0	1	0	0	1	2	0	5
17	Manipur	0	0	0	2	0	4	0	4	0	1	0	0	0	0	0	11
18	Meghalaya	0	2	0	0	0	2	0	0	4	1	0	1	2	2	0	14
19	Mizoram	6	0	0	0	0	1	0	1	0	0	0	0	0	0	0	8
20	Nagaland	2	0	0	0	0	1	0	0	0	0	2	0	0	0	0	5
21	Odisha	0	1	0	2	0	0	0	0	2	2	0	4	1	1	0	13
22	Puducherry	0	0	3	0	0	0	0	1	0	0	0	0	0	0	0	4
23	Punjab	0	0	1	0	0	0	0	0	0	0	0	0	0	1	0	2
24	Rajasthan	0	0	2	0	0	0	2	0	2	1	0	5	1	0	0	13
25	Sikkim	2	0	0	0	0	1	0	0	0	0	4	1	0	0	0	8
26	Tamil Nadu	1	4	0	2	0	0	0	0	0	0	0	2	2	0	0	11
27	Telangana	0	1	0	0	0	0	1	0	0	0	0	4	2	0	0	8
28	Tripura	0	0	2	1	0	1	0	5	4	0	3	3	5	1	0	25
29	Uttar Pradesh	0	0	30	0	0	0	1	0	0	1	0	3	1	24	39	99
30	Uttarakhand	1	0	0	0	0	0	0	0	1	0	0	2	0	0	0	4
31	West Bengal	0	1	4	3	0	0	0	1	2	3	0	5	2	14	2	37
Total number of districts likely for risk of disease		19	27	95	51	0	36	36	46	65	43	46	84	39	88	67	742

3. INTRODUCTION TO NADRES v2

The geographic and seasonal distribution of many infectious diseases are associated with climate and therefore the possibility of using seasonal climate forecasts as predictive indicators in disease early warning system (EWS) became imminent. In this context, ICAR-NIVEDI, in its quest for achieving better livestock health, had developed an interactive web portal named “National Animal Disease Referral Expert System (NADRES)” during early part of the first decade of the millennium. The web portal, which was developed from the financial support of National Agricultural Technology Project, was launched in the year 2005. The portal which is interactive, allows the user/stakeholder to access livestock disease forewarning (n=13) at the district level for entire country two months in advance. The portal which was initially built on oracle platform was later changed to MySQL platform to store the administrator provided disease information and other relevant meteorological and risk factor information. However, with the availability of remote sensed satellite images and the advancement in information technology and statistical algorithms, the upgradation of NADRES became inevitable. To this end, a newer version of NADRES (NADRES V2) has been developed.

How it is different from previous version?

In brief, it can be said that NADRES v2 underwent a sea change not only in its internal structure but also in its physical design. As a result, now the central menu bar consists of Home, about us, Risk factors, Analysis, Livestock disease, post prediction validation and contact details. Risk factors menu comprises of details on resolution, time interval, units and source of 18 meteorological and 5 remote sensing parameters. Analytics menu has various analysis options. The newly created livestock disease menu has the details regarding species affected, clinical signs and preventive measures to be adopted for the 15 economically important diseases. Post prediction validation menu contains the outbreak reports vs prediction. The menu bar on the RHS tabs include online GIS, state wise Livestock disease forecast, district wise Livestock disease forecast, Epi-calculator, download links for mobile app, etc. The website now hosts disease maps in the form of choropleth maps for 15 diseases in two time periods (1990-2000 and 2000-2018). Similarly, disease trends plots exhibit periodic regression plots providing future trend for the disease. On the LHS, Login menu is provided for authorized persons to login and enter disease details and other related parameters. Disease maps provide choropleth maps for 15 diseases in two time periods (1990-2000 and 2000-2018) is presented. Disease trends- Periodic regression plots are exhibited for prediction of the diseases. Auto-messaging option has been created to send the reminders in the form of text messages to concerned PI's and Co-PI's of NADEN centers for submission of outbreak reports. This message is sent weekly to all the concerned officials. Additionally, a message is sent to the concerned veterinary officers in Karnataka for initiation of preventive measures for the forewarned diseases at the block level. Plans are in place to incorporate farmers' and local vets' mobile numbers in to the list so that they may be asked to initiate preventive measures for the forewarned diseases.

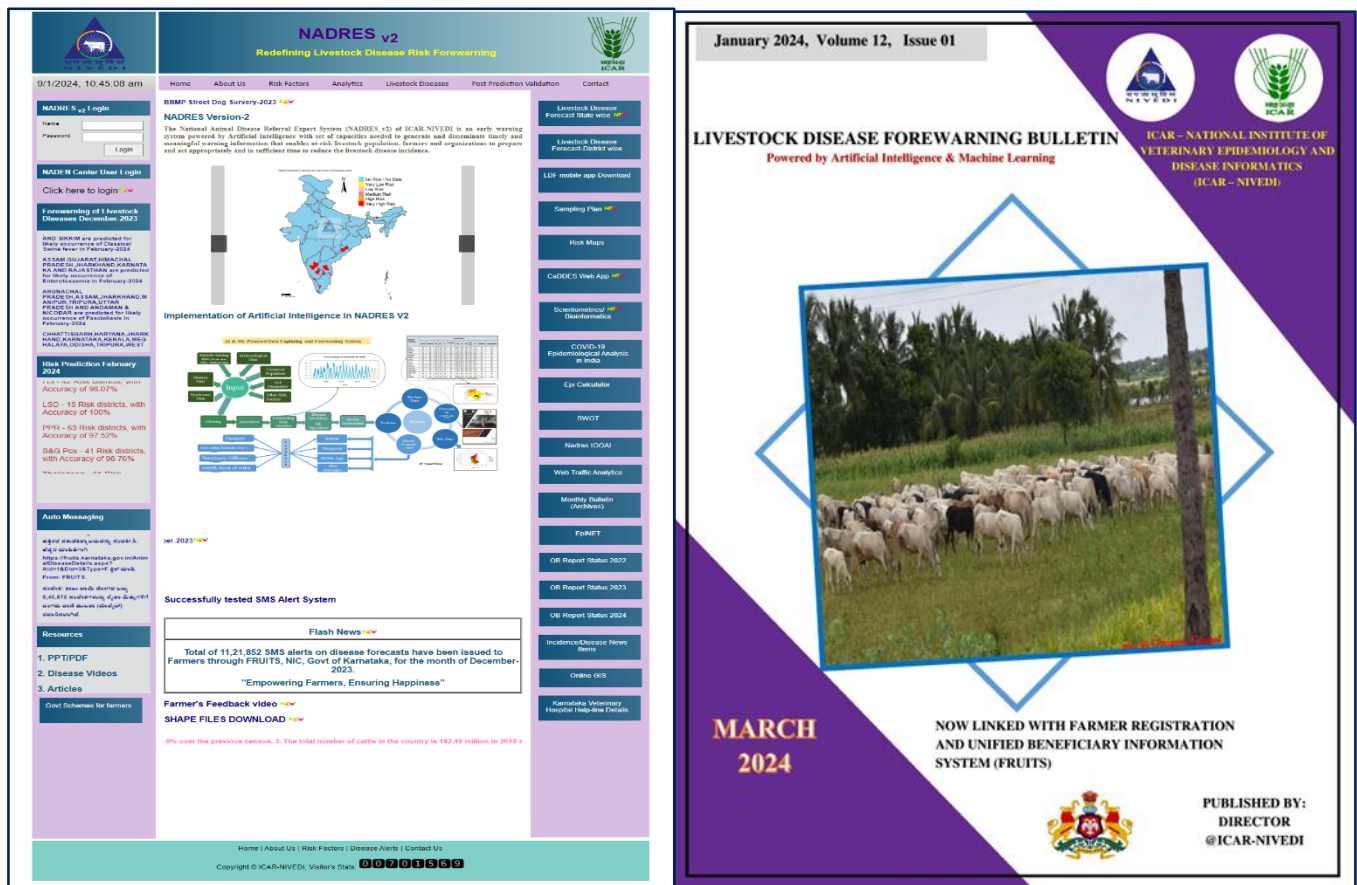


Fig 3.1. NADRES V₂ Home page

The forewarning methodology used is unique and has not been used earlier for livestock disease forewarning in India. Following few paragraphs describe about the forewarning methodology used. It is a well-known fact that weather plays an important role in the precipitation of many diseases and therefore, the climatic parameters such as land surface temperature (LST), precipitation, wind velocity, humidity etc are considered as risk parameters. These parameters along with other non-climatic parameters such as livestock population, density, Normalized Differential Vegetation Index (NDVI), soil moisture constitute the overall risk parameters. A total of 23 such parameters are collected/generated at village level and then aggregated to district level before these are used for analysis.

In addition to the output provided at interactive web portal, the NADRES output are also published in the form of monthly livestock disease forewarning bulletins. The prediction results come with a disclaimer that forewarnings do not take into account of the control measures that already in situ and also may not be realistic for those regions where the data is either unavailable or limited. This bulletin provides the likely occurrence of the 15 shortlisted diseases two months in advance at the district level, disease forewarning maps, prediction accuracy, details on diseases, species affected, clinical signs and its preventive measures.

In summary, it can be said that NADRES v₂ has underwent substantial changes not only in its internal structure but also in its physical design and can be a useful tool for visitors of the website, farmers, vets, policy makers etc.

4. Forewarning Methodology

Preamble

NADRES v2 is an early warning system powered by Artificial Intelligence with set of capacities needed to generate and disseminate timely and meaningful warning information that enables at-risk livestock population, farmers and organizations to prepare and act appropriately and in sufficient time to reduce the livestock disease incidence.

Objectives

- Development of forecasting model for the major livestock diseases and predicting the risk of livestock diseases in advance of two months.
- Development of state of art of communication models to communicate risk of livestock diseases to the stake holders.

I. Materials and data aquisition

Livestock disease data

Previous 10 years' livestock disease outbreak data retrieved from the NADRES database linked with Risk factors data.

Livestock population data

The population data at village level for five major livestock species viz., cattle, buffalo, sheep, goat and pigs were obtained from 20th Livestock census (2019) from Department of statistics, DAHD, GOI.

Species-wise & Category-wise Livestock Population (in thousands)					
Sl No	Species	Category	Population in 2012	Population in 2019	% Change
1	Cattle	Exotic	39732	51356	29.3
		Indigenous	151172	142106	-6
		Total	190904	193462	1.3
2	Buffalo	Total	108702	109852	1.1
3	Sheep	Exotic	3781	4088	8.1
		Indigenous	61288	70172	14.5
		Total	65069	74260	14.1
4	Goat	Total	135173	148885	10.1
5	Pig	Exotic	2456	1897	-22.8
		Indigenous	7837	7159	-8.7
		Total	10293	9056	-12
6	Yaks	Total	77	58	-24.7
7	Mithuns	Total	298	386	29.5
8	Horses & Ponies	Total	625	342	-45.3
9	Mules	Total	196	84	-57.1
10	Donkeys	Total	319	124	-61.1
11	Camels	Total	400	252	-37
Total Livestock			512056	536761	4.8

Meteorological and Remotely Sensed Data:

The parameters such as air temperature ($^{\circ}\text{C}$), perceptible water (mm), Precipitation rate (mm), pressure (millibar), relative humidity (%), Vwind (m/s), Uwind (m/s), Soil temperature ($^{\circ}\text{C}$), Vapour pressure(hPa), Wet day frequency, and sea level pressure (millibar) were extracted from National Centre for environmental prediction (NCEP). The parameters such as potential evapotranspiration (PET), Enhanced Vegetation Index (EVI), Leaf Area Index (LAI), Land Surface Temperature (LST), Normalised Difference Vegetation Index (NDVI) were extracted from remote sensed images from MODIS website (<https://modis.gsfc.nasa.gov/>). In brief, the MODIS products from NASA-TERRA satellite was downloaded for the Indian locations by specifying the tiles (H24V5, H25V6, H24V6, H24V7, H25V7, H25V8, H26V7, H26V6) from 2001 to till date.

The details are given below;

PRODUCT	Science Data Sets (HDF Layers)
MOD15A2H	Lai_500m (Leaf area index) 8 days average
MOD16A2	PET_500m (Total Potential Evapotranspiration) 8 days average
MOD11A2	LST_Day_1km (Daytime Land Surface Temperature) 8 days average
MOD13A1	i. 500m 16 days NDVI (Normalized Difference Vegetation Index) ii. Enhanced Vegetation Index (EVI) 16 days average

The downloaded HDF files (Datasets, which are multidimensional arrays (layers) of a homogeneous type) were converted to GeoTIFF files (single layer data) using R packages, which were later used to extract the parameters by linking it with the sinusoidal values of the Indian villages. The scale factors were multiplied for the extracted values as specified by the MODIS data products to get the values of the parameters. As shown above, the atmospherically corrected NDVI was collected on 16-day interval at 250-meter resolution using MODIS product MOD13A1 and LST was collected on 8-day interval using MOD11A2 at 1 KM resolution.

The parameters such as rainfall, Soil moisture (%), and Wind speed (m/s) were obtained from Global Land DataAssimilation System of NASA (<https://disc.gsfc.nasa.gov>). The remaining parameters were downloaded from climatic research unit (CRU) of University of East Anglia website. It is worth mentioning that the entire process of extraction, assimilation, processing and aligning have been done using R programming language and R environment. After aligning the climatic and non-climatic data with the disease and the livestock population data (aggregated at the district level), the statistical analysis was performed in the R environment.

Delta-Weather parameter

In the context of the National Animal Disease Referral and Export System (NADRES), made significant strides by enhancing database capabilities. original "Static Set" covered a 10-year average of 23 weather parameters from 2011 to 2022. Now, took a bold step forward, introducing a more intricate analysis. The upgraded "Static Set" remains the foundation but with a notable addition. Alongside the initial 23 parameters, we have included another set of 23 parameters known as "Delta" variables. These represent the differences between corresponding weather parameters from 2001 to 2021. This detailed differencing process offers valuable insights into long-term trends in meteorological conditions, especially relevant to understanding animal disease dynamics. In addition, we have introduced the "Dynamic Set" bringing another layer of sophistication. This set focuses on a more recent timeframe, using 2-year averaged parameter values for 2023 and 2024. Derived from data spanning from 2018 to 2023, the Dynamic Set also includes 23 "Delta" parameters, reflecting ongoing changes in climatic trends.

This dynamic approach ensures that NADRES stays updated with the latest meteorological patterns, crucial for timely responses in the early detection of animal diseases. With a total of 46 parameters now, this upgraded database is a cornerstone for NADRES. The integration of static and dynamic perspectives solidifies our position in meteorological analysis. NADRES is a reliable tool for animal disease management and a vital resource for informed decision-making in the intricate landscape of animal disease referral and export processes.

Initially, two regression models and Seventeen machine learning models were applied to test their suitability to fit the data and in all, Fourteen models; two regression model (Generalized Linear Model (GLM), Generalized Additive Model (GAM), and twelve machine learning models, viz., Gradient Boosting Machine Learning Algorithm (GBM), Random Forest (RF), Multivariate Adaptive Regression Splines (MARS), Extreme Gradient Boosting Machine (XGM), Support Vector Machine (SVM), Decision Trees (Tree_prob), Least Absolute Shrinkage and Selection Operator (Lasso), Functional Data Analysis (FDA), Gaussian Process (GP), Neural Network (NN), Multinomial Logistic Regression (Multinom_probs), and Kernel Support Vector Machine (KSVM) which fitted to data well were incorporated for the purpose of disease prediction. The models were trained using the case and control data available at ICAR-NIVEDI. Validation of the models were done by dividing the total observations for a particular disease into marker samples and validation samples and accuracy was tested in terms of discrimination power, which was done using Receiving Operating Characteristics (ROC), Cohen Kappa (Heildke Skill Score) and True Skill statistics (TSS) Accuracy, Error Rate, Precision, Sensitivity, Specificity, F1 score, Log Loss, and Gini-Coefficient. Once the models produce the probability value, it was used for categorizing the risk. Briefly, when all the models produce the p value of more than 0.5, then the highest p value is used for determining the high-risk category. If all the models or any one model produces the p value of less than 0.5, then the lowest p value was used for categorizing lower risk. This was done to minimize the false alert. Thus, the risk predictions based on the probability values ranging from 0-1 are made as follows; Very High Risk ($p=0.81-1.0$), High Risk ($p=0.61-0.80$), Moderate Risk ($p=0.41-0.60$), Low Risk ($p=0.21-0.40$), Very Low Risk ($p=0.0-0.20$) and No Risk ($p=0.0$) for the occurrence of a said disease. It is believed that categorizing districts in to various risk categories will help the stake holders to effectively utilize the available resources (money and manpower).

II.NADRES v2 Data Flow and Data Processing Diagram

A) Data Flow Diagram:

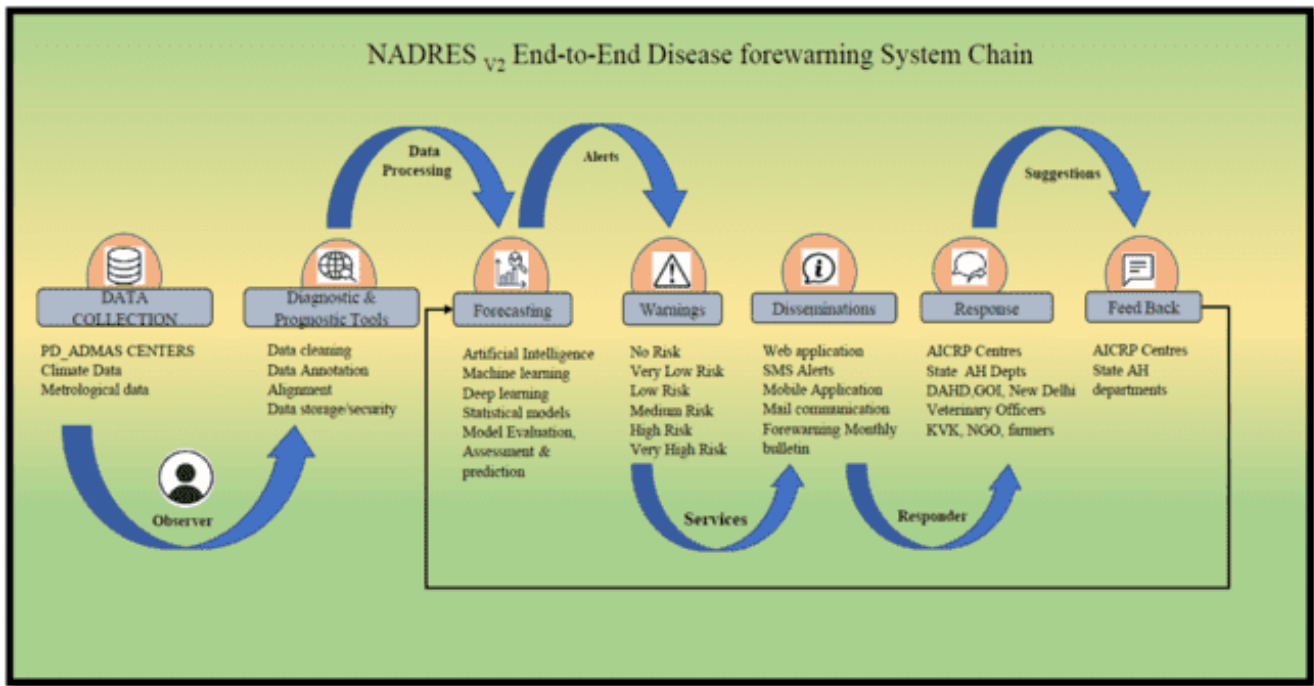


Fig 4.1. NADRES v2 Data Flow Diagram.

B) Artificial Intelligence enabled Data Capturing and Forewarning System:

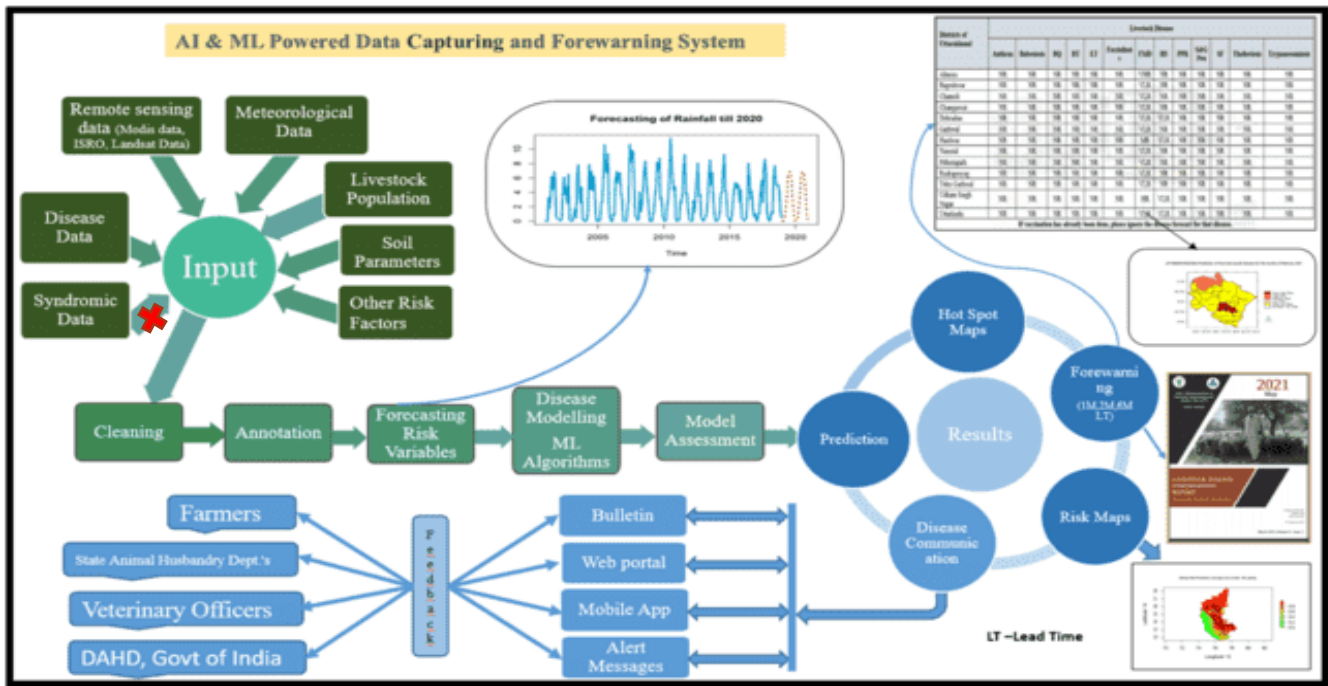


Fig 4.2. Data Capturing and Forewarning system

III. Weighted Outbreak Score

The outbreak data for the month of forecasting is extracted from NADRES database for the period of 10 years from current year. Outbreak data of 15 important livestock diseases are considered. The data is aggregated at district level and the weighted score is defined based on the number of outbreaks for each district in each month considering last 10 years. The weightage score was assigned as 0 for less than three number of outbreaks in the last 10 years for selected month, score 1 for 3–6 number of outbreaks and 2 for more than 6 outbreaks. This weightage score for each district is labelled as risk variable in building the models and risk maps.

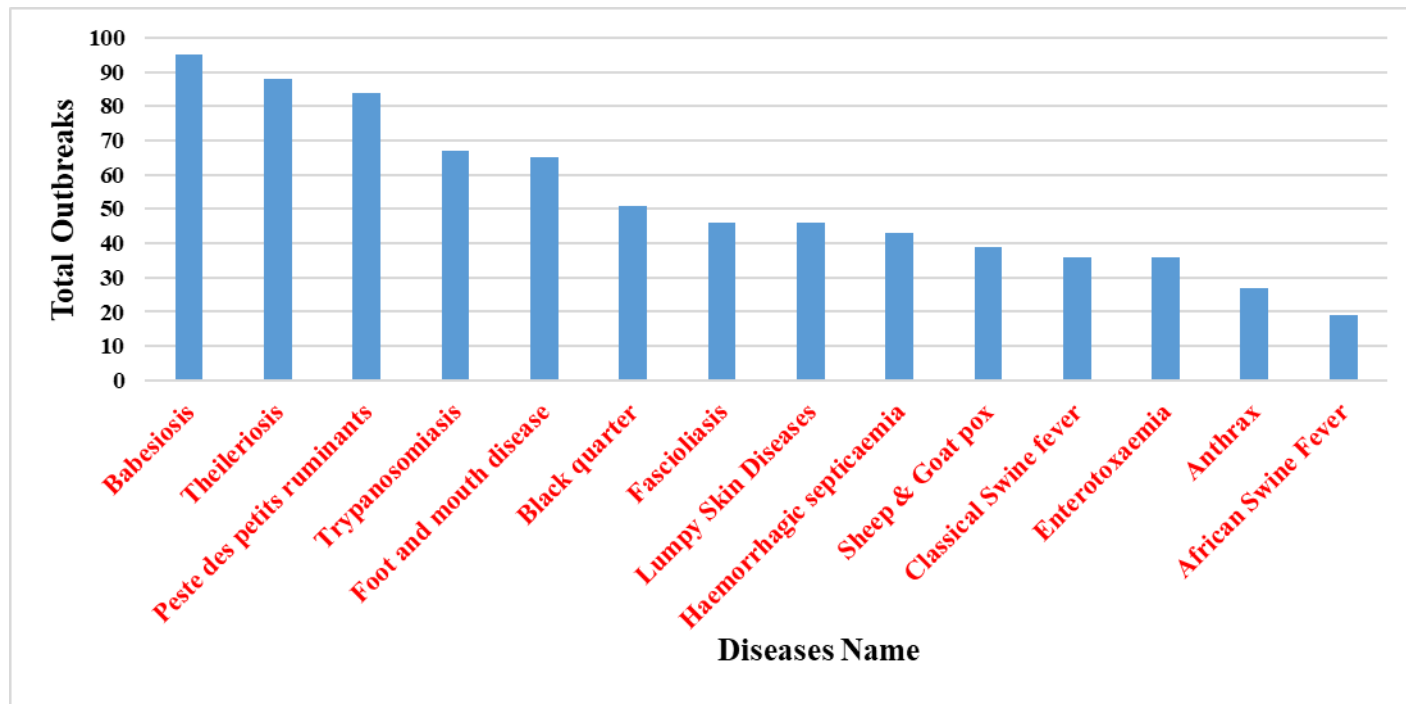


Fig 4.3. Top Fifteen livestock diseases (2024)

IV. Feature Extraction and Data Engineering

Data collection from different sources could be internal and or external to satisfy the objectives of forewarning requirements, data can be of any format, CSV, XML, JSON etc. In this processing of data and feature engineering, we focus mainly on understanding the specified data set and cleaning the dataset, a better understanding of features and their relationships, extracting essential variables, handling missing values and human error, identifying outliers, transforming features if there are outliers so that either truncates a data above a threshold or transform the data using log or any other transformation, scaling the features extracted. This process would be maximising the insights into a dataset.

V. Forecasting of Weather Parameters

Weather forecasting has been one of the most challenging problems around the world because of both its practical value in meteorology and the popular sphere for scientific research. Weather forecast systems are among the most complex equation systems that computer has to solve. A great quantity of data, coming from satellites, ground stations and sensors located around our planet send daily information that must be used to foresee the weather situation in next hours and days all around. Weather forecasts provide critical information about future weather. There are various techniques involved in weather forecasting, from relatively simple observation of the sky to highly complex computerized mathematical models. Further, forecast products by Indian Metrological department were used for validation of our forecasts (https://mausam.imd.gov.in/imd_latest/contents/extendedrangeforecast.php).

Following are the basic steps of forecasting process:

1. Determine the forecast's purpose
2. Establish a time horizon
3. Select a forecasting technique
4. Gather and analyse data
5. Perform the forecast
6. Monitor the forecast and use it in prediction of disease

Statistical Models used for forecasting of weather and remotely sensed variables

ARIMA stands for Autoregressive Integrated Moving Average. ARIMA is also known as Box-Jenkins approach. Box and Jenkins claimed that non-stationary data can be made stationary by differencing the series, Y_t . The general model for Y_t is written as,

$$Y_t = \phi_1 Y_{t-1} + \phi_2 Y_{t-2} \dots \phi_p Y_{t-p} + \epsilon_t + \theta_1 \epsilon_{t-1} + \theta_2 \epsilon_{t-2} + \dots \theta_q \epsilon_{t-q}$$

Where, Y_t is the differenced time series value, ϕ and θ are unknown parameters and ϵ are independent identically distributed error terms with zero mean. Here, Y_t is expressed in terms of its past values and the current and past values of error terms.

The ARIMA Model combines three basic Methods:

- Auto Regression (AR) – In auto-regression the values of a given time series data are regressed on their own lagged values, which is indicated by the “p” value in the model.
- Differencing (I-for Integrated) – This involves differencing the time series data to remove the trend and convert a non-stationary time series to a stationary one. This is indicated by the “d” value in the model. If $d = 1$, it looks at the difference between two-time series entries, if $d = 2$ it looks at the differences of the differences obtained at $d = 1$, and so forth.
- Moving Average (MA) – The moving average nature of the model is represented by the “q” value which is the number of lagged values of the error term.

This model is called Autoregressive Integrated Moving Average or ARIMA (p, d, q) of Y_t . We will follow the steps enumerated below to build our model. ARIMA models were run in 18 combinations of p, d, q. Based on the minimum AIC value, the order of ARIMA model was selected. This order was used for the prediction of all the weather parameters used in developing disease forewarning models.

VI. Implementation of Principal Component Analysis

Large datasets are gradually common and are often difficult to interpret. Principal Component Analysis (PCA) is a technique for reducing the dimensionality of such datasets, increasing the interpretability but at the same time, minimizing the information loss. The PCA is employed in NADRES v2 by creating new uncorrelated variables that successively maximize the variance. This means that `preserving as much variability as possible` translates into finding new variables that are linear functions of those in the original dataset, that successively maximize variance and that are uncorrelated with each other. Determining such new variables, the principal components (PCs) reduce to solve an eigenvalue/eigenvector problem. PCA can be based on either covariance matrix or the correlation matrix and the main use of PCA are descriptive. In the present study, all the meteorological and remote sensing variables are considering for PCA, with correlation matrix, the final output of principal components which are independent of each were considered for further ML modelling and risk estimation.

VII. Machine Learning Models

Disease outbreak data were aligned with generated risk variables to the respective latitude and longitude, which were subjected to climate-disease modelling. A number of models were fit to aligned data and tested for accuracy in terms of discrimination power. Two regression models, Generalized Linear Models (GLM) and Generalized Additive Models (GAM) and Seventeen machine learning algorithms, i.e. Random Forest (RF), Boosted Regression Tree (BRT), Artificial Neural Network (ANN), Multiple Adaptive Regression Spline (MARS), Flexible Discriminant Analysis (FDA), Classification Tree Analysis (CTA), Extreme Gradient Boosting Machine (XGM), Support Vector Machine (SVM), Decision Trees (Tree_prob), Least Absolute Shrinkage and Selection Operator (Lasso), Functional Data Analysis (FDA), Gaussian Process (GP), Neural Network (NN), Multinomial Logistic Regression (Multinom_probs), Kernel Support Vector Machine (KSVM), Ridge Regression, Naive Bayes were employed for disease modelling. Different modelling methods return different types of 'model object' and all these model objects could be used for the predict function to make predictions for any combinations of values of independent variables. Response plots were created to explore and understand model predictions.

The fitted models were assessed for their discriminating power using Receiving Operating Characteristic (ROC) curve, Cohen's Kappa (Heildke Skill Score), True Skill Statistics (TSS) Accuracy, Error_rate, Precision, Sensitivity, Specificity, F1 score, Loglass, Gini-Coefficient. These measures were used to evaluate the quality of predictions based on presence-absence data. Raster Stack was used to combine the results of individual predictions by different model methods. All the models were assessed for overfitting.

The outcome of best fitted models was in probability of disease occurrence and was categorised into 6 risk levels as No risk (NR), Very low risk (VLR), Low risk (LR), Moderate risk (MR), High risk (HR) and Very high risk (VHR) for enabling the stakeholders to take appropriate control measures by suitably allocating available resources.

5. ACCURACY OF PREDICTION

Serial No.	Diseases	Accuracy (%)
1.	African Swine Fever	100
2.	Anthrax	99.17
3.	Babesiosis	97.80
4.	Black quarter	98.90
5.	Bluetongue	99.86
6.	Classical swine fever	98.76
7.	Enterotoxaemia	98.35
8.	Fasciolosis	99.31
9.	Foot and mouth disease	98.76
10.	Haemorrhagic septicaemia	98.62
11.	Lumpy skin disease	99.03
12.	Peste des petits ruminants	97.39
13.	Sheep & Goat pox	99.58
14.	Theileriosis	97.39
15.	Trypanosomosis	99.31

Aggregation and prediction of livestock diseases at district level leading to higher accuracy.

- **Formula Used:** The **Accuracy** of disease prediction was calculated using the following formula.

$$\frac{TP + TN}{Total} * 100$$

TP-True Positive Observations, TN-True Negative Observations, Total- Total observations.

- Internal Accuracy was performed using 10 years of data. Accuracy obtained was >90% for all the diseases predicted.
- Despite the power of climate and disease risk models, considerable uncertainties remain, identifying these uncertainties, highlighting importance of improved data may improve the model accuracy, realism, confidence, together with translating uncertainties in model inputs into uncertainties in model outputs, are important benefits of modelling.

6. MORAN'S I FOR CLUSTERING OF LIVESTOCK DISEASES

Moran's I is a tool that measures spatial autocorrelation (feature similarity) based on both feature locations and feature values simultaneously. Given a set of features and an associated attribute, it evaluates whether the pattern expressed is clustered, dispersed, or random. The tool calculates the Moran's I Index value and both a Z score and p-value evaluating the significance of that index. In general, a Moran's Index value near +1.0 indicates clustering while an index value near -1.0 indicates dispersion.

Autocorrelation tool, the null hypothesis states that "there is no spatial clustering of the values associated with the geographic features in the study area". When the p-value is small and the absolute value of the Z score is large enough that it falls outside of the desired confidence level, the null hypothesis can be rejected. If the index value is greater than 0, the set of features exhibits a clustered pattern. If the value is less than 0, the set of features exhibits a dispersed pattern.

7. R SOFTWARE

R is a programming language and software environment for statistical analysis, graphics representation and reporting. R is a simple and effective programming language, which includes conditionals, loops, user defined recursive functions and input and output facilities. R statistical software version 3.6.2 (version 3.6.2, R Foundation for Statistical Computing, Dark and Stormy Night. <https://www.R-project.org/>) was used as an integrated suite for data mining, calculation and graphical display. Several R packages like *openxlsx*, *raster*, *RMySQL*, *rgdal*, *RColorBrewer*, *spldf*, *sp*, *spdep*, *xlsx*, *plyr*, *randomFores*, *dismo*, *SDMTool*, *dplyr*, *tmap* and *data table* were used for data extraction, data alignment, annotation, analysis, modelling and risk mapping.

8. FOREWARNING OF LIVESTOCK DISEASE FOR THE MONTH OF APRIL, 2024

I. Diseases, Species affected Clinical signs and its preventive measures.

Sl No.	Disease	Species Affected	Clinical Signs	Preventive Measures
1.	African Swine Fever (ASF)	Primarily affects domestic and wild pigs	High Fever, Lethargy and Weakness, Red or Blue Skin Discoloration, Pig display respiratory signs such as coughing, difficulty breathing, and nasal discharge, causes digestive symptoms, including diarrhea and vomiting.	involves strict measures like isolating new pigs, regularly disinfecting facilities, and controlling the movement of pigs and pig-related items. Early detection through surveillance, educating farm personnel, and managing wild boar populations are crucial, while proper disposal of infected material and quick reporting of suspected cases contribute to effective prevention and control. Collaboration among authorities, farmers, and the public, along with clear communication about ASF and its preventive measures, plays a key role in safeguarding pig populations and the swine industry.
2.	Anthrax (AX)	Most of the mammals and ruminants are highly susceptible. Pigs and Horses are moderately susceptible. Carnivores are relatively resistant.	Convulsion and sudden death with oozing of blood from natural orifices such as rectum and nose prior to death. Occasionally oedema develops in the throat and shoulder over a period of one week before death.	Ring vaccination and reporting of the disease is advised. Vaccination to be done in consultation with the veterinarians and as decided by state animal husbandry authorities. Strict biosecurity measures may be followed. Carcass may be disposed by deep burying covered with lime powder. Contaminated area may be disinfected with 4% formalin or 10% caustic soda. Grazing area may be restricted.
3.	Babesiosis (BA)	Cattle. Cross breeds are more susceptible.	High temperature, jaundice like symptoms, yellowish mucosal membrane of eye, rectum and coffee colour urine.	Periodical application of acaricides in and around the animal shed and on the animals. For therapeutic application, Diaminazine or Imidocarb can be useful.
4.	Black Quarter (BQ)	Common disease of cattle and sheep, but occasionally goats and pigs	High fever and lameness followed by swelling in the neck, shoulder, lumbar, gluteal and	Affected animals may be treated with suitable antibiotics. Vaccination to be done in consultation with the veterinarians and as decided by state animal

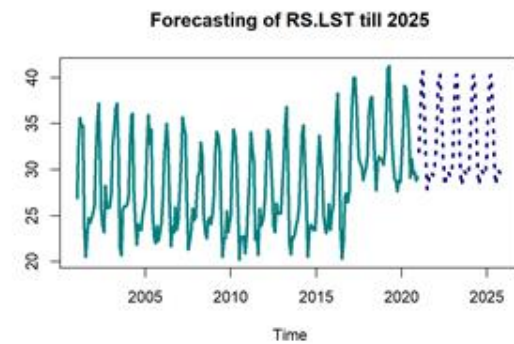
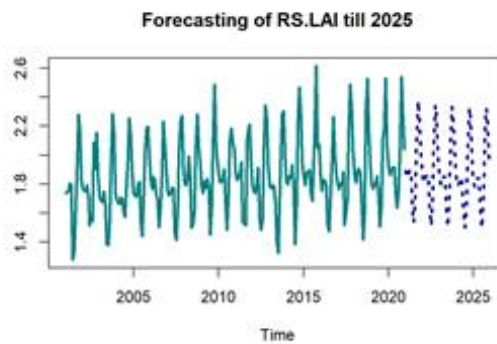
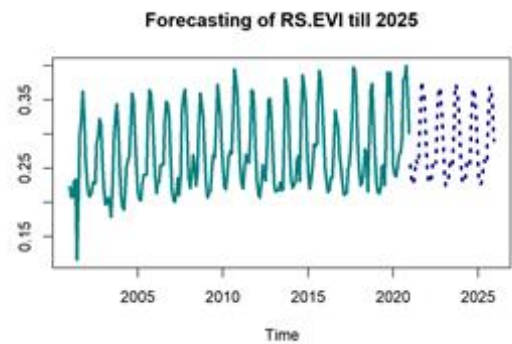
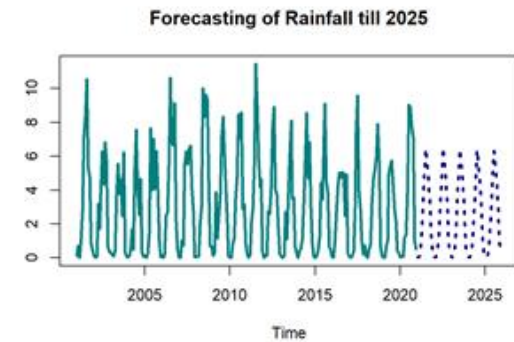
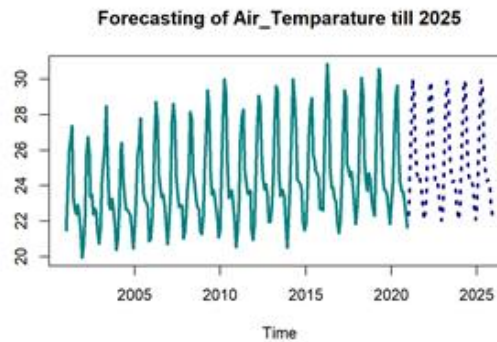
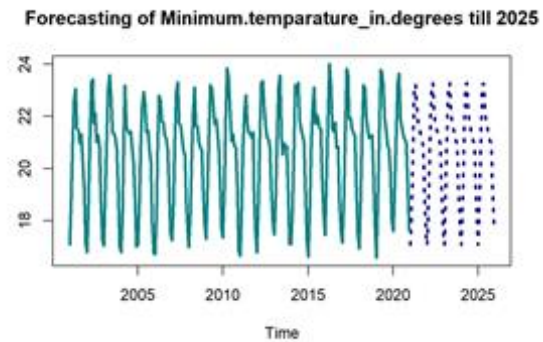
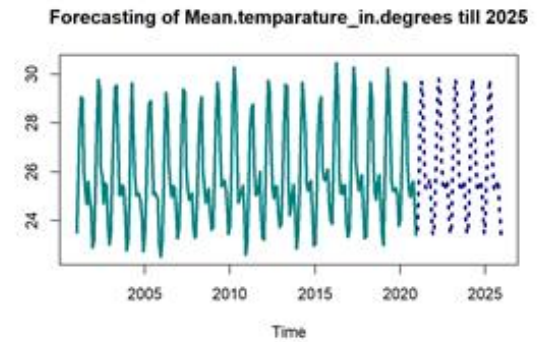
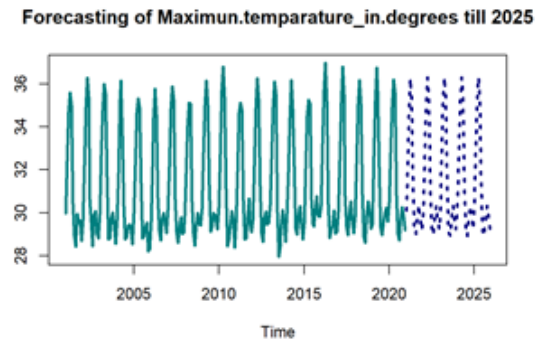
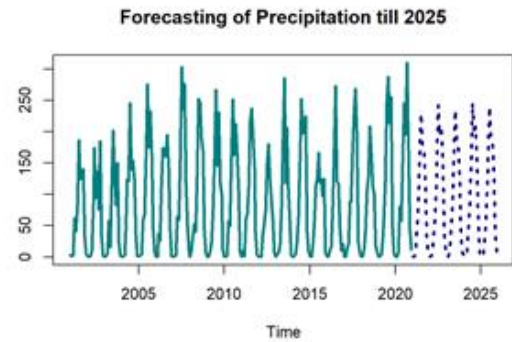
		also suffer from the disease.	sacral regions. Skin over the affected area become dark and crepitate on palpation. Loss of feed intake, colic, lateral recumbency, dyspnoea and death.	husbandry authorities. Strict biosecurity measures may be followed. Grazing area may be restricted. Carcass may be disposed hygienically.
5.	Bluetongue (BT)	Sheep are more susceptible than goats.	Fever, swelling of face, neck, eyelids respiratory distress, nasal discharge, Salivation, necrotic ulcers on tongue, dental pad, gum, lips hyperaemia of muzzle and may bleed at mucocutaneous junction. Affected tongue may become swollen, cyanotic and purple blue in colour – ‘bluetongue’.	Vector control using insecticides and good water management. Vaccination of susceptible animals preferably in the month of May. Do not shear sheep during winter months. Restriction in animal movement, segregation of affected animals and symptomatic treatment. Strict biosecurity measures.
6.	Enterotoxaemia (ET)	Common disease of sheep and goats especially among the young animals.	Dullness, opisthosomas, convulsions, coma and sudden death. Affected adult sheep, which survive for several days May show diarrhoea and staggering.	Affected animals may be treated with suitable antibiotics. Vaccination to be done in consultation with the veterinarians and as decided by State Animal Husbandry Authorities. Strict biosecurity measures may be followed. Carcass may be disposed hygienically. Grazing area to be restricted, stall fed, vitamins and probiotics may be provided.
7.	Fasciolosis (FA)	Cattle, buffalo, sheep and goats.	Progressive anaemia, pale mucous membrane, submandibular oedema (Bottle jaw), loss of appetite, weakness, isolated from flock while grazing, loss in production.	The animal should not be allowed to graze in water stagnant fields or submerged fodder should not be given directly to the animals. The submerged fodder can be processed through hay/silage preparation in order to destroy the metacercariae. The affected animals can be treated with Carbon tetrachloride/ Rafoxanide/Nitroxynil/ Niclofolan/Closantel/Oxyclozanide, under the strict supervision of veterinarian.

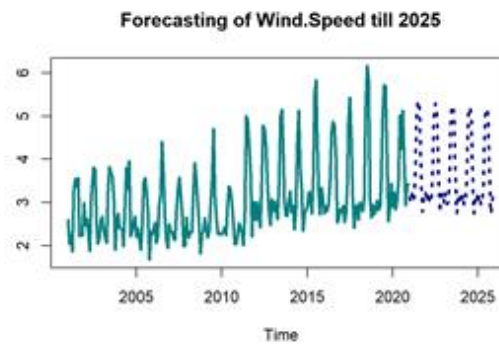
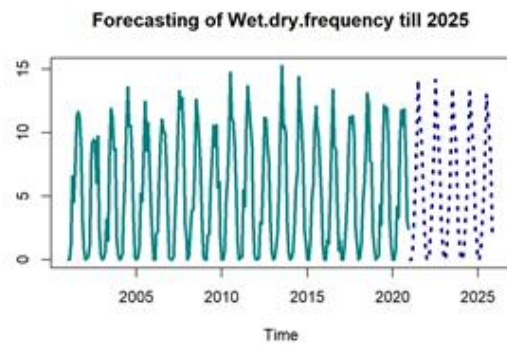
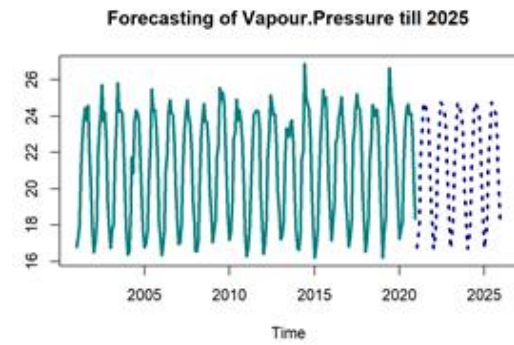
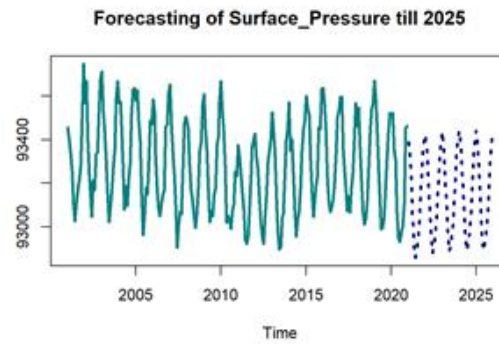
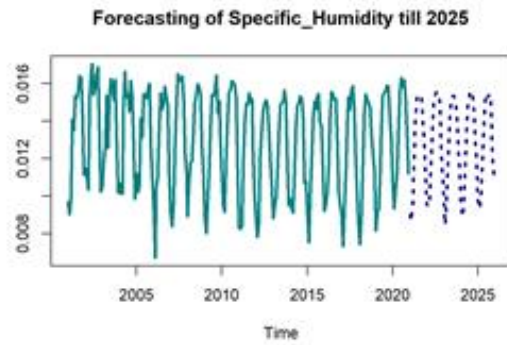
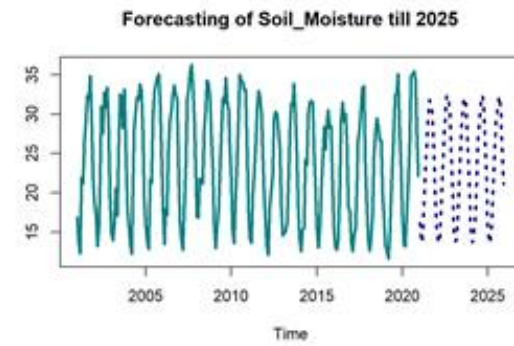
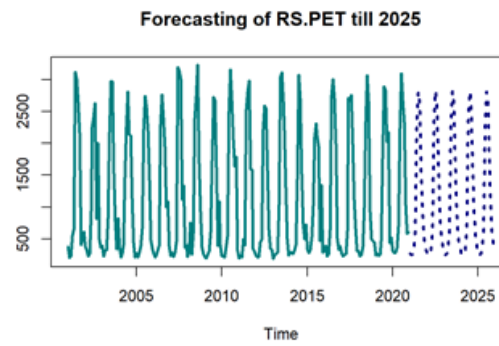
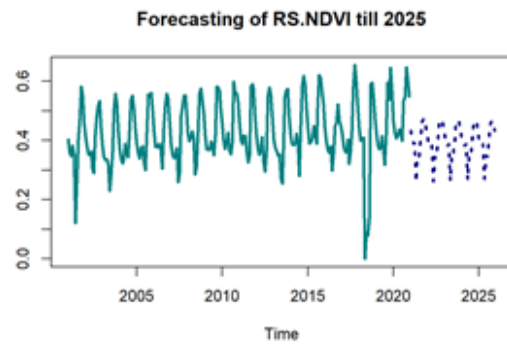
8.	Foot and Mouth Disease (FMD)	Cattle, buffalo, sheep, goats and pigs are often affected domesticated species, but the disease is more severe in cattle and pigs.	Fever, loss of feed intake, drop in milk production, drooling of saliva like ropey string, vesicles develop on the tongue, lips, gums, and palate and eventually rupture. Concurrent to oral lesions, vesicles also appear in inter digital skin and coronary band of the feet. The animal may open and close its mouth with a characteristic smacking sound. Sheep and goats may show lameness. In pigs, lesions may be seen on snout and also on the feet.	Regular vaccination and seromonitoring. Disinfection with sodium carbonate (4%) or 10% washing soda and strict biosecurity measures to be followed and animal movement may be controlled.
9.	Haemorrhagic septicaemia (HS)	Common disease for cattle and buffaloes, but can also occur among other species such as pigs, sheep, goats and many wild animals.	The disease starts with high fever, respiratory distress and haemorrhages maybe seen on the mucous membranes. There is lacrymation, nasal discharge, drop in milk production and anorexia. As the disease progress ear droops and the animals will be prostrated with cyanosis of mucous membranes. There may be oedema along the head, neck, thorax, vulva and anal areas. Sudden death occurs within few hours of clinical signs.	Affected animals may be treated with suitable antibiotics. Vaccination to be done in consultation with the veterinarians and as decided by state animal husbandry authorities. Strict biosecurity measures may be followed. Carcass may be disposed hygienically and stress factors may be reduced by following good animal husbandry practices.

10.	Lumpy skin disease (LSD)	Common disease for Cattle, Buffalo and other domestic animals	Fever, reduced milk production and skin nodules. Mastitis, swelling of peripheral lymph nodes, loss of appetite, increased nasal discharge and watery eyes are also common. Temporary or permanent infertility occur among infected cows and bulls	Vaccination of susceptible animals of above 3 months old age. Restriction on animal movement, strict biosecurity measures and proper disposal of carcass.
11.	Peste des Petits Ruminants (PPR)	Goats and sheep are most affected domestic animals.	Fever, nasal and ocular discharge, respiratory distress, necrotic lesions in buccal mucosa, gum, dental pad, palate, tongue and diarrhoea. Animals may die because of dehydration and pneumonia.	Vaccination of susceptible animals of above 3 months old age. Restriction on animal movement, strict biosecurity measures and proper disposal of carcass.
12.	Sheep and Goat pox (SGP)	Sheep and Goats	Respiratory distress and pock lesions over the non-hairy parts of body, more common in teat, udder, scrotum, head, neck, ear, perineum, inner aspect of thighs and under tail.	Vaccination of susceptible animals of above 3 months old age. Symptomatic treatment of affected animals. Restriction on animal movement, strict biosecurity measures and proper disposal of carcass.
13.	Classical Swine Fever (CSF)	Pigs	Fever, Conjunctivitis, purplish discolouration of snout, ears, abdomen, inner side of the legs and staggering gait.	Vaccination of susceptible animals. Restriction on animal movement, strict biosecurity measures and proper disposal of carcass.

14.	Theileriosis (TE)	Large Ruminants. Crossbreed cattle are more vulnerable.	High temperature, yellowish eye, sometime eyes may be heavily swollen, icteric mucosal membrane of rectum, dark yellowish urine, sometime may reach to coffee colour. Antibiotic is of no use to check the fever.	Periodical application of acaricides in and around the animal shed and on the animals. Therapeutic treatment with Buparvaquone can be useful in both early and advanced stages of the infection.
15.	Trypanosomosis (TR)	Domestic and wild carnivores and herbivores including cattle, buffalo, horse, donkey, camel, dog and cats. Buffaloes are known as carriers.	Fluctuating high fever which is not responded by antibiotics, swollen lymph gland, chronic emaciation and weakness, loss of appetite, gradual loss of production.	The affected animal should be treated with Diaminazine compounds or chloride and sulphate salts of Quinapyramine. Periodical spray of insecticide in and around animal shed to remove the flies.

II. Forecasting of remote sensing and meteorological parameters till, 2025(Ex. Karnataka)





III.SIGNIFICANT WEATHER PARAMETERS TABLE USING DISCRIMINANT FUNCTION ANALYSIS

SI. No	Disease Names	ML Derived Significant Parameters
1	African Swine Fever	Air Temperature, Delta-Precipitation, Delta-Leaf Area Index
2	Anthrax	WET, Maximum Temperature, Wind Velocity, Soil Temperature
3	Babesiosis	Delta-Relative Humidity, Precipitable-water, Delta-Minimum Temperature
4	Black quarter	Wind Velocity, Potential Evapotranspiration, Air Temperature
5	Bluetongue	Wind Velocity, Soil Temperature
6	Classical Swine fever	Air Temperature, Enhanced Vegetation Index, Relative Humidity
7	Enterotoxaemia	Soil Temperature, Precipitable water, Delta-Precipitable water
8	Fascioliasis	Precipitable Water, Air Temperature, Precipitation-rate
9	Foot and mouth disease	Delta-Temperature Minimum, Precipitation-rate, Precipitable water, Soil Temperature
10	Hemorrhagic septicemia	Precipitable Water, Delta-Pressure
11	Lumpy Skin Diseases	Delta-Vapour Pressure, Minimum Temperature, WET, Delta- Enhanced Vegetation Index, Pressure
12	Peste des petits ruminants	Delta-Minimum Temperature, Precipitable water, Delta-Relative Humidity
13	Sheep & Goat pox	WET, Enhanced Vegetation Index, Relative Humidity
14	Theileriosis	Precipitable Water, Delta- Minimum Temperature, Precipitation-rate
15	Trypanosomiasis	WET, Precipitable-water, Delta-soil moisture

Table 7.1: Significant weather parameters govern the Livestock disease incidence (forecast).

9. POST PREDICTION VALIDATION

DIMAPUR | Publish Date: 4/14/2019 AH&VS TEAM VISITS AFFECTED AREAS UNDER MEDZIPHEMA,
Source: <http://www.nagalandpost.com>

Following reports of a good number of buffaloes dying in a recent outbreak of suspected Haemorrhagic septicaemia (HS), a team from Animal Husbandry and Veterinary Services (AH&VS) department visited the affected areas under Medziphema on April 12. (Haemorrhagic septicaemia is a contagious bacterial disease that affects cattle and water buffaloes with a high mortality rate in infected animals).

AH&VS, deputy director & principal investigator, AICRP-ADMAS, Dr S. Amenla Walling, in a press release reported that the team consisted of the department's director, Dr Temsummeren, along with additional director, Dr. Budhi Lama, and other officials from the department. The press release added that the area is prone to such kind of disease outbreaks and the department officials reminded villagers to cooperate with the department and vaccinate their animals against such outbreaks. The team told the villagers that even an outbreak can be contained more effectively if villagers report the matter on time to the nearest Veterinary Health Centre.

The villagers admitted in the meeting that they had not reported the recent outbreak to the department initially. The director appreciated the CVO Dimapur and his Rapid Response Team for their quick action after receiving information and for remaining stationed in the outbreak area to date. Free medicine was also distributed among the villagers. The department, through the press release also appealed to everyone to report such matters to the nearest Veterinary Health Centre (so that qualified staff may intervene quickly), instead of publicizing it in other ways. It stated that the department is prepared to extend services to any outbreak of diseases in animals to control such things.

The press release also pointed out that to control the recent outbreak, the department had to direct its officials to make their own transport arrangements to go to the affected areas because the State Election department did not consider an appeal to exempt the department's emergency duty vehicle from election duty.

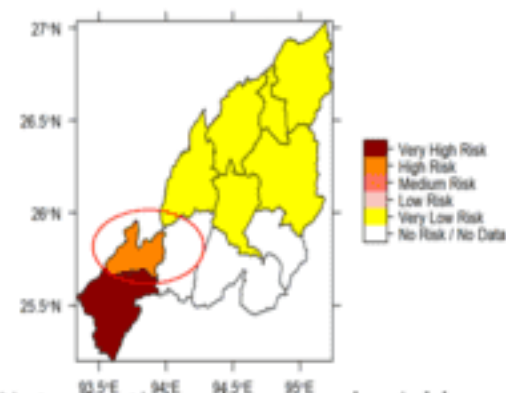
Meanwhile, when contacted, Dr S. Amenla Walling told Nagaland Post that it is difficult to say if the disease has been fully contained since its free grazing season for the animals, but the department is doing its best under the circumstances.

NIVEDI PREDICITONS

Districts of Nagaland	HS prediction for February 2019	HS prediction for March 2019	HS prediction for April 2019
Peren	VLR	VLR	VHR
Dimapur	VLR	NR	HR
Kohima	VLR	VLR	NR
Wokha	VLR	NR	VLR



Risk Prediction of Haemorrhagic septicaemia for the month of April 2019



Note: Spatial and temporal neighbours

Andaman and Nicobar Report June-2020

NIVEDI Prediction

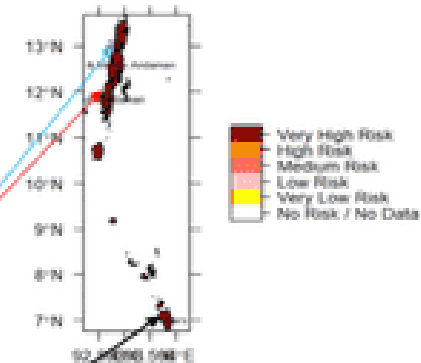
District wise Livestock Disease forwarning for June 2020: Andaman and Nicobar

Districts of Andaman and Nicobar	Livestock Diseases												
	Anthrax	Buberculosis	BQ	BT	ET	Fasciolosis	FM	HN	PP	S&G Fus	SF	Theileriosis	Trypanosomiasis
Nicobar	NR	NR	NR	NR	NR	VLR	NR	NR	NR	NR	NR	NR	NR
North & Middle Andaman	NR	NR	NR	NR	NR	VLR	NR	NR	NR	NR	NR	NR	NR
South Andaman	NR	NR	NR	NR	NR	VLR	NR	NR	NR	NR	NR	NR	NR

If vaccination is already been done please ignore the disease forecast for that disease.

*No risk/No data available (NR), Very low risk (VLR), Low risk (LR), Moderate risk (MR), High risk (HR)/Very high risk (VHR)

NDAMAN & NICOBAR ISLANDS Risk Prediction of Fascioliasis for the month of June 2020



Andaman and Nicobar Report June-2020



Number of cases of of parasitic cases and other diseases reported from A. & N Islands during the month of June 2020

CASES	FASCIOLIASIS	ASCARIASIS	AMPHISTOM	STRONGYLOID	COCCIDIOSIS	MASTITIS	TOTAL
South Andaman	24	48	192	34	2	7	307
N&M Andaman	78	43	14	5	3	10	333
Nicobar	79	31	0	0	0	0	110
TOTAL	361	122	206	39	5	17	750

Dr. Jai Sunder
FI, AICRP-ADMAS
Port Blair

HIMACHAL PRADESH REPORT AUGUST-2020

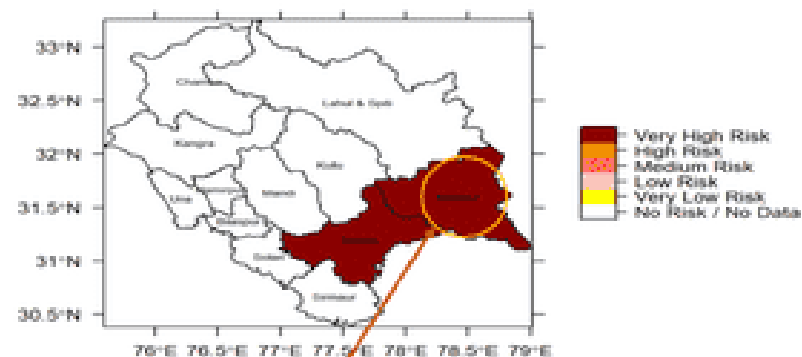
NIVEDI Prediction

District wise Livestock Disease forecasting for July 2020 Himachal Pradesh

Districts of Himachal Pradesh	Livestock Diseases												
	Anthrax	Buberculosis	BQ	BT	ET	Fasciolosis	FMD	HS	PPV	MG	SP	Brucellosis	Typhus
Bilaspur	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR
Chamba	NR	NR	NR	NR	NR	NR	1/1.5	NR	NR	NR	NR	NR	NR
Hamirpur	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR
Kangra	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR
Kasauli	NR	NR	NR	1/1.5	NR	NR	NR	NR	NR	1/1.5	NR	NR	NR
Kullu	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR
Lahul & Spiti	NR	NR	NR	1/1.5	NR	NR	NR	NR	NR	NR	NR	NR	NR
Shimla	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR
Shimla	NR	NR	NR	NR	NR	NR	NR	NR	NR	1/1.5	NR	NR	NR
Sirmaur	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR
Solan	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR
Una	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR

If vaccination has already been done please ignore the disease forecast for that disease.

HIMACHAL PRADESH Risk Prediction of Sheep and Goat pox for the month of July 2020



HIMACHAL PRADESH Report July-2020

Form for submitting livestock disease outbreak data to NIVEDI. (REVISED REPORT-11/07/2020)

NAME OF THE COLLABORATING UNIT: _____

ADDRESS OF THE COLLABORATING UNIT: _____

REPORT FOR THE MONTH OF: _____

DATE OF REPORT: _____

NAME OF THE VILLAGE: _____

Latitude and Longitude of the village: _____

Postal pin code of the village: _____

NAME OF THE DISTRICT: _____

NAME OF THE VILLAGE: _____

SPECIES OF LIVESTOCK: _____

Year: _____

Month: _____

Number of outbreaks: _____

Number susceptible: _____

Number infected: _____

Number of deaths: _____

Number of vaccination: _____

FORMAT FOR SUBMITTING LIVESTOCK DISEASE OUTBREAK DATA TO NIVEDI. (REVISED REPORT-11/07/2020)

NAME OF THE COLLABORATING UNIT: _____

ADDRESS OF THE COLLABORATING UNIT: _____

REPORT FOR THE MONTH OF: _____

DATE OF REPORT: _____

NAME OF THE VILLAGE: _____

Latitude and Longitude of the village: _____

Postal pin code of the village: _____

NAME OF THE DISTRICT: _____

NAME OF THE VILLAGE: _____

SPECIES OF LIVESTOCK: _____

Year: _____

Month: _____

Number of outbreaks: _____

Number susceptible: _____

Number infected: _____

Number of deaths: _____

Number of vaccination: _____

KERALA REPORT JUNE-2021



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SUBMIT INFO SEARCH POSTS

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Anglophone Africa South Asia Middle East/North Africa Antimicrobial Resistance

Published Date: 2021-06-18 03:52:33 IST
Subject: PRO/AH/EDR> Foot & mouth disease - India (02): (KL) cattle, RFI
Archive Number: 20210617.8456180

FOOT & MOUTH DISEASE - INDIA (02): (KERALA) CATTLE, REQUEST FOR INFORMATION

A ProMED-mail post
<http://www.promedmail.org>
ProMED-mail is a program of the
International Society for Infectious Diseases
<http://www.isid.org>

Date: Wed 16 Jun 2021 12:34 IST
Source: The Hindu [edited]
<https://www.thehindu.com/news/national/kerala/steps-to-tackle-fmd-in-alappuzha/article34822360.ece>

The Animal Husbandry Department has ramped up measures to tackle the spread of foot-and-mouth disease (FMD) in Alappuzha.

Around 3000 head of cattle have been affected by the outbreak in the district in the last 1.5 months. At least 193 cattle died of the disease during the period. Around 550 head of cattle have contracted the disease, and 100 perished in the worst-affected Ambalapuzha South and Ambalapuzha North grama panchayats [for Kerala's decentralised administrative structure terminology, see <https://kerala.gov.in/local-self-government-Mod.AS>]. The disease has also been reported from Thakazhi, Thalavady, Edathua, Chettikulangara, Aryad, Pandanad, and Chengannur.

Treatment and vaccination are being carried out to check the spread of the disease. The department has deployed 140 teams in the affected areas. Special teams with doctors have been deployed to the worst-affected Ambalapuzha South and Ambalapuzha North grama panchayats. Doctors have been appointed to Chengannur, Ambalapuzha, and Veliyanad blocks to deal with emergencies during night.

Minister of Animal Husbandry J. Chinchurani said the department had taken measures to bring the outbreak under control.

District Animal Husbandry Officer A.G. Geo said the department had started a vaccination drive on 22 Apr [2021] soon after the FMD cases started to [be reported] in the district. As many as 6140 cattle in 51 grama panchayats have so far been vaccinated against FMD.

The outbreak of the disease was attributed to missed rounds of vaccination due to the COVID-19 pandemic.

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<promed@promedmail.org>

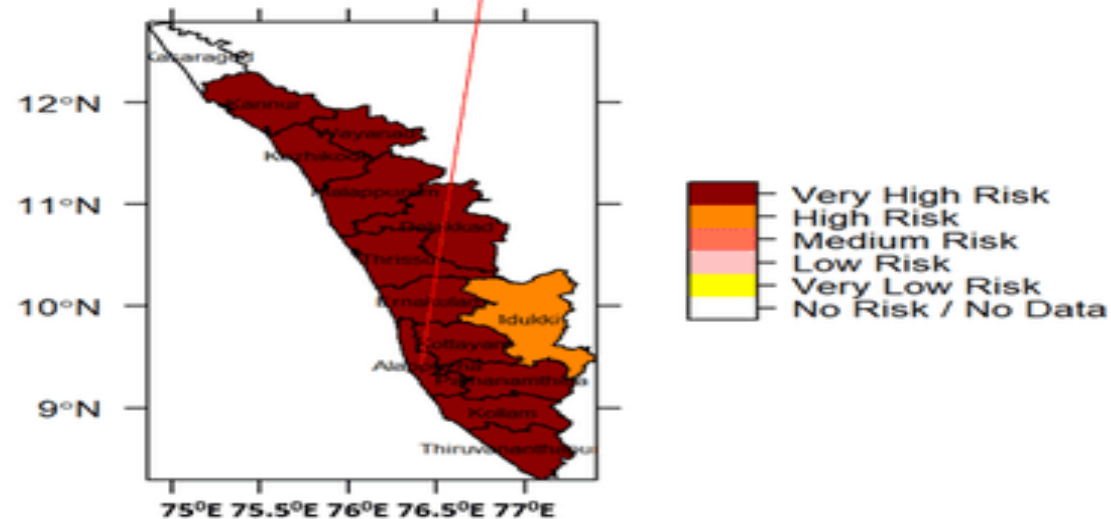
NIVEDI prediction in June 2021

District wise Livestock Disease Risk Forewarning for June 2021: Kerala

Districts of Kerala	Livestock Diseases												
	Anthrax	Babesiosis	BQ	BT	ET	Fascioliasis	FMD	HIS	PPR	S&G Pox	SF	Theileriosis	Trypanosomiasis
Alappuzha	NR	NR	NR	NR	NR	NR	VHR	MR	VLR	NR	NR	NR	NR
Ernakulam	NR	NR	NR	NR	NR	NR	VHR	HR	HR	NR	VHR	NR	NR
Idukki	VHR	VLR	VLR	VLR	NR	NR	HR	NR	VLR	NR	NR	NR	NR
Kannur	NR	VLR	NR	NR	NR	NR	VHR	VHR	VLR	NR	NR	NR	NR
Kasaragod	NR	NR	NR	NR	NR	NR	NR	NR	VLR	NR	NR	NR	NR
Kollam	NR	VLR	VLR	NR	NR	NR	VHR	VHR	HR	NR	NR	NR	NR
Kottayam	NR	NR	NR	NR	NR	NR	VHR	VHR	VLR	NR	NR	NR	NR
Kozhikode	NR	NR	NR	NR	NR	NR	VHR	NR	VHR	NR	NR	NR	NR
Malappuram	NR	VLR	NR	NR	NR	NR	VHR	VHR	VLR	NR	NR	NR	NR
Palakkad	NR	VLR	NR	NR	NR	NR	VHR	HR	HR	NR	NR	VHR	NR
Pathanamthitta	VHR	VLR	NR	NR	NR	NR	VHR	VLR	VLR	NR	NR	NR	NR
Thiruvananthapuram	NR	NR	NR	NR	NR	NR	VHR	VHR	HR	NR	NR	NR	VHR
Thrissur	NR	VLR	NR	NR	NR	NR	VHR	VHR	HR	NR	NR	NR	NR
Wayanad	NR	VLR	NR	NR	NR	NR	VHR	NR	NR	NR	NR	VHR	NR

If vaccination has already been done please ignore the disease forecast for that disease.

KERALA Risk Prediction of Foot and mouth disease for the month of June 2021



KARNATAKA REPORT SEPTEMBER-2021



ProMED
INTERNATIONAL SOCIETY
FOR INFECTIOUS DISEASES

Published Date: 2021-09-30 11:40:53

Subject: PRQ/50AS> Foot & mouth disease - India (30): (Karnataka) cattle

Archive Number: 20210930.8658795

FOOT & MOUTH DISEASE - INDIA (30): KARNATAKA CATTLE

A ProMED mail post:

info@promedmailing.org

ProMED-mail is a program of the
International Society for Infectious Diseases
<http://www.isid.org>

Date: Fri 24 Sep 2021 08:33:57


Source: The Hindu (India)

<https://www.thehindu.com/news/national/karnataka/foot-and-mouth-disease-reported-in-karnataka/article2720274.ece>

District wise Livestock Disease Risk Forewarning for September 2021: Karnataka

Districts of Karnataka	Livestock Diseases												
	Anthrax	Buberculosis	BC	BT	ET	Facidosis	FMD	IT	PPK	MG Fus	ST	Difteria	Cryptosporidiosis
Bellary	NR	NR	11.8	11.8	NR	NR	11.8	11.8	NR	100	NR	NR	NR
Bidar	NR	NR	100	11.8	NR	NR	100	11.8	11.8	100	NR	NR	NR
Bijapur	100	NR	11.8	NR	NR	NR	100	100	NR	100	NR	NR	NR
Bidhar	NR	NR	100	NR	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	100	NR	11.8	11.8	NR	NR	100	11.8	11.8	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	100	NR	11.8	NR	NR	NR	100	11.8	11.8	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
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Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
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Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bijapur	NR	NR	NR	11.8	NR	NR	100	11.8	NR	NR	NR	NR	NR
Bidri													

ARUNACHAL PRADESH REPORT APRIL-2022



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 Antimicrobial Resistance

Published Date: 2021-04-14 00:02:23 IST

Subject: PRO/AH/EDR> Foot & mouth disease - India: (AR)

mithun, RFI

Archive Number: 20210413.8305677

FOOT AND MOUTH DISEASE - INDIA: (ARUNACHAL PRADESH) MITHUN, REQUEST FOR INFORMATION

A ProMED-mail post
<http://www.promedmail.org>
 ProMED-mail is a program of the
 International Society for Infectious Diseases
<http://www.isid.org>

Date: Tue 13 Apr 2021
 Source: Sentinel Assam [abridged, edited]
<http://www.sentinelassam.com/north-east-india-news/arunachal-news/many-mithuns-infected-with-fmd-foot-and-mouth-disease-533689>

A large number of mithuns have been affected by FMD (foot and mouth disease) in various parts of Arunachal Pradesh, and a few have died, said official sources on Monday (13 Apr 2021).

Animal Husbandry & Veterinary Department Deputy director Dr Taba Heli, a top mithun expert in North Eastern region, reported that the disease has taken a severe form in the entire Siang belt, particularly in East Siang, West Siang, and Upper Siang districts [map at <https://tinyurl.com/tft28s3c>].

Though the number of deaths is yet to be known, the disease has spread in

scattered areas of Papum Pare district also. The Department has allocated district funds for procurement of medicines to take all possible steps to contain the disease, he said. Mithun deaths have been reported also from Itanagar and Juliang area.

Pointing out that mithuns are the pride of indigenous people in the state, Nyishi Elite Society president Bengia Tolum has urged the department to take all possible steps to save them.

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Communicated by:
 ProMED from HealthMap Alerts
 <promed@promedmail.org>

(Mithun, also known as "Cattle of Mountain," is a bovine species of the northeastern hill region of India. This massive bovine is reared under free-range conditions in the hill forests at an altitude of 1000 to 3000 meters above mean sea level.

It is important to obtain information on the serotype and genotype of the virus strain involved, as well as of the vaccination history of the affected animals.

FMD is a disease of cloven-hoofed animals, including cattle, buffalo, small ruminants, and swine. It is the most contagious disease of mammal animals, having a great potential for causing severe economic loss in susceptible animals. Mortality, generally, is up to 2-3% but may be significantly higher in young stock.

The disease is regarded as endemic in India; effective mass vaccination, applying potent vaccines -- which include matching vaccine strains -- is essential for its control. Continued circulation of the virus may affect India's exports, such as frozen buffalo meat, of which India is the largest producer and exporter, globally. An example of the said export issue is to be found at <https://agriculture.gov.in/Documents/Report/foot%20and%20mouth%20disease.pdf>. - Mod.A5

NIVEDI prediction in April 2021

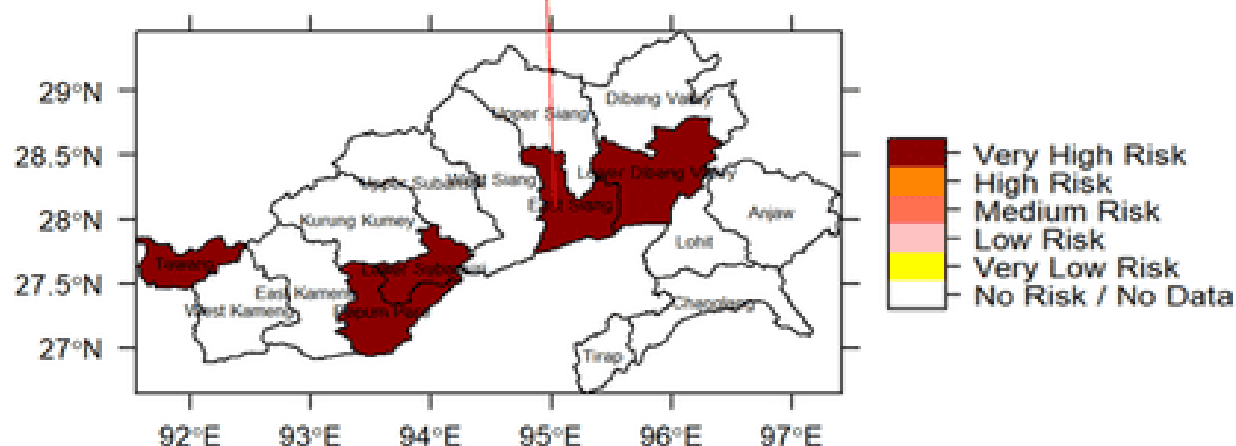
District wise Livestock Disease Risk Forewarning for April 2021: Arunachal Pradesh

Districts of Arunachal Pradesh	Livestock Diseases													
	Anthrax	Babesiosis	BQ	BT	ET	Fascioliasis	FMD	HS	PPR	S&G Pox	SF	Theileriosis	Trypanosomiasis	
Anjaw	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	
Changlang	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	
Dibang Valley	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	
East Kameng	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	
East Siang	NR	NR	NR	NR	NR	NR	VHR	NR	VLR	NR	VHR	NR	NR	
Kurung Kumey	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	
Lohit	NR	VLR	VLR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	
Lower Dibang Valley	NR	NR	VLR	NR	NR	NR	VLR	VLR	VLR	NR	VHR	NR	NR	
Lower Subansiri	NR	NR	NR	NR	NR	NR	VHR	NR	NR	NR	NR	VLR	NR	
Papum Pare	NR	NR	NR	NR	NR	NR	VHR	NR	VHR	NR	VHR	NR	NR	
Tawang	NR	NR	NR	NR	NR	NR	HR	NR	NR	NR	NR	NR	NR	
Tirap	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	
Upper Siang	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	
Upper Subansiri	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	
West Kameng	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	
West Siang	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	NR	

If vaccination has already been done please ignore the disease forecast for that disease.

*No risk/No data available (NR), Very low risk (VLR), Low risk (LR), Moderate risk (MR), High risk (HR), Very high risk (VHR)

ARUNACHAL PRADESH Risk Prediction of Foot and mouth disease for the month of April 2021



UTTARKHAND REPORT JANUARY-2024



Published Date: 2024-01-01 20:19:08 PST

Subject: PRO/SOAS> Peste des petits ruminants - India: (Uttarakhand) goat, sheep, suspected, RFI

Archive Number: 20240102.8714018

PESTE DES PETITS RUMINANTS - INDIA: (UTTARAKHAND) GOAT, SHEEP, SUSPECTED, REQUEST FOR INFORMATION

A ProMED-mail post

<http://www.promedmail.org>

ProMED-mail is a program of the
International Society for Infectious Diseases

<http://www.isid.org>

Date: Sun 31 Dec 2023

Source: The Times of India [edited]

<https://timesofindia.indiatimes.com/city/dehradun/ppr-virus-suspected-cause-of-100-sheep-and-goat-deaths-in-dehradun/articleshow/106413479.cms>

The highly contagious animal disease peste des petits ruminants (PPR), also known as sheep and goat plague, is suspected to have caused the deaths of nearly 100 goats and sheep near Kaffari in Dehradun. The shepherds were following their annual winter migration from high-altitude areas in Uttarakashi to Dehradun when they observed that some of their animals were unwell and a few died.

A team from the animal husbandry department rushed to Kaffari, Motidihar, this week and provided preliminary treatment to the affected ruminants. Sheep and goat rearing is a primary source of income for the shepherds, but with scores of ruminants affected by the virus, they would be adversely impacted, according to them.

Dr Vidyasagar Kapdi, chief veterinary officer of Dehradun, said, "We are waiting for the sample report before attributing the deaths to the PPR virus. However, based on similar symptoms to the PPR virus, treatment of other affected animals has started this week, and now the animals are fine, according to the shepherds."

The impacted sheep and goats exhibited classic symptoms of PPR, which involve fever, nasal and ocular discharge, coughing and sneezing, conjunctivitis, oral lesions, and gastrointestinal symptoms, according to animal husbandry department.

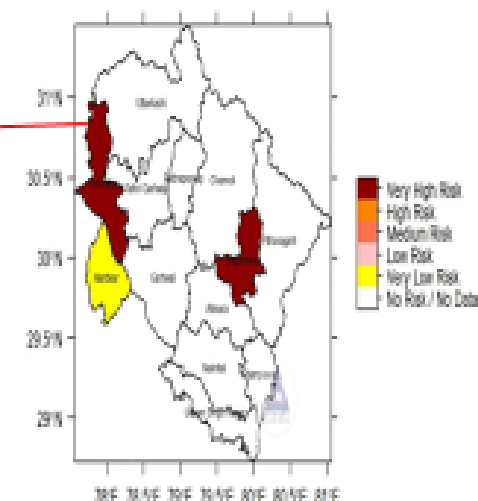
NIVEDI prediction in November 2023

District wise Livestock Disease Risk Forewarning for November 2023

Districts of Uttarakhand	Livestock Diseases													
	Anthrax	Babesiosis	BQ	BT	CSF	ET	Fasciolosis	FMD	HS	LSD	PPR	S&G Fov	Theileriosis	Trypanosomiasis
Almora	NR	NR	NR	VLR	NR	VLR	NR	NR	NR	NR	NR	NR	NR	NR
Bageshwar	NR	NR	NR	NR	NR	NR	NR	VLR	NR	NR	FHR	NR	NR	NR
Chamoli	NR	NR	NR	VLR	NR	NR	NR	VLR	NR	NR	NR	NR	NR	NR
Chamoli	NR	NR	NR	VLR	NR	VLR	NR	VLR	NR	NR	NR	NR	NR	NR
Dehradun	NR	NR	NR	VLR	NR	NR	NR	VLR	VLR	NR	FHR	NR	NR	NR
Garhwal	NR	NR	NR	VLR	NR	NR	NR	VLR	NR	NR	NR	NR	NR	NR
Haridwar	NR	NR	NR	NR	NR	NR	NR	VLR	NR	NR	VLR	NR	NR	NR
Nainital	NR	NR	NR	NR	NR	NR	NR	VLR	VLR	NR	NR	NR	NR	NR
Pithoragarh	NR	NR	NR	NR	NR	NR	NR	VLR	NR	NR	NR	NR	NR	NR
Rudrapur	NR	NR	NR	VLR	NR	VLR	NR	VLR	NR	NR	NR	NR	NR	NR
Tehri Garhwal	NR	NR	NR	NR	NR	NR	NR	VLR	NR	NR	NR	NR	NR	NR
Udham Singh Nagar	NR	NR	NR	NR	NR	NR	NR	VLR	NR	NR	NR	NR	NR	NR
Uttarakashi	NR	NR	NR	VLR	NR	NR	NR	VLR	VLR	NR	NR	NR	NR	NR

If vaccinated, please ignore the disease forecast.

*Negligible risk (NR), Very low risk (VLR), Low risk (LR), Moderate risk (MR), High risk (HR), Very high risk (FHR)



9.1 Correlational Assessment

The number of outbreaks predicted and outbreaks actually reported were mentioned in table 9.1. It has been observed from the table that outbreaks predicted two months in advance and timely alerts were issued that helped the stakeholders to take appropriate preventive measures with in time and accordingly the reported outbreaks were very less. Though the use of artificial intelligence system is more beneficial for accurately predicting the livestock disease outbreaks, there are yet number of limitations, namely, there are expected to be under reporting and also non-reporting cases which created the uncertainties in the model predictions while translating model inputs in to model outputs. However, identifying these uncertainties in the prediction using statistical models and highlighting the importance of quality data may improve the model accuracy and confidence while building the model for livestock disease forecasting.

Table 9.1: Number of districts predicted for livestock diseases risk events and reported outbreaks

Sl No	Livestock diseases	OCT-2023		NOV-2023	
		No. of districts predicted for the disease risk events	No. of districts reported the disease outbreaks*	No. of districts predicted for the disease risk events	No. of districts reported the disease outbreaks*
1	Anthrax	27	1	22	2
2	Babesiosis	85	44	85	2
3	Black Quarter (BQ)	21	1	22	2
4	Bluetongue (BT)	5	0	9	0
5	Enterotoxaemia (ET)	19	7	28	1
6	Fasciolosis	57	16	48	0
7	Foot & Mouth Disease (FMD)	76	30	80	2
8	Haemorrhagic Septicaemia (HS)	26	26	19	1
9	Peste des Petits Ruminants (PPR)	66	9	69	5
10	Sheep & Goat Pox	32	7	43	1
11	Swine Fever	34	2	38	0
12	Theileriosis	82	58	75	1
13	Trypanosomosis	70	32	66	0

*Which takes in to account of action taken for prediction and non-reporting of cases

10 A. LAUNCH OF MOBILE ANDROID APP&LINK TO DOWNLOAD

Livestock forewarning application (LDF) can be downloaded following the link provided:http://www.nivedi.res.in/android_nadres/LDF.apk and google play store link also provided <https://play.google.com/store/apps/details?id=info.androidhive.ldf>

Further launch of LDF application was done, the news provided below.



Livestock Disease Forewarning (LDF Mobile App)

The image displays three sequential screenshots of the Livestock Disease Forewarning (LDF Mobile App) interface on a mobile device.

Screenshot 1 (Home Screen): Shows the app's home screen with a blue header bar labeled "Home". Below the header, there is a "Select a Language" dropdown menu. The main content area features logos for ICAR (Indian Council of Agricultural Research) and NIVEDI (National Institute of Veterinary Epidemiology and Disease Informatics). Below the logos, there is an "About App:" section with text describing the app's purpose and the organization's mandate.

Screenshot 2 (Main Form): Shows the main form for disease forewarning. It has a blue header bar labeled "Livestock Disease Forewarning". The form contains four dropdown menus: "Select disease name" (Anthrax), "Select State name" (KARNATAKA), "Select district name" (Bagalkot), and "Select month name" (December). Below the dropdowns are two buttons: "SUBMIT" (blue) and "RESET" (orange).

Screenshot 3 (Result Pop-up): Shows a "Result" pop-up dialog box. It displays the input details: "Disease name : Anthrax", "State name : KARNATAKA", "District name : Bagalkot", and "Month: December". The result is "Result: Very Low Risk". There is an "OK" button in the bottom right corner of the dialog.

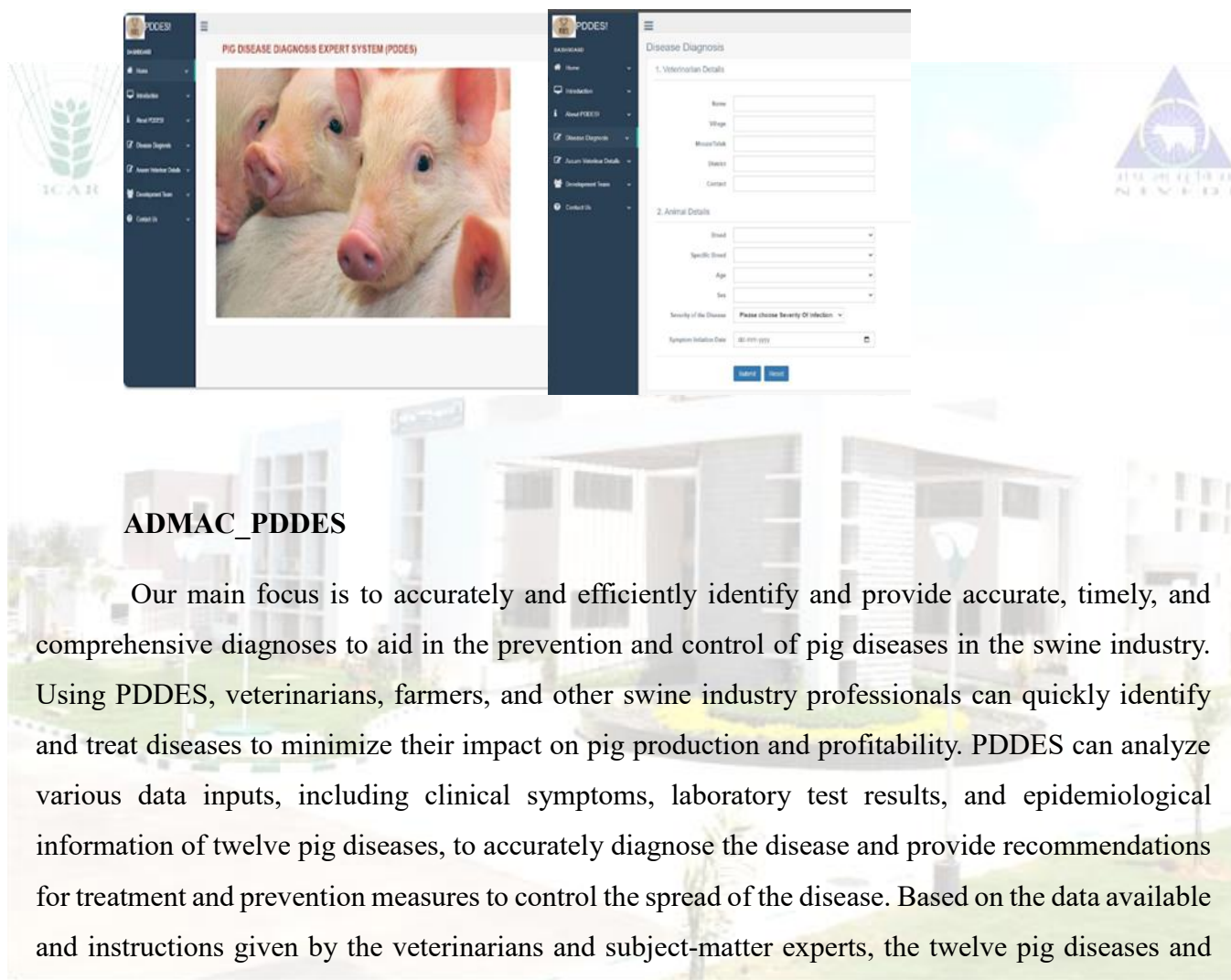
To extend the reach of the NADRES forewarning bulletin among the various stakeholders, a Mobile Application named Livestock Disease forewarning app “LDF-Mobile App” was developed. The forewarning methodology adapted in the “mobile app” remains the same as monthly bulletin; it provides user interface to know the predicted forewarning results stored in NADRES MySQL database. A PHP web-based service is developed in Java to extract the results of forewarning two months in advance by keying state name, district name and disease name and display the same in the mobile app. In addition to forewarning, the LDF-Mobile App also provides the details of clinical samples to be collected in case of outbreaks of the listed diseases for laboratory confirmation and immediate preventive measures to be taken up in case of positive prediction/disease confirmation. The LDF mobile app is available at ICAR-NIVEDI website. It is available on Google play store.

10 B. The ADMAC_PDDES (PIG DISEASE DIAGNOSIS EXPERT SYSTEM) mobile app.

The ADMAC_PDDES mobile app is available on Google play store which takes upto 3.8 MB space.

<https://play.google.com/store/apps/details?id=com.nivedi.myapplication>

Web application : <https://nivedi.res.in/PDDES/>

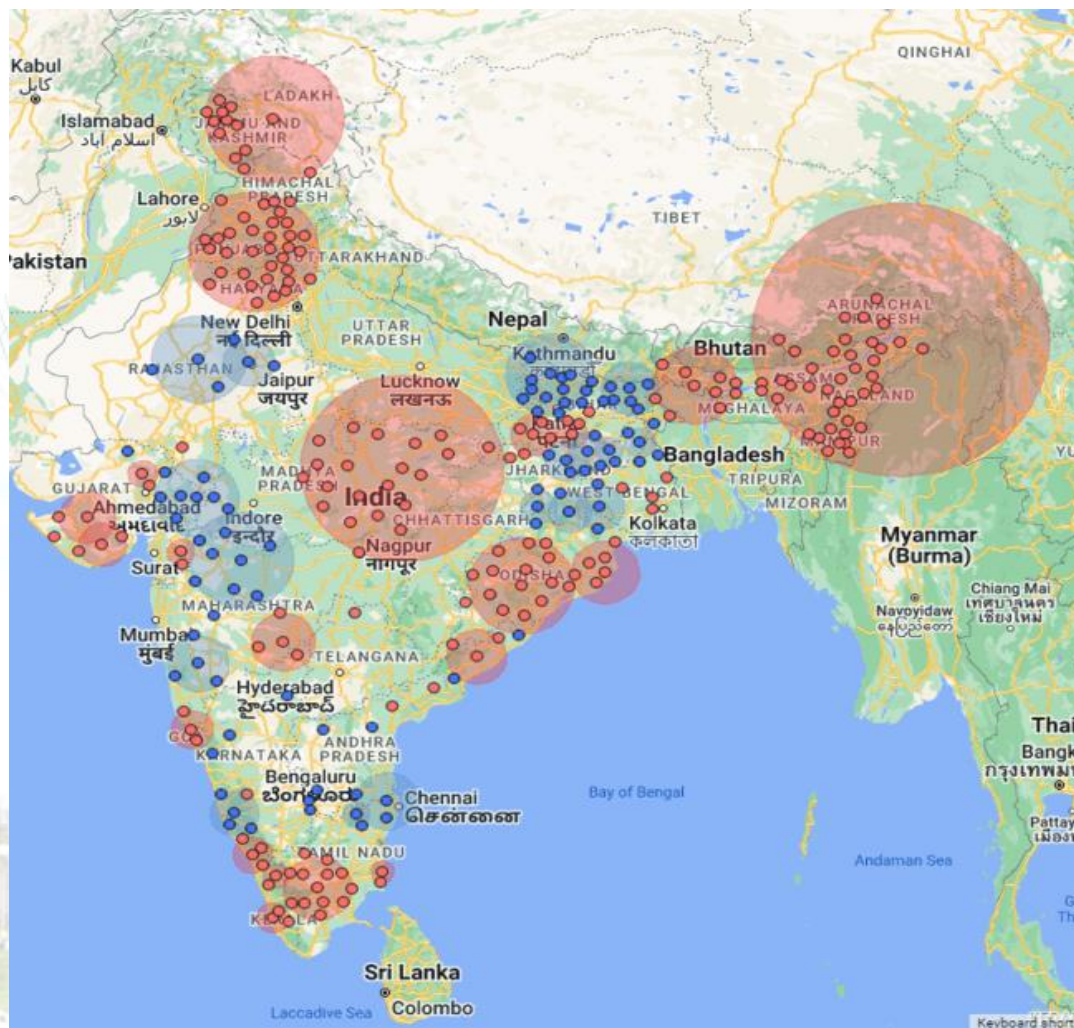


ADMAC_PDDES

Our main focus is to accurately and efficiently identify and provide accurate, timely, and comprehensive diagnoses to aid in the prevention and control of pig diseases in the swine industry. Using PDDES, veterinarians, farmers, and other swine industry professionals can quickly identify and treat diseases to minimize their impact on pig production and profitability. PDDES can analyze various data inputs, including clinical symptoms, laboratory test results, and epidemiological information of twelve pig diseases, to accurately diagnose the disease and provide recommendations for treatment and prevention measures to control the spread of the disease. Based on the data available and instructions given by the veterinarians and subject-matter experts, the twelve pig diseases and their clinical indications or symptoms were determined and Aiken's value index was calculated. The weighted matrix was created for a number of different clinical indicators or symptoms related to swine illnesses. Based on this, an expert system was developed as a mobile application.

11.1. MODEL BASED HOT-SPOT AREA MAPS OF LIVESTOCK INFECTIOUS DISEASES IN INDIA

a) FMD Hotspot by Poisson Model: FMD in India (1989-2022)



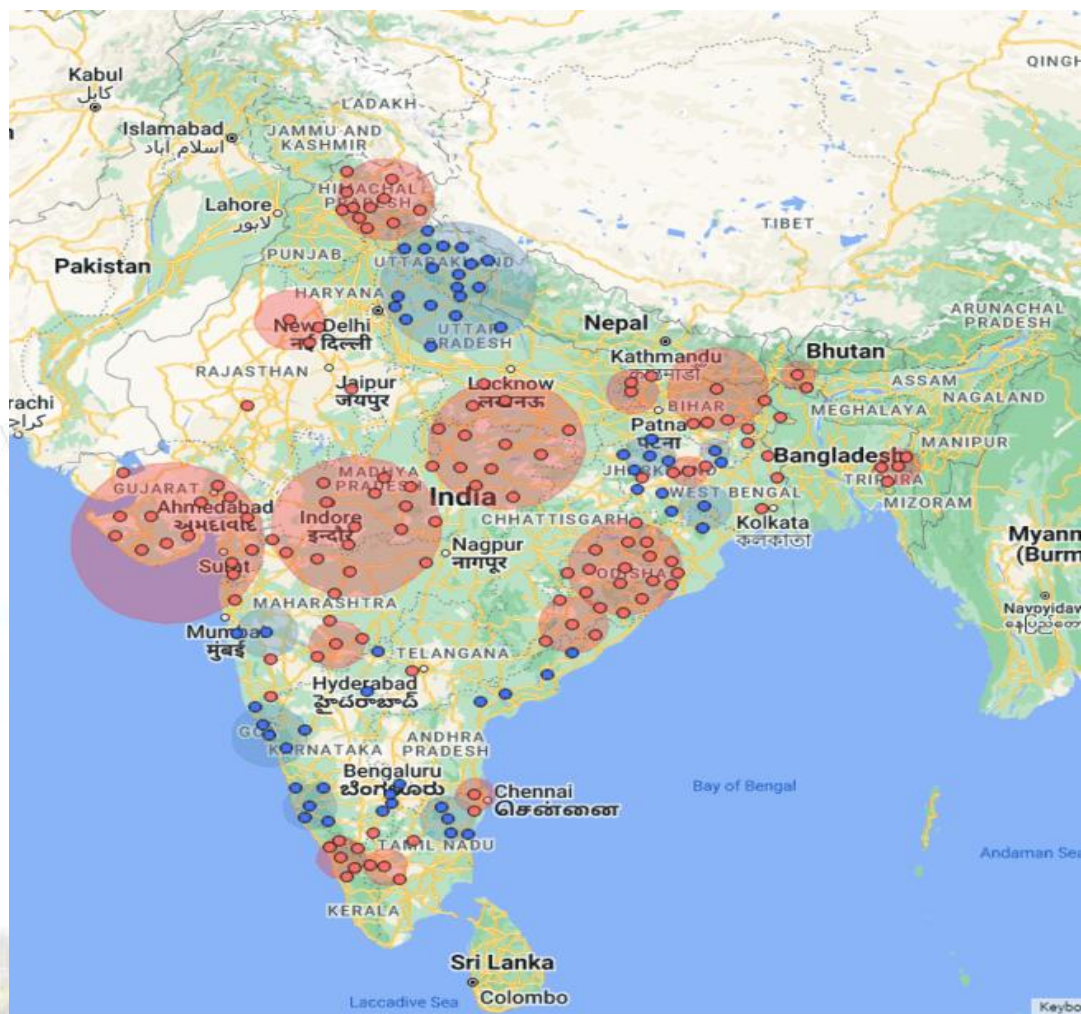
Model Specifications

- **Retrospective Analysis: Space-Time**
- **Probability Model: Space-Time Poisson Model**
 - With the Space-Time Poisson model, the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
 - Circle radius:** 1 km.
 - **Time Aggregation:** 1 year.
 - **Output obtained with no geographical overlap-** Total hotspots–69 (High- 36, Low- 33).
 - The output is generated using SaTScan v9. 6.

Note: Red Dot: High Disease Cluster

Blue Dot: Low Disease Cluster

b) PPR Hotspot by Poisson Model: PPR in India (1995-2022)



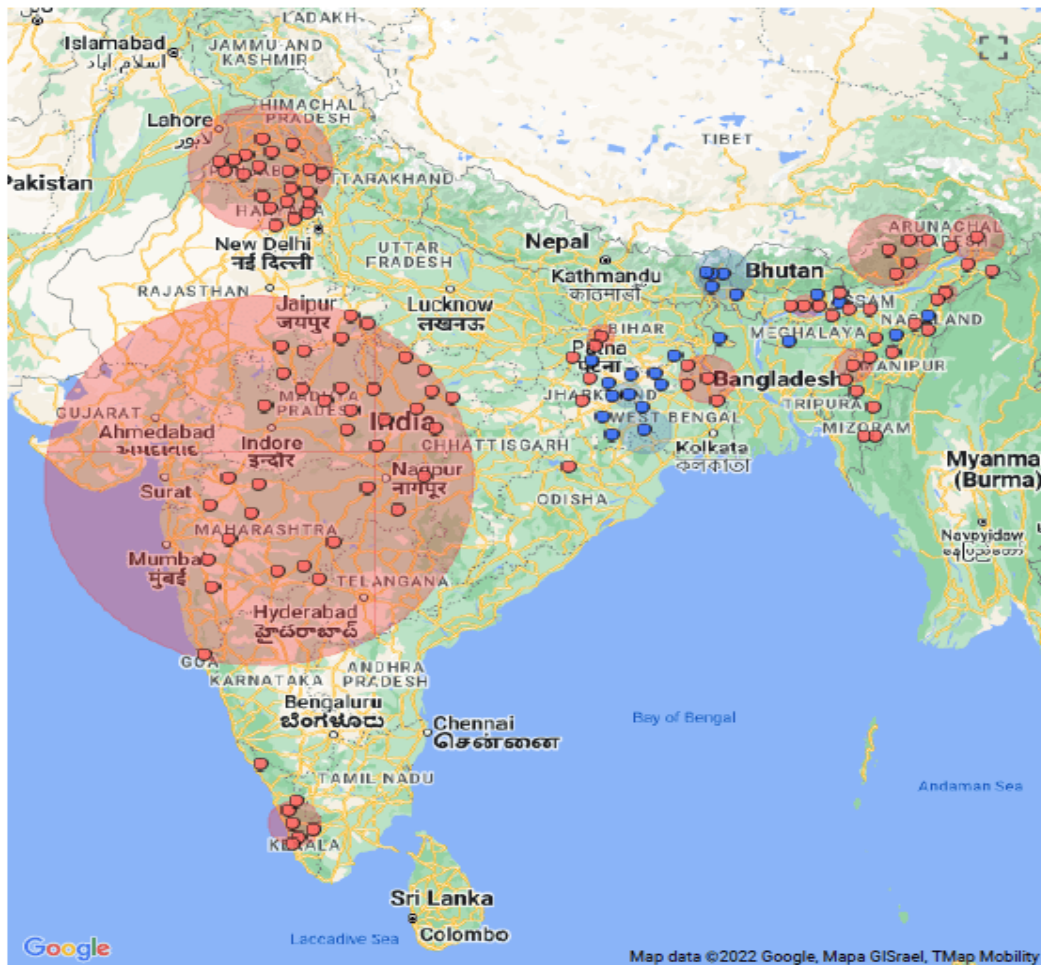
Model Specifications

- **Retrospective Analysis: Space-Time**
- **Probability Model: Space-Time Poisson Model**
 - With the Space-Time Poisson model, the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
- **Circle radius:** 1 km.
- **Time Aggregation:** 1 year.
- **Output obtained with no geographical overlap-** Total hotspots– 51 (High- 30, Low- 21).
- The output is generated using **SaTScan v9. 6.**

Note: Red Dot: High Disease Cluster

Blue Dot: Low Disease Cluster

c) CSF Hotspot by Poisson Model: CSF in India (1992-2022)



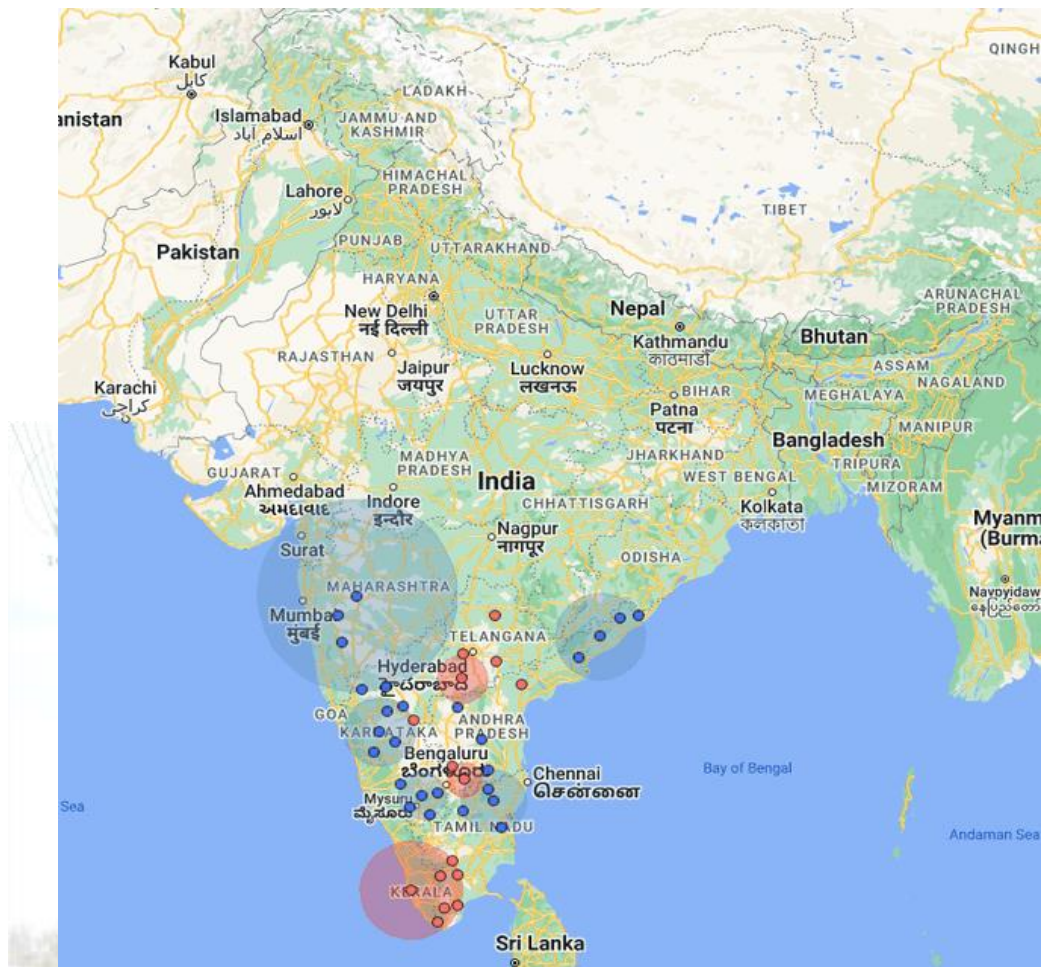
Model Specifications

- **Retrospective Analysis: Space-Time**
- **Probability Model: Space-Time Poisson Model**
 - With the Space-Time Poisson model, the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
 - Circle radius:** 1 km.
 - Time Aggregation:** 1 year.
 - Output obtained with no geographical overlap-** Total hotspots– 46 (High- 29, Low- 17).
 - The output is generated using **SaTScan v9. 6.**

Note: Red Dot: High Disease Cluster

Blue Dot: Low Disease Cluster

d) Bluetongue Hotspot by Poisson Model: Bluetongue in India (2001-2022)



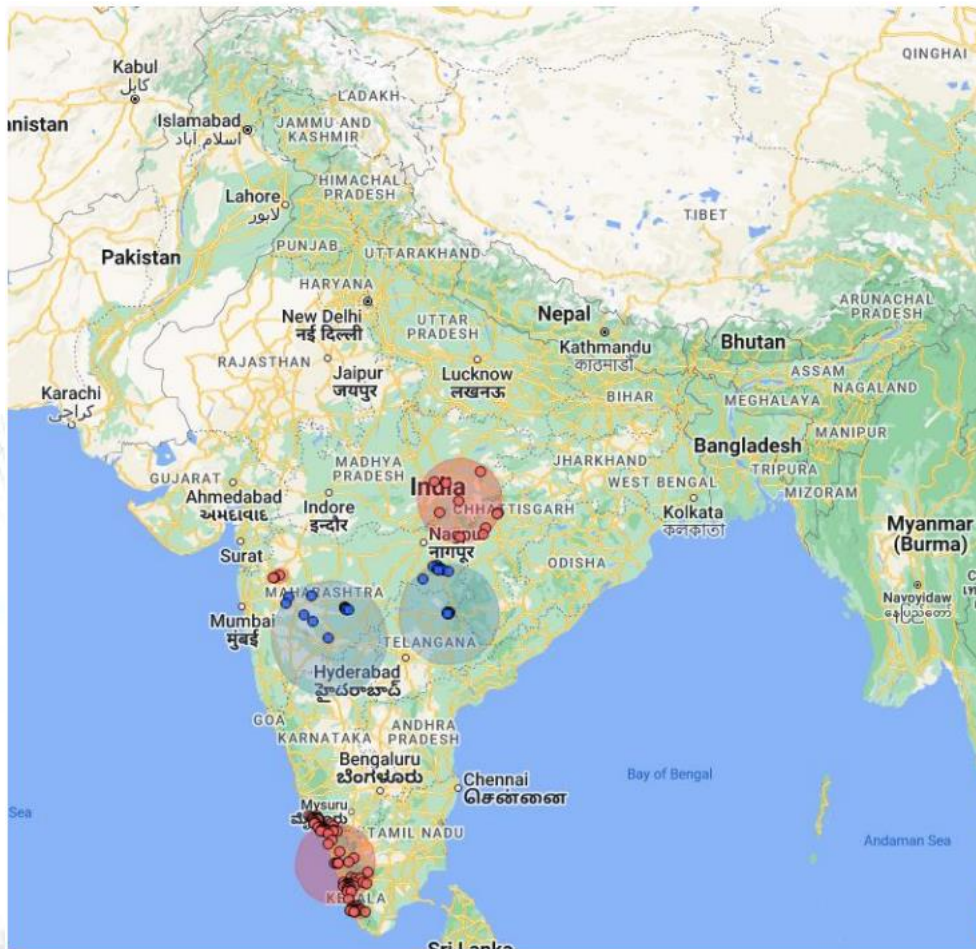
Model Specifications

- **Retrospective Analysis: Space-Time**
- **Probability Model: Space-Time Poisson Model**
 - With the Space-Time Poisson model, the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
 - Circle radius:** 1 km.
 - Time Aggregation:** 1 year.
 - Output obtained with no geographical overlap-** Total hotspots– 14 (High- 7, Low- 7).
 - The output is generated using SaTScan v9. 6.**

Note: Red Dot: High Disease Cluster

Blue Dot: Low Disease Cluster

e) LSD Hotspot by Poisson Model: LSD in India (2019-2022)



Model Specifications

- **Retrospective Analysis: Space-Time**
- **Probability Model: Space-Time Poisson Model**
 - With the Space-Time Poisson model, the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
- **Circle radius: 1 km.**
- **Time Aggregation: 1 year.**
- **Output obtained with no geographical overlap- Total hotspots– 7 (High- 5, Low- 2).**
- **The output is generated using SaTScan v9. 6.**

Note: Red Dot: High Disease Cluster

Blue Dot: Low Disease Cluster

f) Anthrax Hotspot by Poisson Model – Anthrax in India (1989-2022)



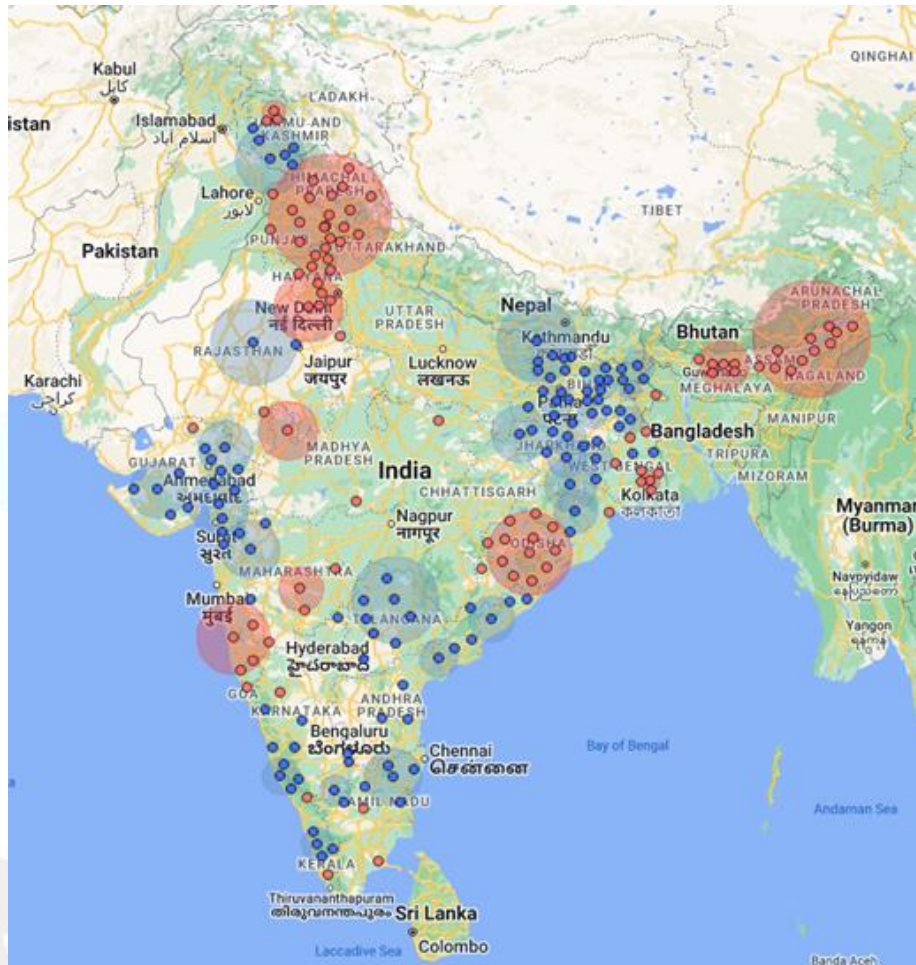
Model Specifications

- **Retrospective Analysis: Space-Time**
- **Probability Model: Space-Time Poisson Model**
 - With the Space-Time Poisson model, the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
- **Circle radius: 1 km.**
- **Time Aggregation: 1 year.**
- The output is generated using **SaTScan v9. 6.**

Note: Red Dot: High Disease Cluster

Blue Dot: Low Disease Cluster

11.2. a) FMD Hotspot by Poisson Model – FMD in India (2000-2004)



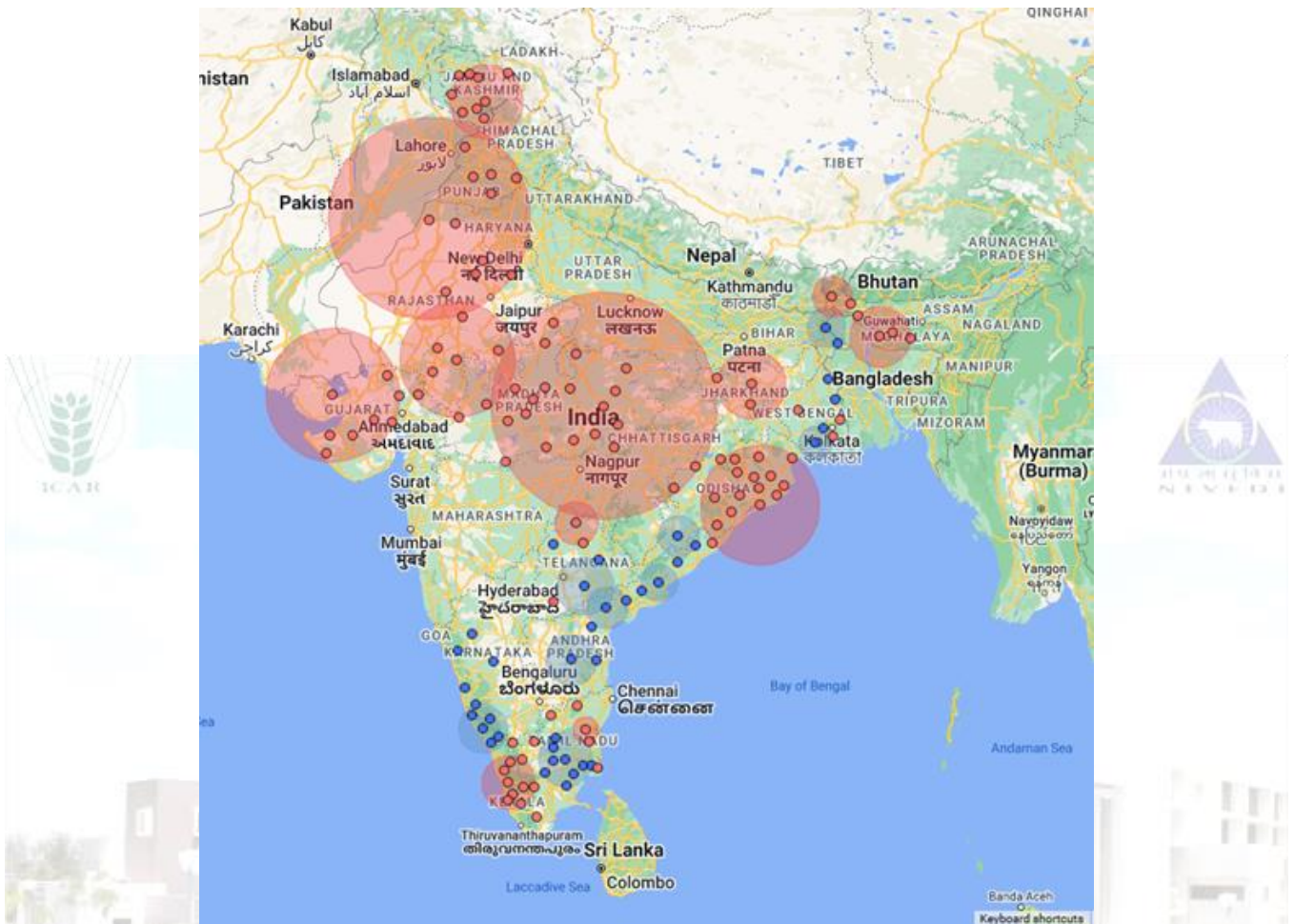
Model Specifications

- **Retrospective Analysis: Space-Time**
- **Probability Model: Space-Time Poisson Model**
 - **With the Space-Time Poisson model**, the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
 - Circle radius:** 1 km.
 - Time Aggregation:** 1 year.
 - Output obtained with no geographical overlap-** Total hotspots– Hotspot: 1 km, 72 clusters, 29 high, 43 low
 - The output is generated using **SaTScan v9. 6.**

Note: Red Dot: High Disease Cluster

Blue Dot: Low Disease Cluster

b) FMD Hotspot by Poisson Model – FMD in India (2005-2009)



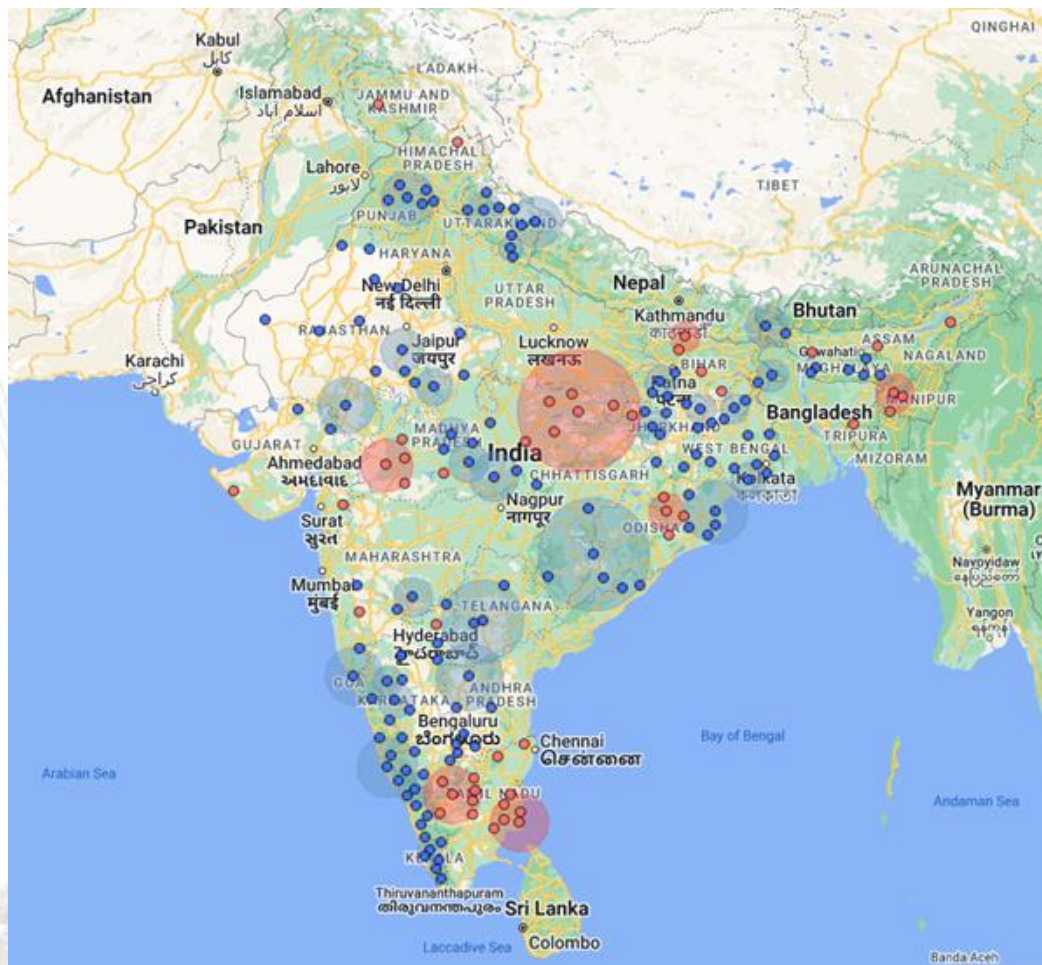
Model Specifications

- **Retrospective Analysis:** Space-Time
- **Probability Model:** Space-Time Poisson Model
 - With the Space-Time Poisson model, the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
 - Circle radius:** 1 km.
 - Time Aggregation:** 1 year.
 - Output obtained with no geographical overlap-** Total hotspots– Hotspot: 1 km, 41 clusters, 23 high, 18 low
 - The output is generated using SaTScan v9. 6.

Note: Red Dot: High Disease Cluster

Blue Dot: Low Disease Cluster

C) FMD Hotspot by Poisson Model – FMD in India (2010-2014)



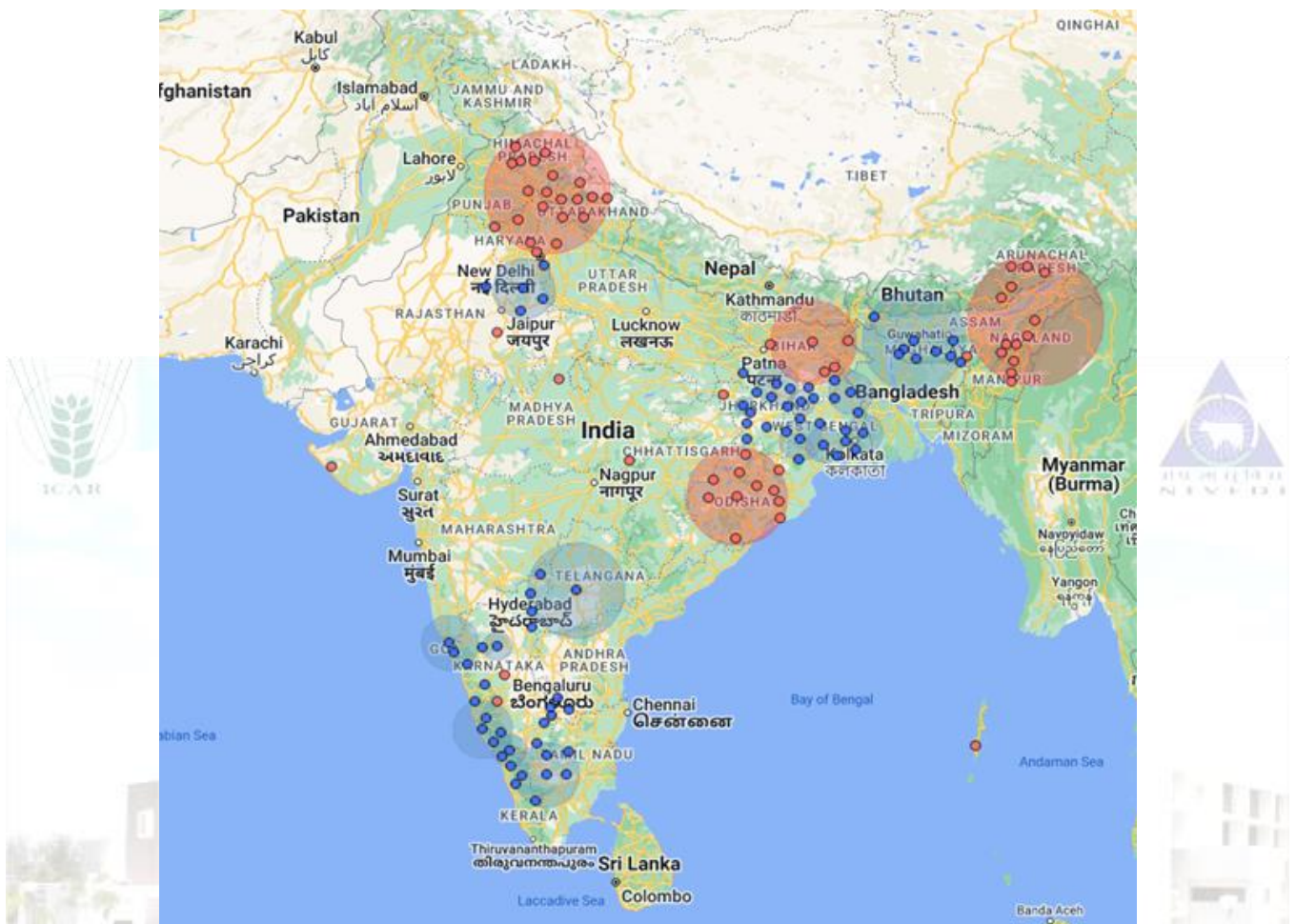
Model Specifications

- **Retrospective Analysis:** Space-Time
- **Probability Model:** Space-Time Poisson Model
 - **With the Space-Time Poisson model**, the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
 - Circle radius:** 1 km.
 - **Time Aggregation:** 1 year.
 - **Output obtained with no geographical overlap-** Total hotspots– Hotspot: 1 km, 103 clusters, 24 high, 79 low
 - The output is generated using **SaTScan v9. 6**.

Note: Red Dot: High Disease Cluster

Blue Dot: Low Disease Cluster

D)FMD Hotspot by Poisson Model – FMD in India (2015-2019)



Model Specifications

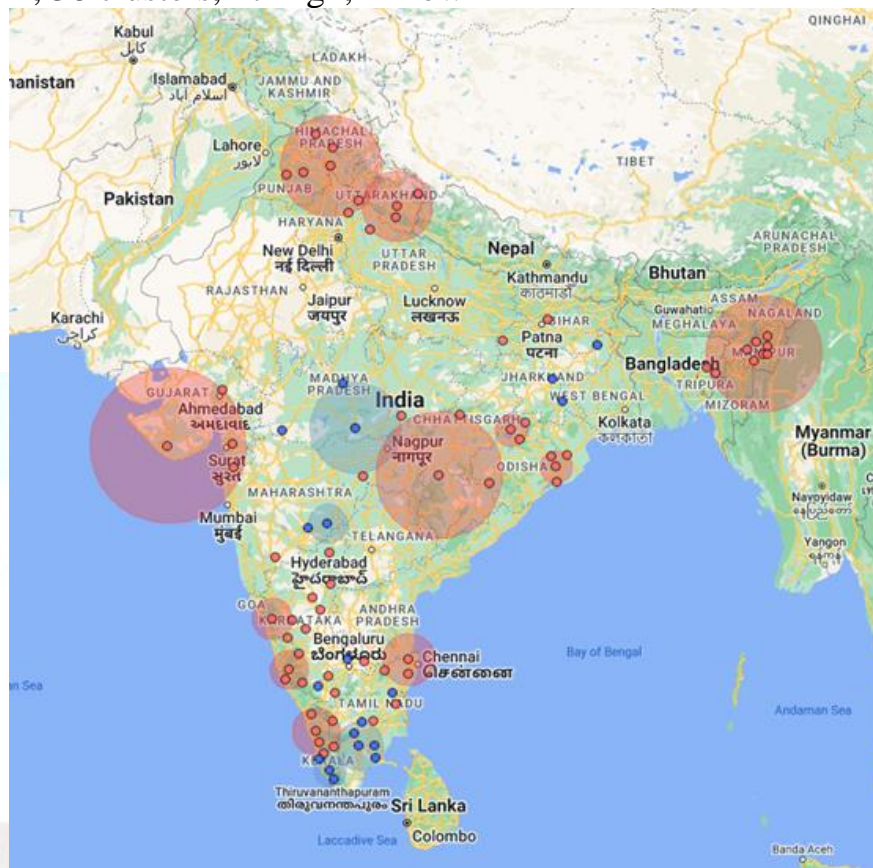
- **Retrospective Analysis:** Space-Time
- **Probability Model:** Space-Time Poisson Model
 - **With the Space-Time Poisson model,** the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
 - Circle radius:** 2 km.
 - Time Aggregation:** 1 year.
 - Output obtained with no geographical overlap-** Total Hotspots-Hotspot: Hotspot: 2 km, 43 clusters, 13 high, 30 low
 - The output is generated using **SaTScan v9. 6.**

Note: Red Dot: High Disease Cluster

Blue Dot: Low Disease Cluster

E)FMD Hotspot by Poisson Model – FMD in India (2020-2023)

Hotspot: 1 km, 38 clusters, 27 high, 11 low



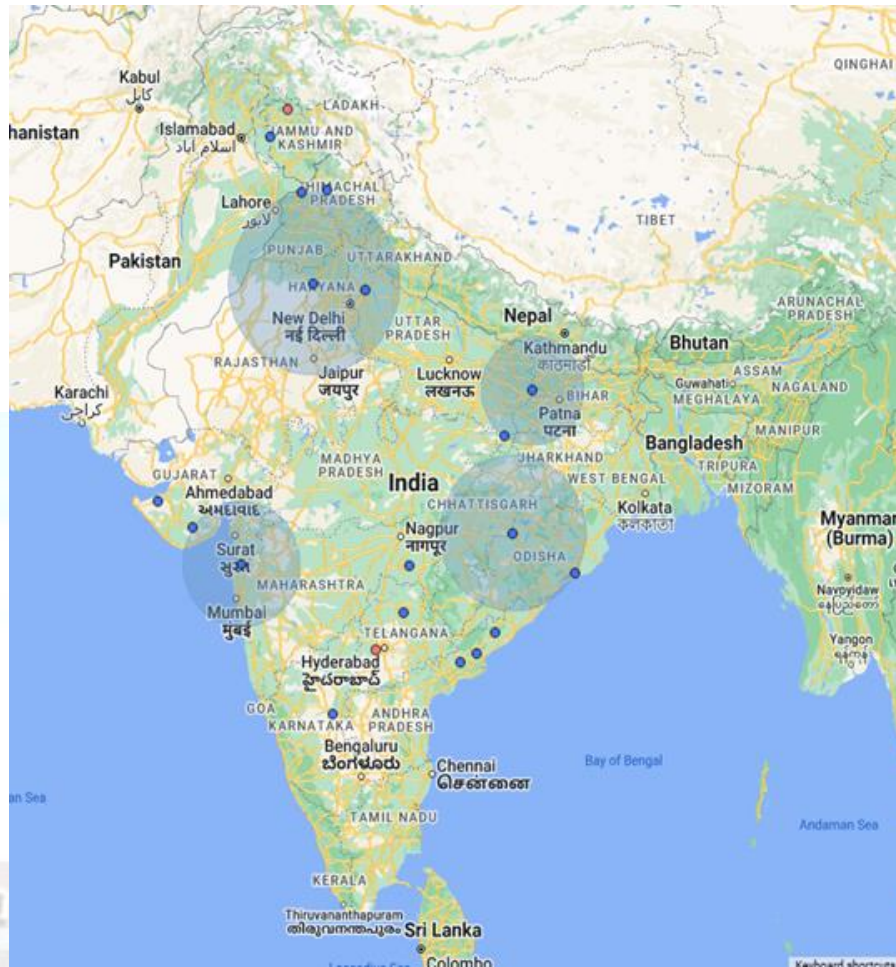
Model Specifications

- **Retrospective Analysis:** Space-Time
- **Probability Model:** Space-Time Poisson Model
 - **With the Space-Time Poisson model**, the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
 - Circle radius:** 1 km.
 - Time Aggregation:** 1 year.
 - Output obtained with no geographical overlap-** Total Hotspots-Hotspot: 1 km, 38 clusters, 27 high, 11 low
 - The output is generated using **SaTScan v9. 6**.

Note: Red Dot: High Disease Cluster

Blue Dot: Low Disease Cluster

11.3. Foot rot district wise Risk map (2001-2019)



Model Specifications

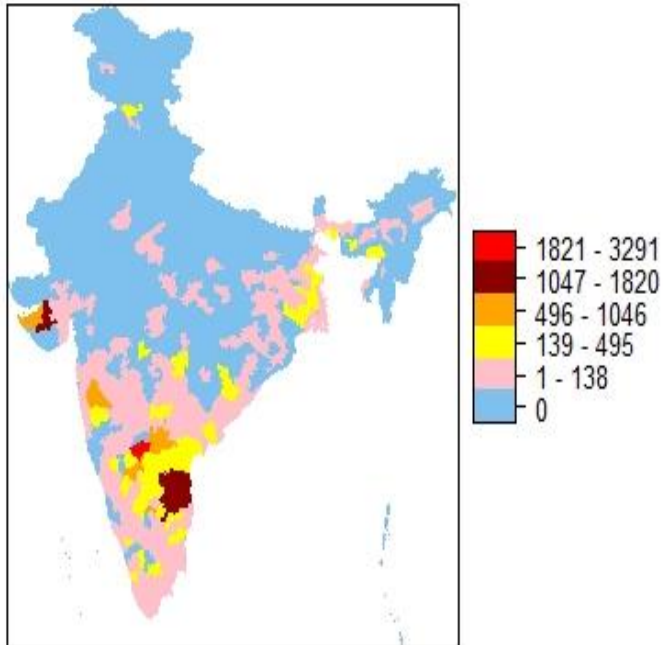
- **Retrospective Analysis:** Space-Time
- **Probability Model:** Space-Time Poisson Model
 - With the Space-Time Poisson model, the number of cases in each location is Poisson-distributed.
 - The model requires case and population counts for a set of data locations and geographical coordinates for each of the locations.
 - SaTScan does a linear interpolation based on the population at the census times immediately preceding and following.
- **Advance Analysis Features:**
 - Circle radius:** 1 km.
 - Time Aggregation:** 1 year.
 - Output obtained with no geographical overlap-** Total Hotspots-Hotspot: Hotspot: 1 km, 15 clusters, 2 high, 13 low
 - The output is generated using SaTScanv9. 6.

Note: Red Dot: High Disease Cluster

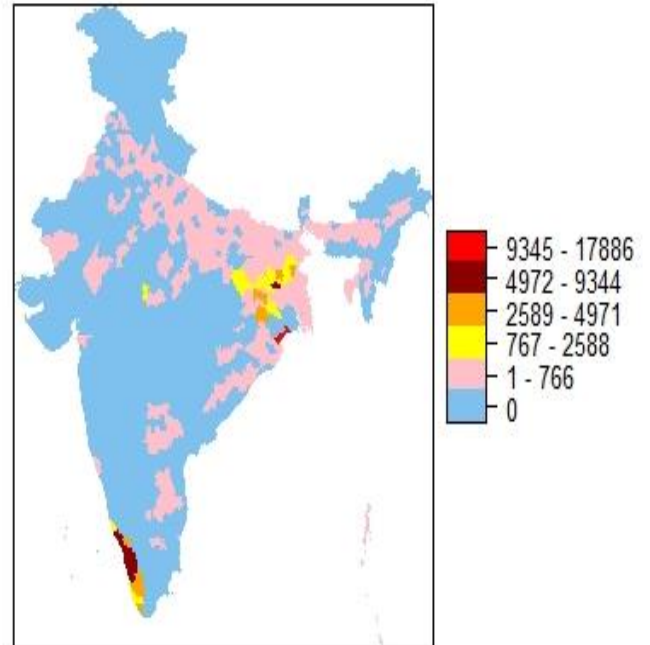
Blue Dot: Low Disease Cluster

11.4. MODEL BASED OUTBREAK/INCIDENCE MAPS OF LIVESTOCK DISEASE IN INDIA

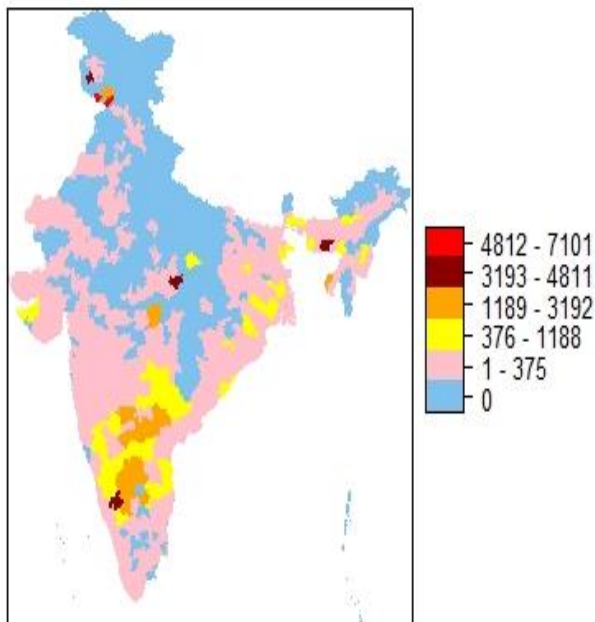
Outbreaks/Incidence- Anthrax(1990-2022)



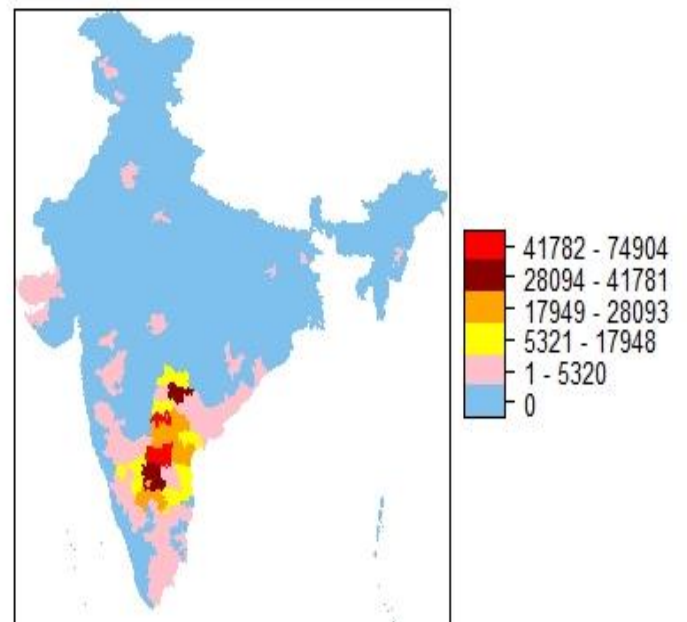
Outbreaks/Incidence- Babesiosis(1990-2022)



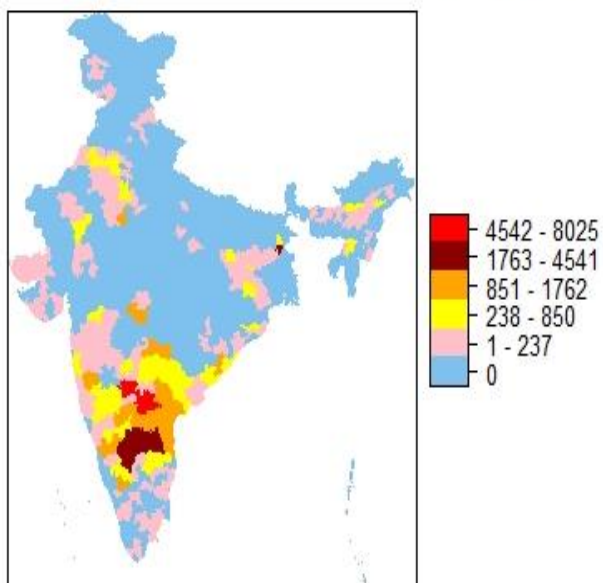
Outbreaks/Incidence- Blackquarter(1990-2022)



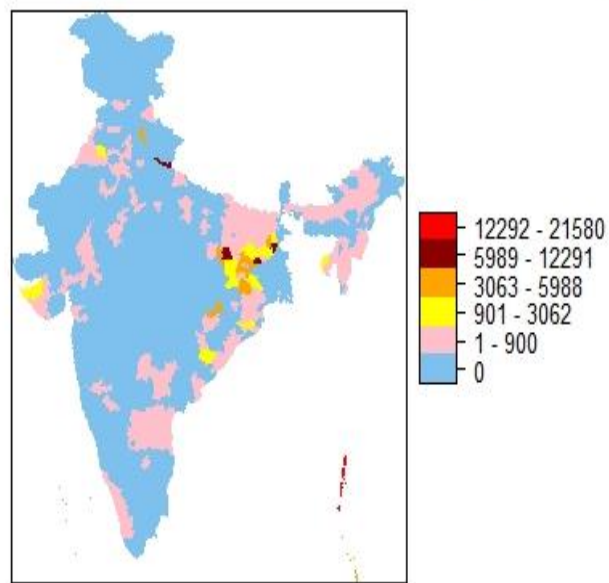
Outbreaks/Incidence- Bluetongue(1990-2022)



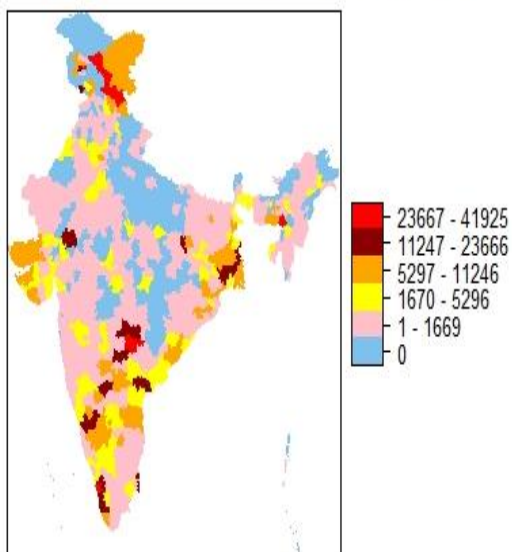
Outbreaks/Incidence- Enterotoxaemia(1990-2022)



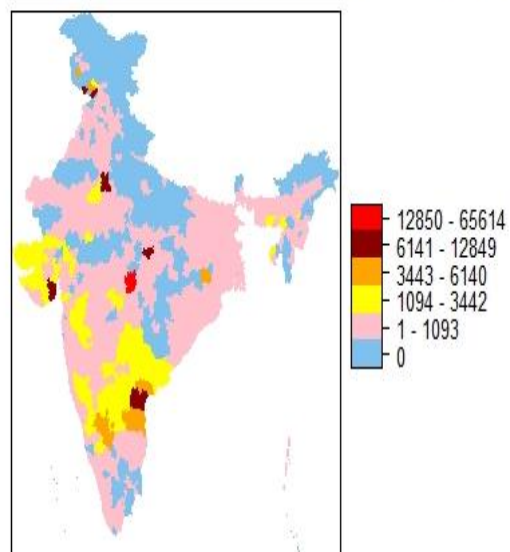
Outbreaks/Incidence- Fascioliasis(1990-2022)



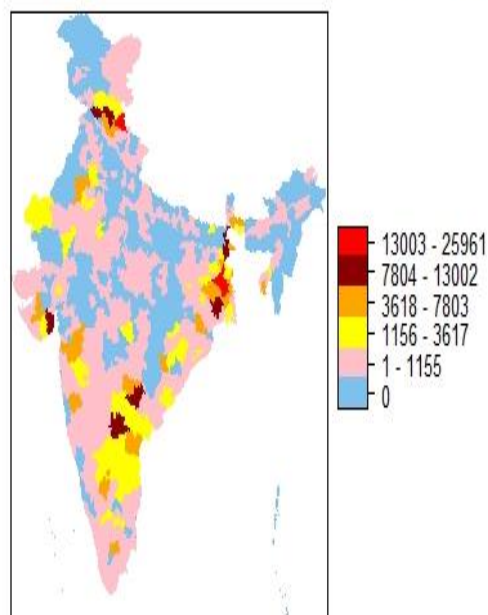
Outbreaks/Incidence-Foot and Mouth Disease(1990-2022)



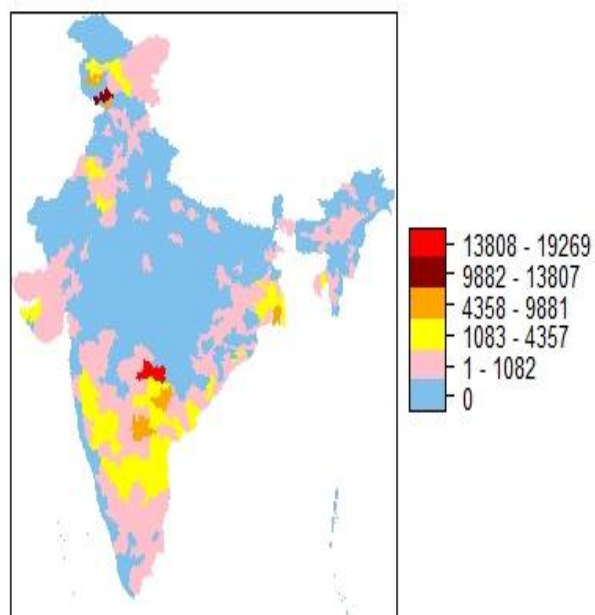
Outbreaks/Incidence-Haemorrhagic Septicaemia(1990-2022)



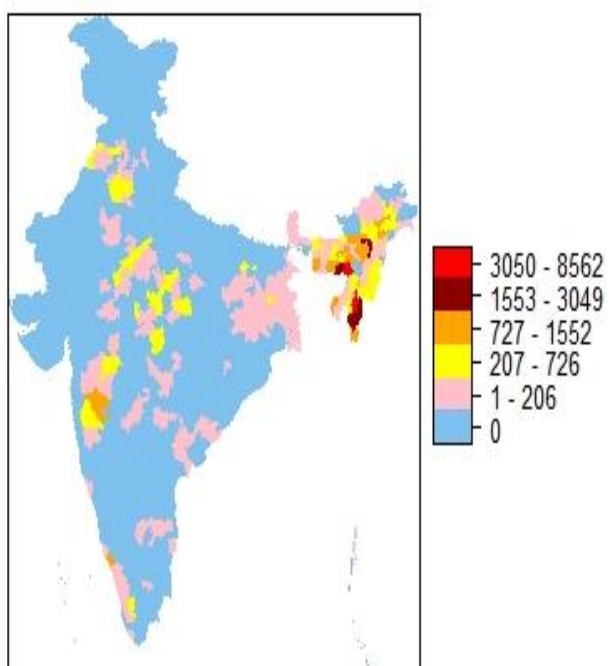
Outbreaks/Incidence-Peste des petits Ruminants(1990-2022)



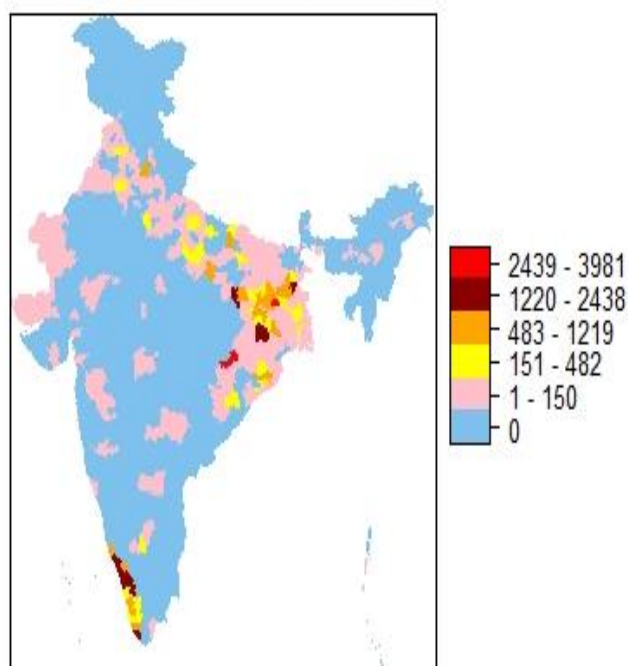
Outbreaks/Incidence-Sheep & Goat Pox(1990-2022)



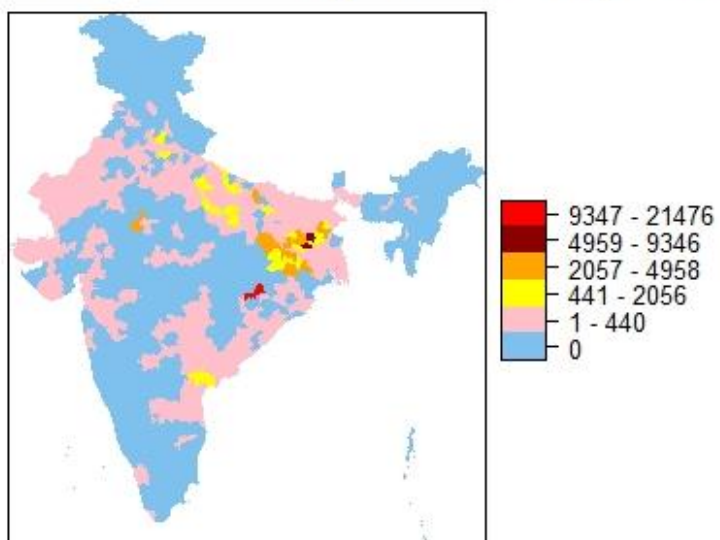
Outbreaks/Incidence-Swine Fever(1990-2022)



Outbreaks/Incidence-Theileriosis(1990-2022)

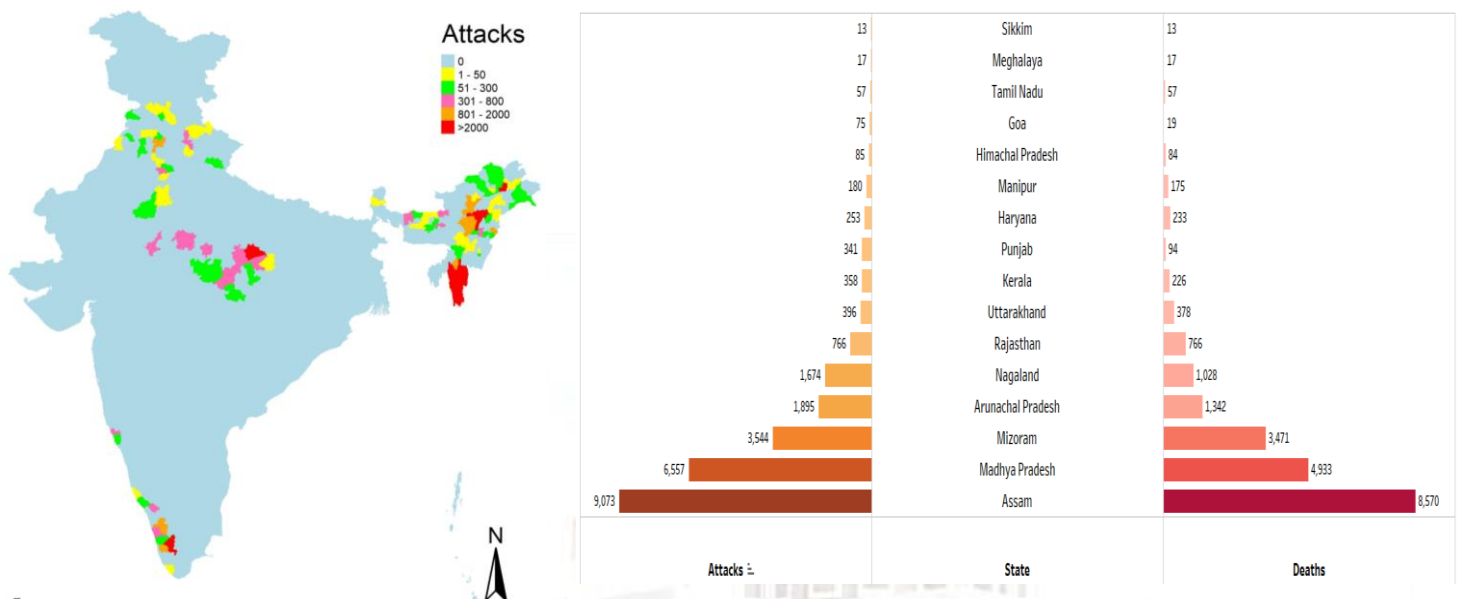


Outbreaks/Incidence-Trypanosomiasis(1990-2022)



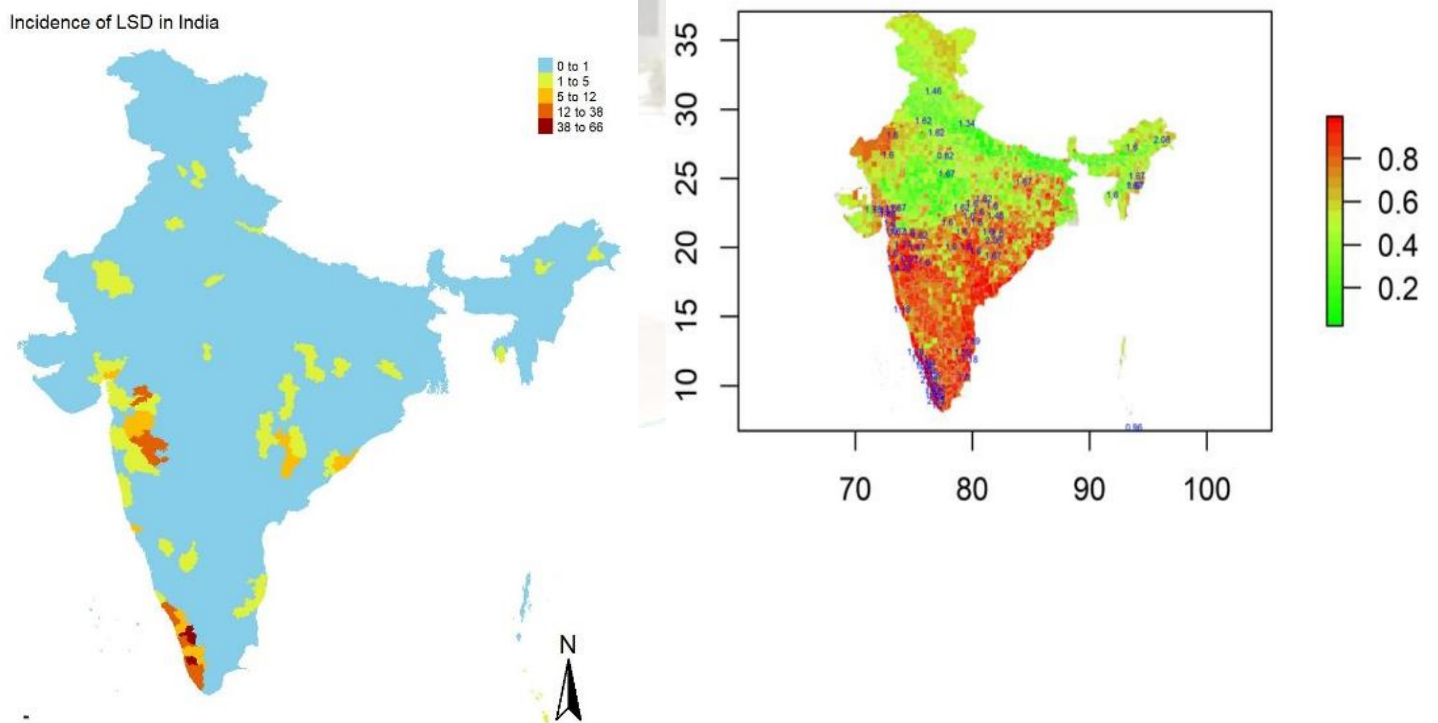
11.5. MODEL BASED RISK MAPS OF LIVESTOCK INFECTIOUS DISEASES IN INDIA

Disease map considering attacks of ASFV in India

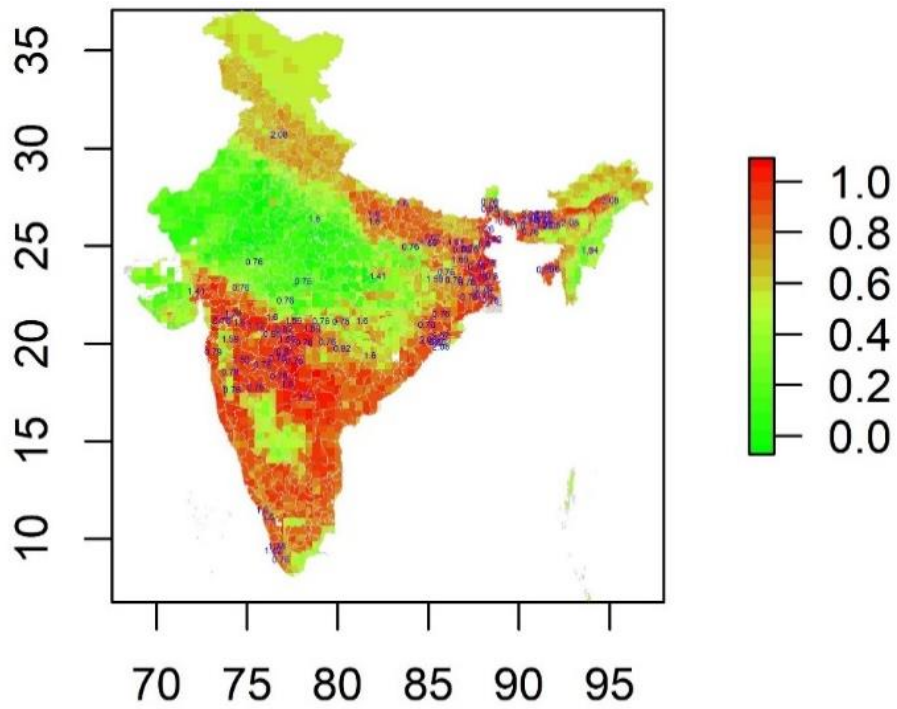


Incidence of LSD in India

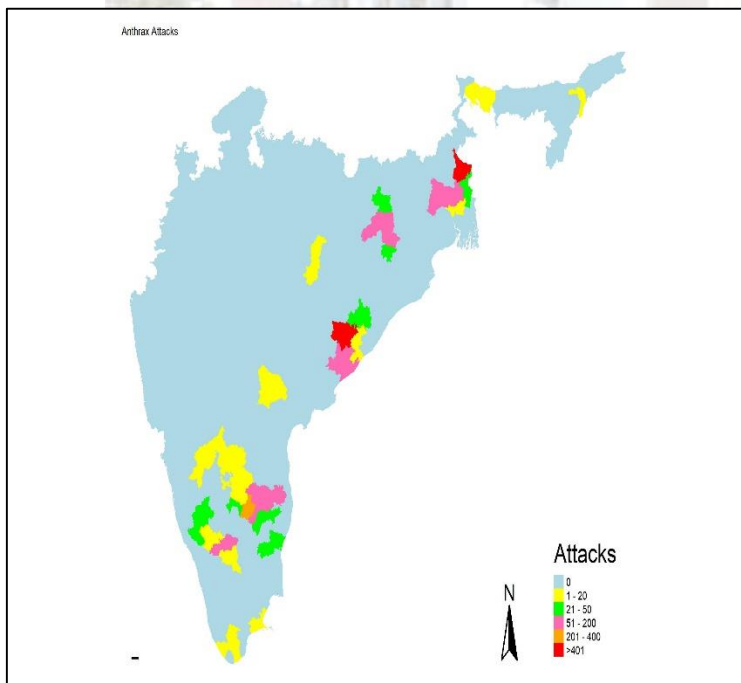
Risk map for LSD in India



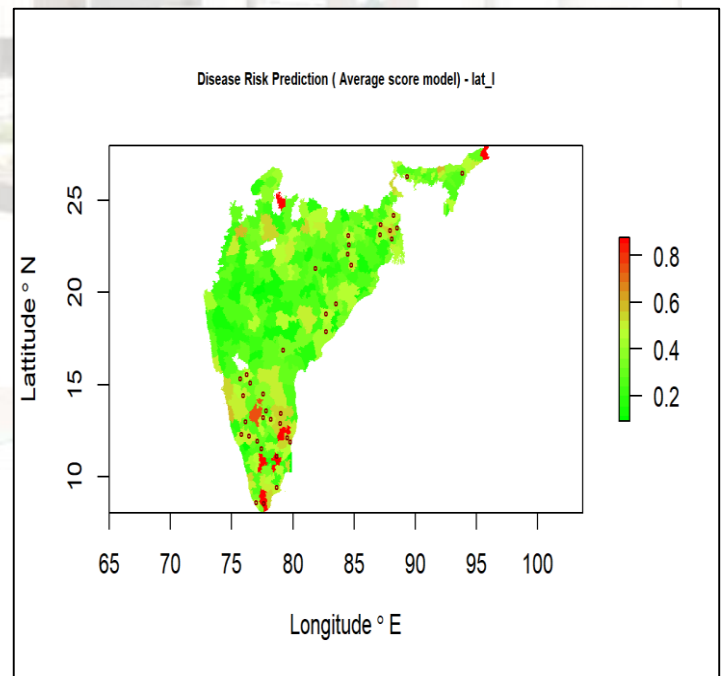
Risk map for Avian Influenza in India



Disease Map for Anthrax on Attacks

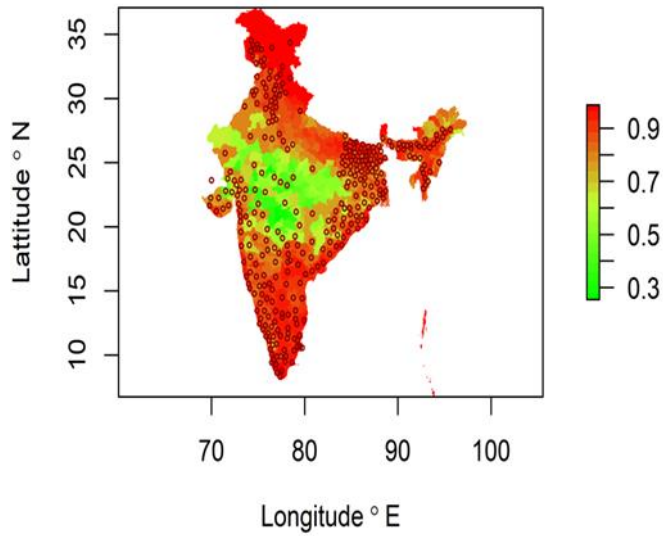


Risk map for Anthrax in India

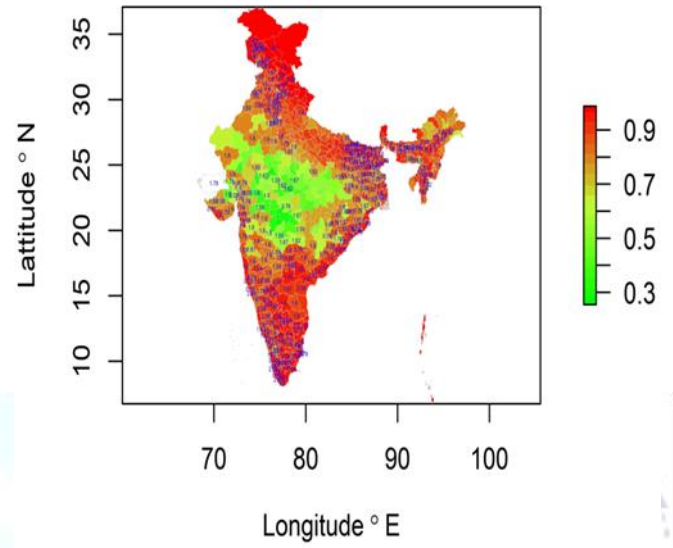


FMD district wise Risk map (2000-2004)

Without R0

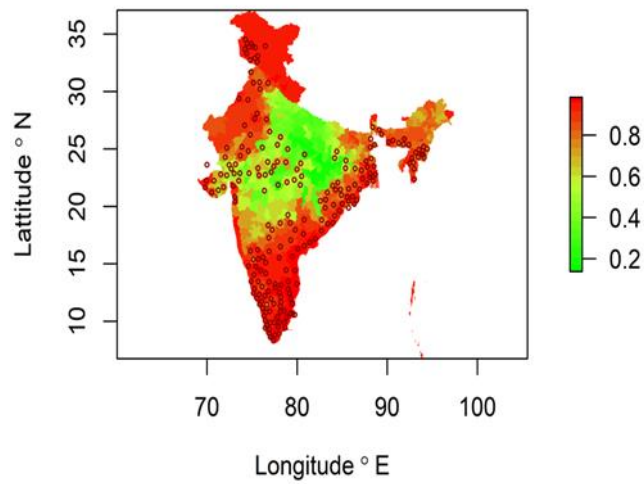


With R0

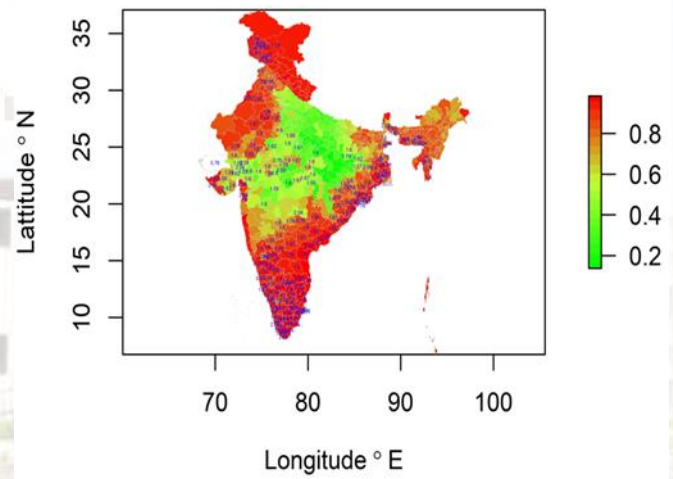


FMD district wise Risk map (2005-2009)

Without R0

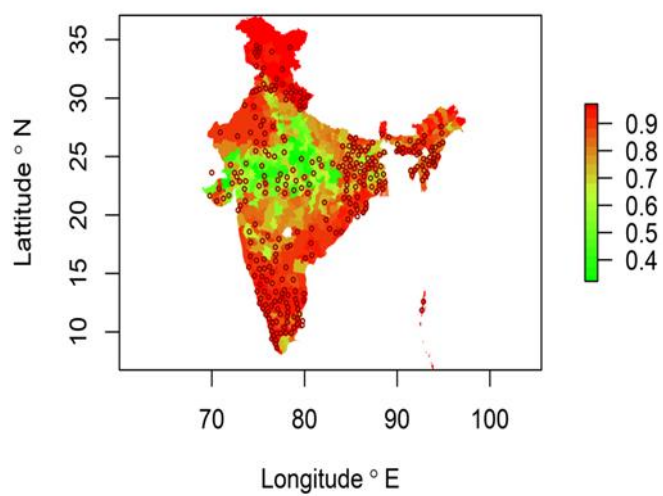


With R0

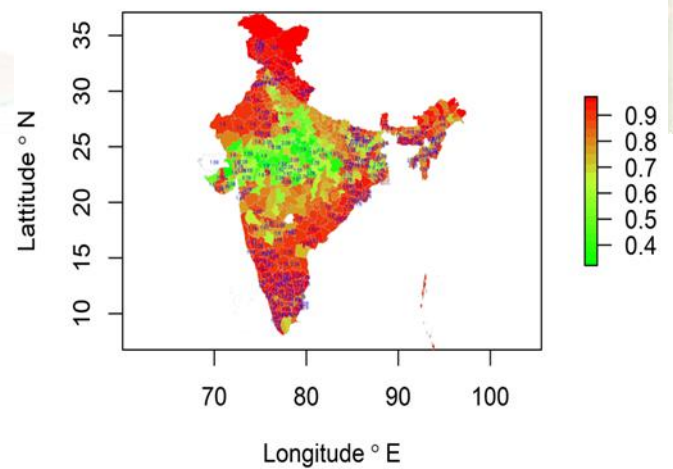


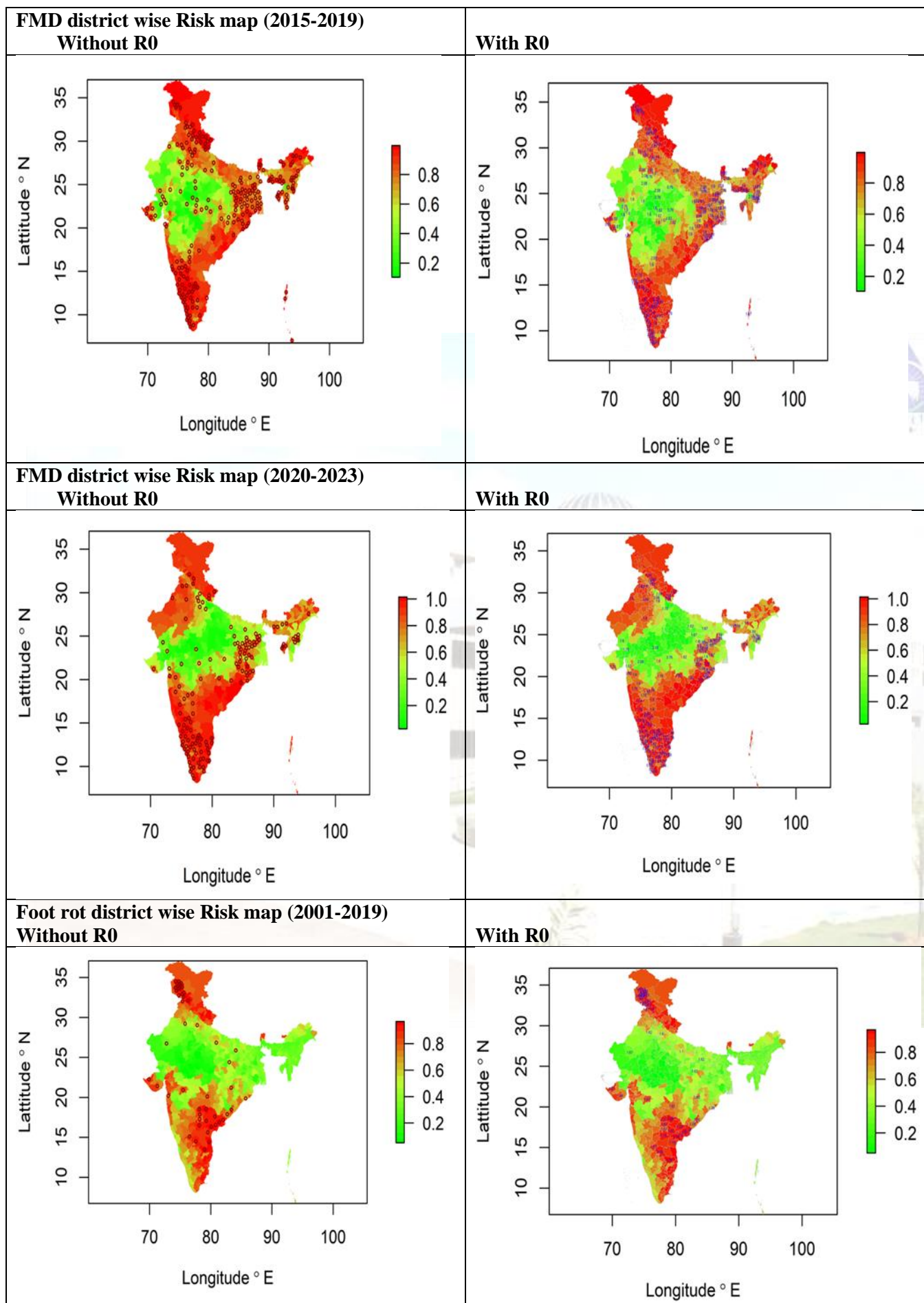
FMD district wise Risk map (2010-2014)

Without R0

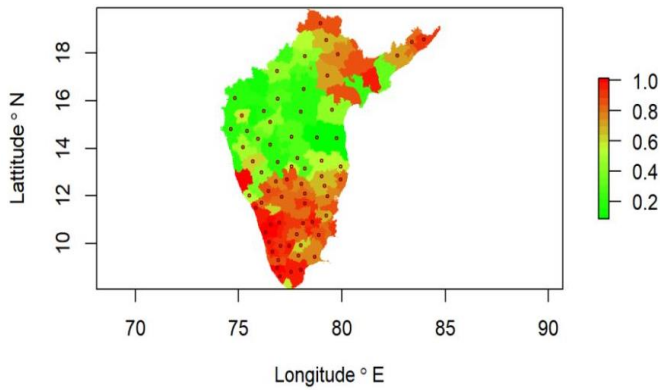


With R0

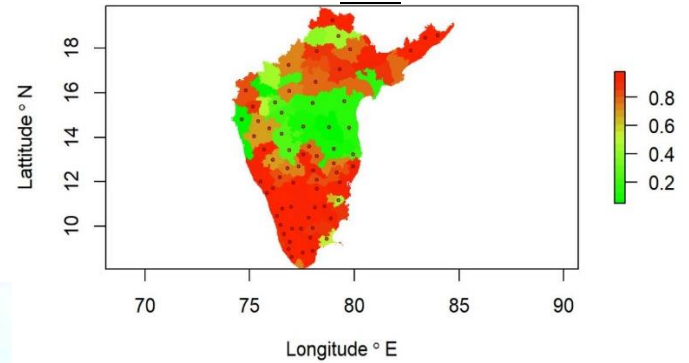




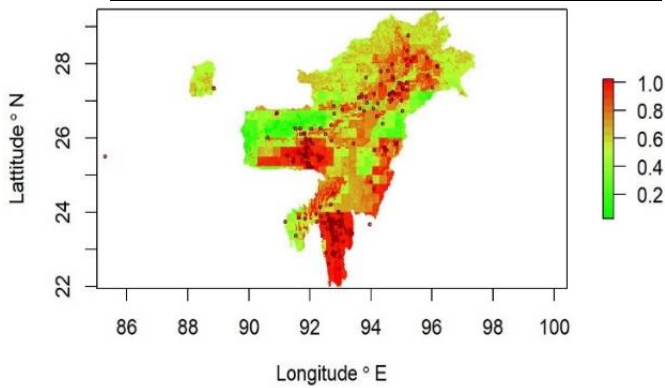
Risk prediction for Anthrax Disease for Southern India



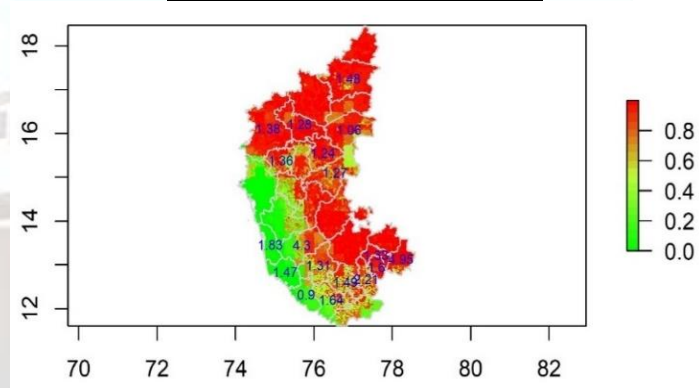
Risk prediction for Enterotoxaemia Disease for Southern India



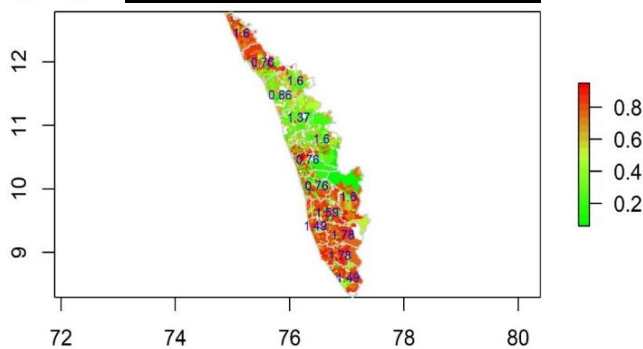
Risk prediction for ASF for North Eastern India



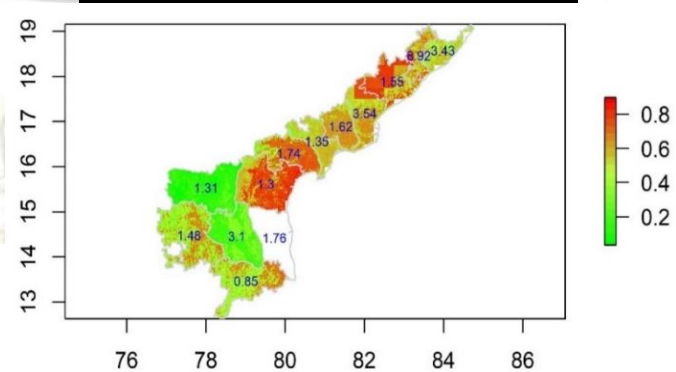
Karnataka Risk map Bluetongue



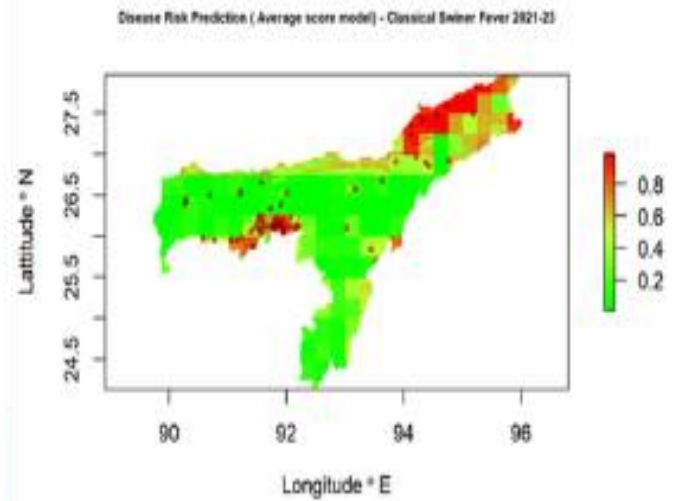
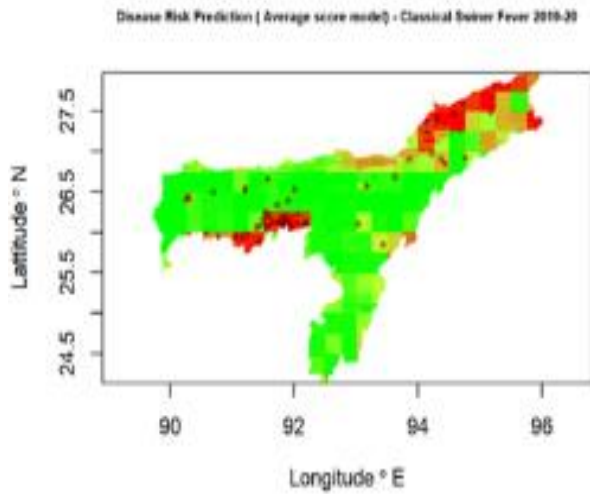
Risk map of Anaplasmosis in Kerala



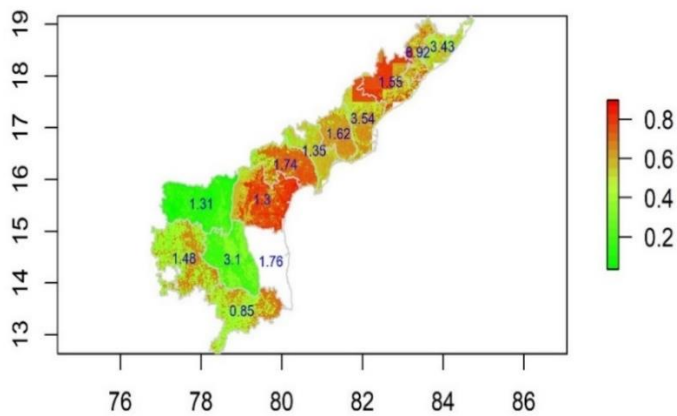
Risk map of Bluetongue in Andhra Pradesh



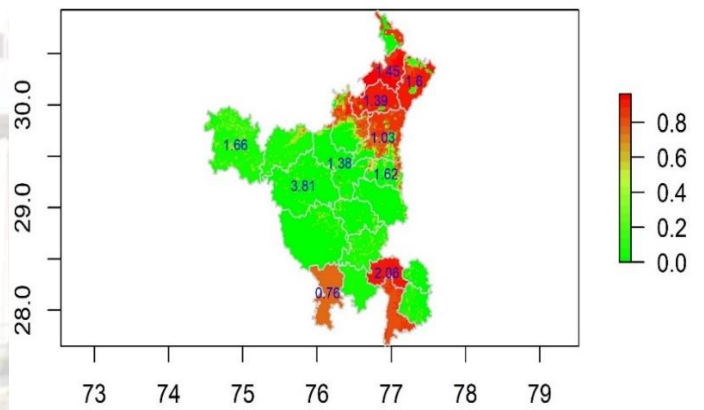
Forecasted risk maps for Classical Swine Fever disease in Assam



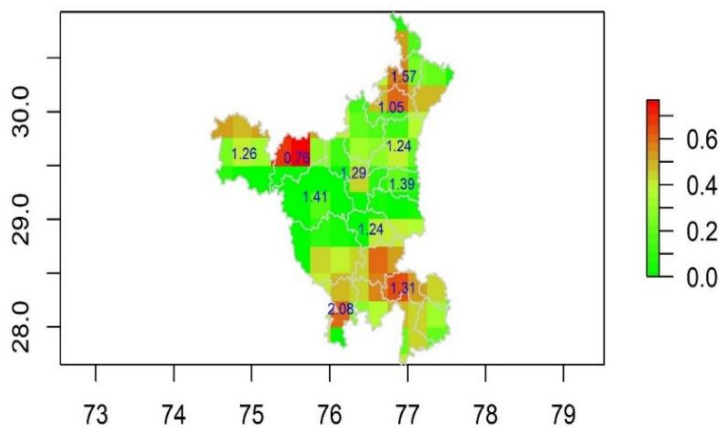
Risk map for Bluetongue in Andhra Pradesh



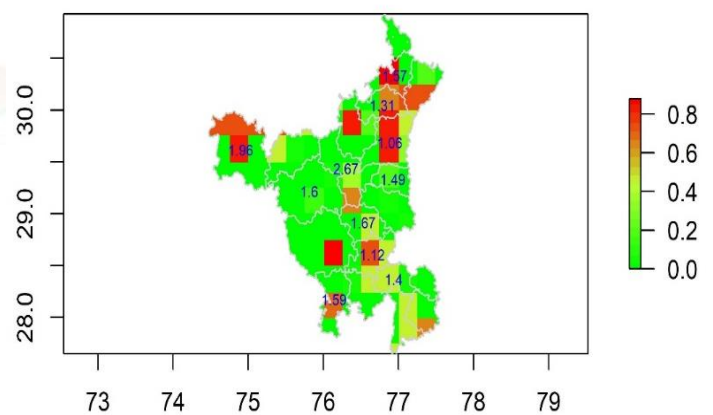
Risk map for Babesiosis in Haryana



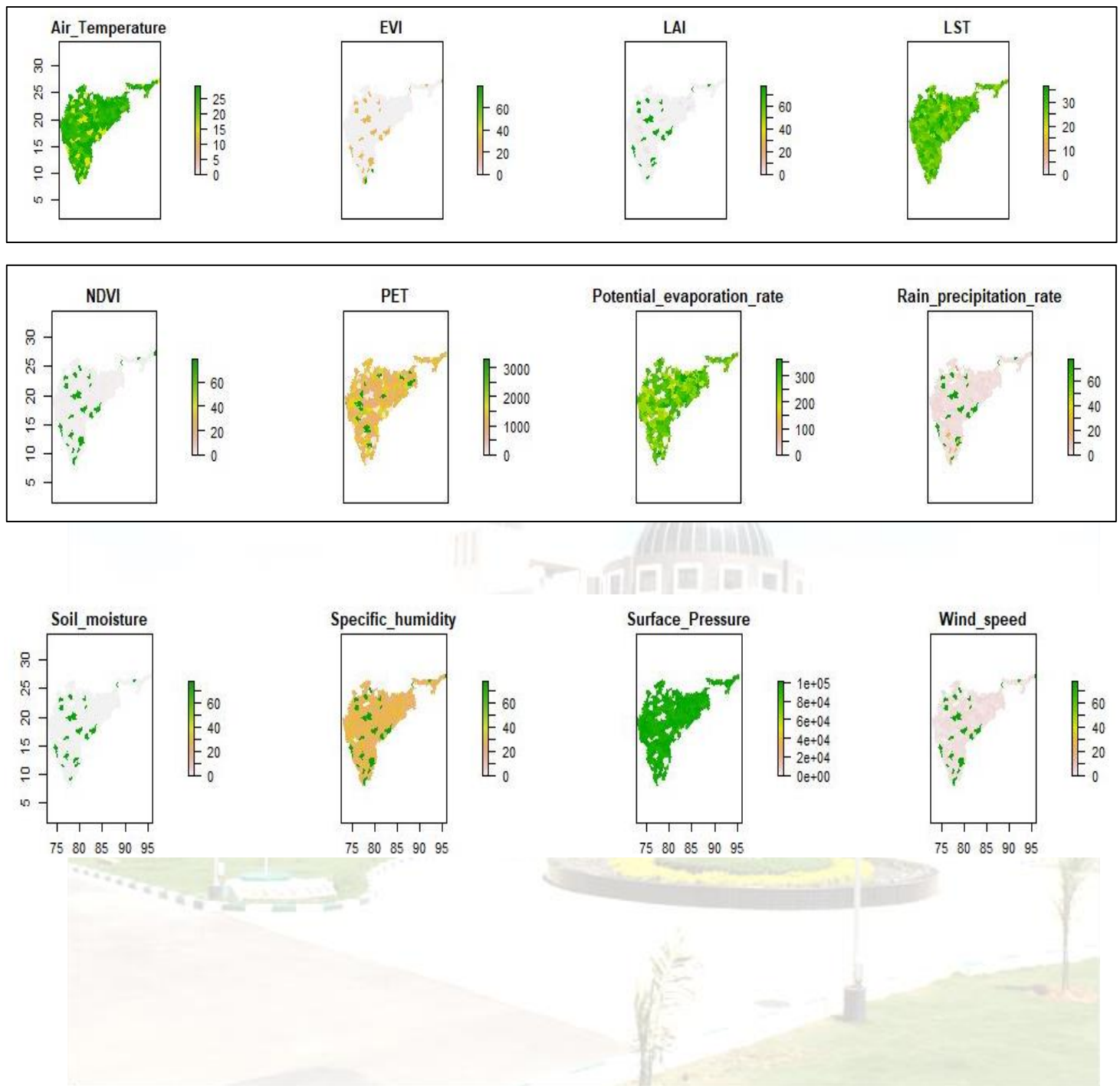
Risk map for Theileriosis in Haryana



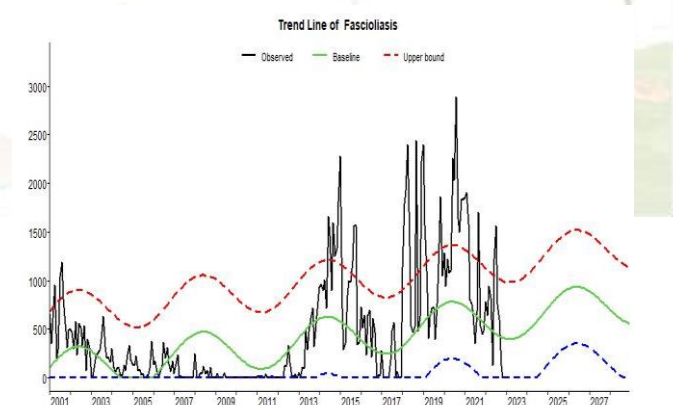
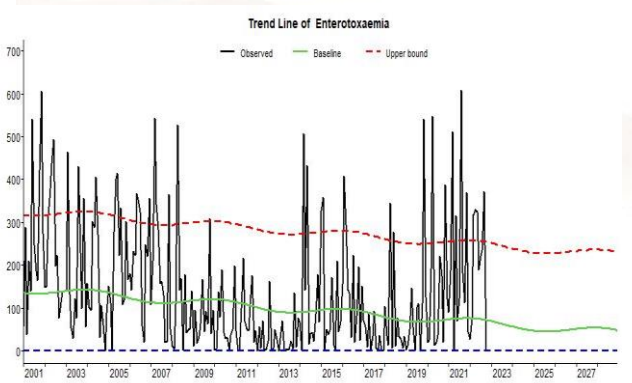
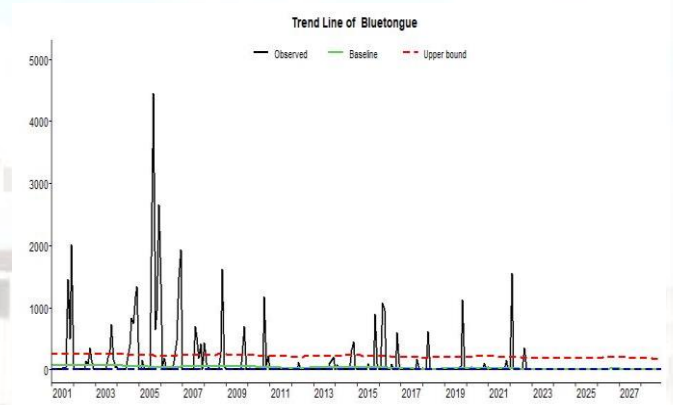
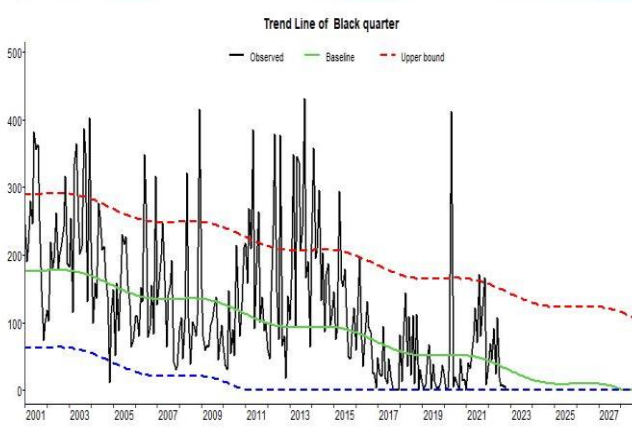
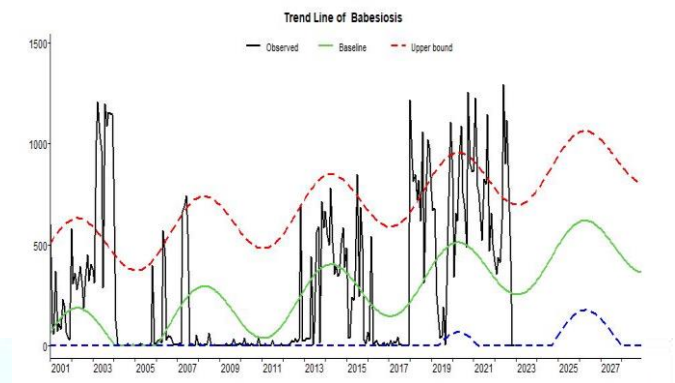
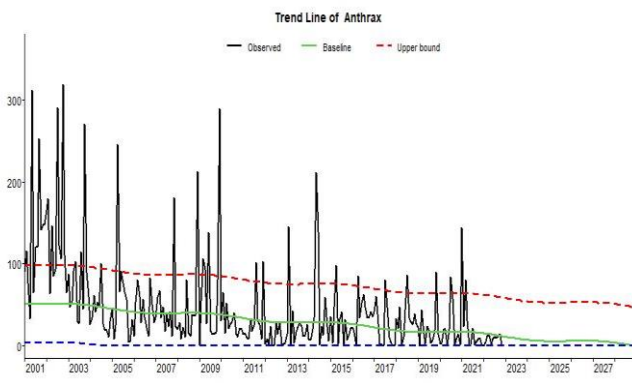
Risk map for Trypanosomiasis in Haryana

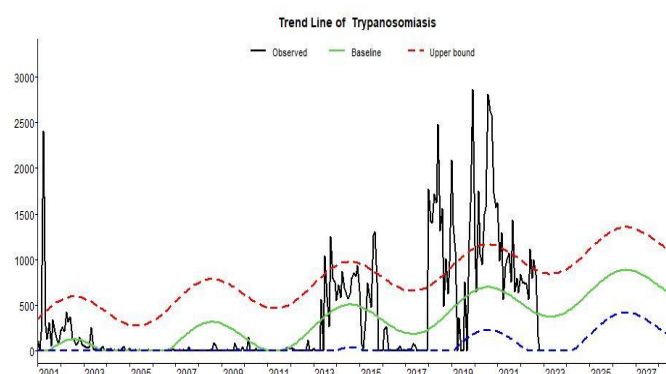
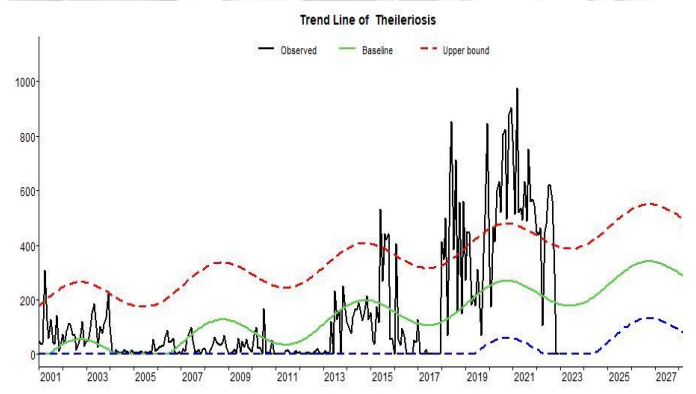
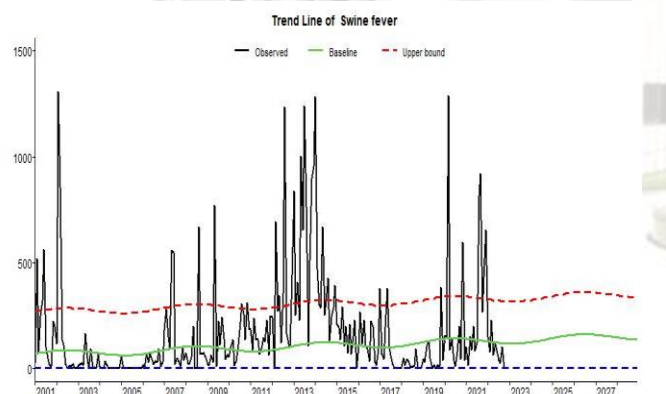
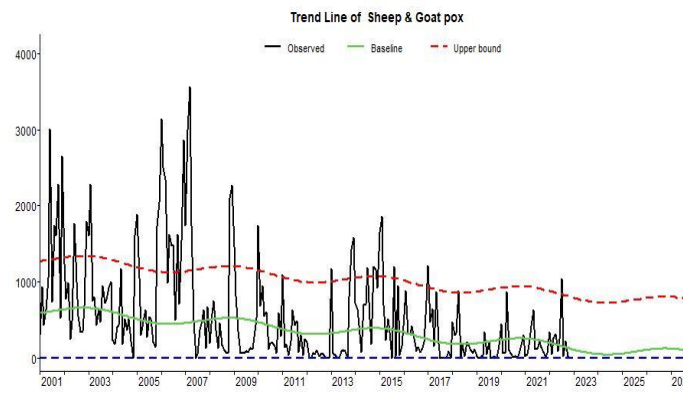
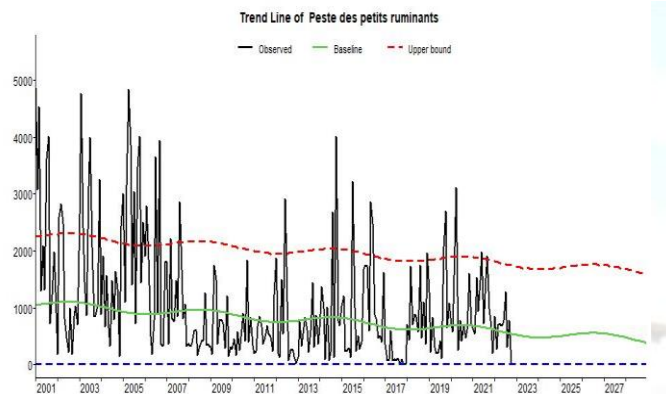
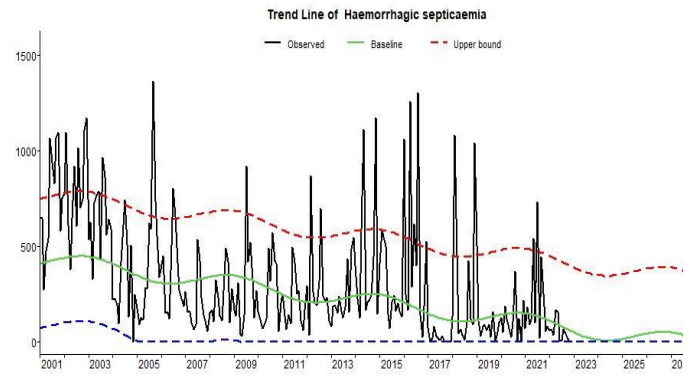
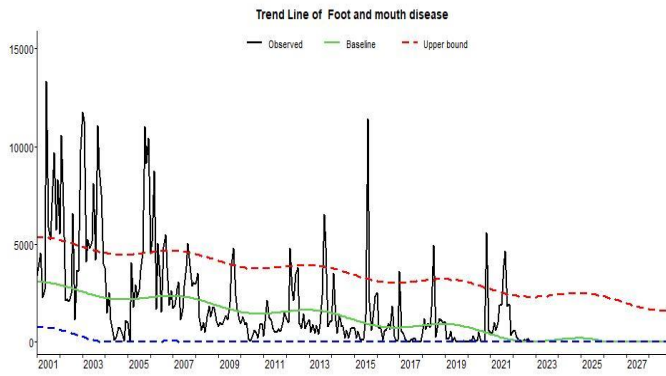


Effects of different parameters on Anthrax disease



11.6. Model based Periodic Regression Model 2023





12. APPLICATION OF META ANALYSIS FOR UNDERSTANDING THE DISEASE PREVALENCE

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RESEARCH ARTICLE

Open Access

Meta-analysis of classical swine fever prevalence in pigs in India: A 5-year study

S. S. Patil, K. P. Suresh, S. Saha, A. Prajapati, D. Hemadri and P. Roy

Indian Council of Agricultural Research - National Institute of Veterinary Epidemiology and Disease Informatics (ICAR-NIVEDI), PBNO-6450, Yelahanka, Bengaluru, Karnataka, India.

Corresponding author: S. S. Patil, e-mail: ss.patil@icar.gov.in

Co-authors: KPS: sureshkp97@gmail.com, SS: sneha.saha88@gmail.com, AP: avi75prajapati@gmail.com, DH: divakar.hemadri@gmail.com, PR: director.nivedi@icar.gov.in

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Abstract

Aim: The aim of the study was to determine the overall prevalence of classical swine fever (CSF) in pigs in India, through a systematic review and meta-analysis of published data.

Materials and Methods: Consortium for e-Resources in Agriculture, India, Google Scholar, PubMed, annual reports of All India Coordinated Research Project on Animal Disease Monitoring and Surveillance, and All India Animal Disease database of NIVEDI (NADRES) were used for searching and retrieval of CSF prevalence data (seroprevalence, virus antigen, and virus nucleic acid detection) in India using a search strategy combining keywords and related database-specific subject terms from January 2011 to December 2015 in English only.

Results: A total of 22 data reports containing 6,158 samples size from 18 states of India were used for the quantitative synthesis, and overall 37% (95% confidence interval [CI]=0.24, 0.51) CSF prevalence in India was estimated. The data were classified into 4 different geographical zones of the country: 20% (95% CI=0.05, 0.55), 31% (95% CI=0.18, 0.47), 55% (95% CI=0.32, 0.76), and 34% (95% CI=0.14, 0.62). CSF prevalence was estimated in northern, eastern, western, and southern regions, respectively.

Conclusion: This study indicates that overall prevalence of CSF in India is much lower than individual published reports.

Keywords: Classical swine fever, India, meta-analysis, pigs, prevalence.



Meta-analysis of the prevalence of livestock diseases in North Eastern Region of India

Nagendra Nath Barman¹, Sharanagouda S. Patil², Rashmi Kurli³, Pankaj Deka¹, Durlav Prasad Bora¹, Giti Deka¹,
Kempanahalli M. Ranjitha², Channappagowda Shivaranjini², Parimal Roy⁴ and Kuralayanapalya P. Suresh³

1. Department of Veterinary Microbiology, College of Veterinary Science, Assam Agriculture University, Guwahati, Assam, India; 2. Department of Virology, ICAR-National Institute of Veterinary Epidemiology and Disease Informatics, Bengaluru, Karnataka, India; 3. Department of Spatial Epidemiology, ICAR-National Institute of Veterinary Epidemiology and Disease Informatics, Bengaluru, Karnataka, India; 4. Director, ICAR-National Institute of Veterinary Epidemiology and Disease Informatics, Bengaluru, Karnataka, India.

Corresponding author: Kuralayanapalya P. Suresh, e-mail: sureshkp97@gmail.com

Co-authors: NNB: nnbarman@gmail.com, SSP: ss.patil@icar.gov.in, RK: rashmikurli@yahoo.in, PD: drpankajaau@gmail.com, DPB: drdpbora@gmail.com, GD: gitideka8906@gmail.com, KMR: ranjuckm10@gmail.com, CS: shivaranjinicveena@gmail.com, PR: director.nivedi@icar.gov.in

Received: 26-07-2019, **Accepted:** 22-11-2019, **Published online:** 11-01-2020

doi: www.doi.org/10.14202/vetworld.2020.80-91 **How to cite this article:** Barman NN, Patil SS, Kurli R, Deka P, Bora DP, Deka G, Ranjitha KM, Shivaranjini C, Roy P, Suresh KP (2020) Meta-analysis of the prevalence of livestock diseases in North Eastern region of India, *Veterinary World*, 13(1): 80-91.

Abstract

Aim: The study aimed to determine the overall prevalence of livestock diseases in North Eastern Region (NER) of India, through a systematic review and meta-analysis of published data.

Materials and Methods: The articles used for the study were retrieved from PubMed, J-Gate Plus, Indian Journals, and Google scholar, R open-source scripting software 3.4.3. Metafor, Meta. The Chi-square test was conducted to assess for the heterogeneity, forest plot (confidence interval [CI] plot) is a method utilized to present the results of meta-analysis, displaying effect estimate and their CIs for each study were used for searching and retrieval of livestock diseases prevalence data in India using a search strategy combining keywords and related database-specific subject terms from 2008 to 2017 in English only.

Results: The prevalence of various livestock diseases are foot-and-mouth disease (21%), bluetongue (28%), brucellosis in bovine (17%), brucellosis in caprine (2%), brucellosis in porcine (18%), brucellosis in sheep and goat (3%), babesiosis (6%), theileriosis (26%), porcine reproductive and respiratory syndrome (1%), porcine cysticercosis (6%), classical swine fever (31%), *Porcine circovirus* (43%), and Peste des petits ruminants (15%). This information helps policymakers to take appropriate measures to reduce the disease burden.

Conclusion: This study indicates that the overall prevalence of various livestock diseases in NER of India.

Keywords: babesiosis, brucellosis, classical swine fever, foot-and-mouth disease, forest plot, livestock, meta-analysis, North Eastern regions, Peste des petits ruminants, *Porcine circovirus*, porcine cysticercosis, porcine reproductive and respiratory syndrome, prevalence, seroprevalence, theileriosis.



Review Article

Peste Des Petits Ruminants in Atypical Hosts and Wildlife: Systematic Review and Meta-Analysis of the Prevalence between 2001 and 2021

Sowjanya Kumari, S^{1,2}, Bhavya, A. P¹, Akshata, N¹, Kumar, K, V¹, Bokade, P. P¹, Suresh, K. P¹, Shome, B. R¹, Balamurugan, V^{1*}

1. Indian Council of Agricultural Research, National Institute of Veterinary Epidemiology and Disease Informatics (ICAR-NIVEDI), Yelahanka, Bengaluru, Karnataka, India

2. Department of Microbiology, Jain University, Bengaluru, Karnataka, India

Received 18 November 2021; Accepted 14 December 2021

Corresponding Author: balavirol@gmail.com

Abstract

Peste des petits ruminants (PPR) or goat plague is considered a leading, highly contagious, and most lethal infectious viral disease of small ruminants affecting the worldwide livestock economy and international animal trade. Although sheep and goats are the primarily affected, the PPR Virus (PPRV) host range has expanded to other livestock (large ruminants) and wildlife animals over the last few decades, resulting in serious concern to the ongoing PPR global eradication program, which is primarily optimized, designed, and targeted towards accessible sheep and goat population. A systematic review and meta-analysis study was conducted to estimate the prevalence and spill-over infection of PPRV in large ruminants (bovine and camel) and wildlife. Published articles from 2001 to October 2021 on the "PPR" were searched in four electronic databases of PubMed, Scopus, Science direct, and Google Scholars. The articles were then selected using inclusion criteria (detection/prevalence of PPRV in bovine, camel, and wildlife population), exclusion criteria (only sheep or goats, lack of prevalence data, experimental trial, test evaluation, and reviews written in other languages or published before 2001), and the prevalence was estimated by random effect meta-analysis model. In the current study, all published articles belonged to Africa and Asia. The overall pooled prevalence of PPR estimates was 24% (95% CI: 15-33), with 30% in Asia (95% CI: 14-49) and 20% in Africa (95% CI: 11-30). The overall estimated pooled prevalence at an Africa-Asia level in bovine and camel was 13% (95% CI: 8-19), and in wildlife, it was 52% (95% CI: 30-74) with significant heterogeneity ($I^2 = 97\%$) in most pooled estimates with a high prevalence in atypical hosts and wildlife across Asia and Africa. Over the last two decades, the host range has increased drastically in the wildlife population, even for prevalent PPR in the unnatural hosts only for a short time, contributing to virus persistence in multi-host systems with an impact on PPR control and eradication program. This observation on the epidemiology of the PPRV in unnatural hosts demands appropriate intervention strategies, particularly at the livestock-wildlife interface.

Keywords: PPR; Bovine; Camel; Wildlife; Systemic Review; Meta-analysis; Prevalence

Web Link https://nivedi.res.in/Nadres_v2/bioinfo.php

13. APPLICATION OF BIOINFORMATICS FOR UNDERSTANDING THE DYNAMICS OF LIVESTOCK INFECTIOUS DISEASES

- a) **An extensive analysis of Codon usage pattern, Evolutionary rate and Phylogeographic reconstruction in Foot and mouth disease (FMD) serotypes (A, Asia 1 and O) of six major climatic zones of India: A Comparative study**

Abstract

Foot and mouth disease (FMD) is a major economically important viral disease of cloven mhoofed livestock globally. The FMD virus (FMDV) spreads widely in confined, cool and humid climatic conditions. Being an RNA virus, FMDV is genetically unstable, and its genome evolution is highly influenced by the mutational pressure. The climatic and environmental conditions have a significant impact on mutational pressure. The present study is a primary effort to establish a comprehensive relationship between climatic factors and molecular evolutionary pattern of serotypes FMDV circulating in India. In this study, isolates of three serotypes (A, Asia 1 and O) were selected from six major climatic zones of India (Montane, Humid subtropical, Tropical wet and dry, Tropical wet, Semi-arid and Arid). Based on the full genome nucleotide sequence data, the codon usage bias, evolutionary and phylogeographic analysis were carried out. The study revealed that the codon use bias indicators in the FMDV serotypes differed significantly depending on the climatic zones. It implies that the selection and mutational pressure influence the codon usage pattern indices (eNC, CAI, RCDI, GRAVY, Aromo), with mutational pressure taking precedence in determining the codon usage bias of the FMDV genome. The tMRCA was estimated to be 1960, 1956 and 1961 for Indian FMD virus serotype-A, Asia 1 and O respectively, which is around 17, 29 and 36 years before its actual identification in the field. Virus transmission across the region was evident from phylogeographic analysis. The integrated analysis of codon usage bias, evolutionary rate and phylogeography analysis signifies the major role of mutational and selection pressure, implying that the in FMD virus co-evolution and adaptations are highly influenced by the climatic/environmental factors.

- b) **Relative Analysis of Codon Usage and Nucleotide Bias between Anthrax Toxin Genes Subsist**

Inpxo1 Plasmid of *Bacillus Anthracis*

Abstract

Anthrax is an ancient and acute illness that affects a large quantity of animal species and is caused by a bacterium *Bacillus anthracis*, which is a rod-shaped, gram-positive and spore forming bacterium. Virulent forms of *B. anthracis* has two large pathogenicity related plasmids pXO1 and pXO2. pXO1 has the different anthrax toxin genes *cya*, *lef*, and *pagA* where as pXO2 has the genes accountable for capsule synthesis and degradation, *capA*, *capB*, *capC*, and *capD*. *B. anthracis* express its pathogenic activity mostly over the capsule and the manufacture of a toxic compound involving three proteins known as edema factor (EF), lethal factor (LF) and protective antigen (PA). These two enormous plasmids of *B. anthracis* are crucial for full pathogenicity, exclusion of either of the plasmids extremely weakens the malignity of *B. anthracis*. In the current study we conducted the relative analysis of the codon usage and nucleotide bias of virulent genes subsist in pXO1 plasmid of *B. anthracis*. Codon usage bias not only plays a substantial role at the extent of gene expression, but also supports to improve the efficacy and accurateness of translation. Codon usage pattern analysis of *B. anthracis* genome is essential for understanding the evolutionary characteristics in the different species. To examine the codon usage arrangement of the *B. anthracis* genome, Nucleotide sequences

of the virulent genes viz *cya*, *lef* and *pag* were collected from National Center for Biotechnology Information (NCBI). The correlations between GC3s, whole GC content, Effective No. of Codons (ENC), Codon Adaptation Index (CAI), Codon Bias Index (CBI), Frequency of Optimal Codons (FOP), General average hydropathicity (Gravy) and Aromaticity (Aroma), of the selected genes were determined. The ENC-plot i.e., ENC values vs GC3s, Pr2 plot i.e., relationship between $A3 / (A3 + T3)$ and $G3 / (G3 + C3)$, Neutrality plot i.e., GC12 versus GC3s, and the RSCU of the genes, all shows codon usage bias existence in all the virulent genes subsists in pXO1 plasmid of *B. anthracis* genome. These results express the codon usage bias existing in the pXO1 plasmid's virulent genes of *B. anthracis* genome could be utilized for further exploration on their evolutionary analysis as in design of primers, design of transgenes, determine of origin of species as well as prediction of gene expression level and gene function.

c) **Reverse vaccinology based *in silico* analysis of Epitope prediction in *cya*, *lef* and *pagA* genes from *Bacillus anthracis* against Anthrax infected species: An Immunoinformatics approach**

Abstract

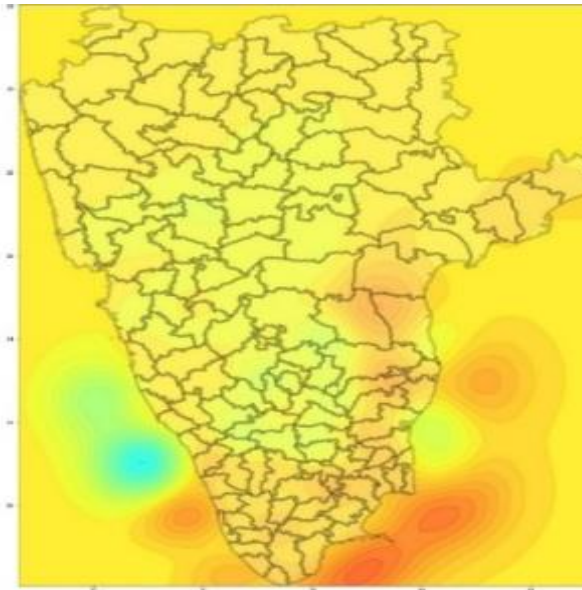
Bacillus anthracis is a Gram-positive spore-forming bacterium that causes the zoonotic disease: anthrax, an abrupt illness that disproportionately impacts grazing livestock and wild ruminants. Anthrax's geographical reach despite years of research on anthrax epizootic and epidemics behaviour, to date, remains to be elucidated. Existing therapeutics, however, are ineffective in combating this infectious disease, necessitating the development of a better vaccine to pause the pandemic using immunoinformatics approaches, this study intended to predict an efficient epitope for a vaccine against anthrax in animals and humans of the toxin genes such as *cya*, *lef* and *pagA* of *B. anthracis* against anthrax. The B-cell and T-cell epitopes were predicted utilizing various bioinformatics tools/software and docking analysis was performed. Consequently, it was found that the evaluated epitopes had no allergenicity, no toxicity and had high antigenicity that provides an effectual and most rapid technique to estimate peptide synthetic vaccines to impede anthrax.



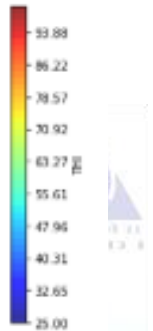
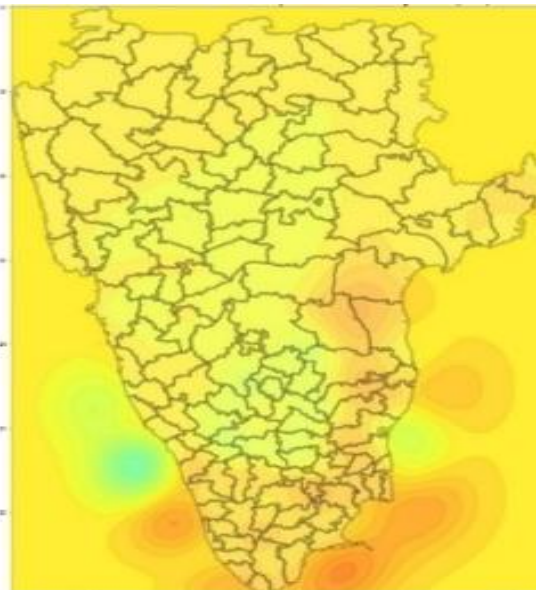
14. IMPACT OF CLIMATE CHANGE ON LIVESTOCK DISEASE INCIDENCE

14.1 Southern States Temperature Humidity Index (THI) 2001-2021

THI for 2001



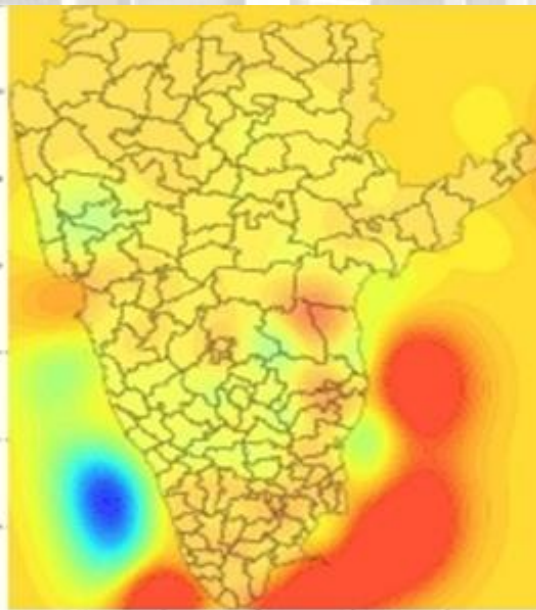
THI for 2011



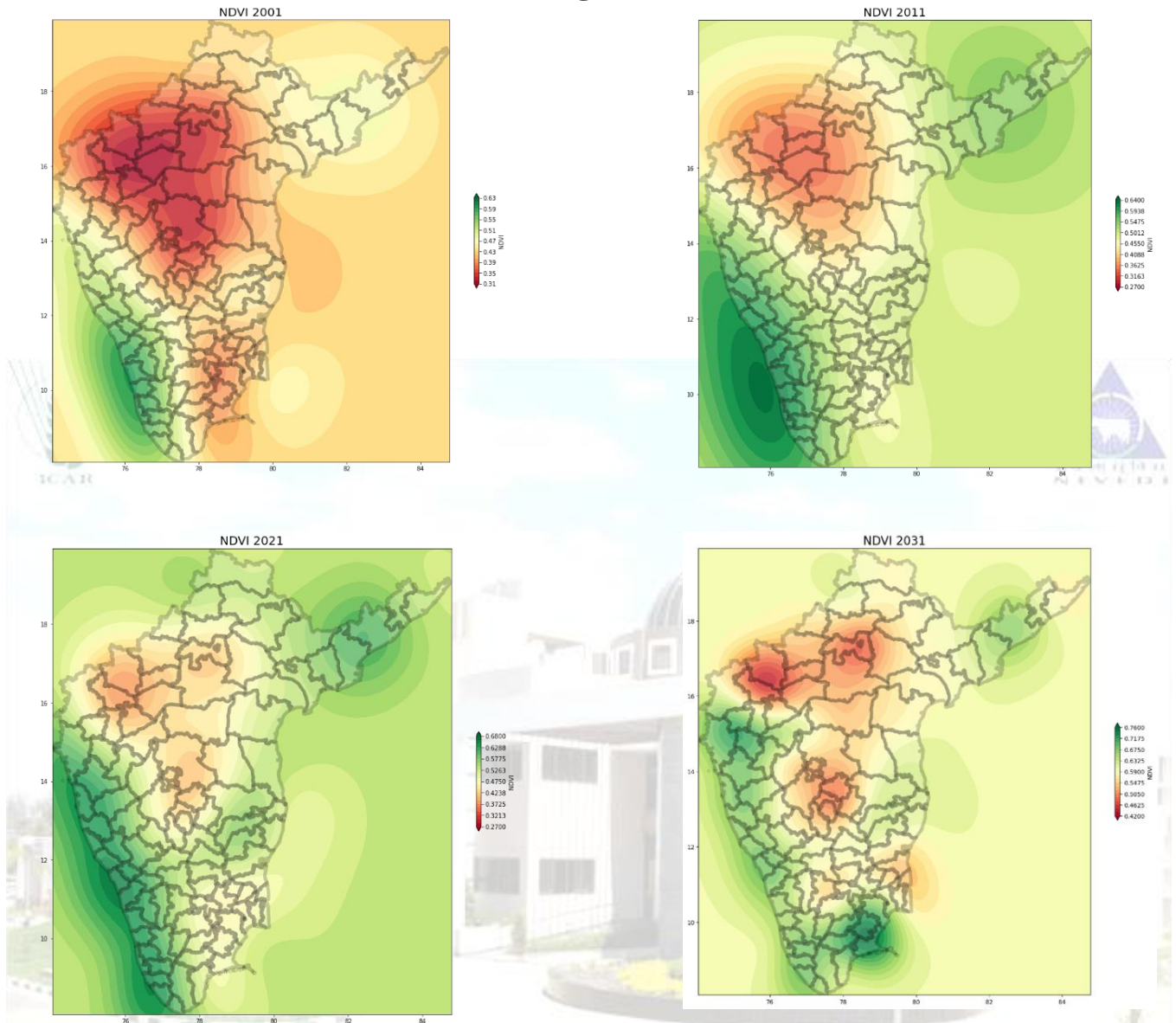
THI for 2021



THI for 2023 (Predicted)

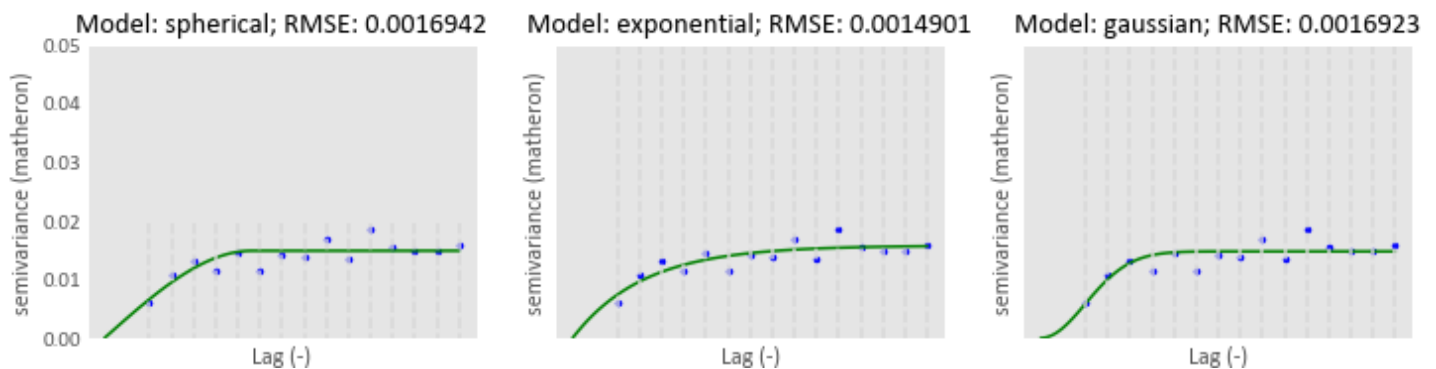


14.2. Southern States Normalized Difference Vegetation Index (NDVI)

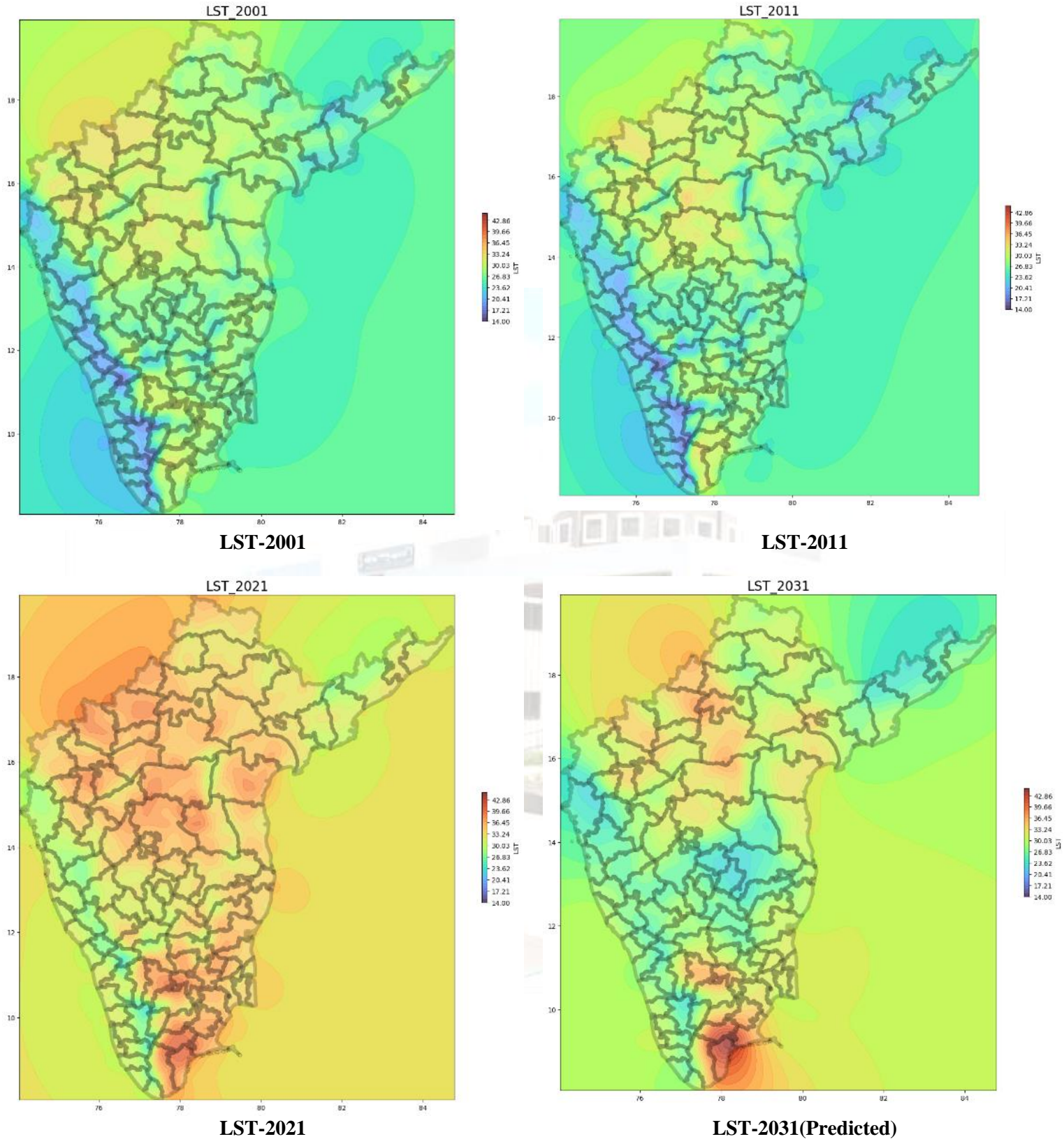


Best fit Variogram model for NDVI:

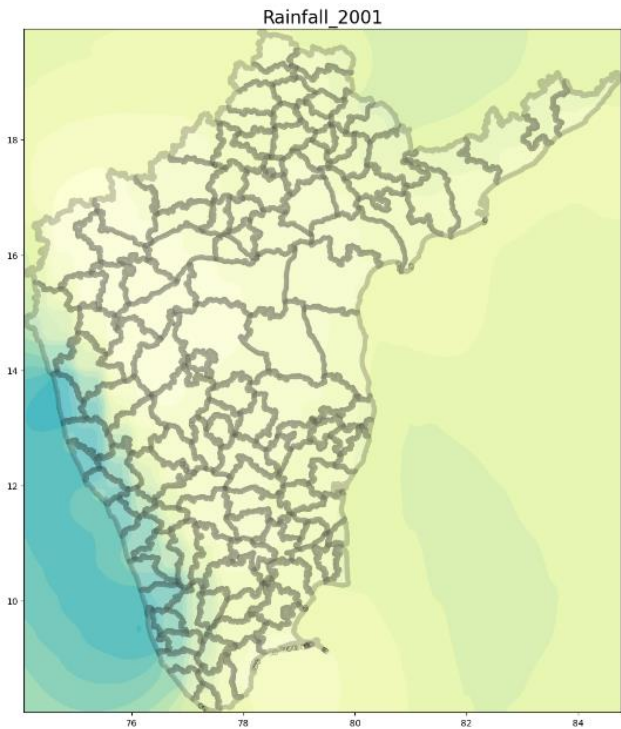
After comparing various variogram models, we could choose the one that would fit kriging the best based on the RMSE value. Based on the result with the lowest RMSE, we may say that the exponential model was best specified.



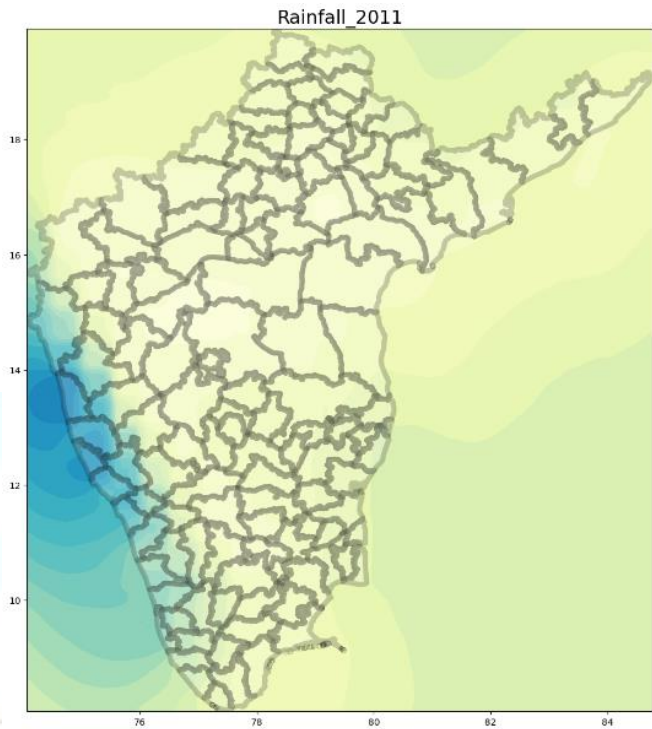
14.3 Land Surface Temperature (LST) - 2001-2031



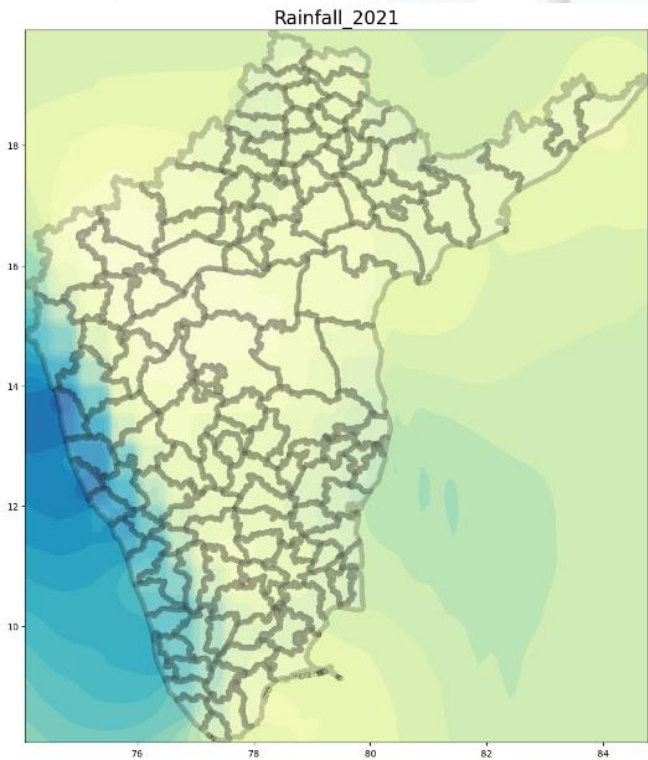
14.4 Rainfall - 2001-2031



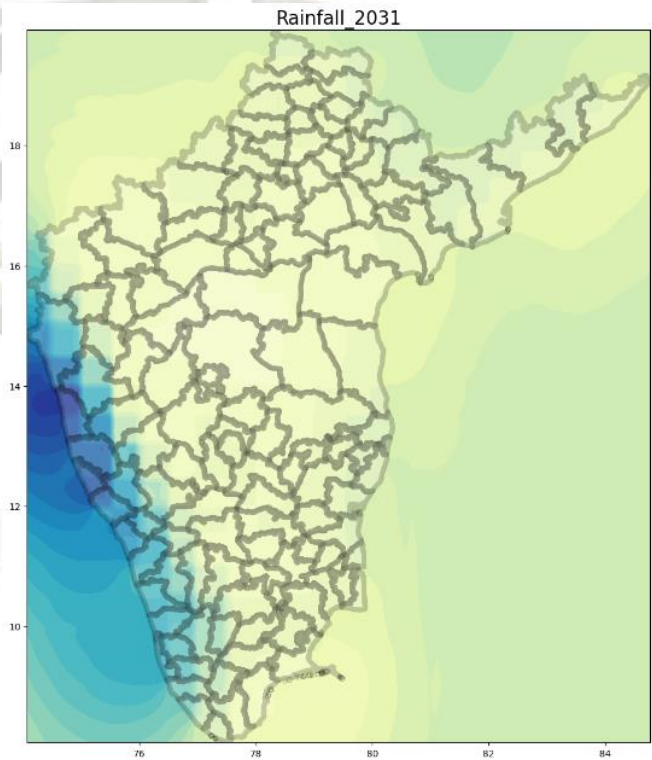
Rainfall 2001



Rainfall 2011

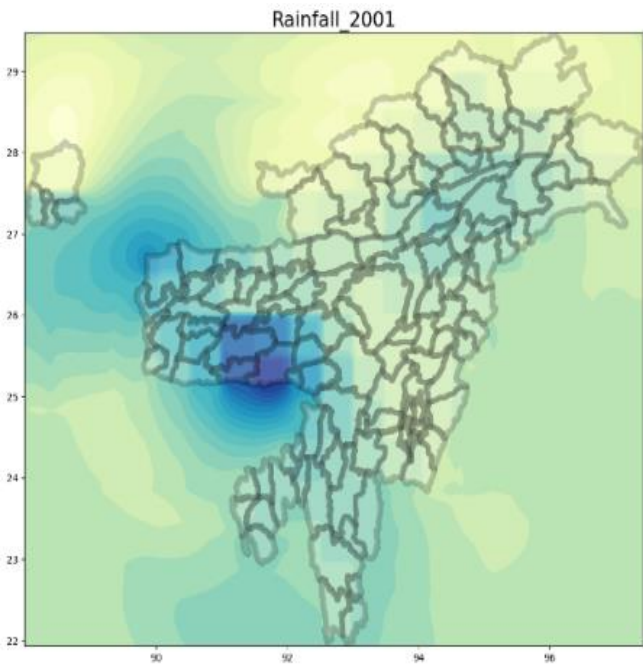


Rainfall 2021

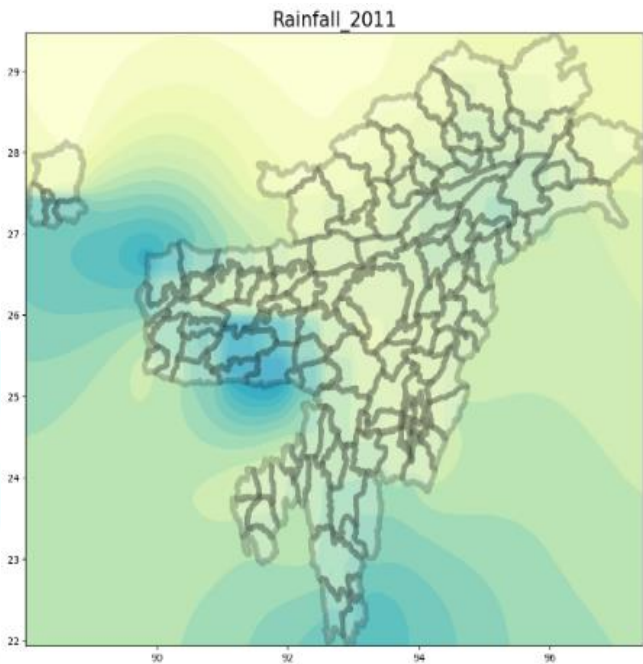


Rainfall 2031 (Predicted)

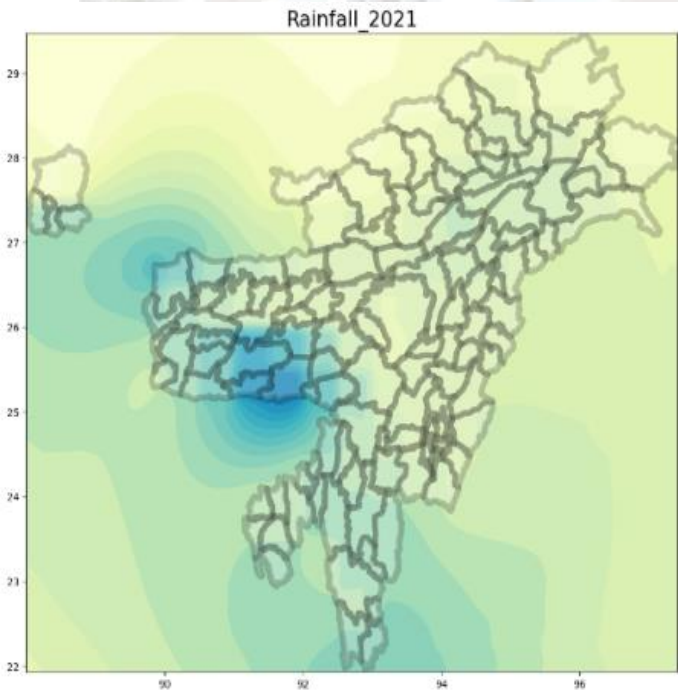
14.5 Rainfal NER 2001-2031



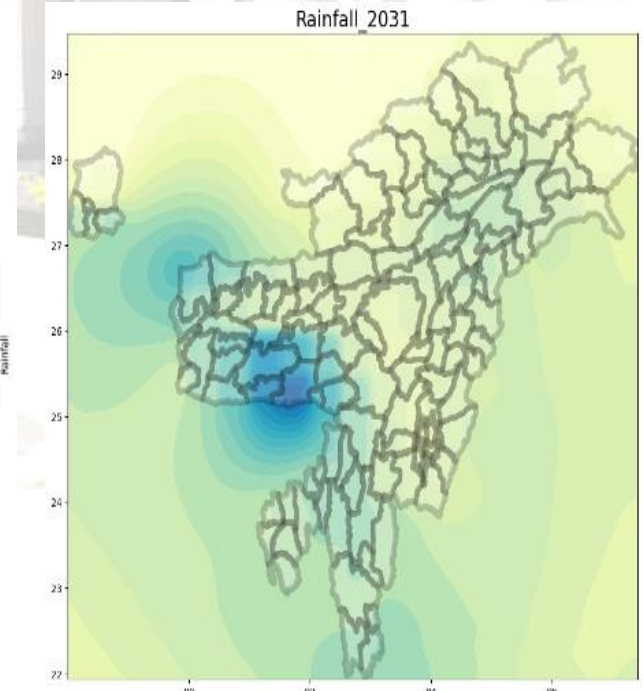
Rainfall 2001



Rainfall 2011

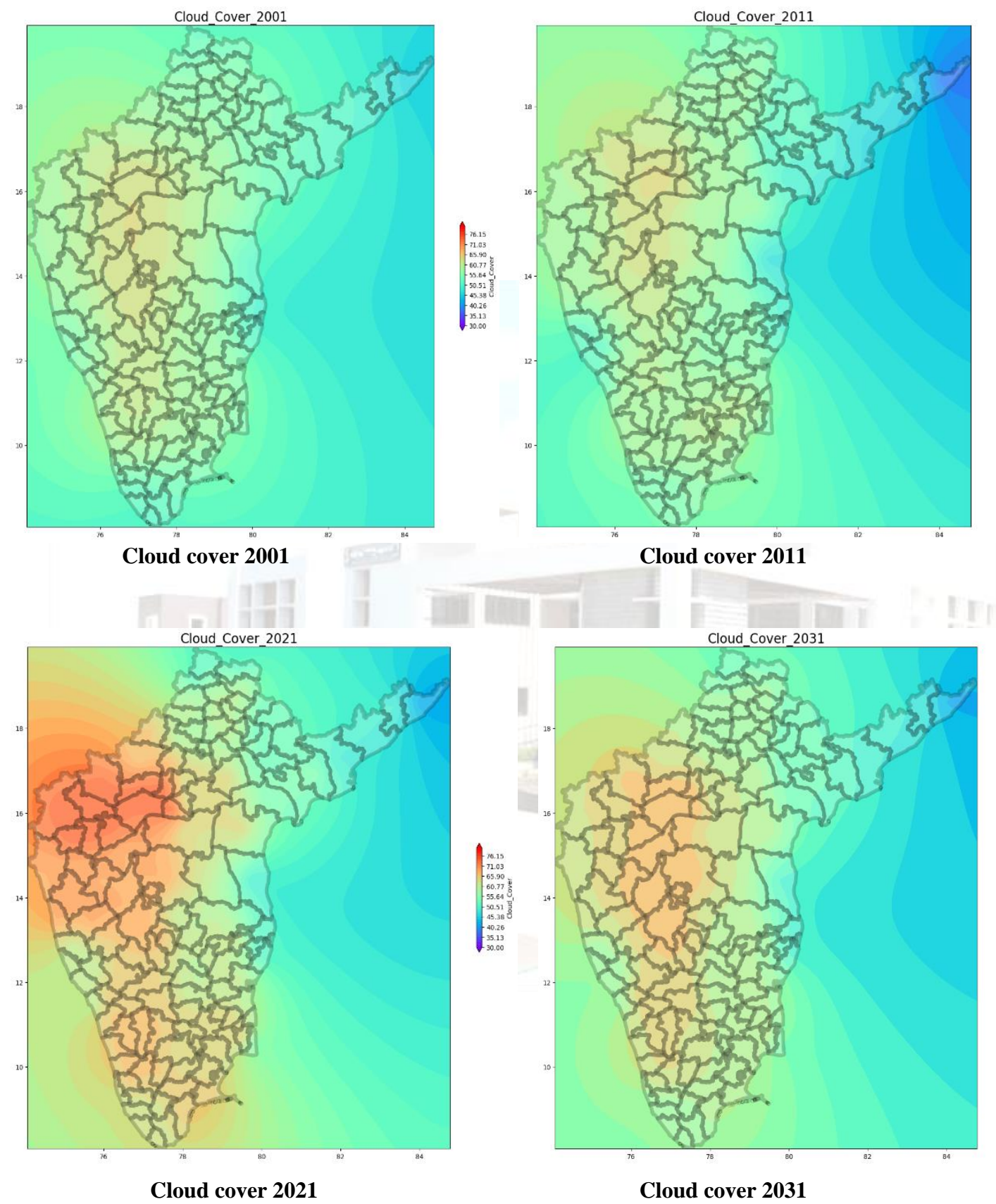


Rainfall 2021

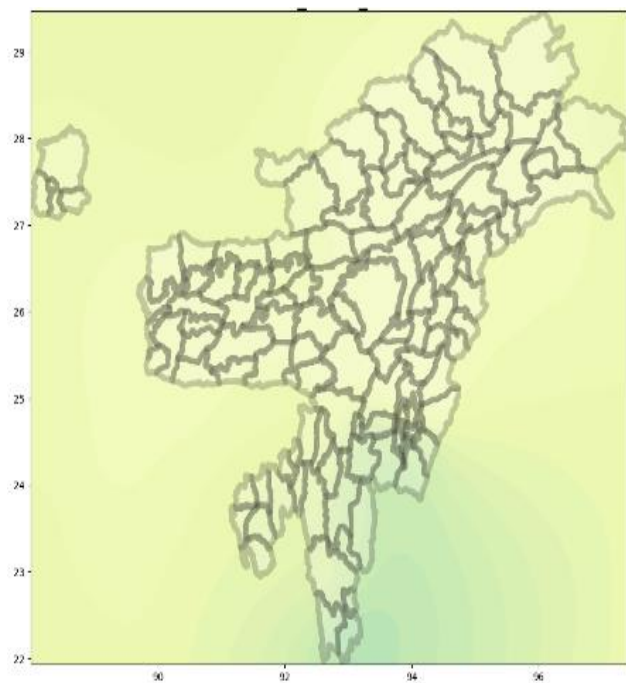


Rainfall 2031(Predicted)

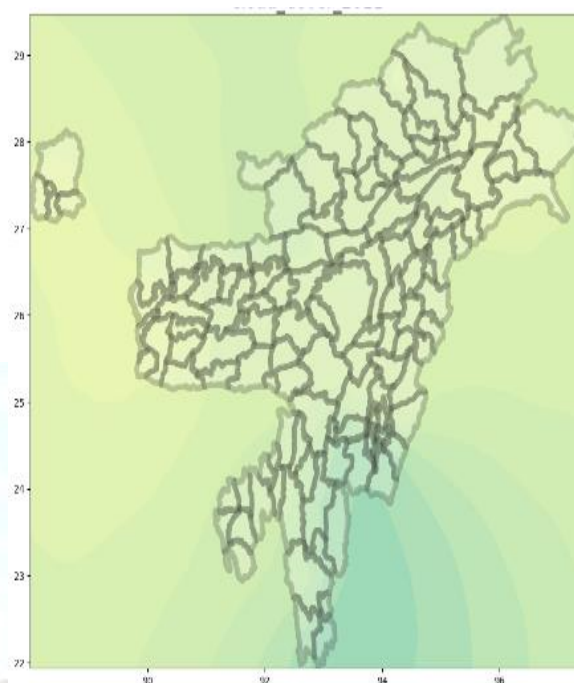
14.6 Cloud cover – 2001-2031



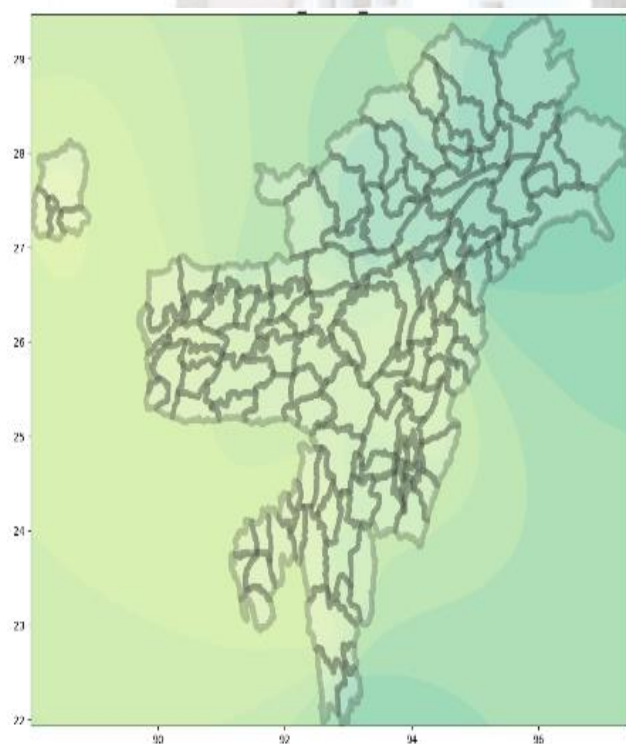
14.7 Cloud cover NER 2001-2031



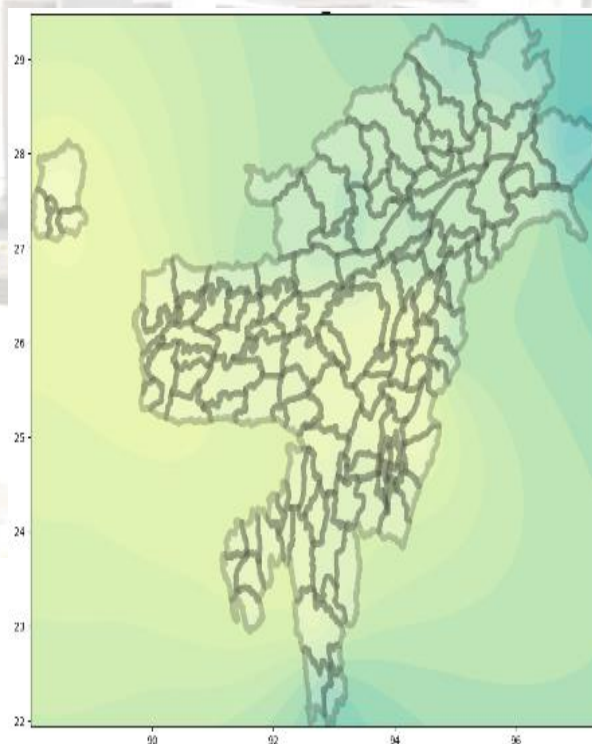
Cloudcover – 2001



Cloudcover - 2011

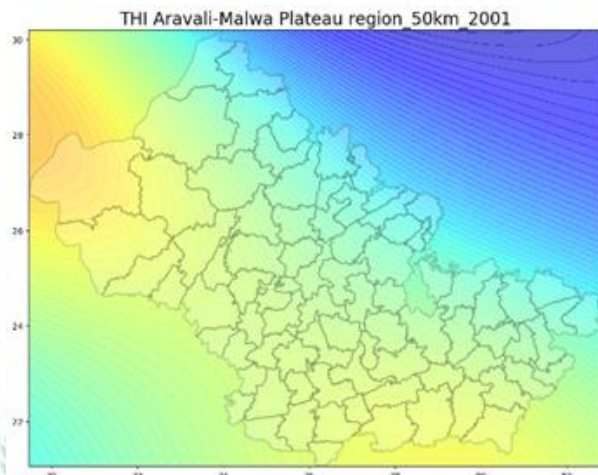


Cloudcover- 2021

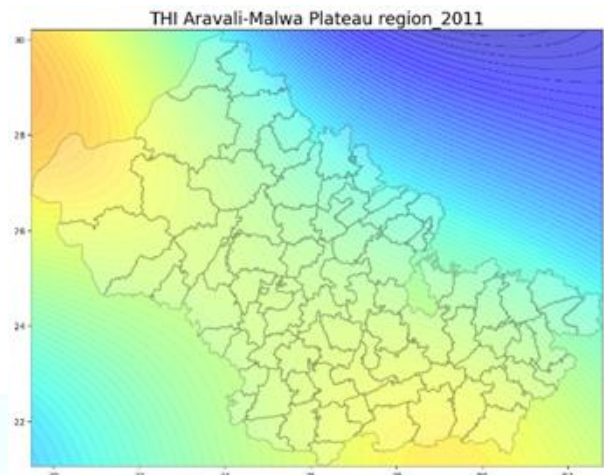


Cloudcover- 2031

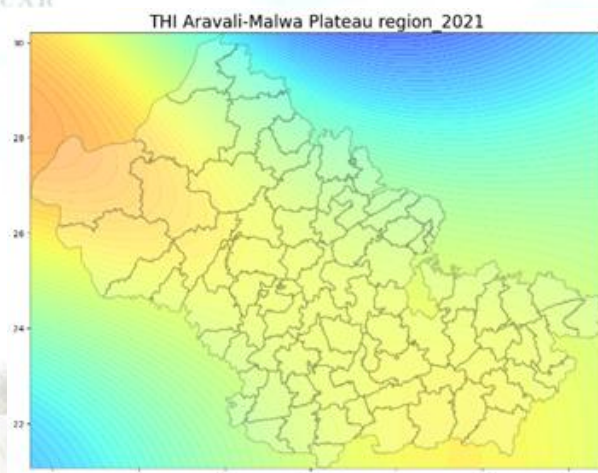
14.8 THI for Aravalli Central Plateau & Hills 2001-2033 (Predicted)



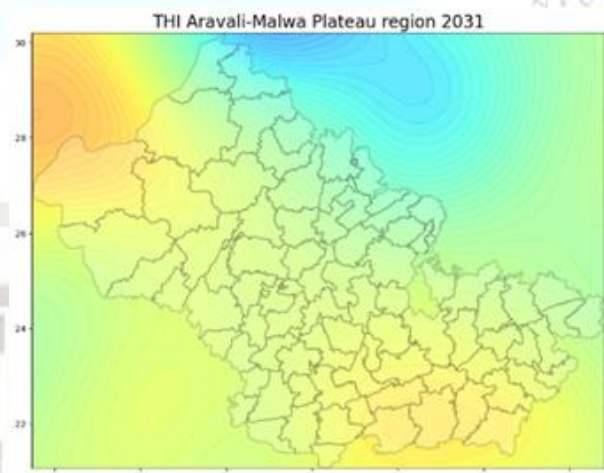
THI for Aravalli Central Plateau & Hills 2001



THI for Aravalli Central Plateau & Hills 2011



THI for Aravalli Central Plateau & Hills 2021



THI for Aravalli Central Plateau & Hills 2031 (predicted)

Zone Name: Aravalli Central Plateau & Hills

Zone No: 08

States: Madhya Pradesh, Rajasthan

No of Districts: 83

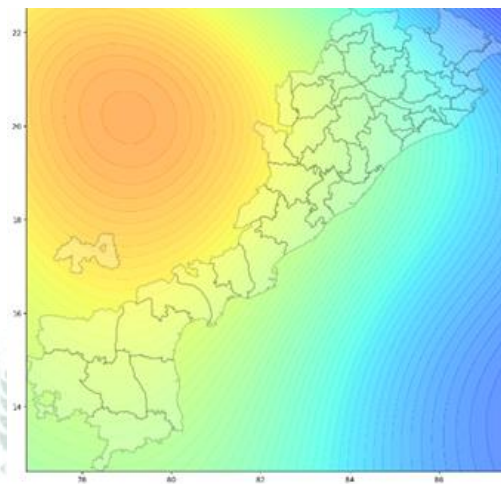
50 KM grids of 0.08 size: No of grids 963

Total Number of Data Points: 374607

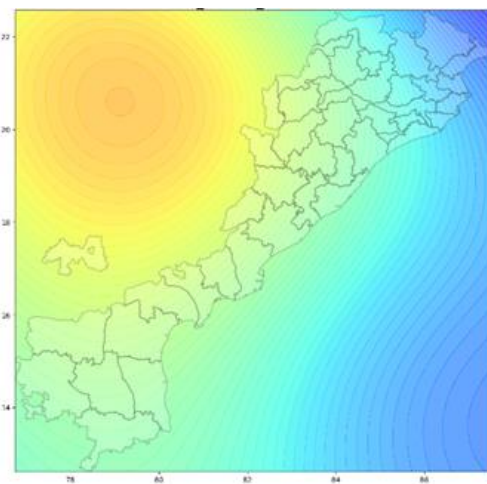
Variogram best fit (least RMSE) Gaussian

Predicted Data: Predicted for next 10 years (2023 to 2033)

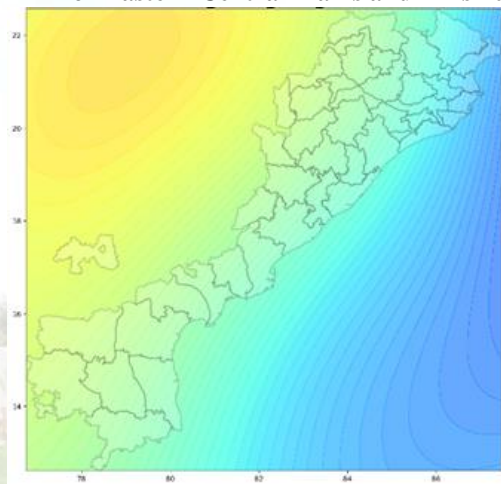
14.9 THI for Eastern Central Plains and Hills 2001-2033 (Predicted)



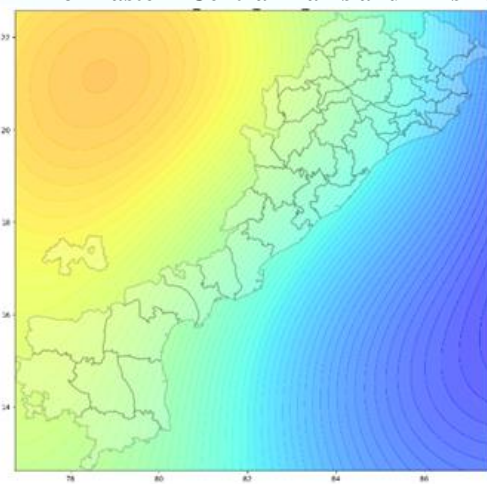
THI for Eastern Central Plains and Hills 2001



THI for Eastern Central Plains and Hills 2011



THI for Eastern Central Plains and Hills 2021



THI for Eastern Central Plains and Hills 2031(Prediction)

Zone Name: Eastern central plains & Hills

Zone No: 11

States: Andhra Pradesh, Odisha

No of Districts: 44

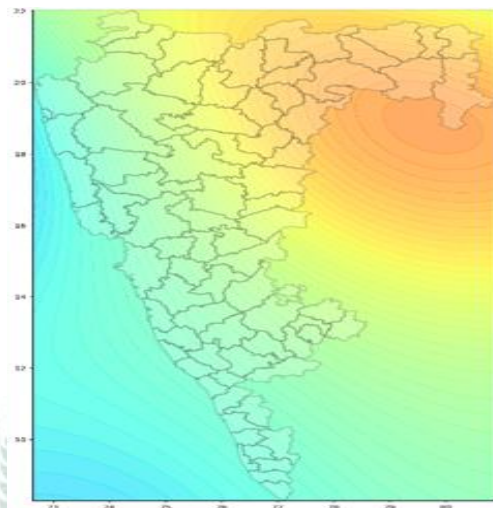
50 KM grids of 0.08 size: No of grids 484

Total Number of Data Points: 188276

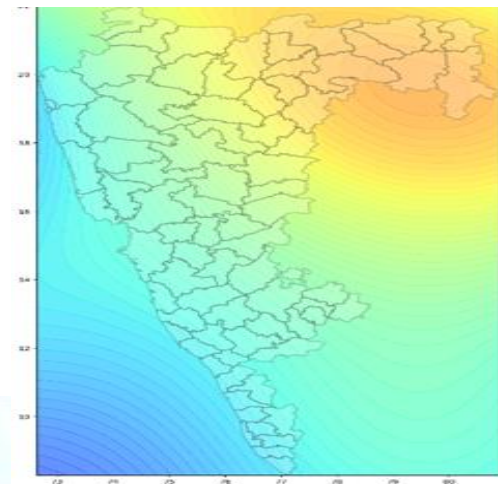
Variogram best fit (least RMSE) Gaussian

Predicted Data: Predicted for the next 10 years (2023 to 2033)

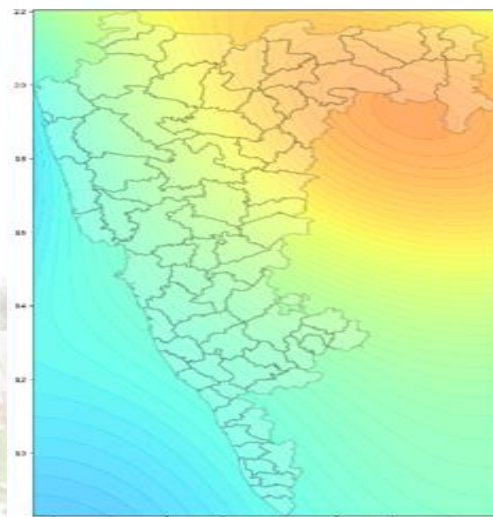
14.10 THI for Western Coastal Plains and Ghats 2001-2033 (Predicted)



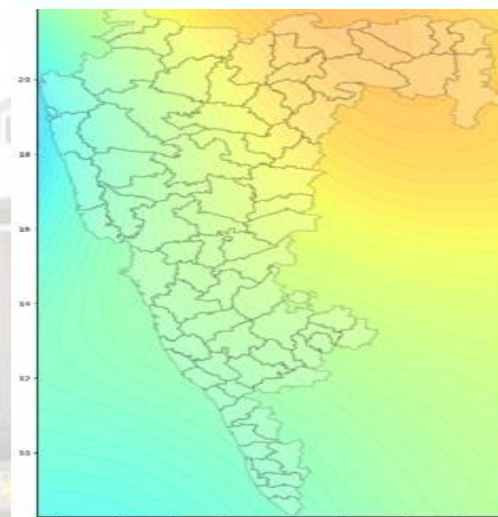
THI for Western Coastal Plains and Ghats 2001



THI for Western Coastal Plains and Ghats 2011



THI for Western Coastal Plains and Ghats 2021



THI for Western Coastal Plains and Ghats 2031(Predicted)

Zone Name: Western Coastal Plains & Ghats

Zone No: 12

States: Karnataka, Kerala, Maharashtra

No of Districts: 80

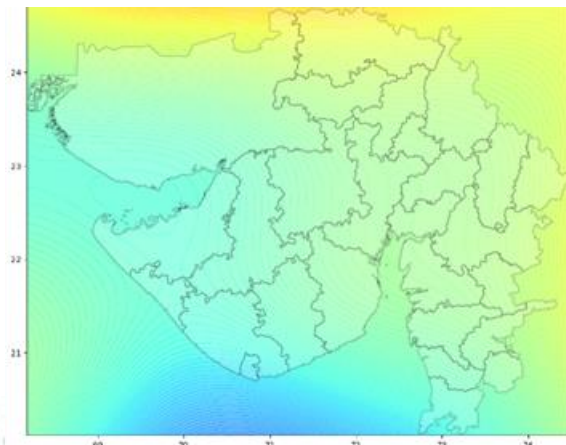
50 KM grids of 0.08 size: No of grids 816

Total Number of Data Points: 317424

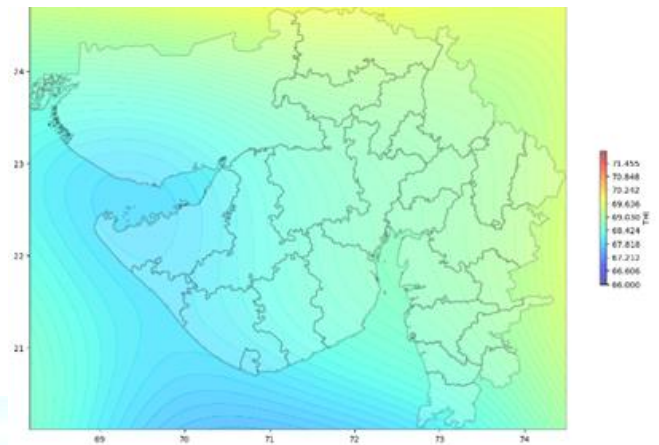
Variogram best fit (least RMSE) Gaussian

Predicted Data: Predicted for the next 10 years (2023 to 2033)

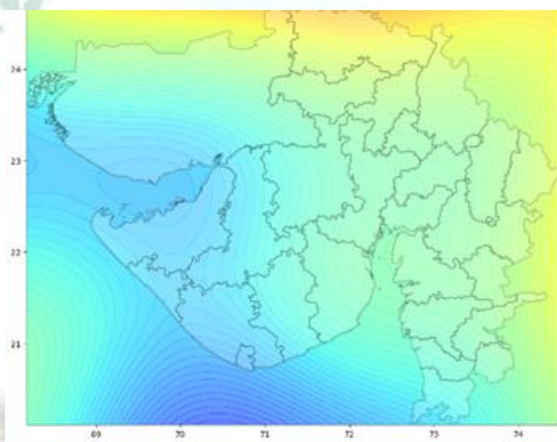
14.11. THI for Gujarat Plains and Hills 2001-2033 (Predicted)



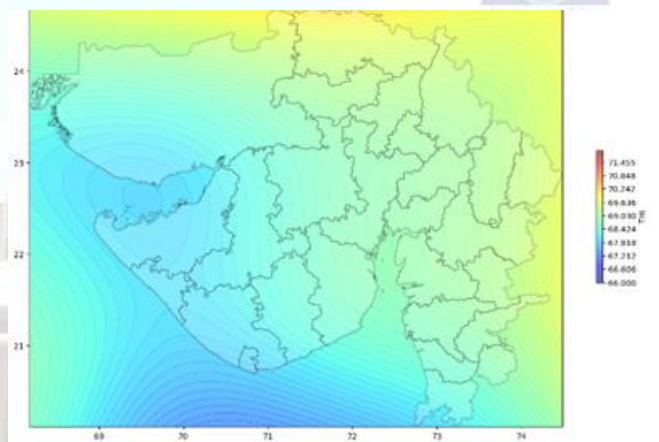
THI for Gujarat Plains and Hills 2001



THI for Gujarat Plains and Hills 2011



THI for Gujarat Plains and Hills 2021



THI for Gujarat Plains and Hills 2033(Predicted)

Zone Name: Gujarat Plains & Hills

Zone No: 13

States: Gujarat

No of Districts: 26

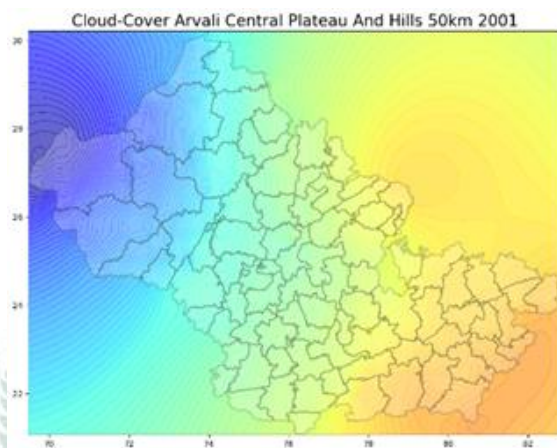
50 KM grids of 0.08 size: No of grids 289

Total Number of Data Points: 112810

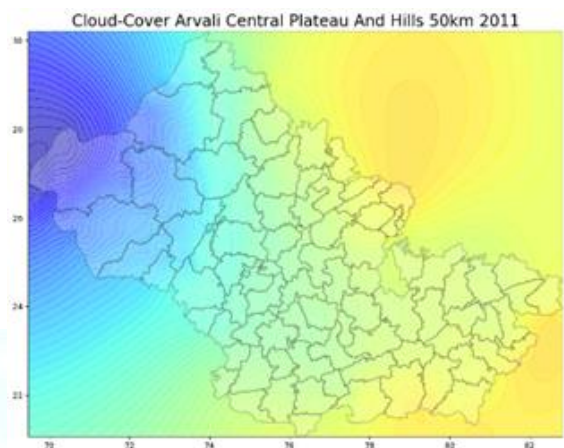
Variogram best fit (least RMSE): Gaussian

Predicted Data: Predicted for the next 10 years (2023 to 2033)

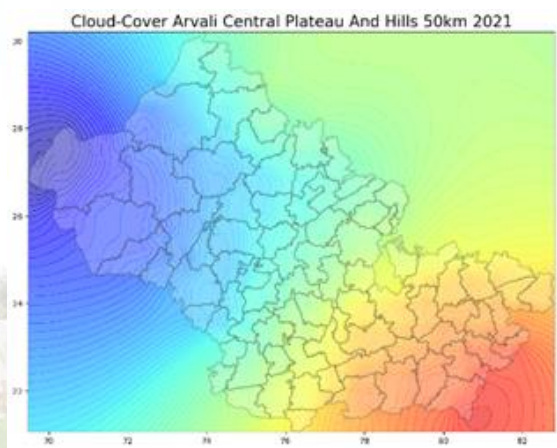
14.12.Cloud-Cover for Aravalli Central Plateau & Hills 2001-2033 (Predicted)



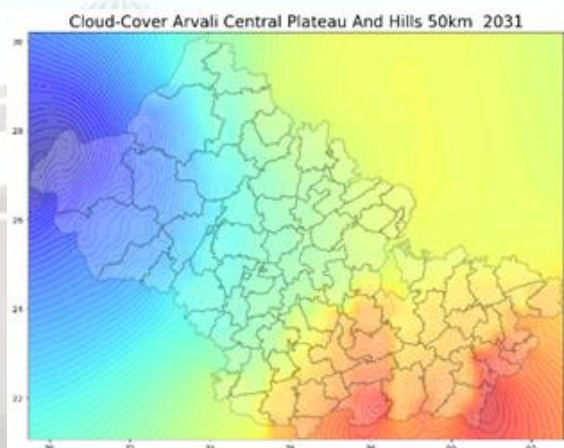
Cloud-Cover for Aravalli Central Plateau & Hills 2001



Cloud-Cover for Aravalli Central Plateau & Hills 2001



Cloud-Cover for Aravalli Central Plateau & Hills 2001



Cloud-Cover for Aravalli Central Plateau & Hills 2031 (predicted)

Zone Name: Aravalli Central Plateau & Hills

Zone No: 08

States: Madhya Pradesh, Rajasthan

No of Districts: 83

50 KM grids of 0.08 size: No of grids 963

Total Number of Data Points: 358236

Variogram best fit (least RMSE): Spherical

Predicted Data: Predicted for next 10 years (2023 to 2033)

Kriging Results

Models:

- **Ordinary Kriging:** Ordinary Kriging is a spatial estimation method where the error variance is minimized

- **variogram_model:** Gaussian Model
 $p \cdot (1 - e^{-d^2/(4r^2)}) + n$

Variables are defined as:

dd = distance values at which to calculate the variogram

pp = partial sill (psill = sill - nugget)

rr = range

nn = nugget

ss = scaling factor or slope

ee = exponent for power model

- **Spatial data:10km/grid**

- **THI (Temperature Humidity Index) data:** The THI data Generate using Air Temperature and Relative Humidity and the Following equation

$$THI = (1.8 \cdot AT + 32) - [(0.0055 - 0.0055 \cdot RH) \cdot (1.8 \cdot AT - 26)]$$

AT: Airtemparture

RH: Relative Humidity

APPENDIX

Abbreviations

NADRES	:	National Animal Disease Referral Expert System
R	:	R environment for statistical computing
ASF	:	African Swine Fever
BQ	:	Black Quarter
BT	:	Bluetongue
CSF	:	Classical Swine Fever
ET	:	Enterotoxaemia
FMD	:	Foot and Mouth disease
HS	:	Haemorrhagic Septicaemia
PPR	:	Peste des Petits Ruminants
SGP	:	Sheep and Goat pox
hPa	:	Hectopascals
NR	:	No risk/No data available
VLR	:	Very low risk
LR	:	Low risk
MR	:	Moderate risk
HR	:	High risk
VHR	:	Very high risk

REFERENCES

1. Bylaiah, S., Shedole, S., Suresh, K.P., Gowda, L., Shivananda, B., Shivamallu, C. and Patil, S.S., 2022. Disease Prediction Model to Assess the Impact of Changes in Precipitation Level on the Risk of Anthrax Infectiousness among the Livestock Hosts in Karnataka, India. *International Journal of Special Education*, 37(3).
2. Suma, A. P., Suresh, K. P., Gajendragad, M. R., Kavya, B.A. (2017). Forecasting Anthrax in Livestock in Karnataka State using Remote Sensing and Climatic Variables, *Int. J. Sci. Res*, 6 (5), 78-96.
3. Suresh KP, Bhavya AP, Chandan S, Raghu R A, Silina E, Stupin V, Shiva P K, Bibek R S and Sharanagouda S P, 2022, Seroprevalence of sheeppox and goatpox virus in Asia and African continent: A systematic review and meta-analysis (Scientometrics), *Veterinary world*, 51(2):455-464
Suresh, K.P., Hemadri, D., Kruli, R., Dheeraj, R. and Roy, P., 2019. Application of Artificial Intelligence for livestock disease prediction. 69(3), 60-62.
4. Nandi, S., Suresh, K.P. and Mondal, S., 2013. Ammonia and urea levels in blood and ovarian follicular fluid in cattle fed with normal and protein rich or imbalanced diet: A meta-analysis. *Indian Journal of Animal Sciences*, 83(5), pp.557-559.
5. SowjanyaKumari, S., Panduranga, B.A., Nayak, A., Kumar, K.V., Bokade, P.P., Suresh, K.P., Shome, B.R. and Balamurugan, V., 2021. Peste Des Petits Ruminants in Atypical Hosts and Wildlife: Systematic Review and Meta-Analysis of the Prevalence between 2001 and 2021. *Archives of Razi Institute*, 76(6), pp.1589-1606.
6. M. Denis, V. Vanderweerd, R. Verbeeke, A. Laudisoit, L. Wynants, D. Van Der Vliet (2020). COVIPENDIUM: information available to support the development of medical counter measures and interventions against COVID-19 (Version 2020-05-05). Transdisciplinary Insights. <http://doi.org/10.5281/zenodo.3782325>
7. Questions and Answers on COVID-19, (consulted on 11/05/2020), <https://www.oie.int/en/scientific-expertise/specific-information-and-recommendations/questions-and-answers-on-2019-novel-coronavirus/>
8. Considerations for sampling, testing, and reporting of SARS-CoV-2 in animals, (consulted on 11/05/2020), https://www.oie.int/fileadmin/Home/eng/Our_scientific_expertise/docs/pdf/COVID-19/Sampling_Testing_and_Reporting_of_SARS-CoV2_in_animals_final_7May_2020.pdf
9. Cohen J. (2020). From mice to monkeys, animals studied for corona virus answers. *Science*, Vol. 368, Issue 6488; pp. 221-222 <https://science.sciencemag.org/content/368/6488/221>
10. CDC, Corona virus Disease 2019 (COVID-19)—pets & other animals (consulted on 29/05/2020) <https://www.cdc.gov/coronavirus/2019-ncov/daily-life-coping/positive-pet.html>

List of NADEN Centres, Principal Investigators and Co-Principal Investigators

Sl.No	State	Center	Principal Investigator	Co-Principal Investigator	Correspondence Address
1	Andaman & Nicobar Islands (ICAR)	Port Blair (ICAR)	Dr. Jai Sunder Mob: 09434281840 Email: jaisunder@rediffmail.com	Dr. T. Sujatha Mob: 9531818976 Email: drsujathaars@rediffmail.com Dr. Debasis Bhattacharya Mob: 8420736888 Email: debasis63@gmail.com	ICAR-Central Island Agricultural Research Institute, Port Blair, Andaman & Nicobar Islands -744105 Ph: 03192-250436, 03192-245983 Fax: 03192-251068
2	Andhra Pradesh	Vijayawada	Dr. L. Ratna Kumari Joint Director(AH) Mob: 09989998027 Email: ratnakumarilam@yahoo.com	Dr. P. Shanila Kumari Mob: 7032443365 Email: shanilapvallapli@gmail.com Dr. M. Sandhya Mob: 8309683137 Email: dmsubrivijayawaada@gmail.com	Veterinary Biological & Research Institute, Veterinary Hospital Campus, Labbipet, Bandar Road, Vijayawada – 520010 Andhra Pradesh Mail: jdvbrvijayawada@gmail.com; Ph: 040-23376016 Fax: 040-23376016
3	Arunachal Pradesh	Itanagar	Dr.Badal Biswakarma Mob: 8638089688 Email: biswkarma@123gmail.com	Dr. Gyamnya Baki Garam, Mob: 8471812963 Email: niyango@gmail.com	Disease Investigation Laboratory, Department of Animal Husbandry & Veterinary, Nirjuli, Itanagar-791109 Ph:0360-2257174
4	Assam	Guwahati	Dr. Durlav Bora Mob: 9435594808, 9954533400 Email: drdpbora@gmail.com	Dr. Arfan Ali Mob: 9101240350 Email: arfan74@gmail.com Dr. Abhijit Deka Mob: 7005578946 Email: docabhijit93@gmail.com Dr. Amitav Chakraborty Mob: 7042736521 Email: vet.amitav@gmail.com	Animal Husbandry & Veterinary Department Government of Assam Khanapara, Guwahati 781022 Ph: +917042736521, +919435106512
5	Bihar	Patna	Dr. Pankaj Kumar Mob: 8084679866 Email: pankajimmunol@gmail.com	Dr. Purushottam Kaushik Mob: 9431260790 Email: drkaushikvet@gmail.com Dr. Pallav Shekhar Mob: 9431060262 Email: Shekhar.medivri@gmail.com	Bihar Veterinary College Campus, Patna-800014, Bihar Email: basu.edn@icar.gov.in
6	Chhattisgarh	Raipur	Dr. Anjana Naidu (Deputy Director) Mob: 9826151076 Email: anju202052@gmail.com	Dr. Varsha Sharma (Veterinary Assistant Surgeon) Mob: 9826650630 Email: varsha14870@gmail.com Dr. Aparna Patel Mob: 9977222214 Email: drpatel7777@gmail.com	Office of the Disease Investigation Laboratory, Animal Husbandry & Veterinary Department, Gau seva aayog campus, in front of GuruTeg Bahadur Uduan GE Road, Govt. of Chhattisgarh Raipur-492001 Gmail: ddvs-sdilrpr.cg@gov.in
7	Goa (ICAR)	Goa (ICAR)	Dr. Susitha Rajkumar Mob: 9502416520 Email: drsusitharajkumar@gmail.com	Dr. Shirish. D Narnaware (Senior Scientist) Mob: 9414052404 Email:Shirish.Narnaware@icar.org.in Dr. Niceta Cunha Costa Mob: 9673584268 Email: niceta.cunhacosta@gmail.com	ICAR-Central Coastal Agricultural Research Institute, Indian Council of Agricultural Research, Ela, Old Goa – 403402, India Ph: 0832-2284678/79 Fax: 0832-2285649
8	Gujarat	Ahmedabad	Dr. Amit Kanani Mob: 9824021874 Email: amit_kanani@hotmail.com	Dr. Dixeeta bhadarka Mob: 6355207010 Email: dixeetabhadarka@gmail.com	Office of the Deputy Director of Animal Husbandry, FMD Typing Scheme, 1st floor, Wing A, Pashupalan Sankul, Makarba Poultry campus, Makarba Ahmadabad – 380015, Gujarat Ph: 079-26304423 Fax: 079-2630442
9	Haryana	Hisar	Dr. Rajesh Khurana Mob: 9416107869 Email: khurana.rajesh866@gmail.com	Dr. Renu Gupta Mob: 9812351344 Email: drenugupta@gmail.com Dr. Ramesh Kumar Mob: 9416435317 Email: rameshkdr@gmail.com Dr. Dinesh Mittal Mob: 9996643737 Email: mittalvet@luvas.edu.in	Veterinary Public Health and Epidemiology, Lala Lajpat Rai University of Veterinary and Animal Sciences (LUVAS), Hisar, -125 004, Haryana E-Mail: dg.ahd@hry.nic.in
10	Himachal Pradesh	Shimla	Dr. Upendar Kumar Mob: 8219216439 Email: dd-epid-hp@nic.in	Dr. Asha Kiran Mob:9817051951, 8988151951 Email: ashakm71@gmail.com, dd-epid-hp@nic.in	State Veterinary Hospital Complex Near lift, Cart road Shimla-171001 Himachal Pradesh Ph./Fax +91-177-2650938
11	Jammu and Kashmir	Srinagar	Dr. Inam Mob : 9797826500 Email : director@jkanimalhusbandry.net, jdiahbp@jkanimalhusbandry.net	Dr. Tariq Pervez Mob: 9469741267 Email: director@jkanimalhusbandry.net	Joint Directorate of Institute of Animal Health and Biological Products Zakura Srinagar Jammu & Kashmir -190024 Ph: 0194-2455091 Fax: 0194-2470518
12	Jharkhand	Ranchi	Dr. Rajesh Kumar Mob:099396 12048 Email: iahp.kankeranchi@yahoo.com	Dr. Shreya Sinha Mob: 9199739351 Email: drshreya1976@gmail.com	Director Institute of Animal Health & Production Kanke, Ranchi-834006 Gmail: iahpkankeranchi@gmail.com
13	Karnataka	Bengaluru	Dr. Asha Mayanna Mob: 08494974621 Email: ashamayanna@gmail.com	Dr. Shivaraj Murag Mob: 09448128283 Email: shivaraj.murag@gmail.com Dr. N. Somasekara Mob: 9845752466 Email: jdeahvs@gmail.com	KVA&FSU Institute of Animal Health & veterinary biologicals, Hebbal Bangalore-560024 Gmail: jdrdiahvb@gmail.com
14	Kerala	Thiruvananthapuram	Dr. Sanjay Devarajan Mob: 9946558727 Email: sanjudee70@yahoo.com	Dr. Soumya Vijayakumar Mob: 9072241650 Email: soumyavijayakumar@gmail.com	Disease Investigation Officer State Institute for Animal Diseases Palode,Pacha(PO), Thiruvananthapuram

					Kerala - 695562 Principal Investigator(NADEN - Kerala) Mob: 9946558272, 8547898727 Gmail: nadenkerala@gmail.com
15	Madhya Pradesh	Bhopal	Dr. Jayant Tapase Mob: 9425022786 Email: jstapase07@gmail.com	Dr.Jha Sai Kiran Mob: 9425022786	Chief Disease Investigation Officer, State Animal Disease Investigation Laboratory, Veterinary Hospital Campus, Jahangirabad, Bhopal-462 008 Ph: 0755-2767583 Fax: 0755-2767583 Gmail: statedilab07@rediffmail.com
16	Manipur	Imphal	Dr. Sabitri Maibam Mob: 9774014506 Email:maibamsabitri@gmail.com	Dr.C. H. Amita Devi Mob: 8575465385 Email:muhindrovvet@gmail.com	Disease Investigation Laboratory, Directorate of Veterinary & Animal Husbandry Services, Imphal-795001 Ph: 0385-2447009 Fax: 0385-2454928
17	Meghalaya (ICAR)	Barapani (ICAR)	Dr. Samir Das Mob: 08730856556 Email: drsamirvph@yahoo.com	Dr. I Shakuntala Mob: 09436166508 Email:ishakuntala92@yahoo.com Dr. Sandeep Ghatak Mob: 09436380401 Email: ghataksnd@rediffmail.com Dr. Rajkumari Sanjukta Mob: 09402196908 Email: rajkumari.sanjukta@gmail.com Dr. K. Puro Mob: 09402134165 Email: akulepuro@rediffmail.com Dr. A.A.P Milton Mob: 8650918630 Email: vetmilton@gmail.com	Division of Animal Health ICAR Research Complex for NEH Region, Umiam, Barapani-793 103 Ph: 0364-2570071 Fax: 0364-2570071 Gmail: dioahvety@bsnl.in
18	Maharastra	Pune	Dr. Y.A. Pathan Mob: 9423324609 Email: dis.pune7@gmail.com	Dr. Madhuri Dharmadhikari Deputy Commissioner of Animal Husbandry Disease Investigation Section, Aundh, Pune-411067 Ph: 7768069912 Email:madhuri.dharmadhikari@gmail.com	Joint commissioner of Animal husbandry WRDDL, Disease Investigation section Aundh,Pune 67 411009 Ph:02025692135 Fax:02025691474 Email: dis.pune7@gmail.com
19	Mizoram	Aizawl	Dr. C. Neithangpuii Mob: 8974206212 Email: puiihualngo@gmail.com	Dr. Tapan Dutta Mob: 09862335294 Email: tapandutta@rediffmail.com Dr. Zotei Mob:8415968680	Directorate of Animal Husbandry & Veterinary, Government of Mizoram, Khatla, Aizawl-796001 Fax: 0389-2333234 Gmail: dilabaizawl@gmail.com
20	Nagaland	Kohima	Dr. Amenla Walling (Deputy Director) Mob: 9612815298 Email: amenlaw2@gmail.com	Dr. Vikulie Mezhatu (VAS) Mob: 8974538691 Email: vikuovet@gmail.com Dr. Khriesamhazou Rhetso (VAS) Mob: 9362611408 Email: Rhetso.k@gmail.com	Animal Husbandry & Veterinary Department, Government of Nagaland, Kohima -797001
21	Odisha	Cuttack	Dr. Jagadish Mohanty	Dr. Jyoti Ranjan Biswal Mob: 9439365472 Email: j.r.biswal2000@gmail.com	Animal Disease Research Institute, Phulnakhara, Cuttack-754 001, Odisha Ph: 0671-2356130 Gmail: adri.odisha@gmail.com
22	Puducherry	Puducherry	Dr.R.Maria Mob: 9443539056 Email: mariamaria14@rediffmail.com	Dr. M. Mohan Mob: 09442486347 Email: mohanpsp@yahoo.co.in	Department of Animal Husbandry & Animal welfare, Government of Puducherry, Maraimalai Adigal Salai, Puducherry-605001 Ph: 0413-2206888
23	Punjab	Ludhiana	Dr. M S Bal (3.12.2021 onwards) Mob: 814661040,9779540624 E-mail: inchargeadre@gadvasu.in	Dr. Gwsimram Filia Mob: 9815804281 Email: harpalfilia@rediffmail.com Dr. Mohinderpal Mob: 9780263576 Email: nrdd12001@mail.com	Animal Disease Research Centre, Guru Angad Dev Veterinary and Animal Sciences University Ludhiana-141004, Punjab Ph: 0161-2414030, 0181-2242335 Fax: 0161-2400822, 0161-24008224
24	Rajasthan	Jaipur	Dr. Khadri	Dr. Lenin Bhatt Mob: 9829791073 Email: leninbhatt@gmail.com	State Disease Diagnostic Centre Department of Animal Husbandry Government of Rajasthan, Panch Batti, Jaipur-302001 Email: ddpathologistraj@gmail.com Fax: +91 141 2374617
25	Sikkim	Tadong	Dr.Seema Rai Mob: 9775136696, 7029635446 Email: seemaravet87@gmail.com	Dr. Tashila Pintso Bhutia (VO - DI) Mob:8918214617 Email: drtashipintosobhutia@gmail.com	Office of Chief Epidemiology Officer, Veterinary Epidemiology Centre (above TNLDA buildings), Veterinary Polyclinic Campus, Saidapet, Chennai-600035. Email: admastn2017@gmail.com;
26	Tamil Nadu	Chennai	Dr.A.Sundaresan Mob: 9884001987 Email id: admastn2017@gmail.com	Dr. K.Karunanidhi Mob: 8056360220 Email: admastn2017@gmail.com	Office of Chief Epidemiology Officer, Veterinary Epidemiology Centre (above TNLDA buildings), Veterinary Polyclinic Campus, Saidapet, Chennai-600035. Email: admastn2017@gmail.com;
27	Telangana	Hyderabad	Dr.G. Sunitha Mob: 9989932774 Email id: jdbp.vbri@gmail.com	Dr. Devender Rao Mob: 7893552900 Email: deshinenid@gmail.com	AICRP ADMAS UNIT, HYDERABAD Telangana State Veterinary Biologicals & Research Institute, Shanthinagar, Hyderabad – 500 028
28	Tripura	Agartala	Dr.Jyotirmoy Roy Mob : 9436124525 Email:sdil.ardd@gmail.com/jyotibu166@gmail.com	Dr. Bina Saikia Mob: 8731069076 Email: binasaikia17@gmail.com	State Disease Investigation Laboratory, P.O.Abhoynagar, Agartala, Tripura West Pin-799005 Gmail: Ahtripura5@gmail.com

29	Uttar Pradesh (ICAR)	Izatnagar (ICAR)	Dr. S. Nandi Principal Scientist Mob: 06395347105 Email: sukdeb@yahoo.in snandi1163@gmail.com	Dr. Vinodh Kumar Mob: 8859784853 Email: vinodhkumar_rajendran@yahoo.com, vinodhkumar.rajendran@gmail.com	Centre for Animal Disease Research and Diagnosis (CADRAD), ICAR-Indian Veterinary Research Institute, Izatnagar – 243122, Bareilly, U.P Gmail:
30	Uttarakhand	Dehradun	Dr. Sunil Kumar Awasthi Mob: 9368036229, 7579407622 Email: Dr. Devendra Sharma (Joint Director) Mob: 8937001273, 9410903526 Email: devendras642@gmail.com	Dr. Chetna Dhapola Mob: 7983962752 Email: chetnadhapola77@gmail.com Dr. Prasoon Dubey Mob: 9411558046 Email: prasoon.dubey@rediffmail.com	Animal Husbandry & Veterinary Department, Government of Uttarakhand, Kedar Puram, Dehradun-248115 Ph: 0135-2450154 Fax: 0135-2450083 Email: kvk.Dehradun@icar.gov.in/ dirahuk@gmail.com
31	West Bengal (ICAR)	Kolkata (ICAR)	Dr. P.S. Banerjee Mob: 9410431207 Email: partha.banerjee@icar.gov.in	-	Collaborating unit of AICRP ADMAS, ICAR-Indian Veterinary Research Institute, Eastern Regional Station, No-37 Belgachia Road, Kolkata-700 037 Tel: 033-25582965 Fax: 033-25565725 Mail: dahvswb@darahwb.org





ICAR-National Institute of Veterinary Epidemiology and Disease Informatics (ICAR-NIVEDI)

P. B. No.6450, Yelahanka, Bengaluru-560064

Phone: +91-80-23093111, Fax: +91-80-23093222, E-mail: director.nivedi@icar.gov.in