

# Sero-Survey in Karnataka State

Summary, Design, and Statistical Methodology



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# Plan of the Talk

- **Part I**

- Survey Design and Implementation
- Probability Model and Statistical Methodology
- Summary of Findings.

- **Part II**

- Nuances of the methodology
- C-optimal Design
- Worst Case Design

# Goal: To Estimate

- **(IgG-Antibody)**--the proportion of people who already have had SARS-CoV2 infection;
- **(Active Infection)**--the proportion of people current active infection;
- **(COVID-19 Burden)**--the prevalence of COVID-19 in the general population of Karnataka state.

# Survey: Units and Implementation

## Geographical Spread

- Each of the 29 districts (other than Bengaluru-Urban) was a unit.
- Each zone of BBMP is considered as a unit;
- There will be total of 38 units. (30 districts+ 8 zones)

## Within Unit

- Locations around District Hospital, Primary/Community Health Care centres were surveyed.
- Counsellors from ICTC and Health Care workers: conducted the survey.
- Survey used existing public health infrastructure.

# Design of Sample at Each Unit

	Risk Subgroups	Samples*	Tests done
<b>Low Risk</b>	Pregnant women presenting for ANC clinic	72	RTPCR and IgG
	Persons attending the outpatient department in the hospitals/ attendees	72	
<b>Moderate Risk</b>	Bus conductors/Auto drivers	28	RAT, RTPCR** and IgG
	Vendors at the vegetable markets	28	
	Healthcare workers	30	
	Individuals in containment zones	28	
	Markets, Malls, Retail stores, Bus stops, Railway stations	30	
<b>High Risk</b>	Elderly (over 60)	72	
	Persons with co-morbid condition	72	
		<b>432</b>	

\*-Divided uniformly across Health Facilities; \*\*-only if RAT was negative.

# Sampling Design

## Participant in each group

- Chosen via systematic sampling and Consent was sought.  
Excluded:
  - if under 18 years of age
  - COVID-19 positive patient
  - Unable to provide informed consent or Not willing to provide sample for test.

## Samples

- Blood was drawn for Antibody test, post centrifugal processing sent to designated IgG Lab.
- Rapid Antigen Test done on spot for Moderate and High Risk group participant.
- Swab was sent to designated Reverse transcription polymerase chain reaction labs.

**290 Health facilities across the state.**

# Data from Survey

## Collection and Compilation

- Surveyor Entered on an specially designed Survey app: participant details.  
( Each participant identified by a Unique S.R.F. id )
- IgG lab sent data to State government.
- RTPCR labs entered directly onto ICMR portal; were downloaded by State government.

## Analysis

- Preparation of one line list for participants (with IgG, RAT, RTPCR results).
- Different Test patterns
- Joint Estimation of IgG and Active Infection



# Model

Table of States and Nominal Test Responses  $M(s,j)$

State of Individual		RAT	RTPCR	IgG
		$j = 1$	$j = 2$	$j = 3$
$s = 1$	Active infection but no IgG antibodies	1	1	0
$s = 2$	IgG antibodies present but no evidence of active infection	0	0	1
$s = 3$	Simultaneous presence of active infection and IgG antibodies	1	1	1
$s = 4$	Neither active infection nor IgG antibodies	0	0	0

# Sensitivity and Specificity

**Sensitivity:** probability of a positive test given that the patient has the disease.

**Specificity:** probability of a negative test given that the patient is well.

$\sigma(m, j)$	RAT	RTPCR	IgG
	$j = 1$	$j = 2$	$j = 3$
Specificities ( $m = 0$ )	0.975	0.97	0.977
Sensitivities ( $m = 1$ )	0.5	0.95	0.921

## Model Needs

- Test pattern taken by individual (E.g. No RAT at low Risk)
- Adjust for Sensitivity and Specificity of each test(s)
- Design effect - Bias in Sample.

# Our First Crude Estimates

# Model- Maximum Likelihood Estimate

- The likelihood at a strata, say  $D$ -district is then given by

$$L(p; t(D), y(D)) := \prod_{n:n \in D} \left( \sum_{s=1}^4 p_s q(y(n)|s, t(n)) \right).$$

- The mapping  $p \mapsto \log L(p; t(D), y(D))$  is concave
- Numerical, iterative, gradient descent procedure is used to arrive at the maximum likelihood estimate for  $p$  in this stratum,

$$\hat{p}(D) = (\hat{p}_1(D), \hat{p}_2(D), \hat{p}_3(D), \hat{p}_4(D)).$$

- IgG-Antibody-Estimate  $\hat{p}_2(D) + \hat{p}_3(D)$ ,
- Active prevalence-Estimate  $\hat{p}_1(D) + \hat{p}_3(D)$ ,
- Total prevalence[ past and active infection]-Estimate -  $\hat{\phi}(D) := \hat{p}_1(D) + \hat{p}_2(D) + \hat{p}_3(D)$ .



# Model-Confidence Interval

- The confidence interval estimations come from the Fisher information matrix associated with this model and a design effect of 3.
- For protocol and logistical reasons, the test patterns differ across the participants.
- So the Fisher information matrices for the different test patterns are averaged in proportion to the observed test patterns. This can be interpreted as a hybrid observation/Fisher information matrix.
- The inverse of this matrix provides an estimate of the covariance of  $\hat{p}(D)$ .

# Karnataka

- The Karnataka estimates are obtained after weighting for district populations:

$$\hat{p} = \sum_D w(D) \hat{p}(D),$$

where  $w(D)$  is the fraction of Karnataka population that lives in district  $D$ .

- Assuming independence across districts, the covariance is

$$\sum_D w(D)^2 \text{cov}(\hat{p}(D)).$$

# Karnataka Numbers

## Seroprevalence of IgG and Active infection in Karnataka state.

Type	Samples	%-IgG against SARS-CoV2	%-Active Infection of COVID-19	%-Prevalence of COVID-19
Crude	15939	2565/15939= 16.1	2363/14132=16.7	4582/15939=28.7
Adjusted	15939	15.4	12.2	26.1
Weighted Adjusted	15624	16.4 (15.1--17.7)	12.7 (11.5--13.9)	27.3 (25.7--28.9)

## Both IgG and Active infection in Karnataka state.

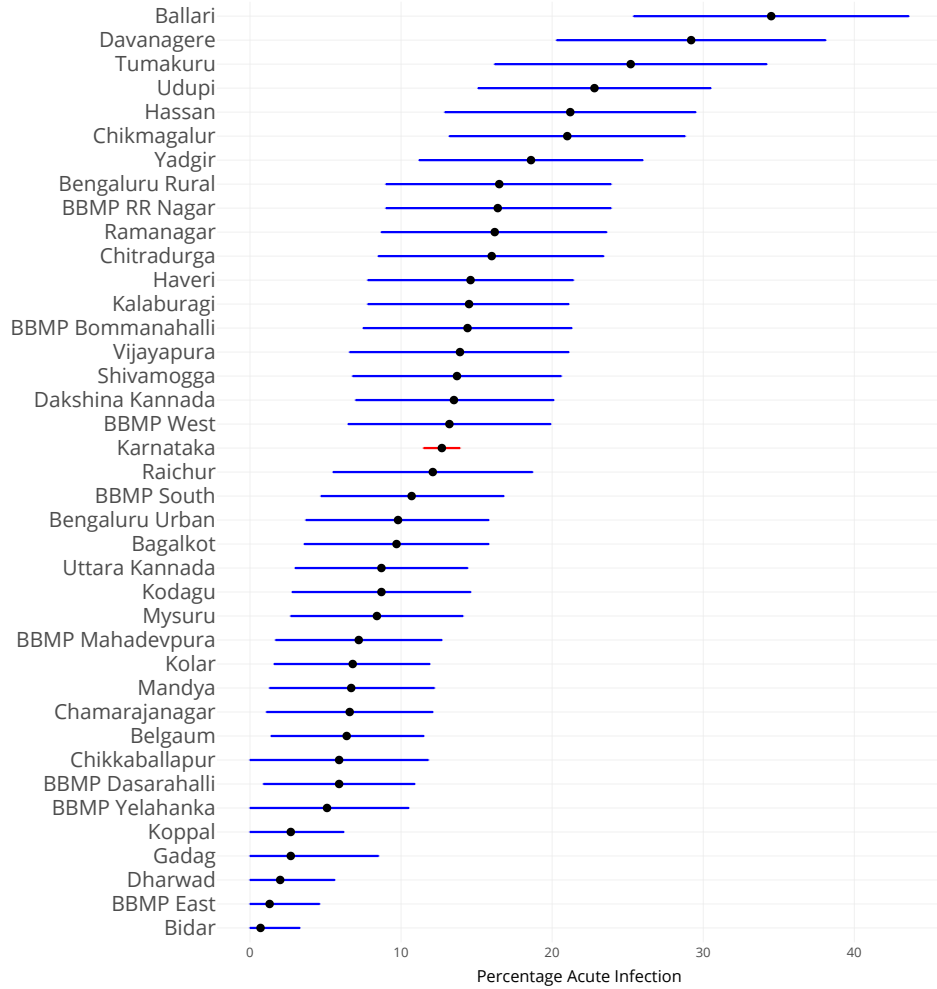
$$16.4 + 12.7 - 27.3 = 1.8$$



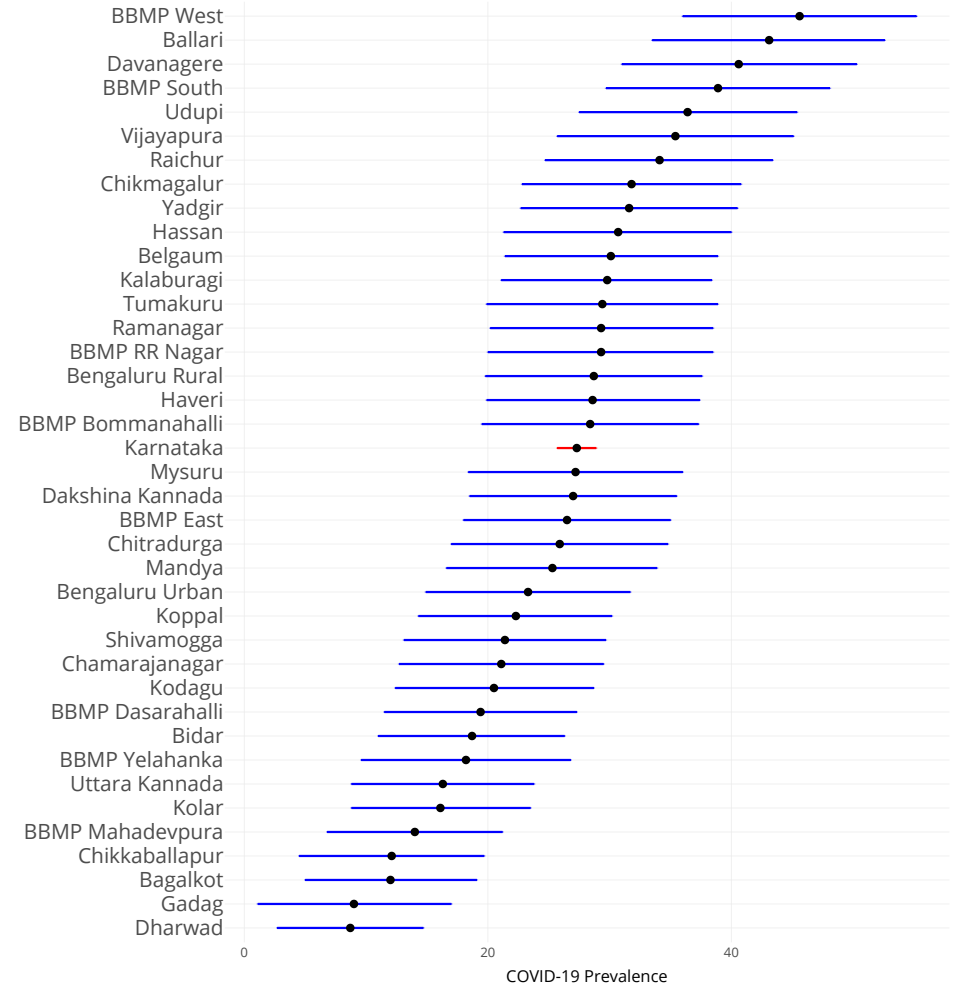
### 95% Confidence Intervals for Percentage IgG Prevalence



95% Confidence Intervals for Acute Infection



95% Confidence Intervals for COVID-19 Prevalence



# Epidemiological Metrics

## Estimated Infection (EI)

$$EI = \text{COVID-19 prevalence} \times \text{Population}$$

## Case to Infection Ratio (CIR)

$$CIR = \frac{EI}{\text{Reported Cases up to 2nd September}}$$

## Past Infection (PI)

$$PI = \text{IgG Prevalence} \times \text{Population}$$

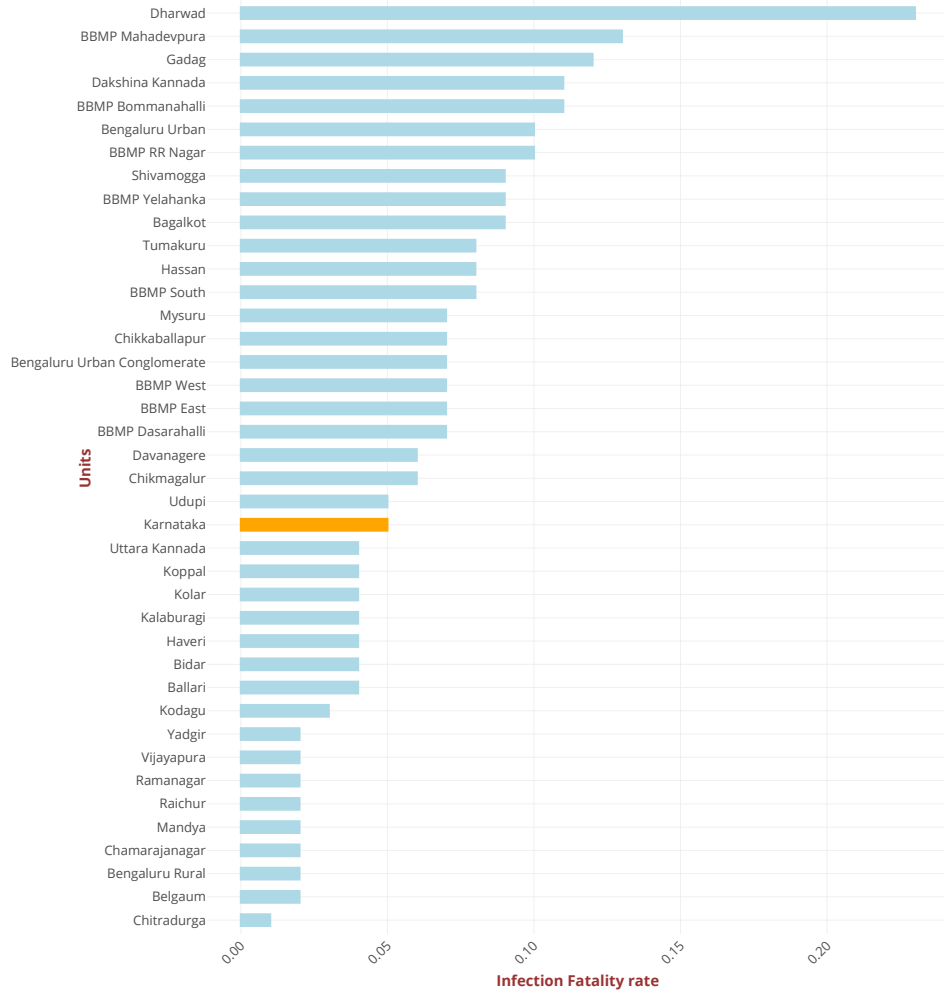
## Infection Fatality Rate (IFR)

$$IFR = \frac{\text{Fatalities Reported up to 2nd September}}{\text{Fatalities Reported up to 2nd September} + PI}$$

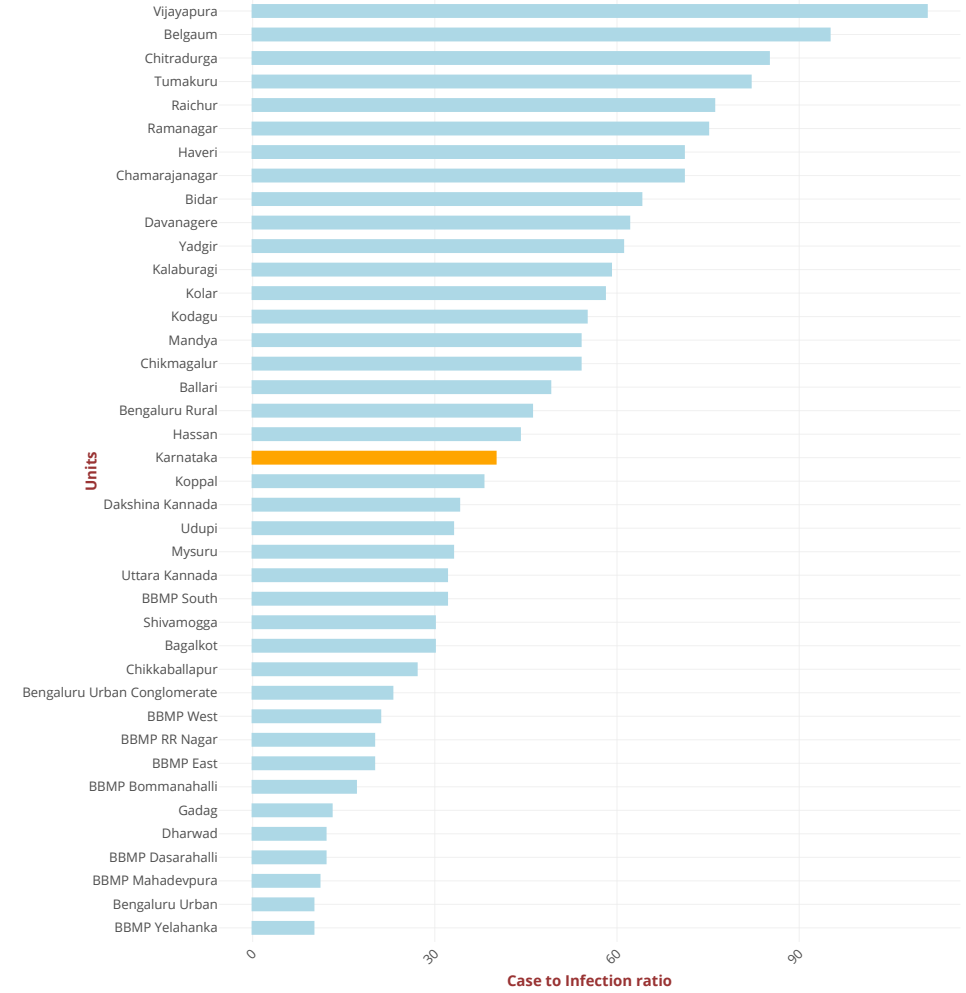
# Karnataka Estimated Numbers

Cases up to 16/09	Estimated total prevalence as on 16/09	CIR	IFR	Active case per day 02-16	Estimated Active Infection
484954	19321334	1:40	0.05%	98028.5	8988313

**Infection Fatality Rate**



**Case to Infection Ratio**



# Karnataka: 16.7 % IgG and 0.05 % IFR

## High IgG and Low IFR

- The district has had surge in cases in the past
- Underreporting of COVID-19 deaths is a possibility

## Low IgG and Low IFR

- District will have surge in cases in the near future
- Low IFR can mean that either there are fewer deaths or there is underreporting.

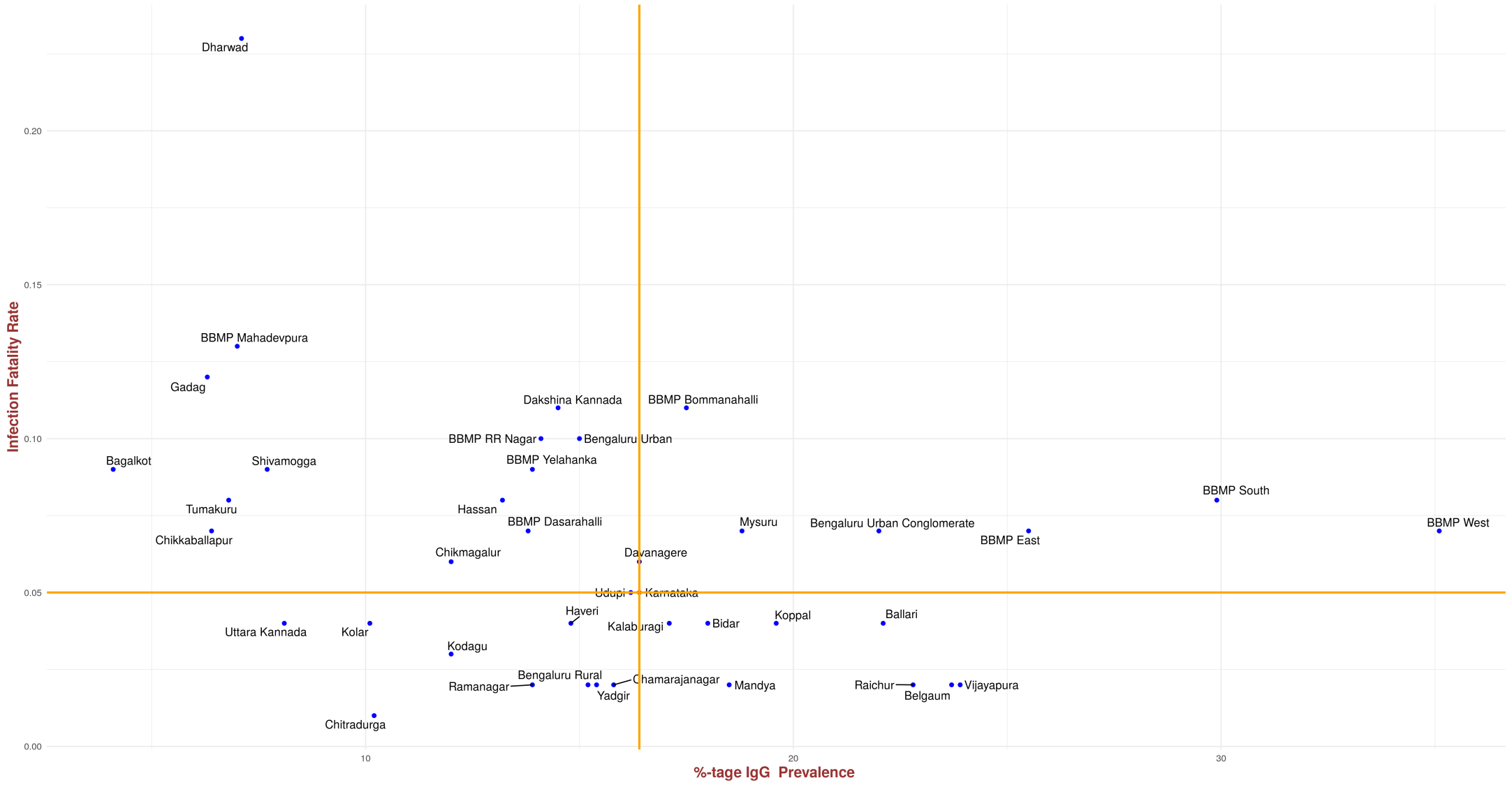
## High IgG and High IFR

- The district has had surge in cases in the past
- The reporting of deaths in the district is relatively reliable

## Low IgG and High IFR

- The district is yet to see surge in cases
- High proportion of deaths due to COVID-19 in the early phase of outbreak

**%-tage IgG Prevalence vs Infection Fatality Rate**



## Karnataka: 16.7 % IgG, 1:40 CIR, and 0.05 % IFR

### High IgG, High CIR, Low IFR

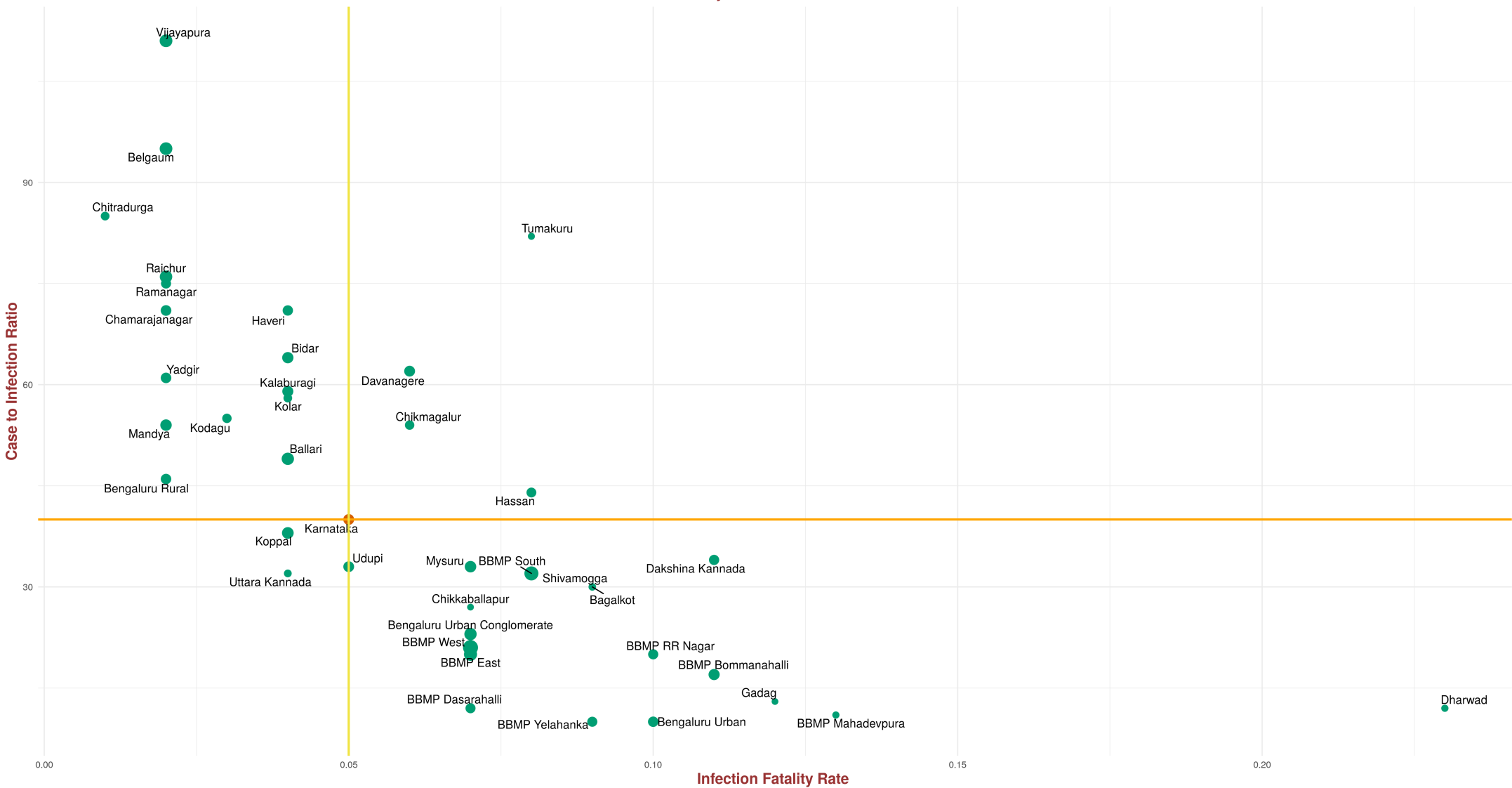
- District has gone through a surge.
- High IgG and high CIR is a clear indication that cases or areas of circulation have been missed.
- High IgG with Low IFR means that there is underreporting of deaths

### High IgG, Low CIR, High IFR

- Despite going through surge in cases in the past, district seems to be detecting cases and areas of circulation
- High IgG with high IFR means that there is reliable reporting of deaths



Infection Fatality Rate vs Case to Infection Ratio



# Survey Findings

- Estimates Across Risk Groups: Age, Gender, Sub-Categories
- RAT Sensitivity amongst : asymptomatic ~ 0.46 and symptomatic ~ 0.67
- Prediction of past and current infection from: Symptoms and Pre Existing Health Conditions
- District wise reports and impacting testing policy
- Prediction of past and current infection from: Symptoms and Pre Existing Health Conditions.  
E.g. Diarrhea and In-contact for past infection.

# Predictions

