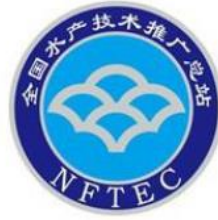




Food and Agriculture  
Organization of the  
United Nations



## FAO/China Intensive Training Course on Tilapia Lake Virus (TiLV)

Sun Yat Sen University, Guangzhou, China

18-24 June 2018

### Session 4

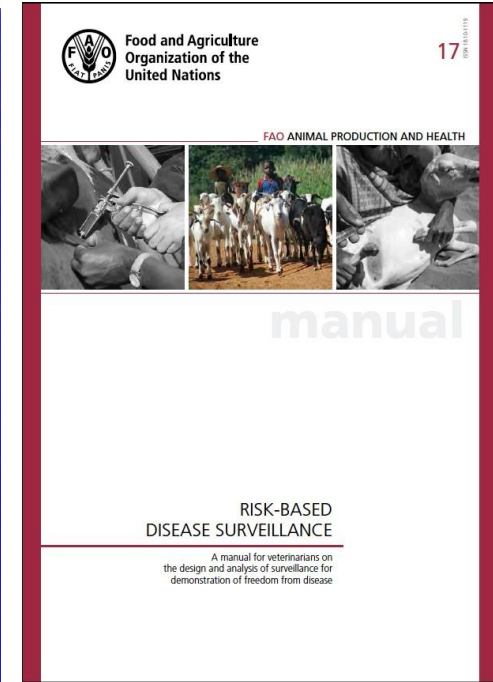
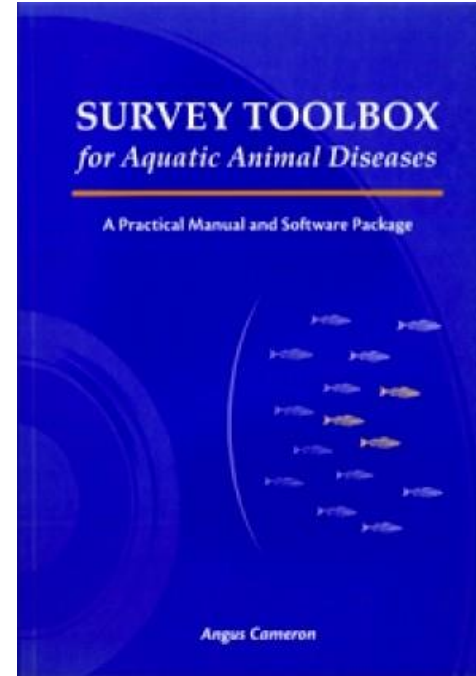
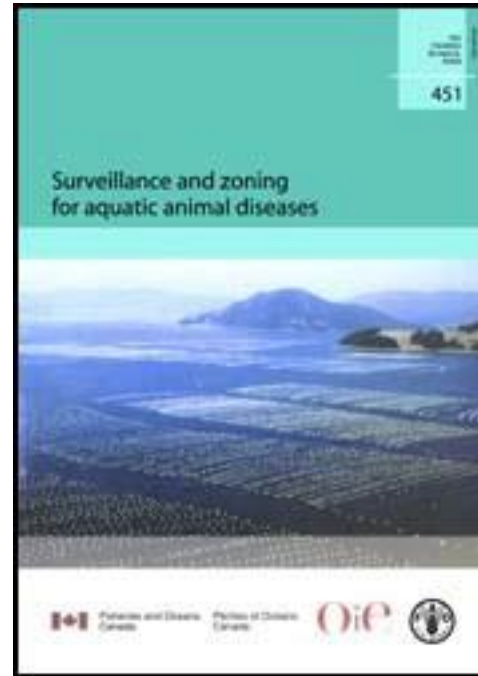
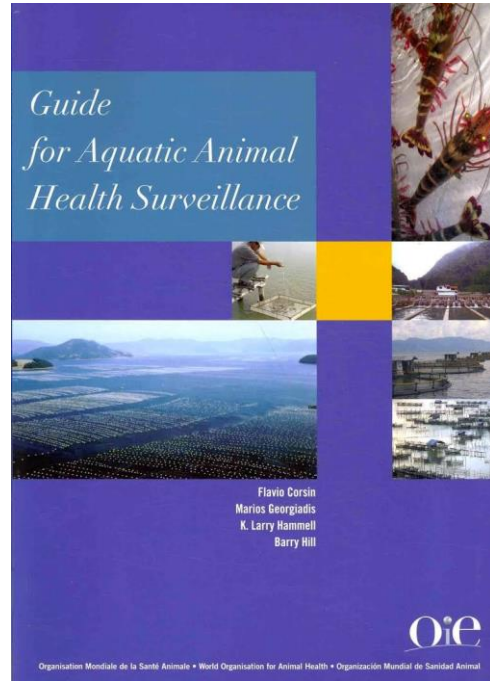
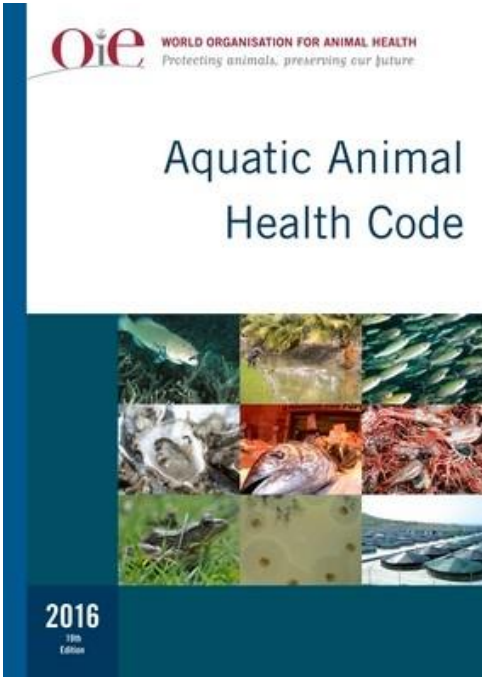
Mona Dverdal Jansen

Epidemiology and surveillance

**Principles of surveillance**

# Learning objectives

- Gain basic knowledge on surveillance systems
- Understanding the keys steps for setting up a surveillance system



# What is surveillance?

- A planned, continuous activity for collecting, collating, analysing and disseminating disease and health data information from a defined population

- OIE International Aquatic Animal Health Code:

“the **systematic series of investigations** of a given population of aquatic animals to detect the occurrence of disease **for control purposes**, and which may involve testing samples of a population”

# What is surveillance?

- A planned, continuous activity for collecting, collating, analysing and disseminating disease and health data information from a defined population

- FAO Surveillance and zoning for aquatic animal diseases:

“a **mechanism** applied to **collect and interpret data on the health of animal populations**, to accurately describe their health status with respect to specific diseases of concern”.

# Survey vs monitoring

- Survey
  - The structured collection of health information from a specified population
  - Targets only specific health problems -> can supplement surveillance.
  - A single survey rarely provides sufficient evidence for disease status
- Monitoring
  - The systematic series of investigation of a given population of aquatic animals
  - May involve testing samples of a population
  - E.g. to detect
    - Changes in the prevalence
    - Changes in geographical distribution of disease
    - Emerging diseases

# What is the overall purpose of surveillance?

- To inform stakeholders and assist in decision-making on the planning and implementation of control measures
  - Trade
  - Support safe movement at farm, zone or national level
  - Eradication
  - Prioritization of resources
  - Socio-economic impact
  - .....
- Should be cost-efficient!

# Objectives of surveillance

- Objectives defined to achieve the purpose
  - Substantiate claims of absence of infection (or disease)
  - Detect emerging (exotic) disease
  - Monitor endemic diseases (distribution, prevalence, incidence)
  - Monitor the success of intervention (e.g. eradication)
- Must be clearly stated to help surveillance system design
  - Consistency with disease situation
  - Recognition of stakeholders needs
  - Types of surveillance system



# Stakeholders

- Surveillance => a multifaceted activity => many potential stakeholders
  - Competent authority
  - Industry (aquaculture, service)
  - Consumers
  - Wildlife (organizations, public)
  - Trading partners
  - Decision makers
- Who will benefit?
- How will results be disseminated/communicated?



# Various surveillance approaches

- **Passive surveillance**
- **Active surveillance** (Targeted surveillance)
- **Risk-based surveillance**
- Sentinel surveillance
- Syndromic surveillance
- Proxy surveillance
- Post-harvest processing
- ...and more



# Passive surveillance

- Not targeted for a specific disease
  - Notifications by farmers
  - Routine diagnostic samples
- Will mainly detect disease (rather than infection only)
  - Non-requested agent investigation based on history and other laboratory findings
- Usually underestimates the prevalence of disease
- Recommended by the OIE as a first step in the surveillance effort

# Active surveillance

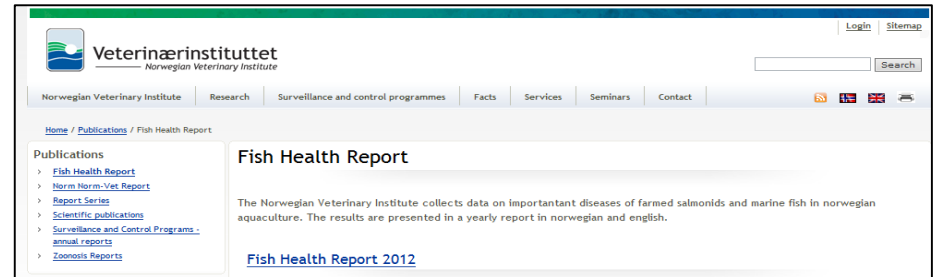
- Specifically designed sampling of a defined population
- Can achieve a “reasonably high” confidence level
- Can be designed to identify infection or disease
- Usually very resource intensive
  - People
  - Laboratory
  - Money

# Risk-based surveillance

- Most commonly recommended surveillance design (e.g. the EU)
- Aims to increase the cost-efficiency
  - Disease probability
  - Severity of consequences of disease
- Several areas of inclusion, e.g.
  - Populations
  - Sample size calculation
  - Data analyses
  - Data interpretation

# Agent/disease information

- Aetiological agent/disease
- Case definition
- Clinical signs
- Epidemiology – e.g. transmission routes
- Diagnostics – availability, validity



# Population definitions (1)

- Population of interest
  - aim to get a representative sampling of this susceptible population
  - -> affects agent entry, agent spread and likelihood of agent/disease detection
- Population structure (demography)
- Locations
- Disease histories
- Exposure
- Susceptibility

# Population definitions (2)

- Population of interest



- Target population

- population to which the conclusions (absence/presence) will be applied



- Study population

- population from which the surveillance data are gathered
- May = target population or be a subset of it

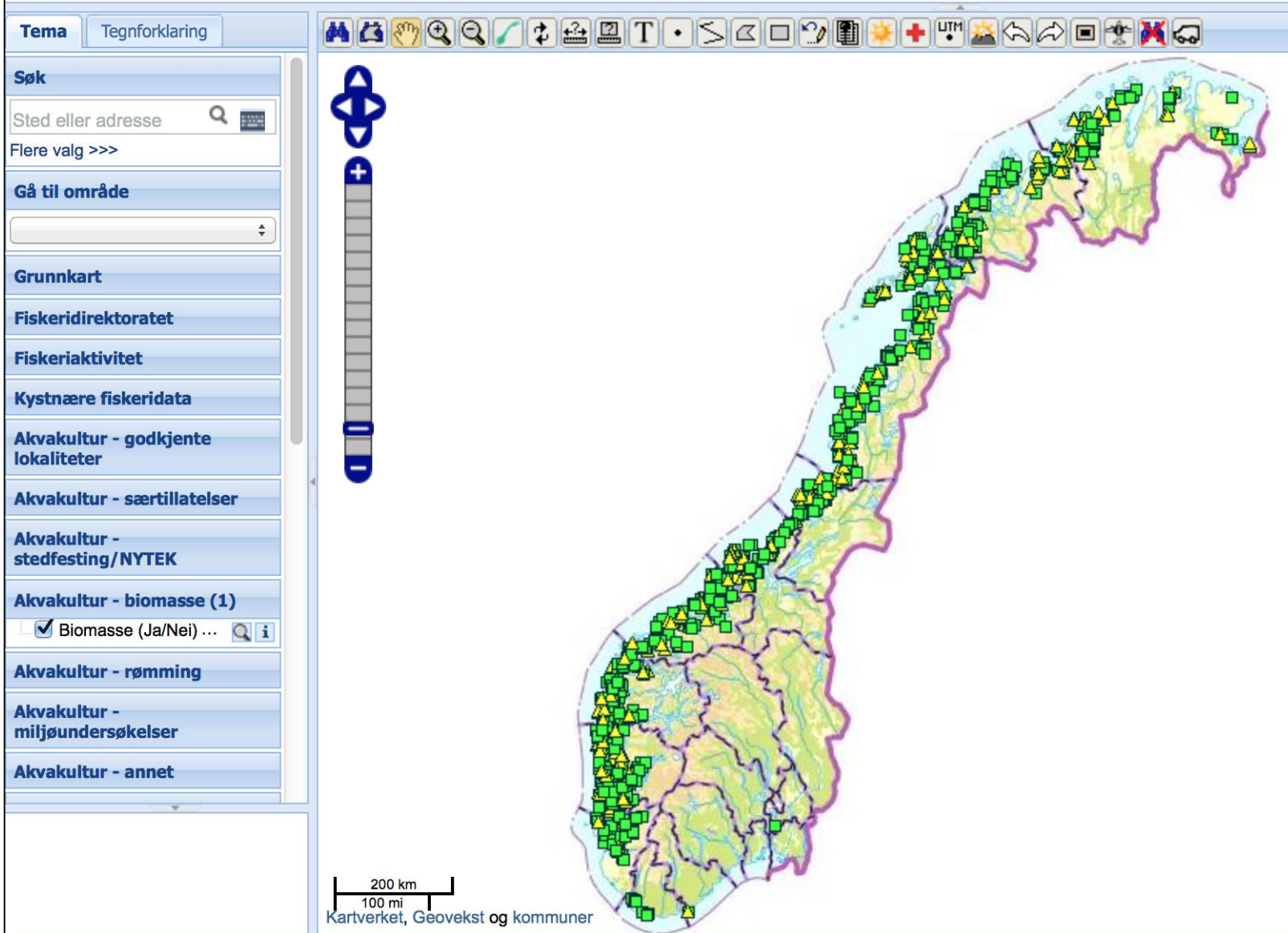


# Epidemiological unit

In the context of surveillance:

- = units selected when sampling
- E.g. animals, ponds/cages, farms, villages, districts





https://www.barentswatch.no/fiskehelse/locality/24315/2018/25

Gjøsingen uke 25 2018 | Fisk...

Fil Rediger Vis Favoritter Verktøy Hjelp

Lokalitetsnavn/id eller brønnbåt

Gjøsinga  
Gjøsinga  
Gjøsinga

**Gjøsinga**

Seiskjæra  
Kvernøya  
Måøya  
Brandsøya  
Berfjorden  
Sørkråkøya  
Nes  
Hongsand  
Kiran  
Hagadalen  
Kiransfjellet  
Skjærafjorden  
Pålodden  
Kavsodden  
Stokken  
Harbak  
Harbaks-  
Svønningsneset  
Håvikneset  
Skjøra  
Nordskjø  
Tøssvika

**0,01**  
Hunnlus per fisk  
(Lusegrense: 0,5)

**PD**  
Pankreassykdom  
Mistanke 11. juni 2018

**Lus per fisk (alle stadier) 0,09**

- Hunnlus 0,01
- Lus i bevegelige stadier 0,08
- Fastsittende lus 0

Sjøtemperatur: 10,5 °C

Lokaliteten er i den nasjonale PD-sonen

Ingen brønnbåter ved lokaliteten denne uken

**Hunnlus - utvikling i 2018**

Hunnlus per fisk 2018  
0,50

Uke 24 2017 UKE 25 | Juni, 2018

TA KONTAKT

OIE\_chapitre\_salmonid\_alphavirus.pdf - Adobe Acrobat Reader DC

Fil Rediger Vis Vindu Hjelp

Hjem Verktøy Epidemiology Secti... OIE\_chapitre\_salm... x

Logg på

1 / 14 125%

## CHAPTER 2.3.6.

# INFECTION WITH SALMONID ALPHAVIRUS

### 1. Scope<sup>1</sup>

For the purpose of this chapter, infection with salmonid alphavirus (SAV) means infection with any subtype of SAV of the genus *Alphavirus* of the family *Togaviridae*.

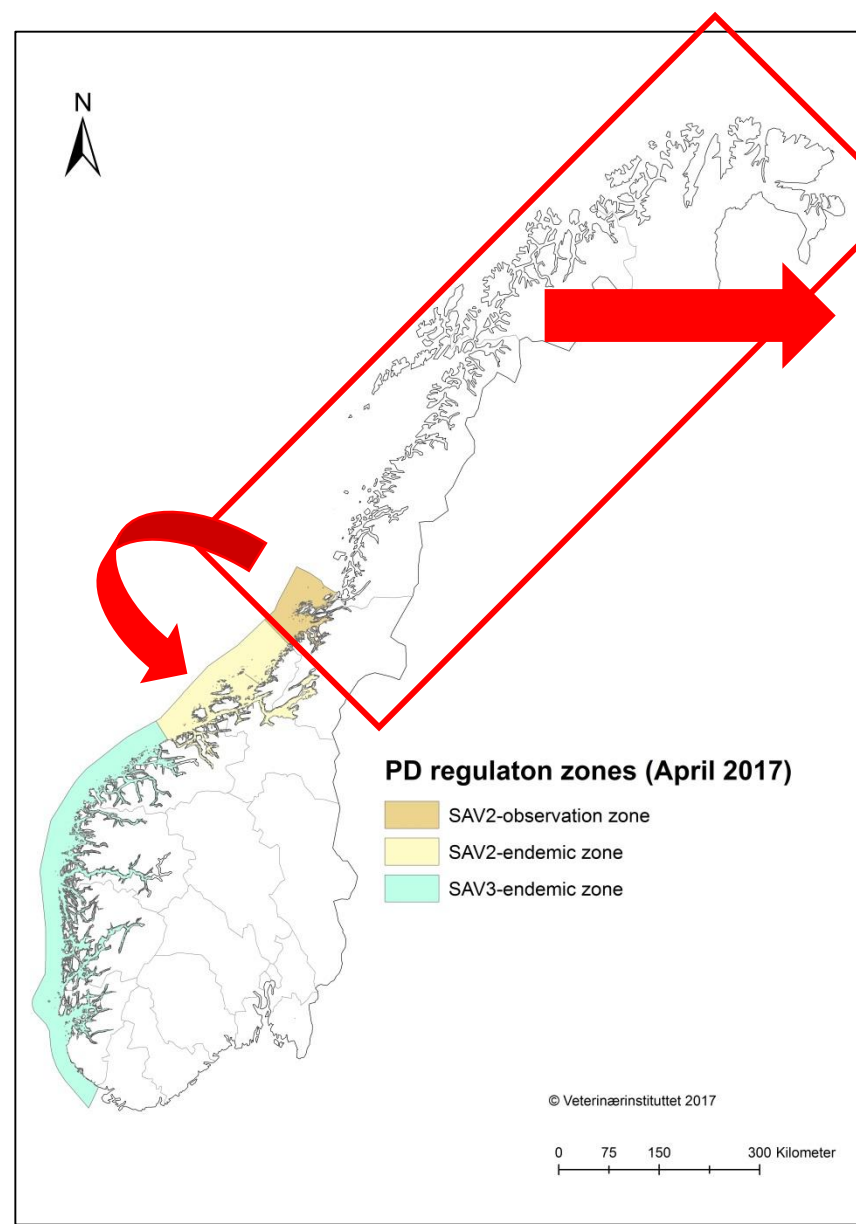
Infection with SAV may cause pancreas disease (PD) or sleeping disease (SD) in Atlantic salmon (*Salmo salar* L.), rainbow trout (*Oncorhynchus mykiss*) and brown trout (*Salmo trutta* L.) (Boucher et al., 1995; McLoughlin & Graham, 2007). The virus is horizontally transmitted, and the main reservoirs of SAV are clinically diseased or covertly infected fish (Viljugrein et al., 2009). The disease is a systemic disease characterised microscopically by necrosis and loss of exocrine pancreatic tissue, and heart and skeletal muscle changes. The mortality varies significantly, from negligible to over 50% in severe cases, and up to 15% of surviving fish will develop into long, slender fish ('runts') (McLoughlin & Graham, 2007).

### 2. Disease information

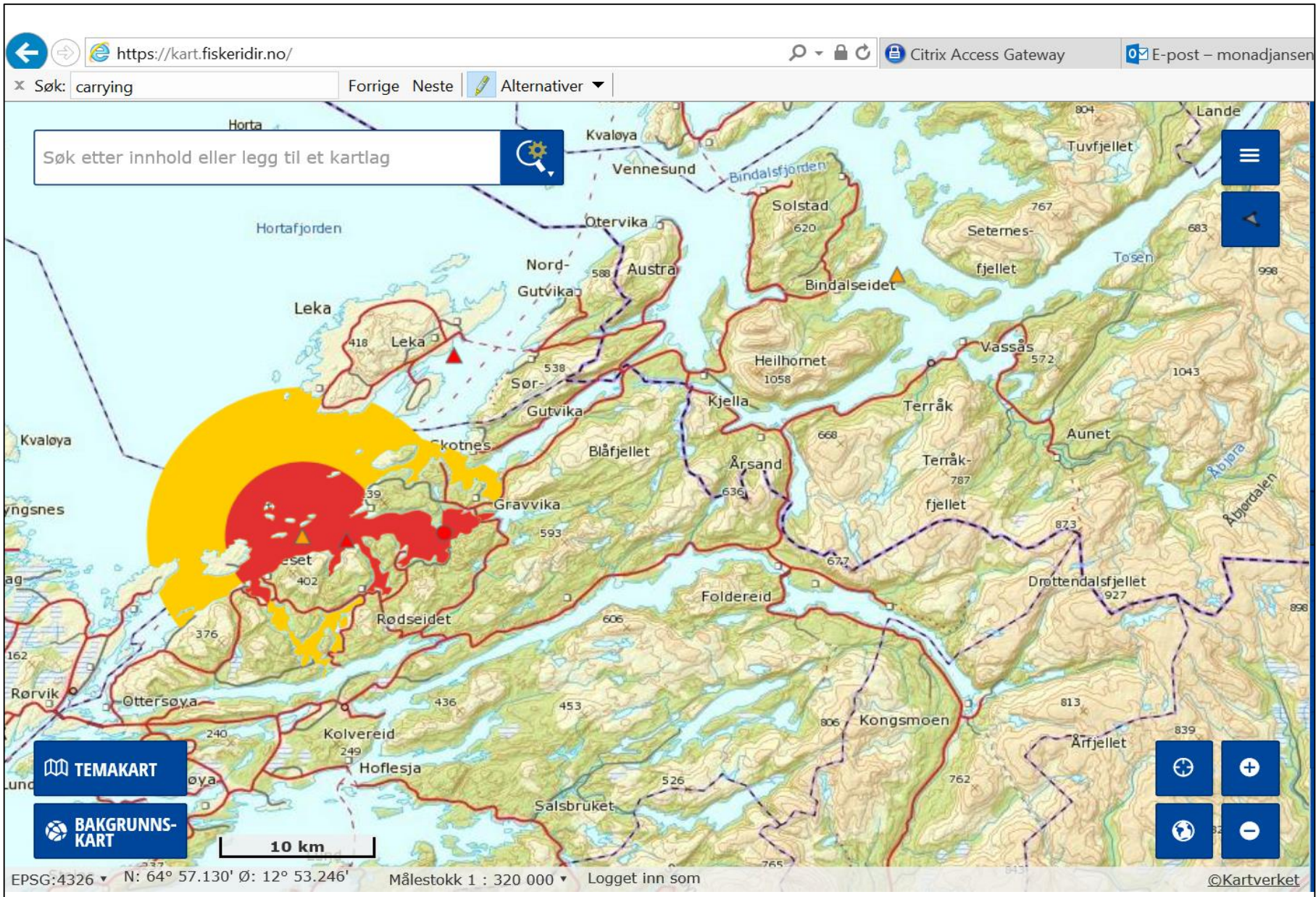
#### 2.1. Agent factors

##### 2.1.1. Aetiological agent, agent strains

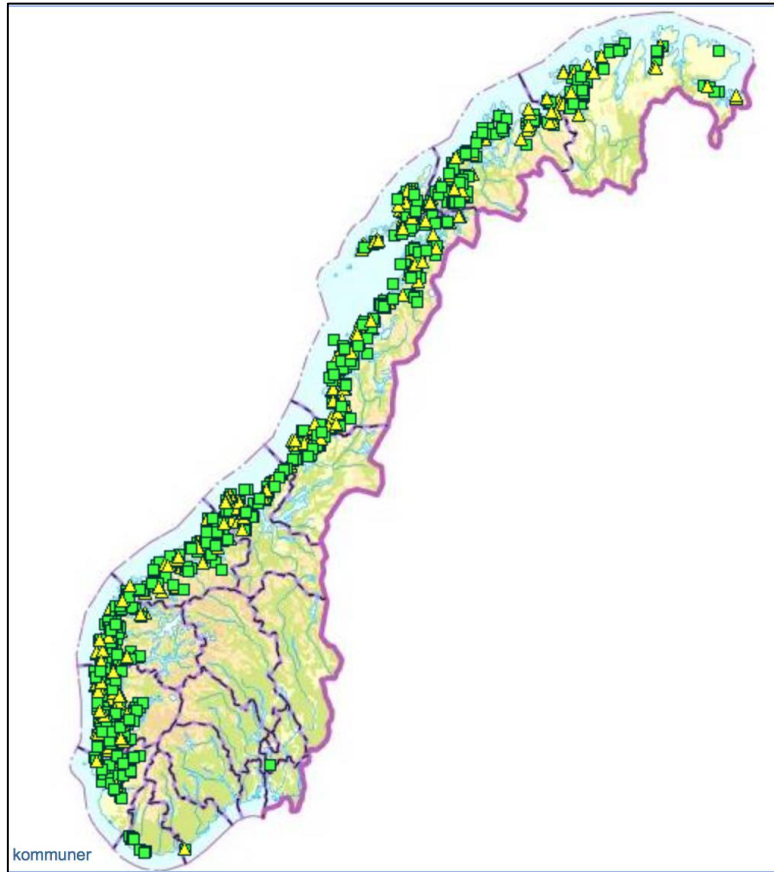
Northern zone:  
tested negative for 2  
consecutive years  
=>  
free status application



All sites:  
Risk-based sampling  
twice/year  
+  
Passive surveillance



# New surveillance system



- All sites
- Risk-based

Main purpose:

- Improved overview
- Trade

Industry bears the cost!

# Sampling issues (1)

- For surveillance in the aquaculture industry we commonly need to be able to select:
  - Fish farms - within country/zone
  - Fish within farms
- Need to determine a sample size to (randomly) sample enough farms and fish to optimize your estimate
  - not too many → waste of resources
  - not too few → won't give you the results you want



# Sampling issues (2)

- Is there access to individuals that are representative of the population?
  - Random samples are rarely possible in fish farms
    - Dead, sick, moribund, apparently healthy, healthy
    - Different ease of sampling
  - Fully representative samples are rarely achieved

# How to sample?

- Several methods available:

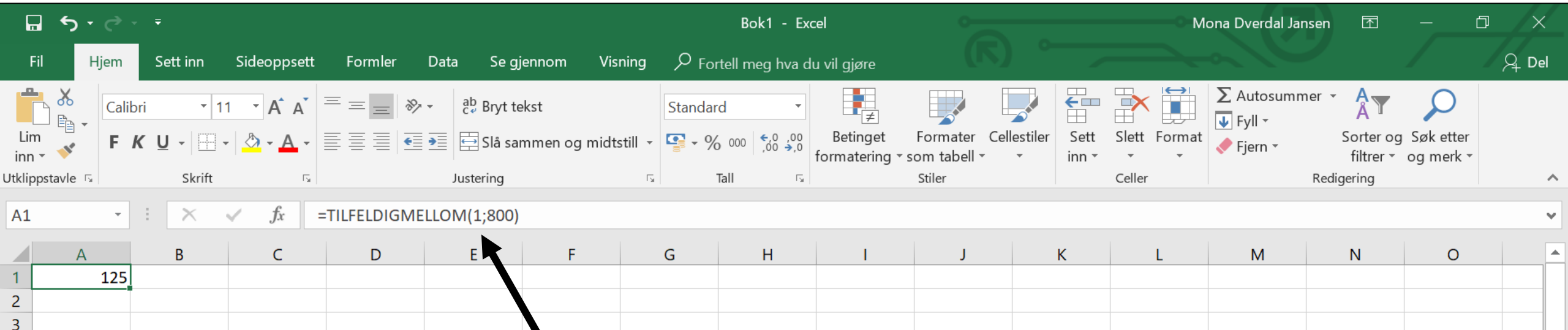
- Non-probability sampling (not random, e.g. convenience - only cooperating farmers)
- Simple random sampling
- Systematic random sampling
- Stratified random sampling (random sampling within groups)
- .....



# Simple random sampling

- Sampling from a population (e.g. farms) that is homogeneous in relation to disease distribution
- Requires a sampling frame
  - complete list of all sampling units available in the source population
  - requires individual identification for each sampling unit
- Each individual is selected using a random process so that each has a equal chance to be selected
- Collect the units with ID indicated by the random process

# Random number generator



The screenshot shows the Microsoft Excel interface. The title bar indicates the file is named "Bok1 - Excel" and the user is "Mona Dverdal Jansen". The ribbon is set to "Hjem" (Home). The formula bar shows the formula `=TILFELDIGMELLOM(1;800)` in cell A1. The spreadsheet grid shows the value 125 in cell A1. A black arrow points from the text `=RANDBETWEEN(1;800)` below to cell E1 in the grid.

	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O
1	125														
2															
3															

`=RANDBETWEEN(1;800)`

Bok1 - Excel

Mona Dverdal Jansen

Fil Hjem Sett inn Sideoppsett Formler Data Se gjennom Visning Fortell meg hva du vil gjøre

Utklippstavle Skrift Justering Tall Betinget formatering Formater Cellestiler Sett inn Slett Format Autosummer Fyll Fjern Sorter og Søker etter filter og merk

A1 =TILFELDIGMELLOM(1;800)

	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O
1	495														
2	452														
3	743														
4	114														
5	480														
6	142														
7	594														
8	438														
9	441														
10	526														
11	455														
12	754														
13	40														
14	728														
15	173														
16	644														
17	690														
18	609														
19	96														
20	652														
21															
22															

Ark1

Klar Gjennomsnitt: 463,3 Antall: 20 Summer: 9266 100 %

# Systematic random sampling

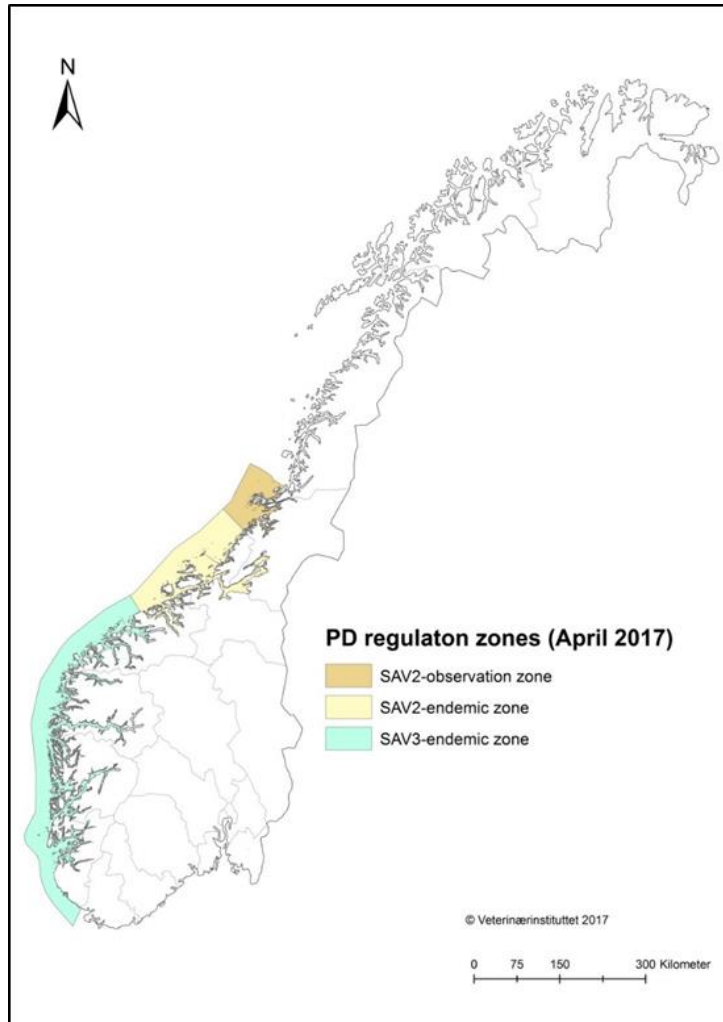
- Sampling from a population that is homogeneous in relation to disease distribution
- Units are sampled at a regular interval after a random start
  - No need for a sampling frame
- All units have to be sequentially accessible and each unit has an equal chance of being selected
  - Farms
  - Fish e.g. at slaughter, during vaccination
- Subject to bias in the random start, the interval or in the listing order

# Systematic random sampling - example

1. Calculate the Interval  $j = \text{Total number of animals} / \text{Sample size}$ 
  - We want 100 samples from 10 000 fish unit
  - $j = 10\,000/100 = 100$
2. Randomly pick a number from within  $j$ 
  - Starting with random in first 1-100 e.g. 27
3. Sample every  $j^{\text{th}}$  animal
  - E.g. 27, 127, 227,....., 9 927



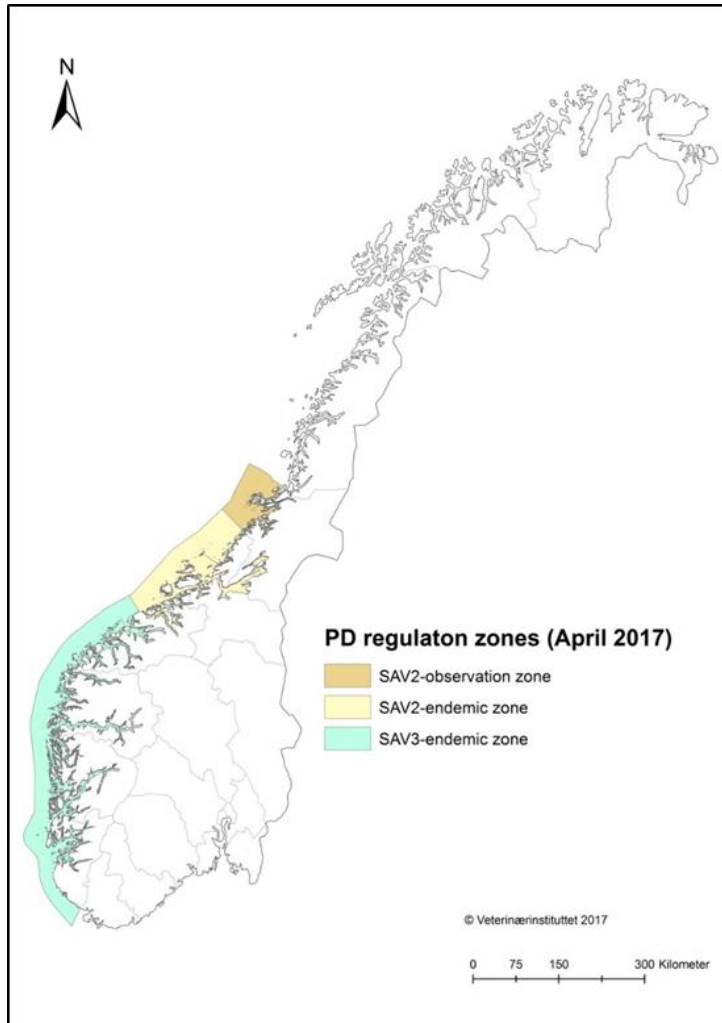
# Stratified random sampling



- Sampling from a population that is heterogeneous regarding the disease distribution
- The population can be categorized into groups where the units are more homogeneous than between than between groups



# Stratified random sampling



- Separate all units into mutually exclusive strata -> no overlap between them
  - Units within each stratum are assumed to be homogeneous in the disease distribution
- Include all strata
- Within each stratum, randomly select units (simple random or systematic random)
  - The same number of units in each strataor
  - **A proportionate sample to the number of units in each stratum**

# Now we know how we want to sample....

..... how many should be included?

- How many farms?
- How many fish?



# Sample size – general considerations

- Rarely possible to measure the entire population
- -> Parameters of interest calculated from subset of population
- Important to have enough samples to have sufficient certainty to aid decision-making
- Almost always restricted by the available budget

# Sample size – common approach

- Need to get sufficient sensitivity at farm level (or other unit-level)
  - Enough fish at each site
- Often result in:
  - Calculation of the cost per site
  - This then determines the maximum number of sites that can be sampled

# Sample size determination

## Key inputs

1. Design prevalence
  - expected prevalence of infection in the sampled group (e.g. moribunds)
2. Confidence level required for the result
  - typically use 95% (99% if aim at higher certainty)
3. Sensitivity of the test used
  - Usually assume perfect specificity
4. Population size
  - E.g. number of farms, number of fish






# Design prevalence



- Minimum expected prevalence of infection in the study population
  - Dynamics of infection for the agent in question
- How to determine the value?
  - **Between 1% and 5%** if only a small proportion of the population is expected to be infected (e.g. early in the outbreak)
  - **>5%** if expecting high prevalence (**10%** - often used with risk-based sampling)
  - **2%** = standard if no reliable information is available
- (Where available: see relevant disease chapter of the OIE Aquatic Manual)

# Sample size calculation


## Use

- Free software available at:
  - <http://www.winepi.net/uk/index.htm>
  - <http://epitools.ausvet.com.au/content.php?page=home>
- Published tables and equations in epidemiology text books, manuals

← →  http://www.winepi.net/uk/index.htm     

EpiTools  WinEpi: Working IN EPIdemi... x 

# WinEpi Working in Epidemiology



**Sample size**

- Detection of Disease
- Maximum possible Prevalence
- Estimate Percentage
- Estimate Mean
- Estimate Differences between Percentages

[ Start ]

---


## Sampling: Detection of Disease (1)

---

Confidence level :  ▾

Population size : -

Detection level :  ▾

Next 

“Population” can be the country’s tilapia farms, the fish in a farm and so on



# Working in Epidemiology

## Sampling: Detection of Disease (2)

Confidence level % :

Population size :

Minimum expected prevalence (%) :

# Working in Epidemiology

## Sampling: Detection of Disease (3)

Sample size needed to detect a disease (or infection) in a population:

Confidence level % :	95%
Population size :	40000
Expected minimum prevalence (%) :	10.00%

N. of infected animals to detect :	4000
<b>Needed sample size :</b>	<b>29</b>
Sampling fraction :	0.07%

Assuming  
perfect  $Se$  &  $Sp$

Risk-based  
sampling

# Working in Epidemiology

## Sampling: Detection of Disease (3)

Sample size needed to detect a disease (or infection) in a population:

Confidence level % :	99%
Population size :	40000
Expected minimum prevalence (%) :	10.00%

N. of infected animals to detect :	4000
<b>Needed sample size :</b>	<b>44</b>
Sampling fraction :	0.11%

Assuming  
perfect  $Se$  &  $Sp$

Risk-based  
sampling

# Working in Epidemiology

## Sampling: Detection of Disease (3)

Sample size needed to detect a disease (or infection) in a population:

Confidence level % : 95%

Population size : 40000

Expected minimum prevalence (%) : 5.00%

N. of infected animals to detect : 2000

**Needed sample size : 59**

Sampling fraction : 0.15%

Assuming  
perfect Se & Sp

Risk-based  
sampling

# What about wild fish populations?

- More difficult due to many unknowns
  - Best available information to estimate population size
  - Is it possible to get risk-based samples?
  - If not, random sampling or systematic random sampling of caught fish
  - Harder to define design prevalence so use a conservative estimate



# Working in Epidemiology

## Sampling: Detection of Disease (3)

Sample size needed to detect a disease (or infection) in a population:

Confidence level % :	95%
Population size :	40000
Expected minimum prevalence (%) :	1.00%

N. of infected animals to detect :	400
<b>Needed sample size :</b>	<b>297</b>
Sampling fraction :	0.74%

Assuming perfect  
Se & Sp

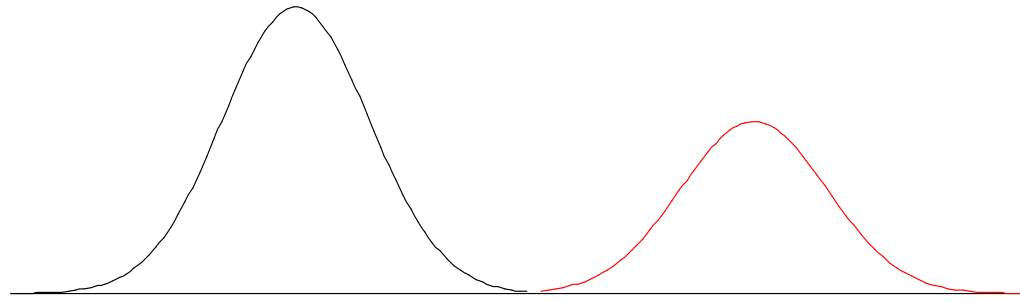
Risk-based  
sampling possible?

Large effect of a  
low design  
prevalence!

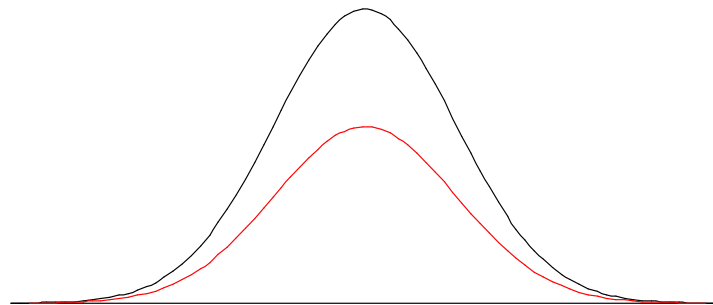
# Diagnostic tests



# The ability of the test to distinguish between diseased and non-diseased



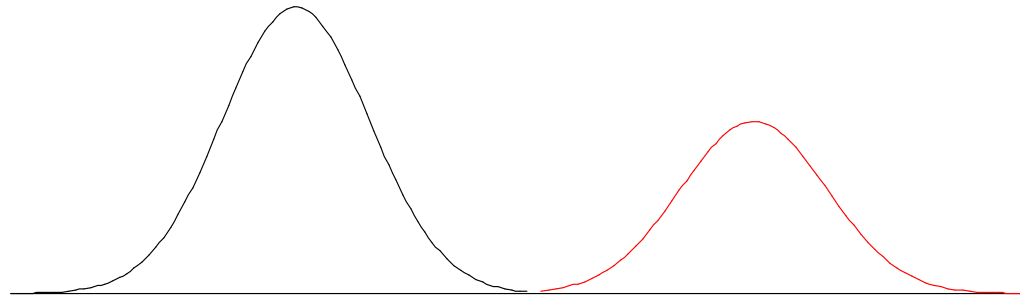
Perfect test  
(no tests are perfect)



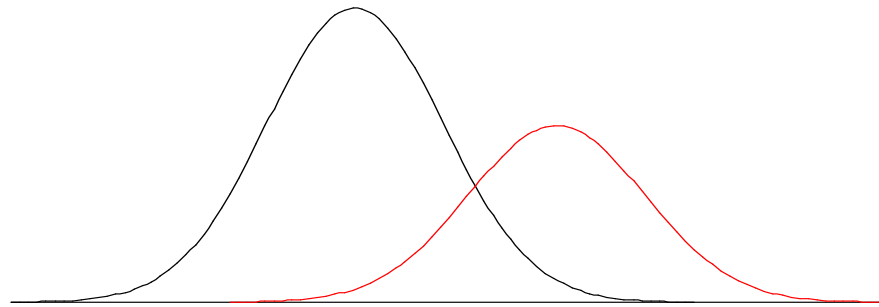
Useless



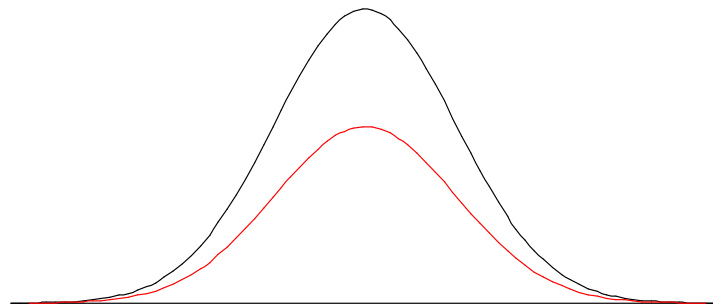
# The ability of the test to distinguish between diseased and non-diseased



Perfect test  
(no tests are perfect)

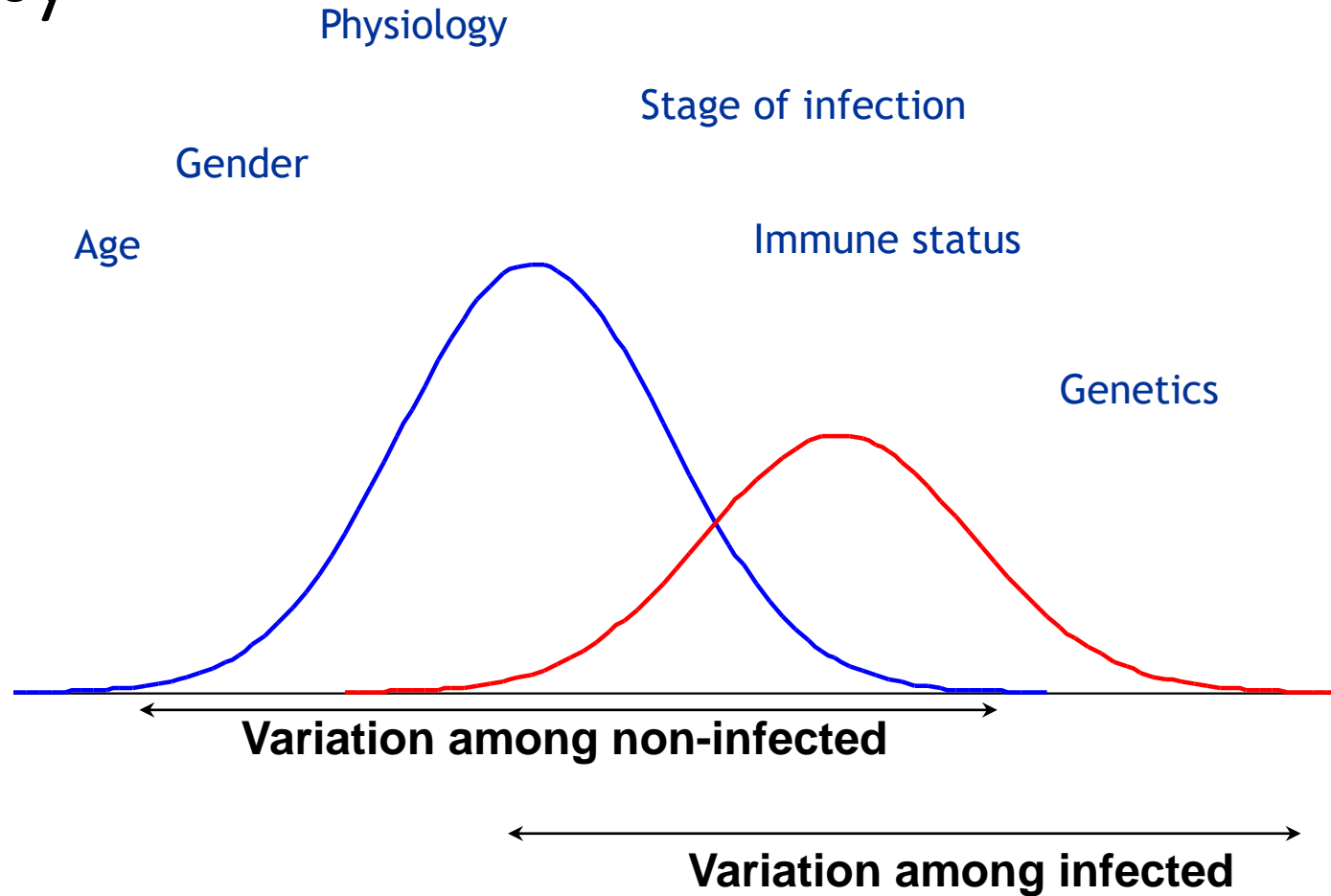


Normal situation



Useless

# Variability



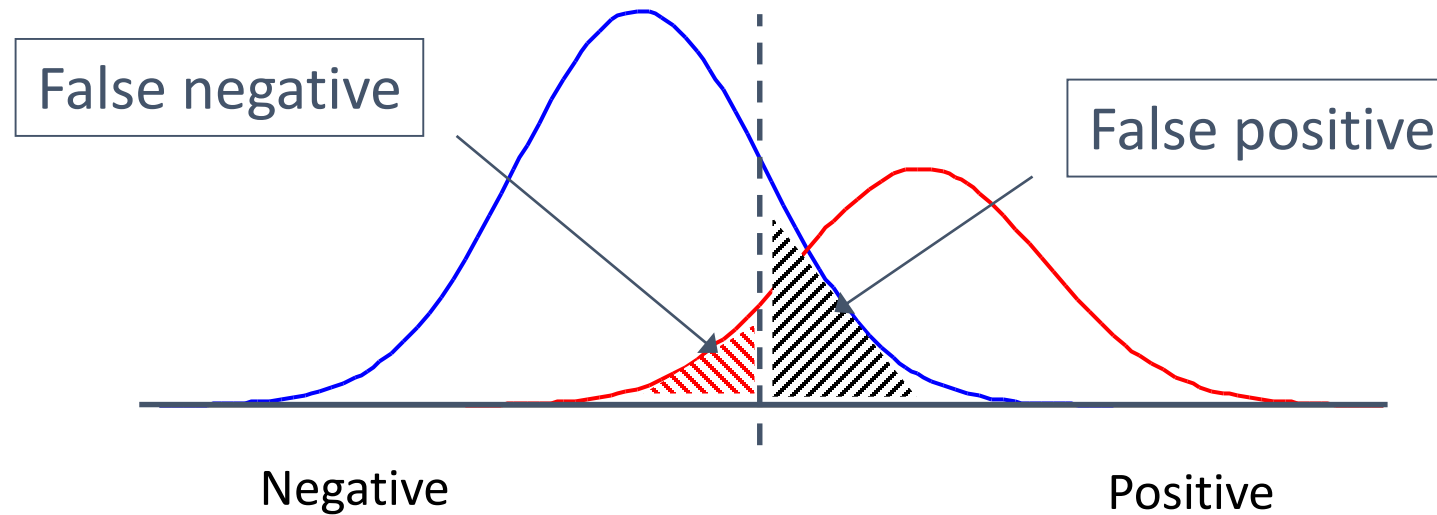
# Variability

- Biological variation
  - age, stage of infection, immune status, physical condition,...
- Variation due to differences in case definition
  - e.g. Histopathology vs PCR
- Technical variation
  - within lab variability (repeatability)
  - between lab variability (reproducibility)

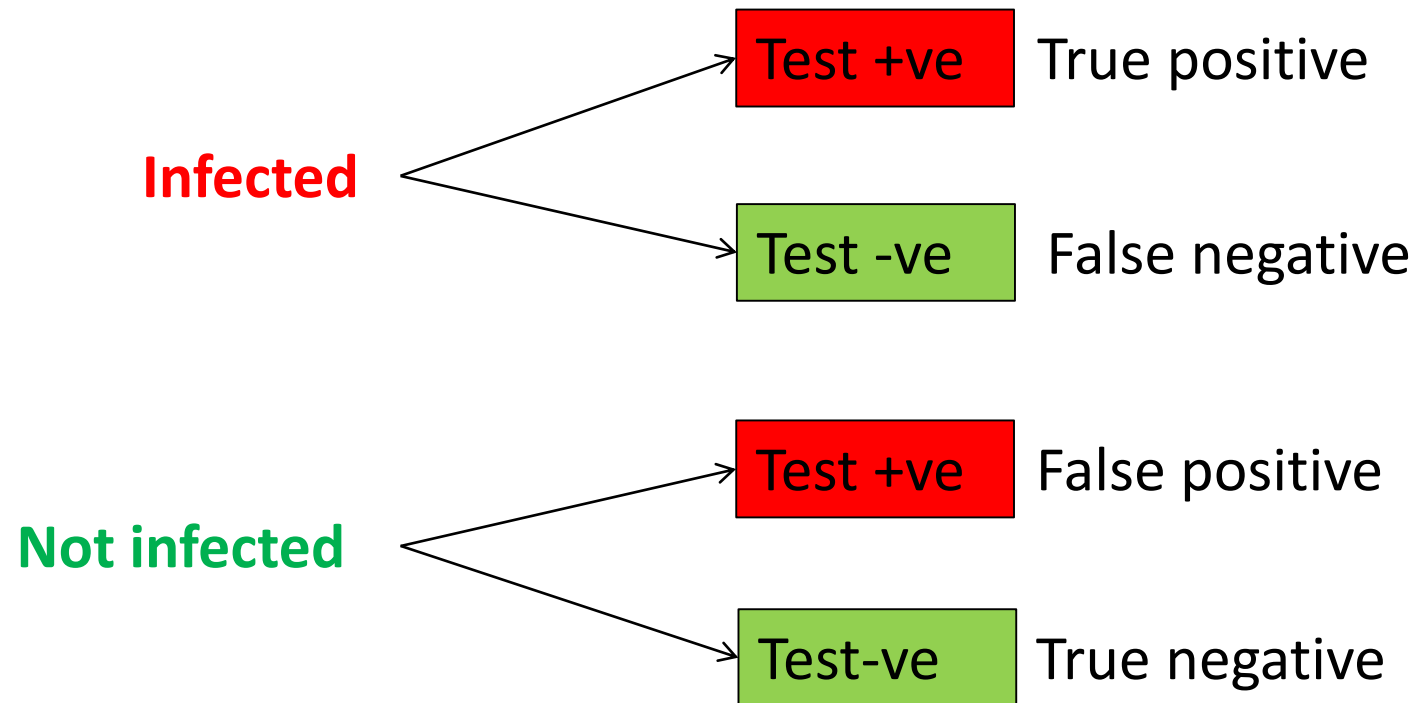
# Some consequences of variability

- Different individuals or populations with same status can give a different response
- Overlap in the distributions will usually occur
  - Due to overlap in distributions and means
- Need of “Cut-off values” : False positive – false negative

# False positives and false negatives



# Binary test output – no/yes; 0/1



# Sensitivity and specificity

	True disease status		
	D+	D-	
Test +			
Test -			

# Sensitivity and specificity

	D+	D-	
Test +	a	b	$g = a + b$
Test -	c	d	$h = c + d$
	$a + c$	$b + d$	$g + h$

$Se = a / (a + c)$

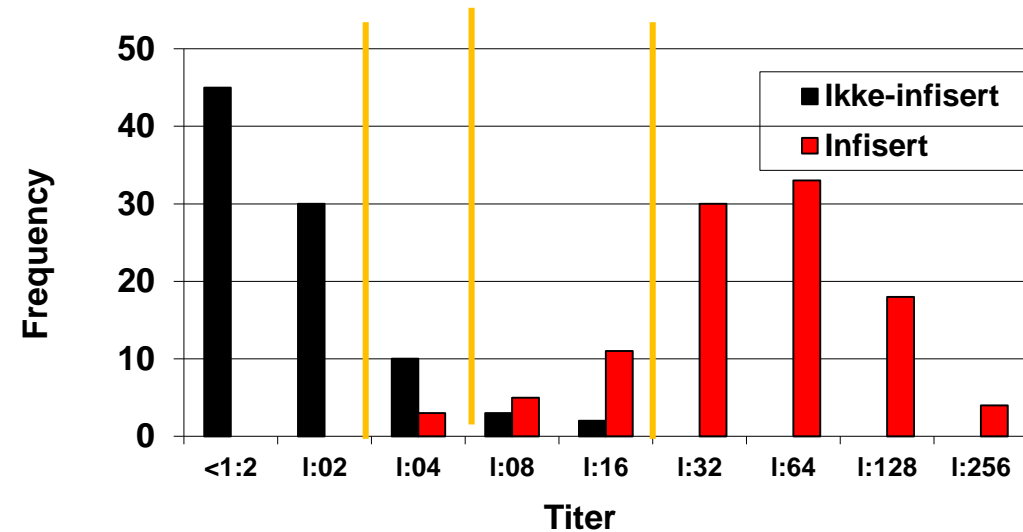
$Sp = d / (b + d)$

**Sensitivity (Se):** probability of testing positive if truly infected

**Specificity (Sp):** probability of testing negative if truly non-infected



# Sensitivity and spesificity



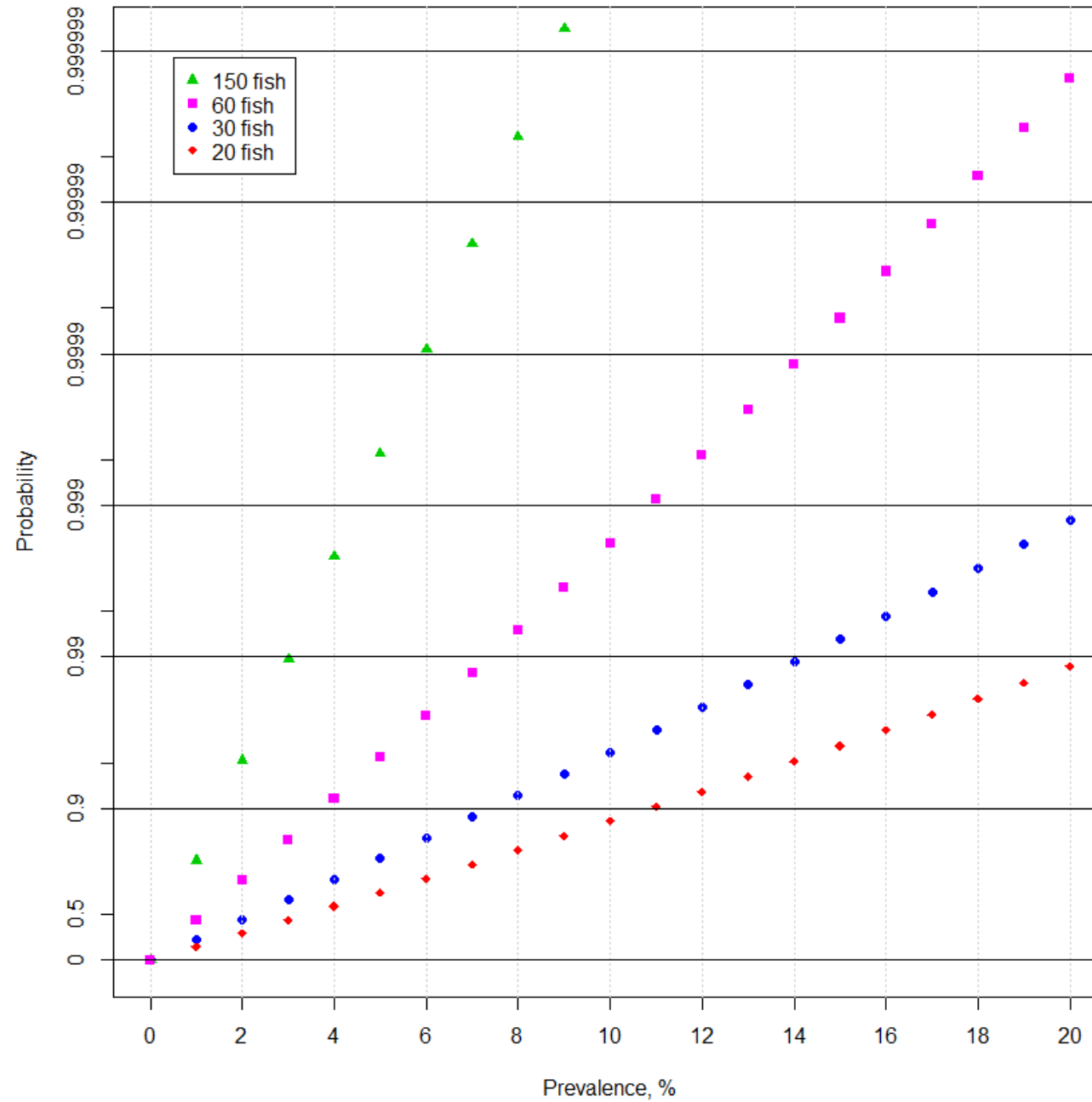
Cut-off  
 $\geq 1:82$

	D+	D-
Test +	104	15
Test -	0	75

$$Se = \frac{851}{1041} = 0,817$$

$$Sp = \frac{75}{90} = 0,83$$

### Probability of positive result perfect tests



Source: Norwegian  
Veterinary Institute

# Aspects to be considered when interpreting diagnostic test results

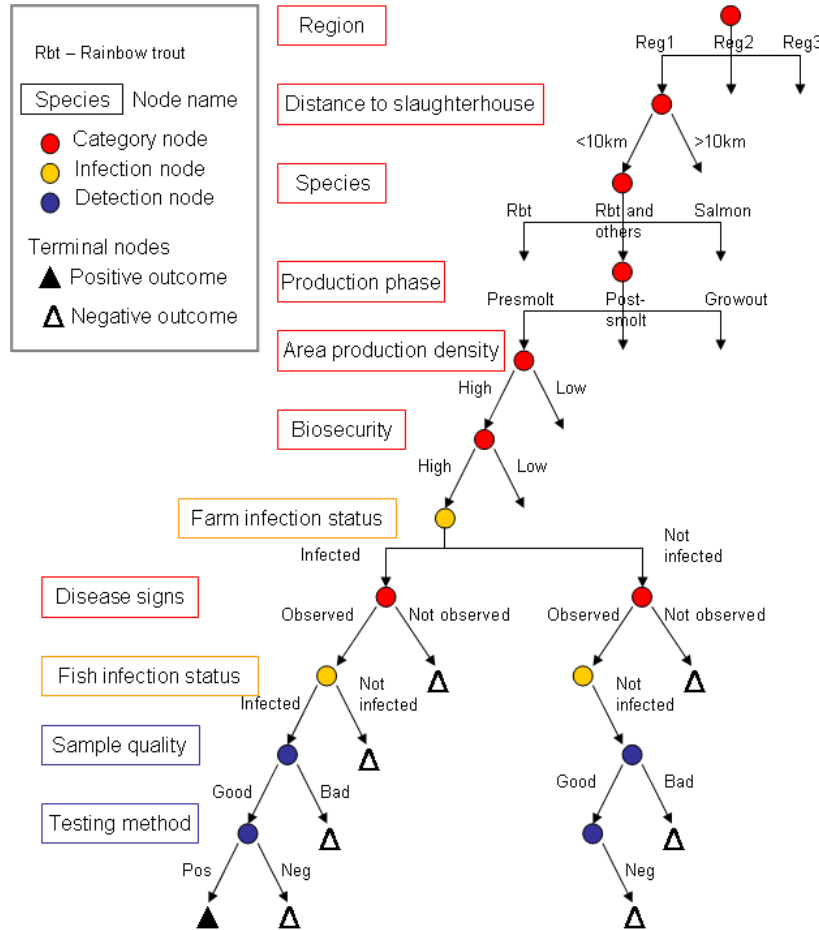
- Pooling of samples
  - How does that affect the Se (/Sp)?
- When utilizing several tests
  - How to combine the results?
  - How does this affect the Se (/Sp)?
- Aggregate testing
  - Testing several individual animals in order to classify herds
  - How does that affect the Se (/Sp)?

Design prevalence	Sensitivity (%)	Specificity (%)	Sample size	Maximum number of false positives if the population is free
2	100	100	149	0
2	100	99	524	9
2	100	95	1,671	98
2	99	100	150	0
2	99	99	528	9
2	99	95	1,707	100
2	95	100	157	0
2	95	99	542	9
2	95	95	1,854	108
2	90	100	165	0
2	90	99	607	10
2	90	95	2,059	119
2	80	100	186	0
2	80	99	750	12
2	80	95	2,599	148
5	100	100	59	0
5	100	99	128	3
5	100	95	330	23
5	99	100	59	0
5	99	99	129	3
5	99	95	331	23
5	95	100	62	0

5	95	99	134	3
5	95	95	351	24
5	90	100	66	0
5	90	99	166	4
5	90	95	398	27
5	80	100	74	0
5	80	99	183	4
5	80	95	486	32
10	100	100	29	0
10	100	99	56	2
10	100	95	105	9
10	99	100	29	0
10	99	99	57	2
10	99	95	106	9
10	95	100	30	0
10	95	99	59	2
10	95	95	109	9
10	90	100	32	0
10	90	99	62	2
10	90	95	123	10
10	80	100	36	0
10	80	99	69	2
10	80	95	152	12

Source: OIE Guide for Aquatic Animal Surveillance (p. 34)

# Surveillance system sensitivity



Lyngstad *et al* (2016) Preventive Veterinary Medicine, 124, 85-95



## EpiTools epidemiological calculators

This site is developed and maintained by Ausvet. The site is intended for use by epidemiologists and researchers involved in estimating disease prevalence or demonstrating freedom from disease through structured surveys, or in other epidemiological applications.

### Surveillance utilities

- 1-Stage representative freedom surveys
- 2-Stage representative freedom surveys
- Risk-based freedom surveys
- Random Sampling from a population
- Estimating true prevalence
- Pooled prevalence calculator
- Survey Toolbox for livestock diseases and freedom in finite populations
- HerdPlus module for herd-sensitivity and freedom in finite populations

### Case study data

- GIS case study data from *Epidemiology for Field Veterinarians* text
- *Epidemiological Problem Solving* case studies and model answers

### Epidemiological studies

- Sample size calculations
- Summarise categorical or continuous data
- Statistical significance testing
- Probability distributions
- Bioequivalence analysis

### Diagnostic tests

- Application of diagnostic tests

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ausvet EpiTools epidemiological calculators

Home Prevalence Freedom Studies Diagnostics Sampling English

### Simple risk-based surveillance

### Input Values

Relative risk :

Population proportion in high risk group :

Design prevalence :

Test sensitivity :

Number of high-risk units sampled :

Number of low-risk units sampled :

### Sample Size - single level

### Sensitivity - single level

- The relative risk: this measures the risk of animals being infected in the high-risk group, relative to the risk of animals being infected in the low-risk group. For risk-based surveillance, this should be greater than 1. If analysing biased surveillance (for instance abattoir testing), the animals tested may have *lower* probability of being infected than the rest of the population.
- The population proportion: this is the proportion of animals from the entire population that are in the high-risk group.

In addition, the following parameters are required:

- The design prevalence: this is the assumed prevalence of disease, if the disease is present in the population. It is used as a standard by which the sensitivity of the surveillance can be evaluated.
- The individual animal test sensitivity: this is the sensitivity of the test performed on individual animals.
- The number of animals tested in each of the high and low risk groups.
- The prior confidence of freedom before the surveillance was undertaken. This is combined with System sensitivity to calculate posterior confidence of freedom.

The results include:

- the sensitivity of the surveillance system, or in other words, the probability that the surveillance system would detect at least one infected animal was present at the specified design prevalence.

## Simple risk-based surveillance - calculation of surveillance sensitivity

### Input Values

This page calculates the surveillance sensitivity for simple risk-based surveillance, for instance, a survey in which a high risk population is targeted.

This analysis assumes that there is no clustering of disease (for instance, we are working at the herd level), and that the effective specificity of the surveillance system is equal to one (all positives are followed up to ensure that they are not false positives).

Relative risk :

One risk factor is considered, for which the following information is required:

- The relative risk: this measures the risk of animals being infected in the high-risk group, relative to the risk of animals being infected in the low-risk group. For risk-based surveillance, this should be greater than 1. If analysing biased surveillance (for instance abattoir testing), the animals tested may have *lower* probability of being infected than the rest of the population.
- The population proportion: this is the proportion of animals from the entire population that are in the high-risk group.

Population proportion in high risk group :

Design prevalence :

In addition, the following parameters are required:

- The design prevalence: this is the assumed prevalence of disease, if the disease is present in the population. It is used as a standard by which the sensitivity of the surveillance can be evaluated.
- The individual animal test sensitivity: this is the sensitivity of the test performed on individual animals.
- The number of animals tested in each of the high and low risk groups.
- The prior confidence of freedom before the surveillance was undertaken. This is combined with System sensitivity to calculate posterior confidence of freedom.

Test sensitivity :

Number of high-risk units sampled

Number of low-risk units sampled

The results include:

- the sensitivity of the surveillance system, or in other words, the probability that the surveillance system would detect at least one infected animal if disease was present at the specified design prevalence.
- the level of confidence that the population is free of disease (at the design prevalence).
- For comparison, the sensitivity of the system and confidence of freedom if representative sampling were used are also shown, along with the sensitivity ratio. This indicates how much more sensitivity the risk-based approach achieves, relative to a representative approach.
- the effective probability of infection (EPI) for both high and low risk groups.

Prior confidence of freedom :

Submit



## Simple risk-based surveillance - calculation of surveillance sensitivity

Analysed : Wed Jun 20, 2018 @ 02:22

### Inputs

Relative risk	2
Population proportion in high risk group	0.1
Design prevalence	0.05
Test sensitivity	0.95
Number of high-risk units sampled	25
Number of low-risk units sampled	5
Prior confidence of freedom	0.5

### Results

	Surveillance sensitivity	Confidence of freedom
Risk-based	<b>91.6%</b>	<b>92.3%</b>
Representative	<b>76.8%</b>	<b>81.2%</b>
Sensitivity ratio	<b>1.19</b>	
EPI in high-risk group	<b>9.1%</b>	

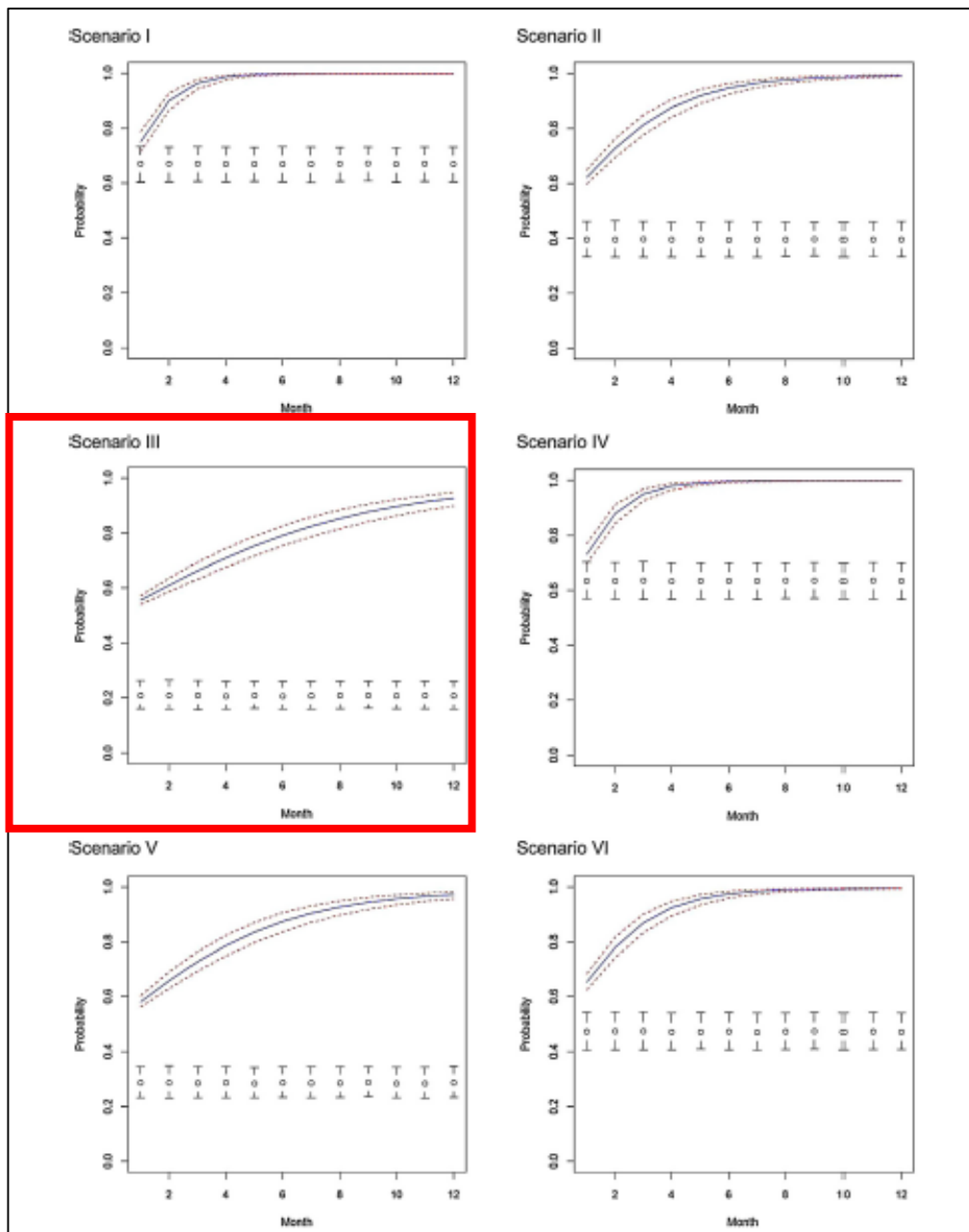
### Inputs

Relative risk	2
Population proportion in high risk group	0.1
Design prevalence	0.1
Test sensitivity	0.95
Number of high-risk units sampled	25
Number of low-risk units sampled	5
Prior confidence of freedom	0.5

### Results

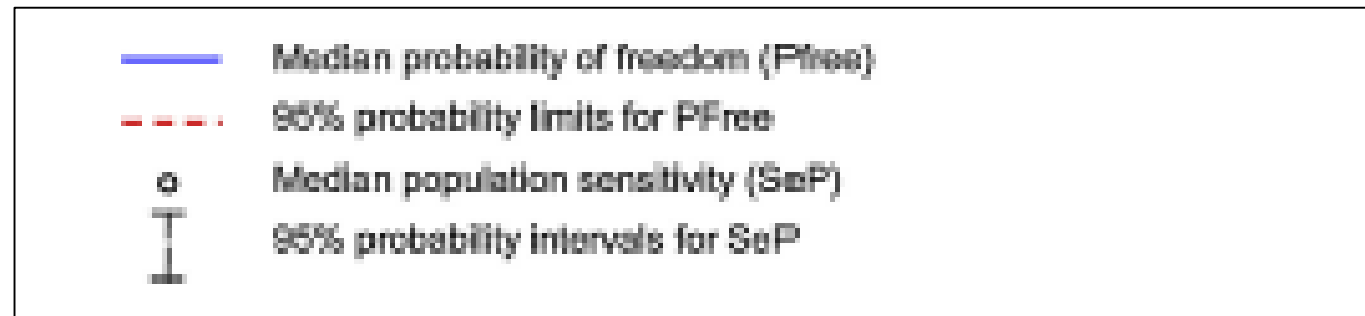
	Surveillance sensitivity	Confidence of freedom
Risk-based	<b>99.4%</b>	<b>99.4%</b>
Representative	<b>95.0%</b>	<b>95.2%</b>
Sensitivity ratio	<b>1.05</b>	

Can also be calculated for other levels of clustering e.g. 2-stage sampling



## Yearly inspection frequency:

- I. 6 inspections for both Atlantic salmon (AS) & rainbow trout (RT)
- II. 3 inspections (AS & RT)
- III. 6 inspections RT, 1 inspections AS
- IV. 6 inspections (southern Norway), 3 inspections northern Norway
- V. 6, 4 or 0 inspections depending on biosecurity level
- VI. 6 or 3 inspections depending on production intensity



# Conclusions

- There is an exciting variety of surveillance options
  - Purpose
  - Cost
- New approaches tend to be more complex
  - Start simple and build up depending on the requirements
- Use professional guidance when appropriate
- Good biosecurity and procedures
  - -> possibility for less intensive surveillance over time

# References

- *OIE Aquatic animal health code* (current, online version)
- *OIE Guide for Aquatic Animal Health Surveillance*, 2009.
- *FAO Surveillance and zoning for aquatic animal diseases*, 2004
- Cameron, *et al* . (2002) *Survey toolbox for aquatic animal diseases – a practical manual and software package*. Australian Centre for International Agricultural Research (ACIAR).



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