

Faculty of Agricultural and Nutritional Science

NETWORK ANALYSIS Interruption of the chain of infection by removal of the most central premises

Kathrin Büttner, Joachim Krieter and Imke Traulsen

Institute of Animal Breeding and Husbandry Christian-Albrechts-University, Kiel

63rd Annual EAAP Meeting Bratislava, Slovakia August 27th to 31st, 2012 CAU

Christian-Albrechts-University Kiel Institute of Animal Breeding and Husbandry











Introduction

- Animal diseases like classical swine fever cause extensive economic losses in the livestock industry
- The transport of live animals is a major risk factor for the spread of infectious diseases
- Source of classical swine fever virus infection in German domestic pig herds from 1993 – 1998





Introduction

- To interrupt the chain of infection during an epidemic it is important to know the underlying structure of trade networks
- Network analysis
 - Characterisation of network topology
 - Detection of central or important farms in the network



Introduction

- To interrupt the chain of infection during an epidemic it is important to know the underlying structure of trade networks
- Network analysis
 - Characterisation of network topology
 - Detection of central or important farms in the network

Aim of the study

- To characterize the changes in the network topology by successive removal of the most central farms in the trade network
- To evaluate which centrality parameter is the most suitable measure for a rapid fragmentation of the trade network
- Interruption of the chain of infection



- Trade network of the pork supply chain from a producer community in Northern Germany
- Observation period: June 2006 to May 2009
- Transported livestock: Piglets, pigs, sows and boars



• Network properties: Directed & static



- **Degree:** Number of **direct** trade contacts
 - Ingoing trade contacts: In-degree
 - Outgoing trade contacts: Out-degree
- Infection chain: Number of direct and indirect trade contacts regarding the chronological order of the trade contacts
 - Ingoing trade contacts: Ingoing infection chain
 - Outgoing trade contacts: Outgoing infection chain





- Components: Two farms are part of the same component if they are connected by at least one path through the network
 - Number of components
 - Size of components
- Fragmentation: Number of components in relation to the number of farms in the network
 - Fragmentation = 0 (totally connected network)
 - Fragmentation = 1 (every farm is isolated)





Number of components: Size of largest component: Fragmentation:





Number of components: 1 Size of largest component: 21 (100 %) Fragmentation: 0





Number of components: 12 Size of largest component: 7 (33.3 %) Fragmentation: 0.89





Number of components: 16 Size of largest component: 2 (9.5 %) Fragmentation: 0.99











Reduction of the size of the largest component of more than 75% Number (Proportion) of removed farms Farm type n In-Outdegree degree 5 16 Multiplier 29 (17.2%)(55.2%)Farrowing 24 11 34 (70.6%)(32.4%)farm Finishing 77 153 (50.3%)farm Farrow-to-114 4 267 (42.7%)(1.5%)finishing farm 31 220 Total 483 (45.5%) (6.4%)







Reduction of the size of the largest component of more than 75%

Farm type	n	Number (Proportion) of removed farms	
		Ingoing infection chain	Outgoing infection chain
Multiplier	29	5 (17.2%)	18 (62.1%)
Farrowing farm	34	29 (85.3%)	9 (26.5%)
Finishing farm	153	129 (84.3%)	-
Farrow-to- finishing farm	267	199 (74.5%)	5 (1.9%)
Total	483	362 (74.9%)	32 (6.6%)





Conclusion

- The parameters regarding the ingoing contacts are not suitable for a rapid fragmentation of the trade network
- The successive removal of the most central premises regarding the parameters
 - out-degree
 - outgoing infection chain

is an appropriate method to interrupt the chain of infection during an epidemic

- Only 6% of the farms have to be removed
 - to get a reduction of the largest component of more than 75%
 - to get a fragmentation of more than 0.9







This project is kindly financed by the DFG.