

National Institute for Public Health and the Environment Ministry of Health, Welfare and Sport

# Wastewater based surveillance of AMR in the Netherlands

Hetty Blaak, Merel Kemper, Ana Maria de Roda Husman, Heike Schmitt



#### Environmental surveillance

- > Human feces and urine ends up in sewers
- Sewage consists of wastewater from:
  - households (71%)
  - industry and institutions (29%)
  - among which health care institutions (3.3%)
  - and agricultural, hunting and foresting industries (0.6%)
- Sewage is collected and treated in wastewater treatment plants (WWTP)
- > Effluent is released to surface water
- Wastewater can be used to gain insight in the prevalence of fecal pathogens in the general population and their geographical distribution



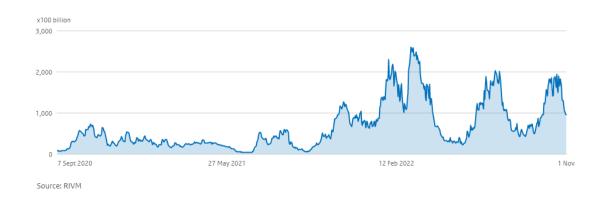


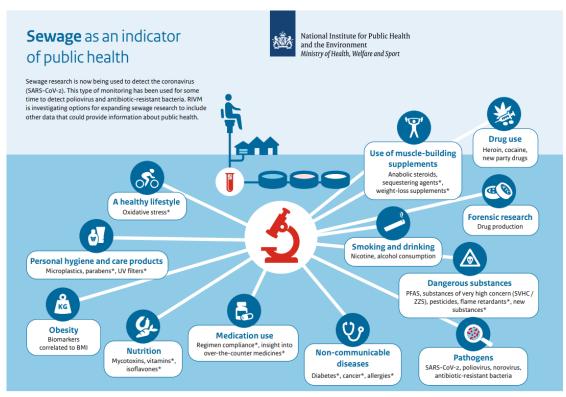




### Environmental surveillance – now and further

 AMR surveillance since 2016 within Aanpak Antibioticaresistentie





\*These parameters are not yet used in sewage research. Further research is needed to determine whether this is possible



## Environmental AMR surveillance – current goals

- 1. Prevalence of BRMO and specific BRMO types
- 2. Distribution across the country
- 3. Trends in time
- 4. Estimates of carriage
- 5. Evaluation of usage for local healthcare institutions

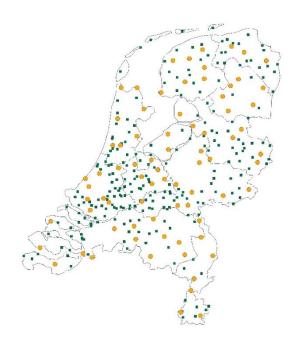


..This for the case of specific rare BRMO, such as carbapenemase producing Enterobacterales



### Surveillance of antibiotic resistance - method

**RWZI** selection



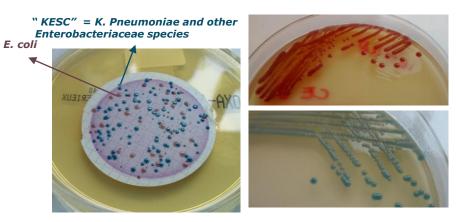
Sampling



Filtration



#### Selective culture



#### **CPE** isolates:

Identification of carbapenemase-gen by PCR: OXA-48, NDM, KPC, IMP, VIM, GES

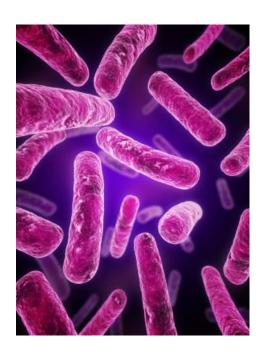
Genetyping: sanger sequencing (GES, OXA-48) and Wi

Genotyping: sanger sequencing (GES, OXA-48) and WGS (KPC en NDM)



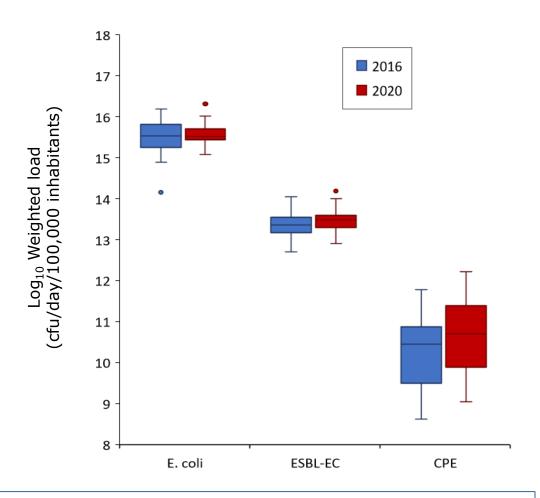
# CPE (carbapenemase producing Enterobacterales)

- Low prevalence in NL (~1:10.000 persons)
- Carriage associated with international travel and international hospital admission
- Resistance encoded by different genes (OXA-48, NDM, KPC, VIM)





#### CPE load- trends over time



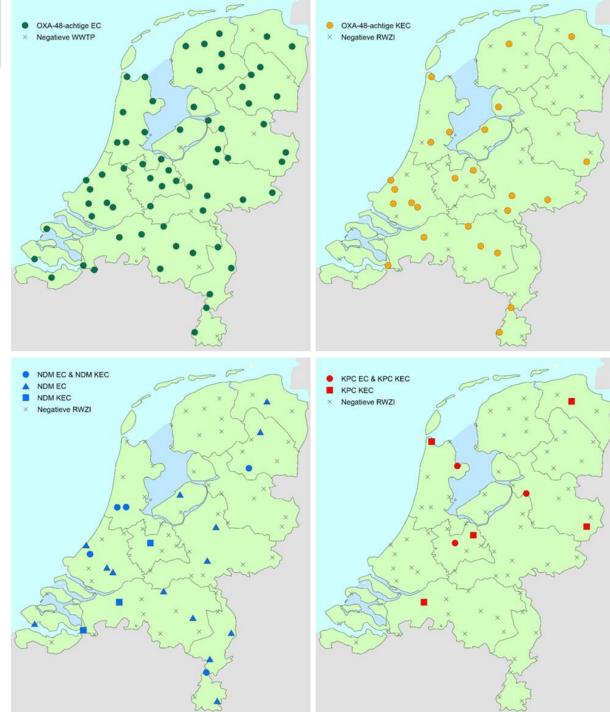
- CPE load 10<sup>2</sup>-10<sup>3</sup> times lower than ESBL-EC, and 10<sup>4</sup>-10<sup>5</sup> times lower than *E. coli* load
- Constant loads for E. coli and ESBL-EC between 2016 and 2020
- Slight increase in CPE load (2.5fold difference in geomean), p=0.093

Load = concentration x 24h volume of influent (cfu/day)



# Regional distribution

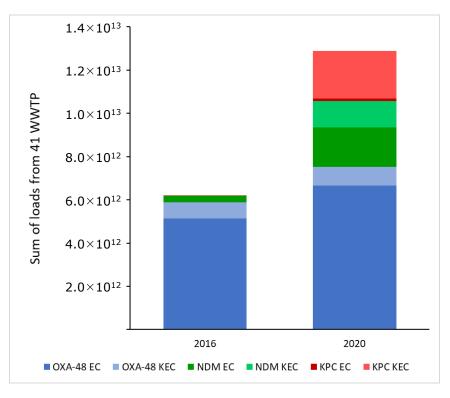
 No signs of difference in regional distribution of 4 CPE genotypes





### Population carriage estimates

Sum of CPE loads of 41 WWTP  $-4.4 \times 10^6$  inhibitants



#### Point estimate of CPE carriage

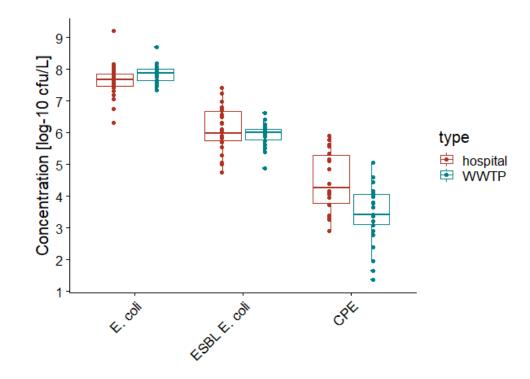
	2016	2020
CPE	1043 (0.02%)	2078 (0.05%)
NDM/KPC CPE	55 (0.001%)	860 (0.02%)

- Based on ΣCPE-load relative to ΣESBL-load  $(1.3 \times 10^{15})$ ,
- Based on 5% ESBL-carriers in 2014-2016 (vd Bunt et al. 2019) ,
- Assumed: same fecal load/carrier for ESBL and CPE carriers



# In-depth studies: hospitals as source of CPE

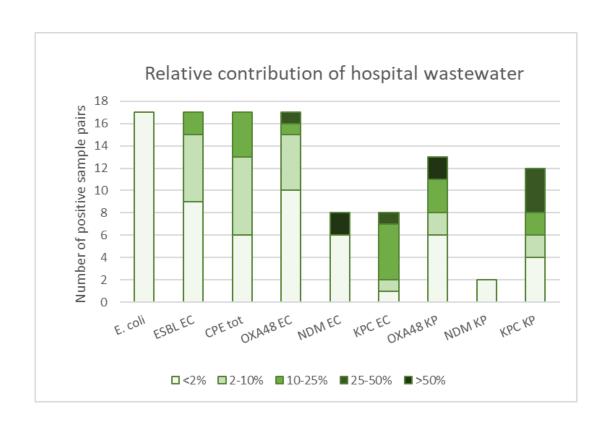
- > 8 hospitals, sampled 3-4 times, paired samples from hospital and WWTP
- Significantly higher concentrations of CPE found in hospital effluent





# Contribution of hospitals to total WWTP - load

- Hospitals on average 1% of the total wastewater volume
- Load: total number of bacteria (=concentration \* wastewater volume)
- Hospitals do contribute to CPE load
- > But mostly much less than 25%
  - Apart from single gene types





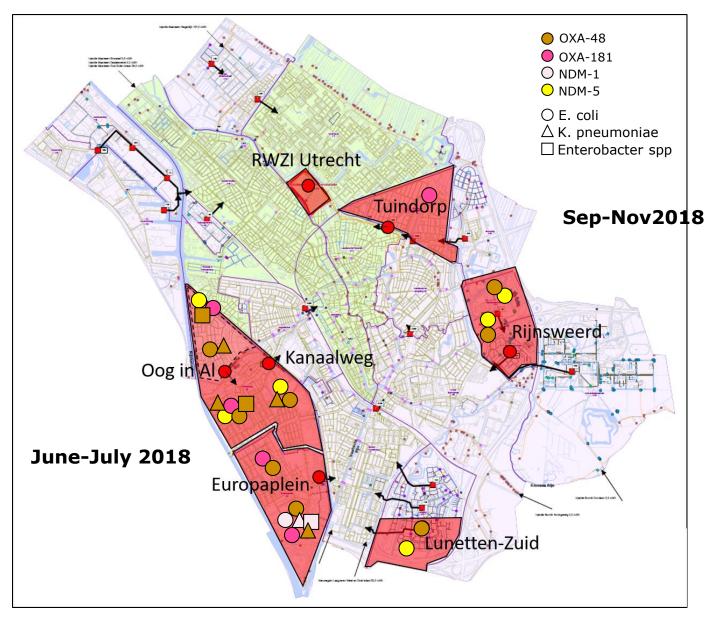
### Healthcare institutions and wastewater

- No major point source for emissions to the environment
- Wastewater treatment at institutions: no major contribution to emission reduction

Contribution of environmental surveillance to human surveillance?

- Outbreak detection
- Overall prevalence in HCI





# CPE in district sewers

- CPE are detected in district sewers, further substantiating carriage in the community
- Multiple CPE types are found in different districts of the same city
- Detection of CPE in small districts suggests high sensitivity (likely restricted numbers of carriers)



#### Other BMRO

Colistin resistant *E. coli* 

• presence/absence analyses rather than concentrations

CPE method applicable to global analysis

• JPE TRIuMPH

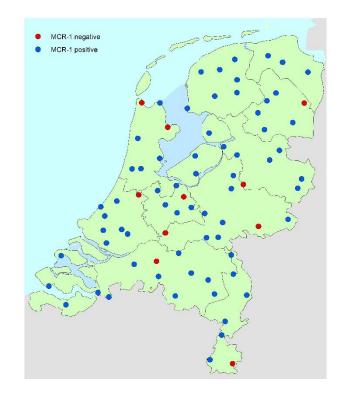
Carbapenem resistant *Pseudomonas aeruginosa* 

JPE SAMPAN

Metagenome, Acinetobacter baumannii

Wish list

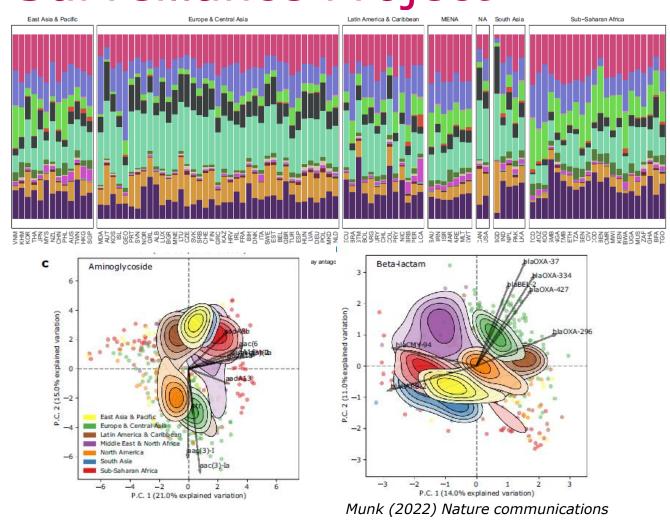
	Samples	Isolates
Total	76	296
MCR-1 pos	66	203
	87%	69%





# Global Sewage Surveillance Project

- Metagenomic analyses of sewage samples from >100 countries
- Differences in types of resistance genes across continents
- Correlation not only with antibiotic usage, but also health spendings / sanitation





#### Conclusions



Wastewater based surveillance (WBE)

- Most promising application: prevalence of rare antibiotic resistant bacteria in the population and their geographical distribution
- Trends in prevalence of CPE: No decrease despite measures to control Covid-19 pandemic (longitudinal measurements still ongoing)
- Single resistotypes: Overlap of wastewater isolates and isolates from human surveillance - WBE provides additional information
- WBE can also be used at the level of districts or institutions



#### Thanks to

- Hetty Blaak, Merel Kemper, Ka Yin Leung, Ana Maria de Roda Husman
- Many people at all Dutch waterboards, Waternet, Delfluent, Aquon, Aqualysis, WBL
- > CBS
- Min VWS

