Molecular typing data collection of STEC isolates from food and animals

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Human side: ECDC to collect molecular typing data from foodborne pathogens isolated from human cases

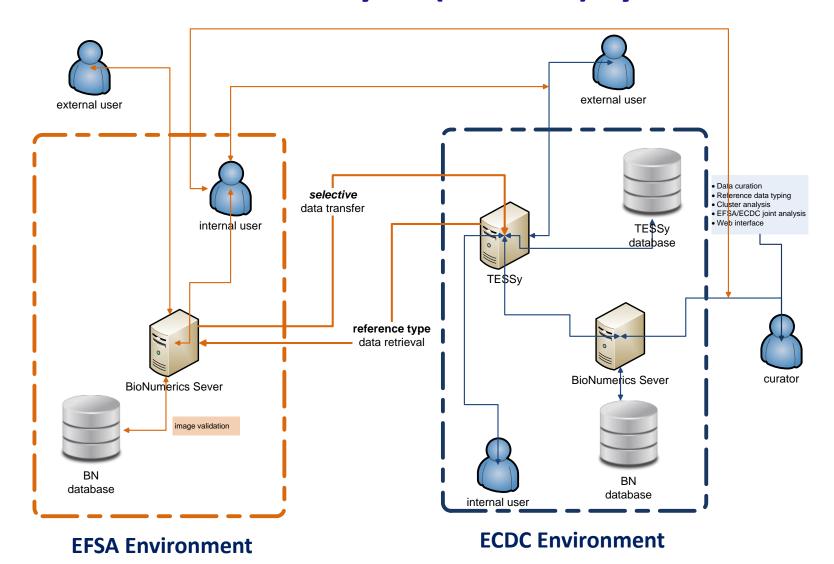
<u>Veterinary side</u>: EFSA, in collaboration with relevant EU Reference Laboratories (EURLs), to collect similar data from food, feed and animal isolates

A pilot project is going to be run, starting at the end of this year

The data collection will cover initially Salmonella, VTEC and Listeria with **PFGE** method (and MLVA only for S. Typhimurium) – other methods and pathogens can be taken aboard later

Regular joint analyses of the data by ECDC, EFSA and EURLs – integration at EU level

Structure of the Molecular Typing Data Collection and Analysis (MTDCA) system



MTDCA system

	EPIDEMIOLOGICAL DATA (PARTLY SENSITIVE)				Typing Data (not sensitive by itself)		
ID	Country	Animal/Food Type	Area in the country	Date of Sample	Serotype	PFGE_Xbal	AST_NAL
EFSA- 001	XX	Smoked salmon	X	2012-06-01	STANLEY		S
EFSA- 002	XX	Turkey	Υ	2012-07-01	STANLEY		R

- **Epidemiological Data**: data about an isolate in terms of sampling time and place, origin (food/ animal type, country of origin, producer) etc. partly sensitive data
- **Typing Data**: the results of typing experiments performed on an isolate. e.g. serotype, PFGE pattern, antimicrobial susceptibility results, MLVA results non sensitive data
- **Joint EFSA-ECDC analyses** to be carried out only with non-sensitive or anonymised data.

Molecular typing data collections in the EU

Standard Operating Procedures (PFGE, image aquisition, image quality assessment, curation) are under development and will be made available

Member states/NRLs will be asked to participate in the pilot phase, providing molecular typing data

In line with the orientations provided in the Vision paper, a **Scientific Collaboration Agreement** will be signed by all parties to define data ownership, access, use, publication, procedures and confidentiality.