





# Development and validation of a method for quantification of 15 antiviral drugs against influenza in poultry muscle using LC-MS/MS

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#### Influenza

■ Three types of influenza: A, B and C.

- Symptoms: fever, headache, dry cough, nasal congestion.
- Seasonal epidemic form: between 300k and 500k deaths globally every year.
- Pandemic form from zoonotic origins every 10-50 years: deaths in millions.



## **Avian Influenza**

- Regular avian influenza outbreaks in poultry industry.
- Constant mutation: new strains (e.g. H10N3).
- Combination of avian influenza and human influenza can cause the next pandemic.



# Antiviral drugs against influenza

- Antiviral drugs against influenza developed for human use.
- It is suggested that illegal use of antiviral drugs in farm causes development of drug-resistant influenza strains.
- Consequently: Chinese and US authorities (FDA) have banned use of certain antiviral drugs in poultry farms.
- Necessity of developing an analytical methodology to monitor antiviral drugs residue in poultry.

# **Existing methods**

- Existing fast screening techniques.
- LC-MS/MS methods with very long/complicated procedures (SPE, two analytical columns) and/or not including all compounds of interest.
- Necessity to improve existing methods.
- Goal : develop a fast LC-MS/MS method including all relevant antiviral drugs against influenza.



#### **Antiviral drugs selected**







### **Antiviral drugs : properties**

- Broad range of polarity: from highly polar to non-polar compound.
- Acid, basic and neutral compounds.
- Broad range makes analytical method challenging.



### LC-MS/MS

- Method was developed on a LC-MS/MS instrument: Waters Acquity I-Class coupled with a Xevo TQ-XS.
- Provide suitable selectivity and sensitivity for confirmatory quantitative analysis of residue traces.



# **Mass Spectrometry optimisation**

- Electrospray source of ionisation was compared to Unispray.
- Electrospray gave better sensitivity at the desired flow and chromatography mode



- Source parameters were optimised and most suitable MS transitions were selected.
- Positive mode and protonated molecules [M+H]<sup>+</sup> were chosen for this method

# **Chromatography: C18 columns**

- Wide range of C18 columns experimented.
- Little to no retention for most polar compounds (Zanamivir, Ganciclovir, Ribavirin).

■ Not adapted to polar analytes.



# **Chromatography: alternative to C18**

- Alternative reversed-phase columns screened.
- Best retention for polar compounds: Selectra PFPP, better retention factor.
- Still very low retention, improved by use of ion pairing (HFBA) but was not practical.



# **Chromatography: HILIC**

- HILIC is the best chromatography mode for polar compounds.
- Best results were found with columns driven by partitioning mechanism.
- BEH Amide (A) gave more appropriate retention to non-polar or neutral compounds than HILIC-Z (B).



### **Chromatography: ion suppression**

- HILIC gave better ion suppression results than reversedphase for polar compounds.
- Very high ion suppression zone around the dead time in reversed-phase mode.



### **Sample preparation: sample extraction**

- QuEChERS is widely used in drug residue food analysis but was found ineffective at extracting highly polar compounds.
- Acetonitrile, methanol and isopropanol were tried.
- Best compromise was found with [acetonitrile:water] 80:20 v:v, best overall extraction efficiencies (>40%) and relatively clean extracts.



# **Sample preparation: clean-up**

- Various standard clean-up procedure were tested and found inefficient:
  - Ultrafiltration did not provide additional clean-up.
  - Freezing step trapped polar compounds in ice.
  - Several dspe sorbent were not beneficial.
- Dilute and shoot was chosen prior to LC-MS/MS analysis. Best final solvent mixture was [acetonitrile:methanol:water] 50:25:25 v:v:v.

### **Method performance: Validation**

- Method was validated according to EU guidelines 2021/808.
- All criteria were validated.
- Reproducibility trueness were ranging from 84% to 127% and reproducibility precision from 2.8% to 22.7%.



### Method performance: LOQ/LOD

- Eight matrix matched calibrants and the zero level were used to build a calibration curve, all R<sup>2</sup> were ≥0.988.
- LOQ ranged from 0.03 to 10.46 µg/kg.

Analyte	Curve range (µg/kg)	LOD (µg/kg)	LOQ (µg/kg)
Arbidol Sulfoxyde	0.1 - 0.8	0.0	
Arbidol	0.1 - 0.8	0.0	
Arbitol Sulfone	0.5 - 4	0.0	
Acyclovir	0.5 - 4	0.0	
Rimantadine	0.5 - 4	0.0	
Oseltamivir	0.5 - 4	0.0	
Amantadine	1-8	0.0	
Zanamivir	2 - 16	0.1	
Peramivir	2 - 16	0.1	
Ganciclovir	2 - 16	0.1	9 0.62
Viramidine	2 - 16	0.0	8 0.26
Oseltamivir Acid	2 - 16	0.2	8 0.92
Laninamivir	10 - 80	0.4	4 1.47
Ribavirin	25 - 200	2.4	8 8.26
Favipiravir	25 - 200	3.1	4 10.46

# **Occurrence in Irish market**

- A total of 120 poultry products were selected from Irish market.
- Mix of EU/UK and outside UK/EU origin and mix of cooked and raw products.
- No trace of antiviral drug against influenza was found.



Cooked/Processed Raw

# Conclusion

LC-MS/MS method for the confirmatory and quantitative analysis of 15 antiviral drug residues in avian muscle.

■ Fast and easy method.

- Method to monitor antiviral drug residues to better control the risk of resistant avian influenza strains generation.
- Sensitive method with LOQ ranging from 0.03 to 10.46 µg/kg.









### Thank you for your attention Any questions?



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