# Identification and Differentiation by PCR and MALDI-TOF-MS

## of Brachyspira spp. Isolated from Dogs





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#### Introduction

Brachyspira are known to cause severe symtomatic infection in poultry and characterized by diarrhea, weight loss and reduced egg prodution. Porcine and dog isolates of *Brachyspira pilosicoli* are very similar to those of humans. Therefore a possible role as zoonotic agent is under discussion. Presently insufficiently studied is the prevalence, distribution pathgenic potential in dogs. The aim of this study was the charactarization of Brachyspira subpopulations in asymtomatic, clinically healthy dogs by PCR and MALDI-TOF-MS.

#### **Materials and Methods**

Rectal swab samples (n = 6) were taken in a first screening round. In a second screening round 10 samples were taken from 10 dogs living in one kennel. In a third screening round 2 samples were taken from 2 dogs living without contact to the other experimental animals. All samples were cultured on trypticase-soja agar (TSA) enhanced with antibotics (A) in addition to blood agar. The putative Brachyspira isolates in pure culture were differentiated using MALDI-TOF-MS analysis (B) and Multiplex-PCR (C).

### **Agars were Created with the**

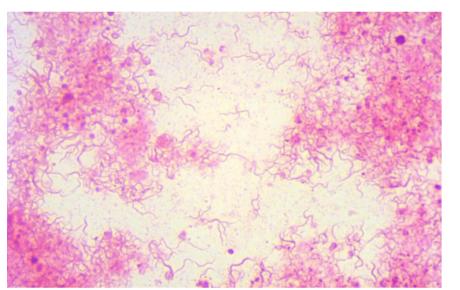
**Following Antibiotics** 

TSA 3: Colistin Rifampicin

Spectinomycin

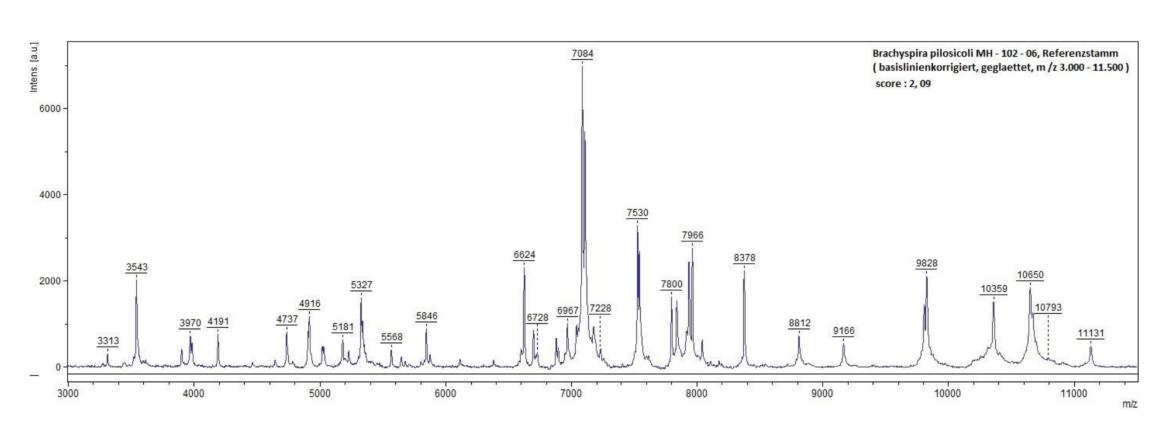
Colistin Rifampicin Spectinomycin Spiramycin Vancomycin

TSA 5:



Gram stain 07.06.17 from first screening round

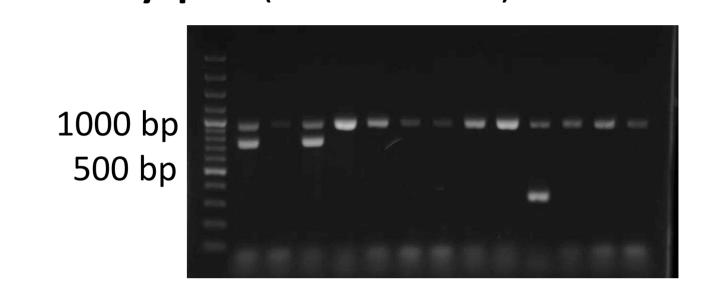
### **MALDI-TOF-MS Spectrum** (Matrix-Associated-Laser-Disorption-Ionisation Time of Flight Massspectrometry)



**Experimental Approach** 

Representitive spectrum of *Brachyspira pilosicoli* 

### **MP-PCR Differentiation of Pathogenic Brachyspira** (K.K. 06.01.17)



- 1) nox PCR
- nox gene 939 bp amplicon
- Brachyspira sp.
- 2) abgB PCR
- hippurat-hydrolase gene
- **B.** pilosicoli 744 bp amplicon
- 3) tnaA PCR
- tryptophanase gene
- 325 bp amplicon
- B. intermedia
- > B. hyodysenteriae

### Results

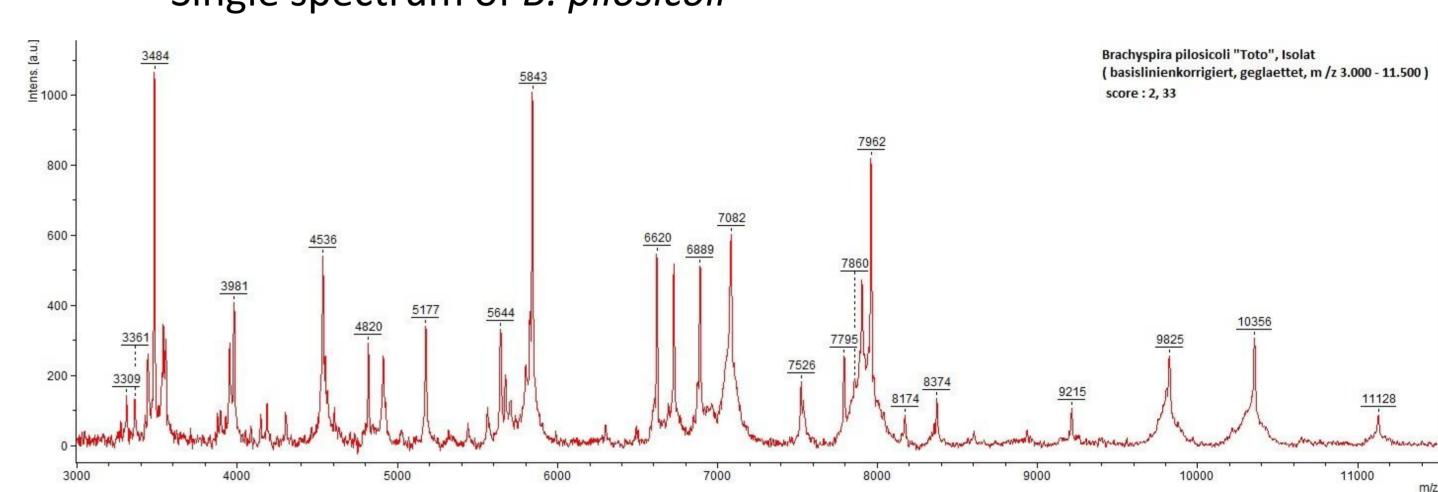
#### Comperative Growth on TSA 3 and TSA 5

Lola'. Representative cultures 07.06.17 taken from dog ,Lola'.



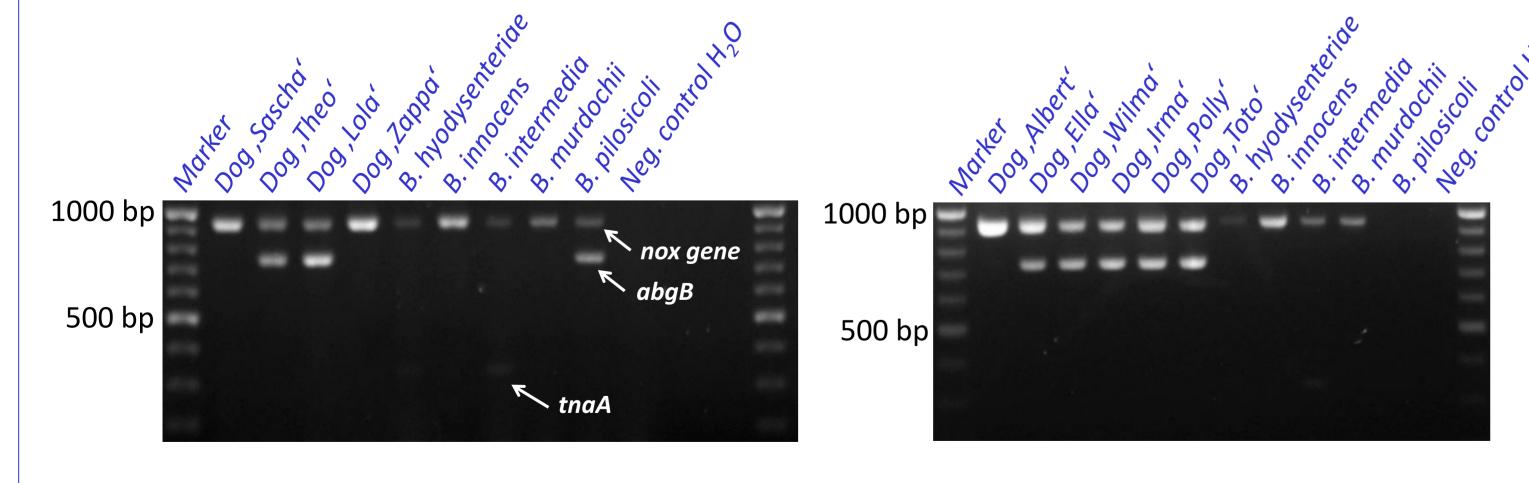
Brachyspira growth was much enhanced on TSA 3. Semiquantative analysis showed 3,5 cm for TSA 3 vs. 0,5 cm for TSA 5 from central starting point.

#### D MALDI-TOF-MS Single spectrum of *B. pilosicoli*



B. pilosicoli could be identified in 6 isolates.

	<b>Multiplex</b>	PCR
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Brachyspira could be proven to be present in all isolates. The abgB-PCR (744 bp), specific for *B. pilosicoli*, could be identified in 7 isolates.

Isolat	Dog	MALDI-	MALDI-	MP-PCR			Remark
	IVAIIIC	Result	Score	nox-Gen	abgB	tnaA	
VA II <sub>1,2,3</sub>	Toto	B. pilosicoli	2.33	+	+	-	
VA II <sub>2,3</sub>	Zappa	B. species		+	_	_	
VA II <sub>1,2</sub>	Sascha	B. species		+	-	-	
VA II <sub>1,3</sub>	Theo	B. species		+	+	_	
VA II <sub>1,2</sub>	Albert	B. species		+	-	-	
VA <sub>7,8,10</sub>	Irma	B. pilosicoli	2.06	+	+	-	
	Polly	B. pilosicoli	2.15	+	+	-	
VA II	Lola	B. pilosicoli	2.13	+	+	-	
VA II <sub>2,3</sub>	Ella	B. pilosicoli	1.53	+	+	-	
VA II <sub>1,2,3</sub>	Wilma	B. pilosicoli	2.01	+	+	-	
-	Gina	-	-	-	-	-	neg. culture
-	Eila	-	-	-	-	-	neg. culture
	VA II <sub>1,2,3</sub> VA II <sub>1,2</sub> VA II <sub>1,2</sub> VA II <sub>1,3</sub> VA II <sub>1,2</sub> VA <sub>7,8,10</sub> VA <sub>8,9,10</sub> VA II VA II VA II <sub>2,3</sub>	$\begin{array}{c c} \textbf{Name} \\ \hline \textbf{VA II}_{1,2,3} & \textbf{Toto} \\ \textbf{VA II}_{2,3} & \textbf{Zappa} \\ \hline \textbf{VA II}_{1,2} & \textbf{Sascha} \\ \hline \textbf{VA II}_{1,2} & \textbf{Shert} \\ \hline \textbf{VA II}_{1,2} & \textbf{Albert} \\ \hline \textbf{VA}_{7,8,10} & \textbf{Irma} \\ \hline \textbf{VA II} & \textbf{Lola} \\ \hline \textbf{VA II} & \textbf{Lola} \\ \hline \textbf{VA II}_{2,3} & \textbf{Ella} \\ \hline \textbf{VA II}_{1,2,3} & \textbf{Wilma} \\ \hline \textbf{-} & \textbf{Gina} \\ \hline \end{array}$	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	Name         TOF-MS Result         TOF-MS Score         nox-Gen           VA II <sub>1,2,3</sub> Toto         B. pilosicoli         2.33         +           VA II <sub>2,3</sub> Zappa         B. species         +           VA II <sub>1,2</sub> Sascha         B. species         +           VA II <sub>1,3</sub> Theo         B. species         +           VA II <sub>1,2</sub> Albert         B. species         +           VA <sub>7,8,10</sub> Irma         B. pilosicoli         2.06         +           VA <sub>8,9,10</sub> Polly         B. pilosicoli         2.15         +           VA II         Lola         B. pilosicoli         2.13         +           VA II <sub>2,3</sub> Ella         B. pilosicoli         1.53         +           VA II <sub>1,2,3</sub> Wilma         B. pilosicoli         2.01         +           -         Gina         -         -         -	$\begin{array}{ c c c c c c c c } \textbf{Name} & \textbf{TOF-MS} \\ \textbf{Result} & \textbf{Score} & \textbf{nox-Gen} & \textbf{abgB} \\ \hline \\ VA II_{1,2,3} & Toto & B. pilosicoli & 2.33 & + & + \\ VA II_{2,3} & Zappa & B. species & + & - \\ \hline VA II_{1,2} & Sascha & B. species & + & - \\ \hline VA II_{1,3} & Theo & B. species & + & + \\ \hline VA II_{1,2} & Albert & B. species & + & + \\ \hline VA_{7,8,10} & Irma & B. pilosicoli & 2.06 & + & + \\ \hline VA_{8,9,10} & Polly & B. pilosicoli & 2.15 & + & + \\ \hline VA II & Lola & B. pilosicoli & 2.13 & + & + \\ \hline VA II_{2,3} & Ella & B. pilosicoli & 1.53 & + & + \\ \hline VA II_{1,2,3} & Wilma & B. pilosicoli & 2.01 & + & + \\ \hline & Gina & - & - & - & - \\ \hline \end{array}$	$ \begin{array}{c ccccccccccccccccccccccccccccccccccc$

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#### Conclusions

Culturing of Brachyspira species was achieved in all dogs on TSA 3. PCR analyses of the nox- and abgB -PCR showed presence of Brachyspira in all isolates (nox +) and B. pilosicoli in 7 isolates (abgB +). Interestingly, two dogs, with no contact to the kennel dogs, were tested negative already in culture, suggesting spread among the kennel dogs. The clinical implications for asymptomatic, apparently healthy dogs with positive Brachyspira culture are presently unknown. Whether Brachyspira carriers need treatment for prevention of public health problems also warrants further research.