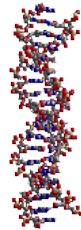




# Molecular epidemiology of the neglected meat-borne pathogen *Sarcocystis* spp. in pigs and cattle: an update

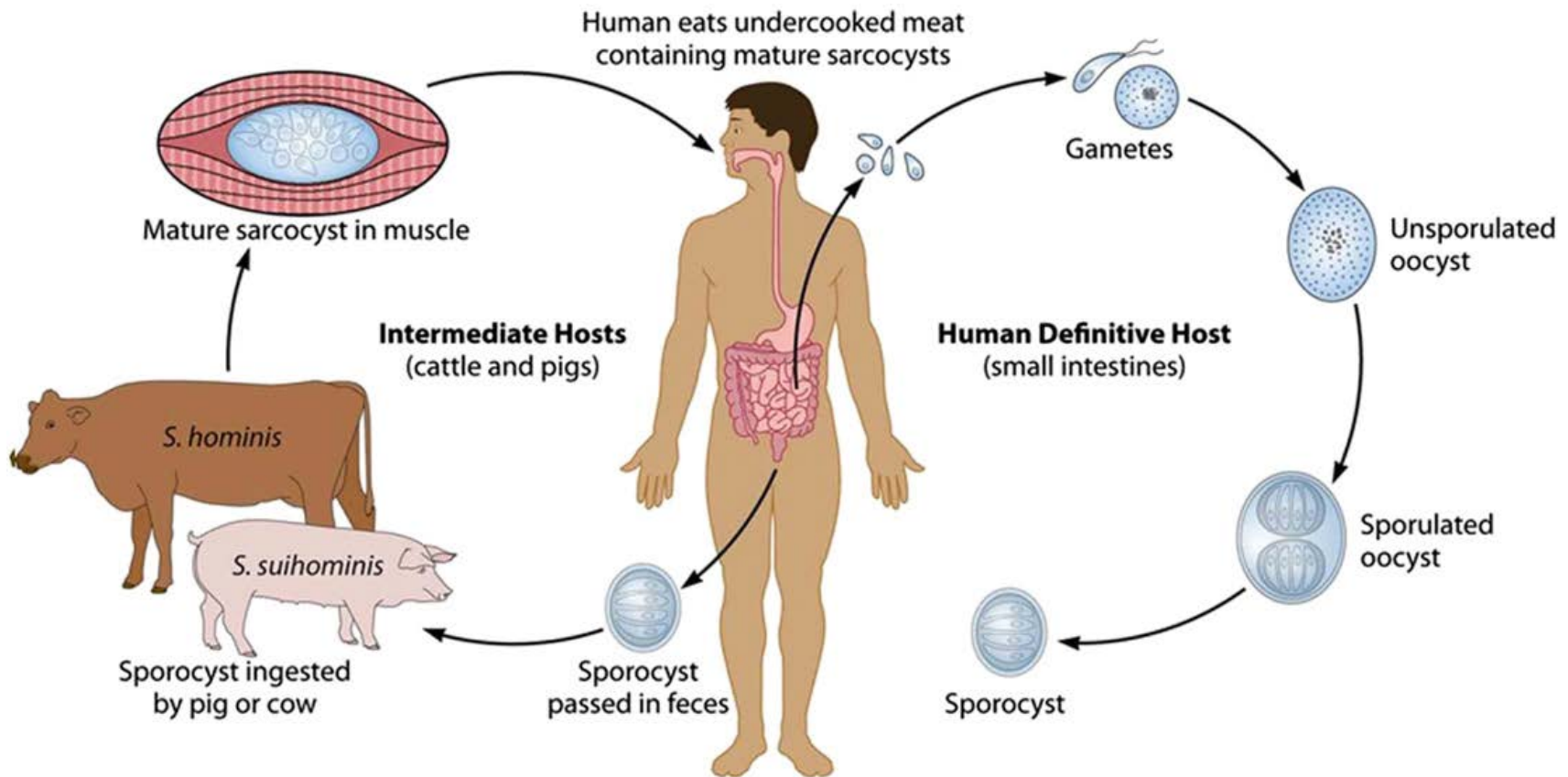


**Kálmán IMRE, Adriana MORAR, Mirela IMRE, Claudia SALA**

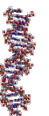
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Faculty of Veterinary Medicine,  
Banat's University of Agricultural Sciences and Veterinary Medicine "King  
Michael I of Romania" from Timișoara, Romania

# General considerations

- Sarcocystis* spp. are considered cyst-forming intracellular coccidian (Apicomplexa: Sporozoa) parasites



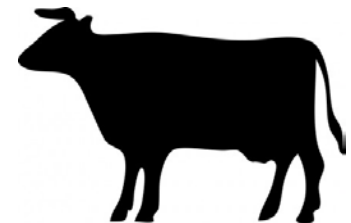
# General considerations



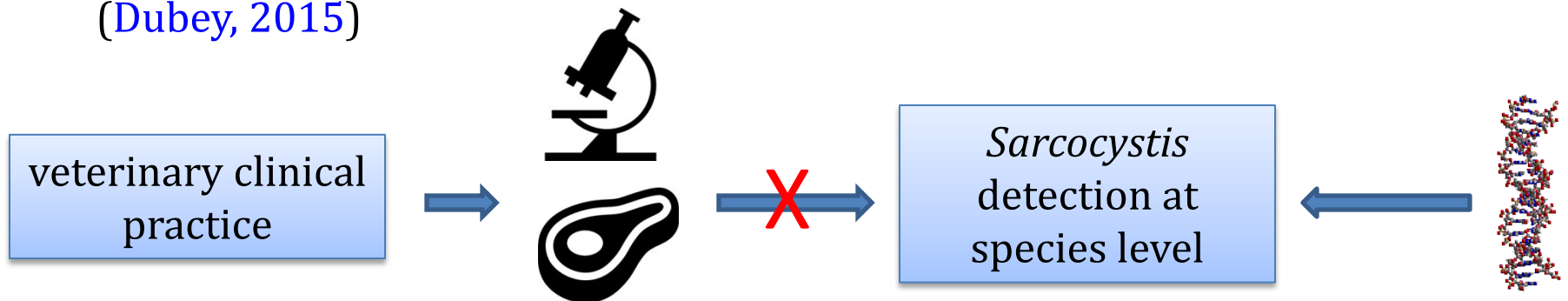
Consequences of the illness in humans (**sarcocystosis**):  
appetite loss, vomiting, nausea, diarrhea, abdominal  
pain, respiratory disorders or tachycardia



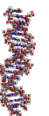
- *Sarcocystis* spp. in **cattle** (*Bos taurus*):  
*S. cruzi*, *S. hirsute*, ***S. hominis***, *S. rommeli*, ***S. heydorni*** and  
*S. sinensis* – nomen nudum (Dubey, 2015)  
*S. bovifelis*, *S. bovini* (Gjerde, 2016)



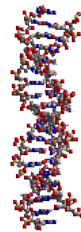
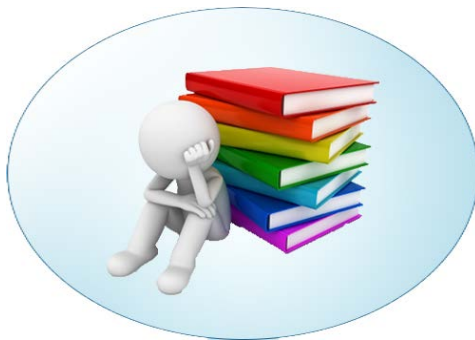
- *Sarcocystis* spp. in domestic (*Sus scrofa domesticus*)  
and feral **pigs** (*Sus scrofa*):  
*S. miescheriana*, ***S. suis***, and *S. porcifelis*  
(Dubey, 2015)



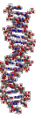
# Aim



- review the **worldwide distribution** and **molecular epidemiology** of *Sarcocystis* spp. in cattle and pigs, processing **PubMed** retrieved relevant scientific papers, published in the new century and based on molecular tools
- prevalence values of the infections according to countries and geographical regions
- incriminated *Sarcocystis* spp. and their isolation sources together with the targeted genes and used primers
- public health significance of the isolates

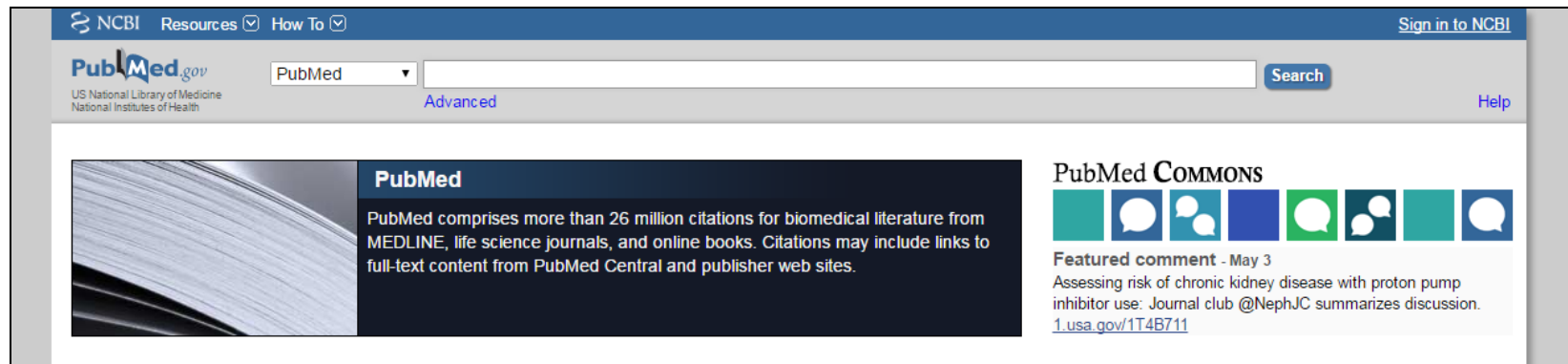


# Study design



The data for the current update covers the period from 2009 to nowadays.

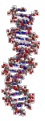
Pub Med searching using “*Sarcocystis*” and “sarcocystosis” terms combined with “pig” “wild boar” and “pork” or “cattle” “bovine” “beef”



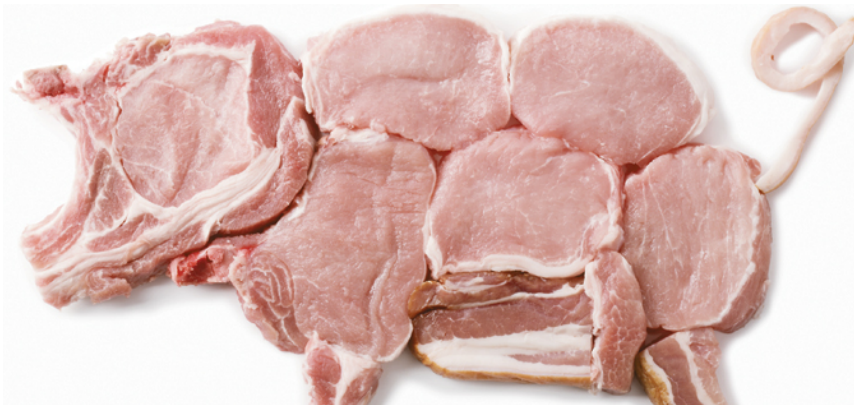
a total of 18 representative papers were selected

more than 80% out from the available published information from the scientific literature in the last 6 years.

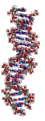
# Domestic pigs – epidemiological aspects



Host	County (geographical region)	No. of <i>Sarcocystis</i> positive samples / examined (%prevalence)	Isolation source (tissue)	Targeted gene region/primer set	Identified species (%)/no. of sequenced isolates	Reference s
<i>Sus scrofa domesticus</i>	Switzerland	1* – case study	myocardium	18S rRNA / COC1 and COC2	<i>S. miescheriana</i>	Caspari et al. 2011
	China (Henan Province- Central region)	4/4 (100%) – isolates for slaughtered pigs	diaphragm	18S rRNA / S18SF and S18SR	<i>S. miescheriana</i> /4	Yan et al. 2013
	India (Punjab)	182/250 (72.8%)	myocardium	18S rRNA /	<i>S. miescheriana</i> (n=1), <i>S. sui hominis</i> (n=6)/7	Kaur et al. 2016



# Feral pigs (wild boar) – epidemiological aspects

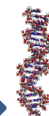


Host	County (geographical region)	No. of <i>Sarcocystis</i> positive samples / examined (%prevalence)	Isolation source (tissue)	Targeted gene region/primer set	Identified species (%)/no. of sequenced isolates	Referenc es
<i>Sus scrofa</i>	Portugal (North- Eastern)	76/103 (73.8)	diaphragm	18S rRNA / <b>Sar-F, Sar-R</b> , SmiesF and SsuihR	<i>S. miescheriana</i> /19	<a href="#">Coelho et al. 2015</a>
	United States of America	44/147 (29.9)	myocardium	18S rRNA / <b>2L</b> and <b>3H</b>	<i>S. miescheriana</i> /31	<a href="#">Calero - Bernal et al. 2015</a>
	Spain	8/25 (32)	myocardium	18S rRNA / 2L , 2H and 3H	<i>S. miescheriana</i> (n=7) <i>S. suihominis</i> (n=1)/8	<a href="#">Calero - Bernal et al. 2015</a>
	Iran	1*	thigh muscle	18S rRNA / <b>SarcoF</b> and <b>SarcoR</b>	<i>S. miescheriana</i>	<a href="#">Kia et al. 2011</a>





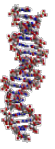
# Cattle– epidemiological aspects



Host	County (geographic al region)	No. of <i>Sarcocystis</i> positive samples / examined (%prevalence)	Isolation source (tissue)	Targeted gene region/primer set	Identified species (%)/no. of sequenced isolates	References
<i>Bos taurus</i>	Hungary	100/151 (66)	oesophagus myocardium	18S rRNA / COC1 and COC2; BJ1 and BN2	<i>S. cruzi</i> (64), <i>S. hominis</i> (19), <i>S. sinensis</i> (17)/36	<a href="#">Hornok et al. 2015</a>
	Germany	174/257 (67.7)	minced (muscle) beef	18S ribosomal DNA/ SarcoFext and SarcoRext	<i>S. sinensis</i> (n=22), <i>S. hominis</i> (n=5) and <i>S.</i> <i>hirsuta</i> (n=1)/28	<a href="#">Moré et al. 2014</a>
	Argentina	135/380 (35.5)	loin	18S rRNA / SarcoF and SarcoR	<i>S. cruzi</i> (n=16), <i>S. hominis</i> (n=13)	<a href="#">Moré et al. 2011</a>
	Italy (North- Western)	279/384 (72.7) 228/384 (59.4) 225/384 (58.6)	oesophagus diaphragm myocardium	18S rRNA / -	<i>S. cruzi</i> (285), <i>S. hominis</i> (164), <i>S. hominis</i> -like (71), <i>S. hirsuta</i> (7)	<a href="#">Domenis et al. 2011</a>
	Iran	90/101 (89.1)	diaphragm	18S rRNA / designed primers	<i>S. cruzi</i> (89), <i>S. hominis</i> (48)	<a href="#">Akhlaghi et al. 2016</a>
		40/123 (32.5)	oesophagus diaphragm myocardium intercostals muscle	18S rRNA / 18S1H and 18S9L	<i>S. cruzi</i>	<a href="#">Hamidinejat et al. 2015</a>
		1*	hamburger	18S rRNA / SarF and SarR	<i>S. hominis</i>	<a href="#">Ahmadi et al. 2015</a>
			diaphragm	18S rRNA	<i>S. hominis</i>	<a href="#">Hajimohammadi et al. 2014</a>
	Malaysia	49/77 (63.6)	myocardium	18S rRNA	<i>S. cruzi</i> (49)	<a href="#">Latif et al. 2015</a>
	Vietnam	63/101 (62.4)	oesophagus, diaphragm, tongue and cervical muscle	18S rRNA / 18S9L and 18S1H	<i>S. cruzi</i> (55), <i>S. hominis</i> (54), <i>S. hirsuta</i> (28)	<a href="#">Jehle et al. 2009</a>



# Cattle– epidemiological aspects

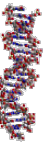


Host	County (geographical region)	No. of <i>Sarcocystis</i> positive samples / examined (%prevalence)	Isolation source (tissue)	Targeted gene region/primer set	Identified species (%)/no. of sequenced isolates	Referen ces
<i>Bos taurus</i>	Argentina	147 excised individuals sarcocysts from 12 samples of cattle meat	beef muscle tissue	Partial <i>Cox1</i> /SF1 and SR9 and for 7 isolates of <i>S. bovifelis</i> SR8D and and SR9; and Ss1R, Ss2R for <i>S. bovifelis</i> , Sb1R and Sb2R for <i>S. bovini</i>	<i>S. bovifelis</i> (n=67) <i>S. bovini</i> (5), <i>S. hirsuta</i> (1) <i>S. cruzi</i> (n=21)	Gjerde, 2016
	New Zealand				<i>S. bovifelis</i> (n=38), <i>S. bovini</i> (19) <i>S. hirsuta</i> (19)	
	Brazil				<i>S. bovifelis</i> (n=15), <i>S. hirsuta</i> (15)	
	Uruguay				<i>S. bovifelis</i> (n=1) <i>S. cruzi</i> (1)	
	Germany				<i>S. bovifelis</i> (n=26) <i>S. hirsuta</i> (n=21)	



# Conclusions

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- **Scanty background** of molecular studies, **especially in European countries**, even if the food safety regulations require the surveillance of protozoa in meat and derivate products by each member state.
- In domestic pigs and wild boar: **widespread** occurrence of *S. miescheriana*, a narrower appearance of *S. sui hominis* (only 2 studies) and the lack of *S. porcifelis* with uncertain taxonomic status
- In cattle: *S. cruzi* and the zoonotic *S. hominis* seems to be the **most prevalent** species, followed by *S. sinensis* (currently recognized as *nomen nudum*) and *S. hirsuta*. In addition, the presence of a considerable number of *S. bovis* and *S. bovini* has been reported in a single study, and currently they can be considered genetically distinct species.
- The relatively reduced number of publications presenting *Sarcocystis* molecular epidemiologic background in livestock meat intended for human consumption, pointed out the **necessity of the increasing the number of surveys** dedicated to this topic at worldwide level and special emphasis on European countries.

# Thank you for your attention!

