Effects of maternal nutrition on immune competence and microbiota composition of piglets

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Introduction

- Programming intestinal immune system via maternal intervention
 - Transmission of mucosal immune memory
 - Chickens: Rebel et al. (2006); Comp. Biochem. Physiol. 145:502-508
- Microbial colonisation initiates maturation and programming of intestinal immune system
- Program intestinal immune system piglets via maternal intervention on microbial colonisation e.g. via:
 - Transmission via vaginal flora
 - Transmission via faeces
 - Transmission via colostrum and milk





Experimental set-up

AIM:

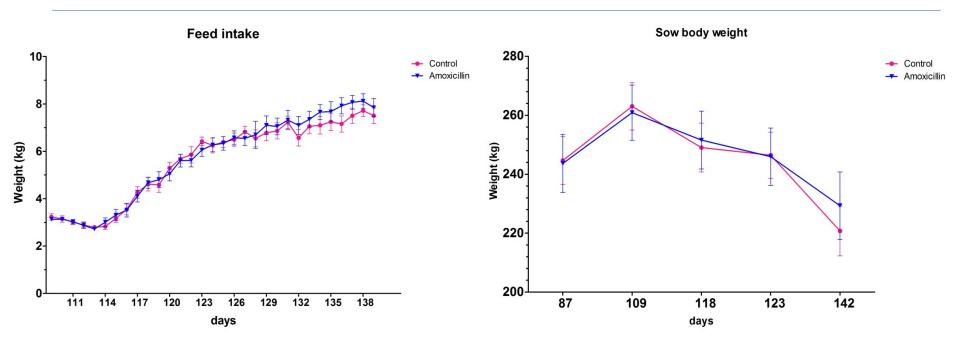
Study the effect of in feed amoxicillin treatment in sows on microbial composition and intestinal development of piglets

- Sows divided over 2 treatments:
 - Regular lactation feed (n = 16)
 - Lactation feed with amoxicillin for 1 week (n = 15)
- Effects of treatment on sows and offspring determined





Zootechnical parameters: Sow lactation feed intake + weight



RA2

- No significant differences in feed intake nor in body weight due to amoxicillin addition
- No significant differences in reproduction due to amoxicillin treatment

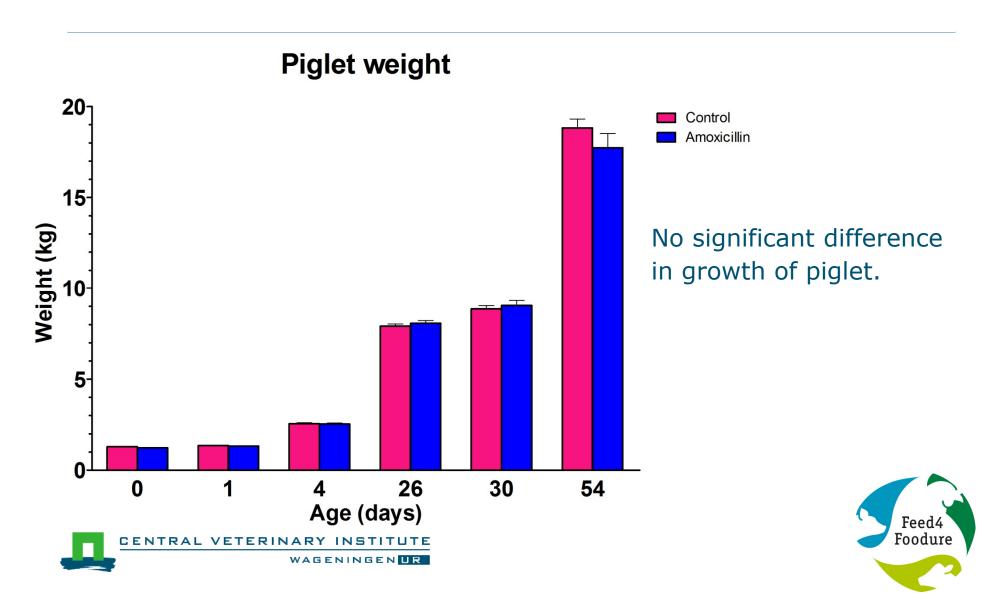


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mag dit of kan dit er beter Rebel, Annemarie, 12/08/2014 RA2

Zootechnical parameters:

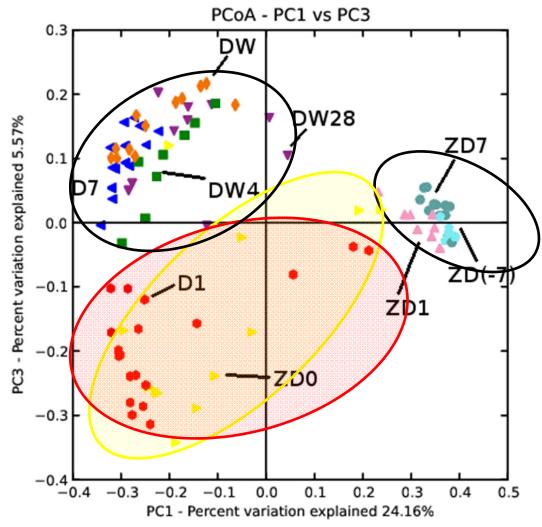
Piglet weight



Microbiota composition

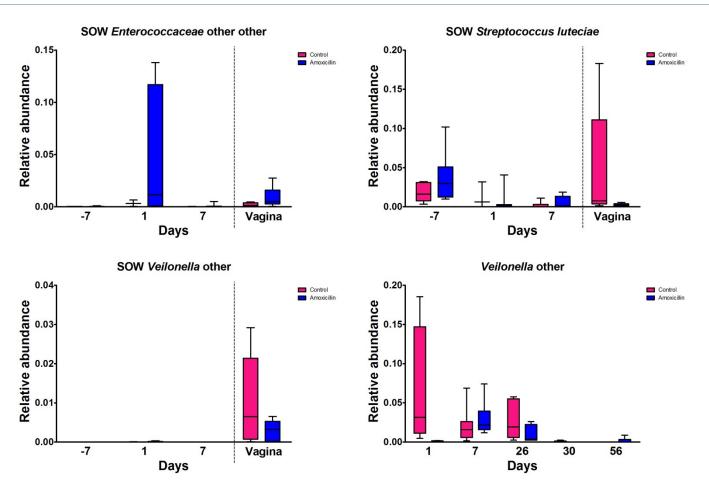
Transmission route sow - piglets

- Microbiota composition
 - Faeces sows around farrowing (ZD-7, ZD1, ZD7)
 - Vagina sows after farrowing
 (ZD0)
 - Ingesta proximal jejunum piglets
 (D1, D7, DW, DW4, DW28)
- Microbiota composition piglets day 1 overlaps microbiota sow vagina
- No significant differences between treatment groups



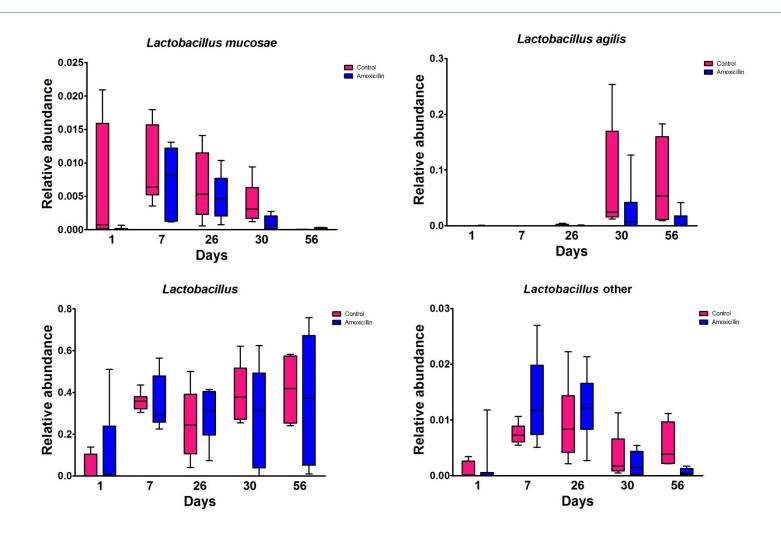


Microbiota composition Effect of amoxicillin treatment





Specific changes in microbiota piglets





Intestinal microbiota

Conclusions

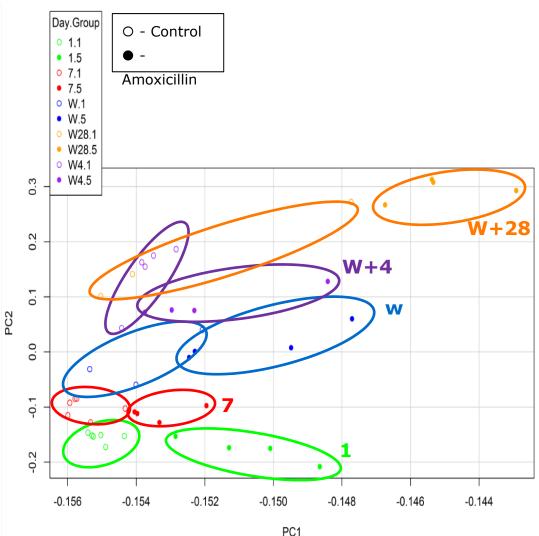
- Significant differences between treatment groups at level of specific microbiota in sow and piglets
- Microbiota composition piglets day 1 overlaps microbiota sow vagina

Intestinal gene expression offspring

Can maternal feed interventions affect immune competence?

- Transcriptional analysis of proximal jejunum offspring
- Principle component analysis (PCA) demonstrates differences between control and amoxicillin piglets
- PCA shows development as function of time in both groups





Intestinal gene expression offspring Statistics of regulated probes and genes in offspring

Table 1. Descriptive statistics of regulated probes/genes between treatments on day 1, 7, W, W+4, and W+28.

Contrast	 Day	Regulated Probes ¹		Regulated Annotated Genes ¹	
		Up	Down	Up	Down
Amoxicillin vs control	1	159	28	52	10
Amoxicillin vs control	7	0	1	0	0
Amoxicillin vs control	28	194	301	95	148
Amoxicillin vs control	32	11	153	1	84
Amoxicillin vs control	56	154	272	67	156

¹ logFC>|1.5| and adjusted p-value < 0.05

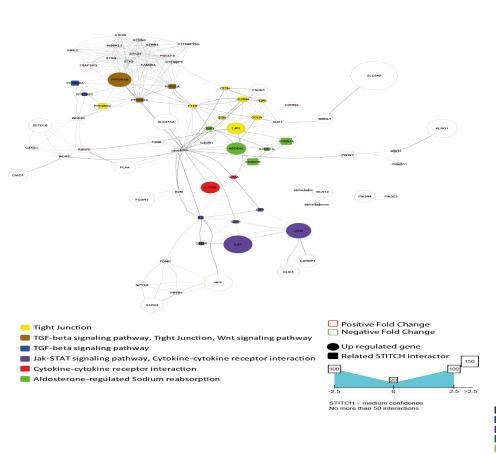
- Most pronounced changes at weaning
- Only part of regulated probes has been annotated (30 50%)
- Limited regulation at day 7





Intestinal gene expression offspring day1

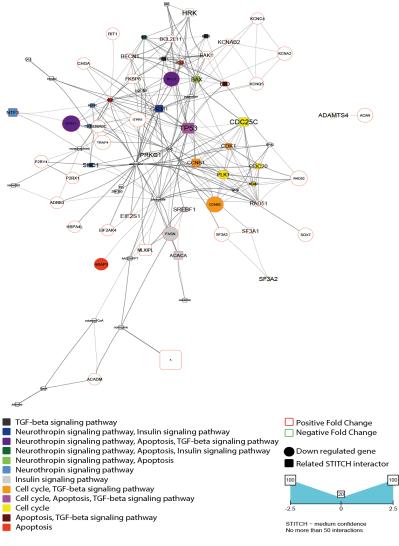
Up regulated genes



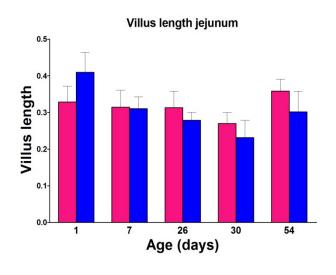
CENTRAL VETERINARY INSTITUTE

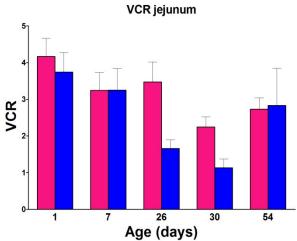
WAGENINGEN UR

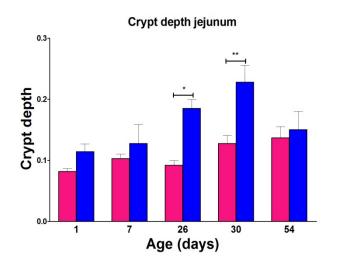
Down regulated genes

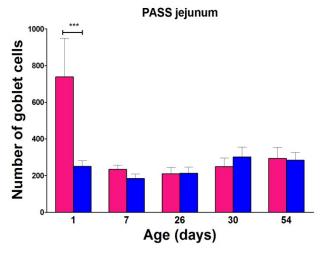


Parameters intestine piglets











Intestinal development offspring Conclusions

- Transcriptional differences in offspring due to maternal feed intervention
- Transcriptional differences translates in intestinal cellular parameters
- Treatment of sows with amoxicillin leads to a difference in immune processes at day 1 with decreased number of goblet cells
- Treatment of sows with amoxicillin leads to difference apoptosis processes at weaning with an increase in crypt depth







Conclusion

- Treatment sows with amoxicillin no effect on performance piglets until day 42.
- Treatment of sows with amoxicillin has effect on microbiota and gene expression of piglets as measured
 - Either due to changes in microbiota composition of sows via vaginal microbiota or oral-fecal transmission.
- Possibility to modulate the intestinal development as well as microbiota of piglets by maternal feed intervention
- Follow up: determine correlation microbiota with gene expression

Credits

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- Astrid de Greeff
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- Alex Bossers
- Freddy de Bree
- Frank Harders
- Ralph Kok

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