



# Dangerous liaisons between gene and microbiota: the example of Card9 in IBD

Harry SOKOL



Equipe AVENIR U1157

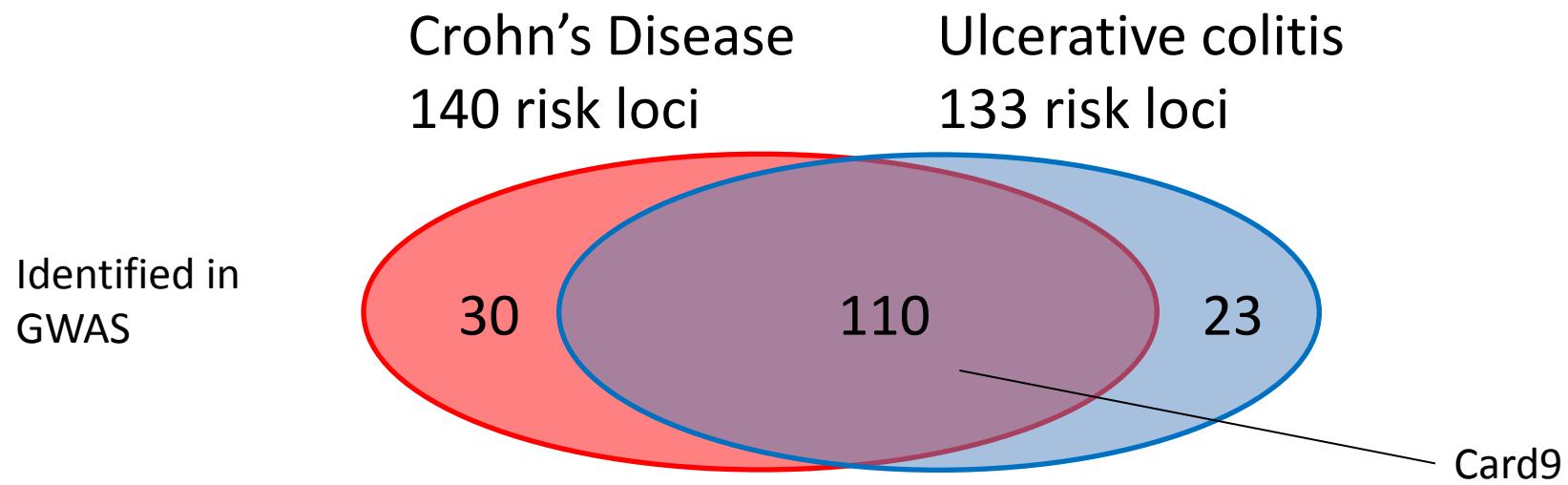


Gastroenterology Department  
Saint-Antoine Hospital

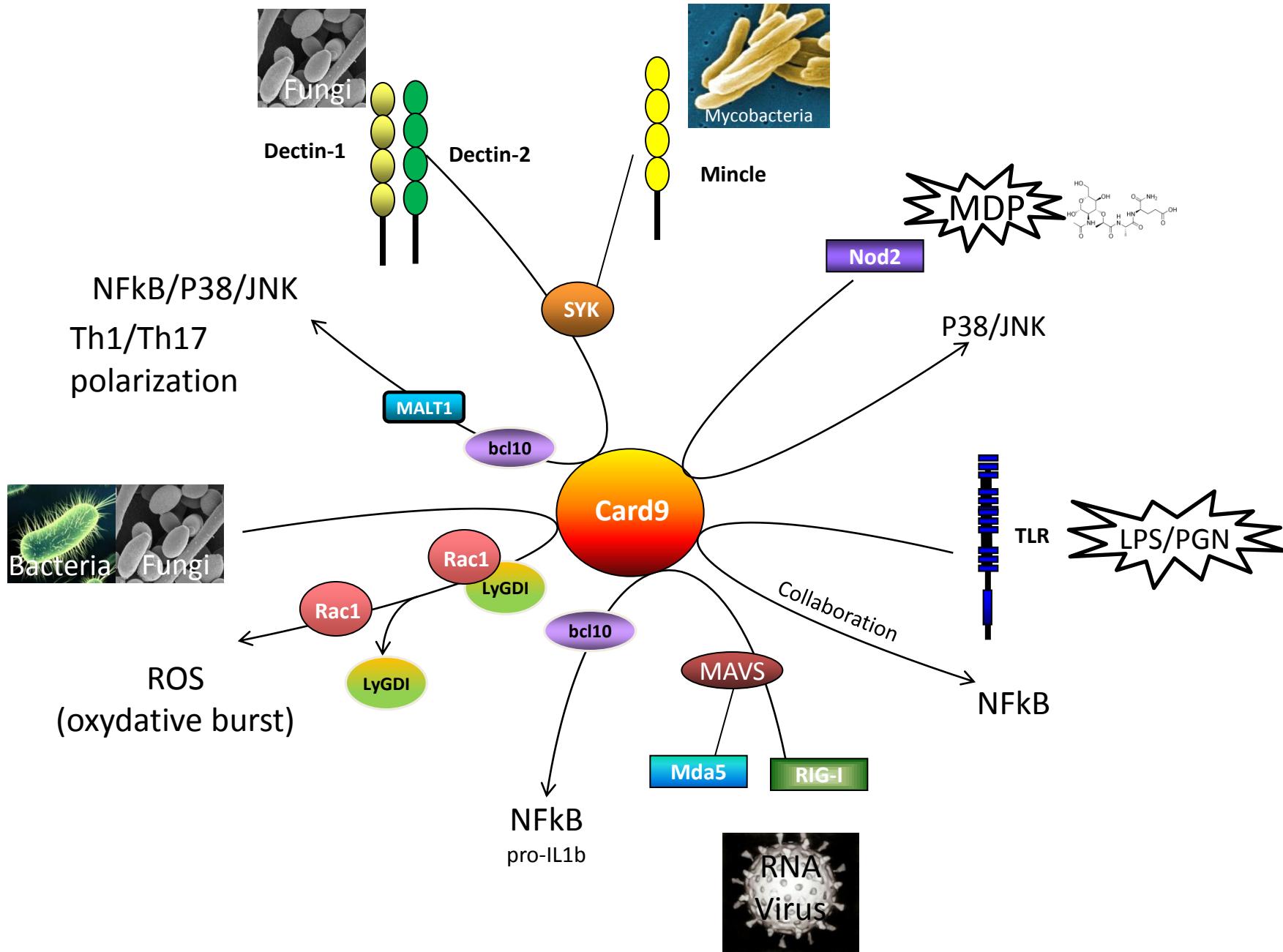


ProbiHote

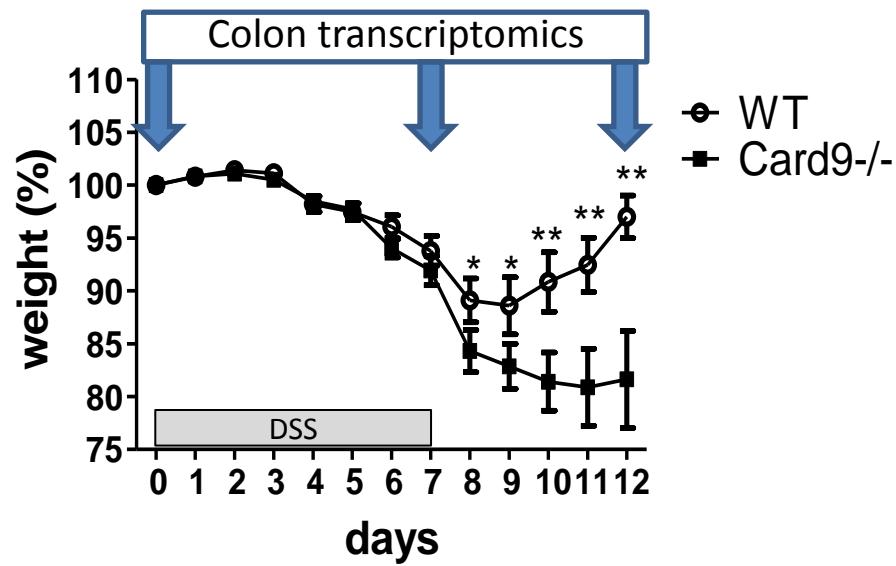
# Genes in IBD



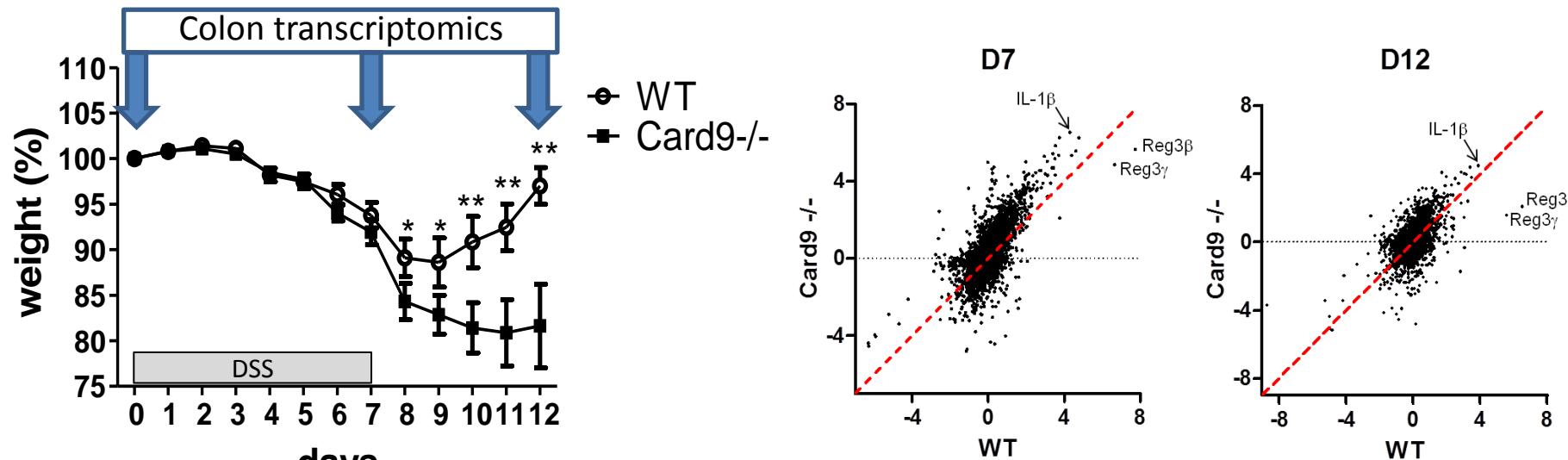
# Card9 : a central adaptor molecule in innate immunity



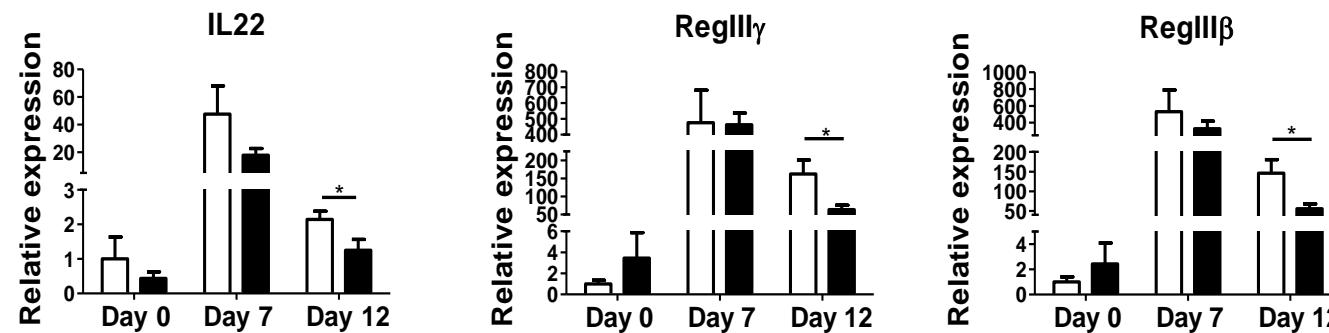
# Card9 KO mice are more susceptible to DSS-induced colitis



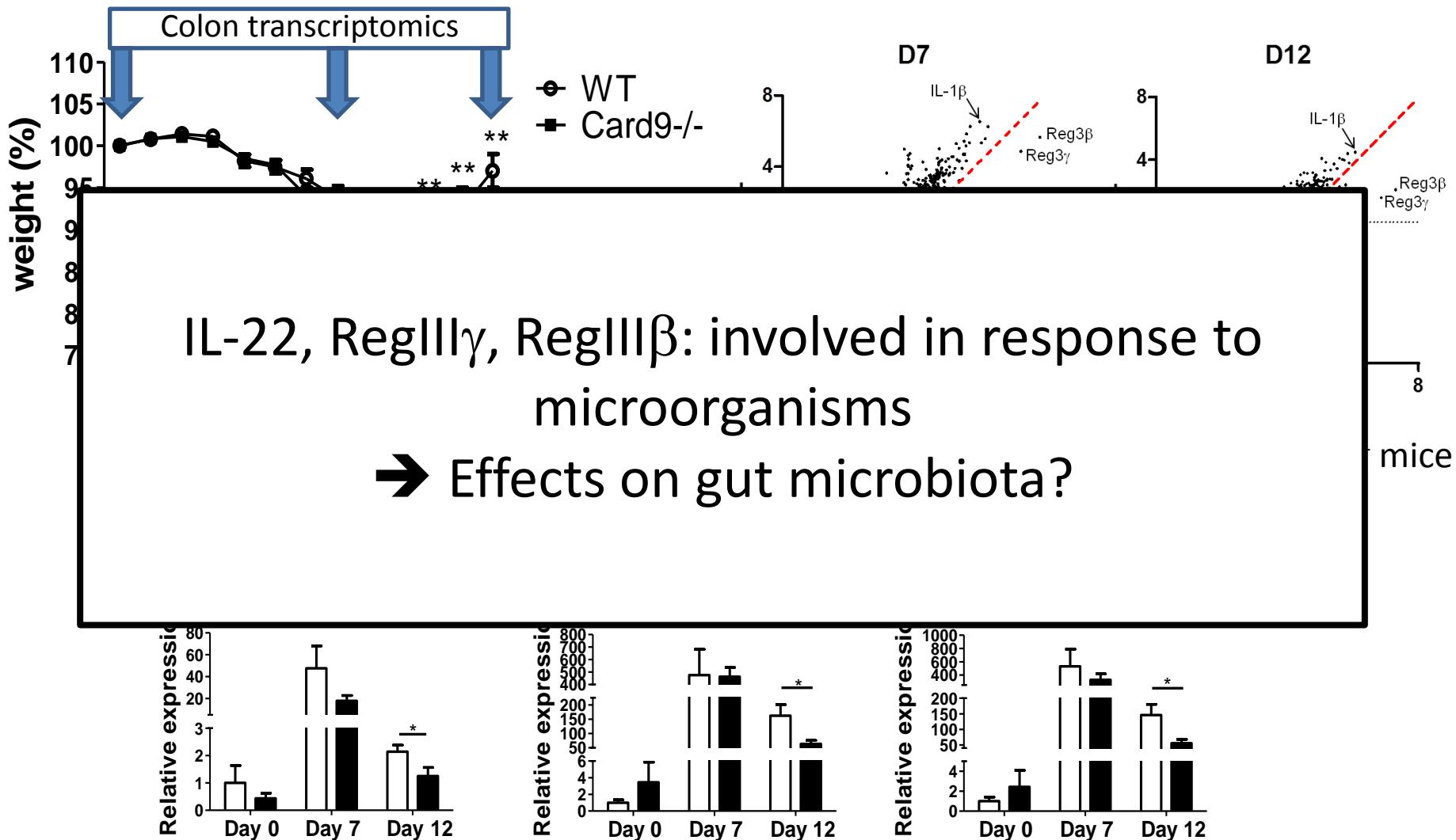
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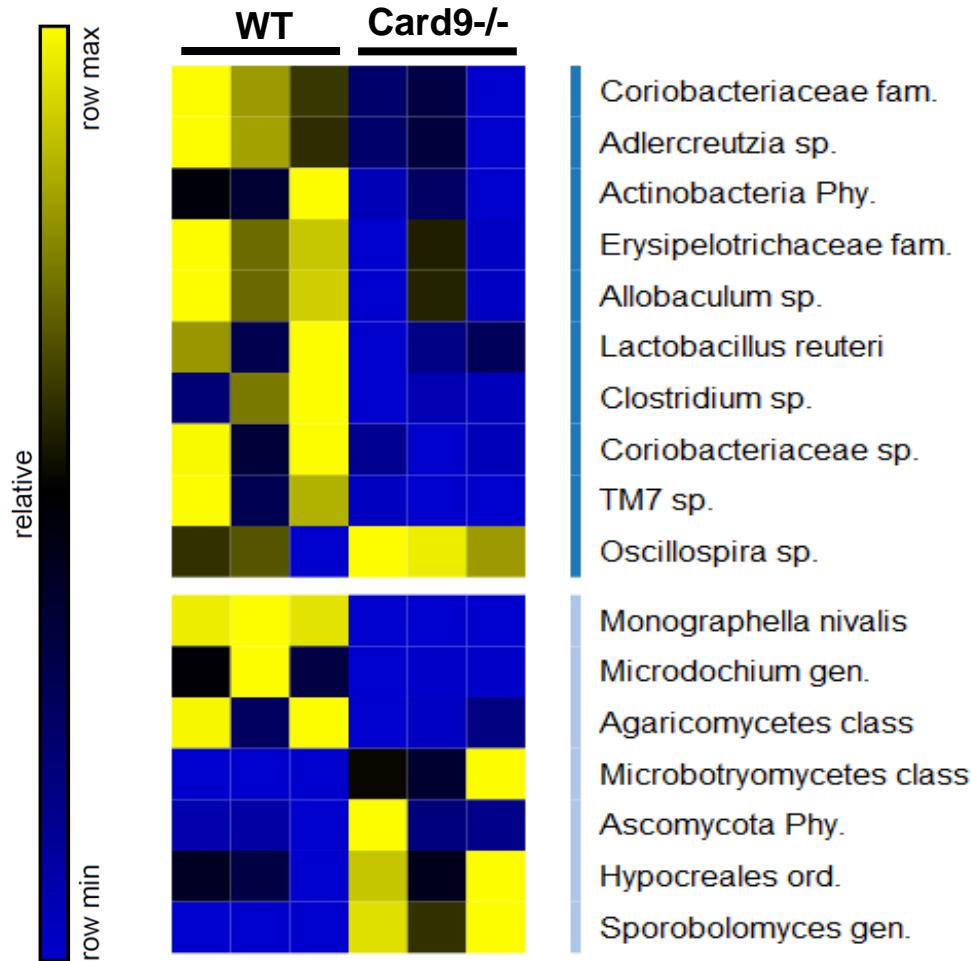
Reg3 $\gamma$  and Reg3 $\beta$  down regulated in Card9<sup>-/-</sup> mice



# Card9 KO mice are more susceptible to DSS-induced colitis

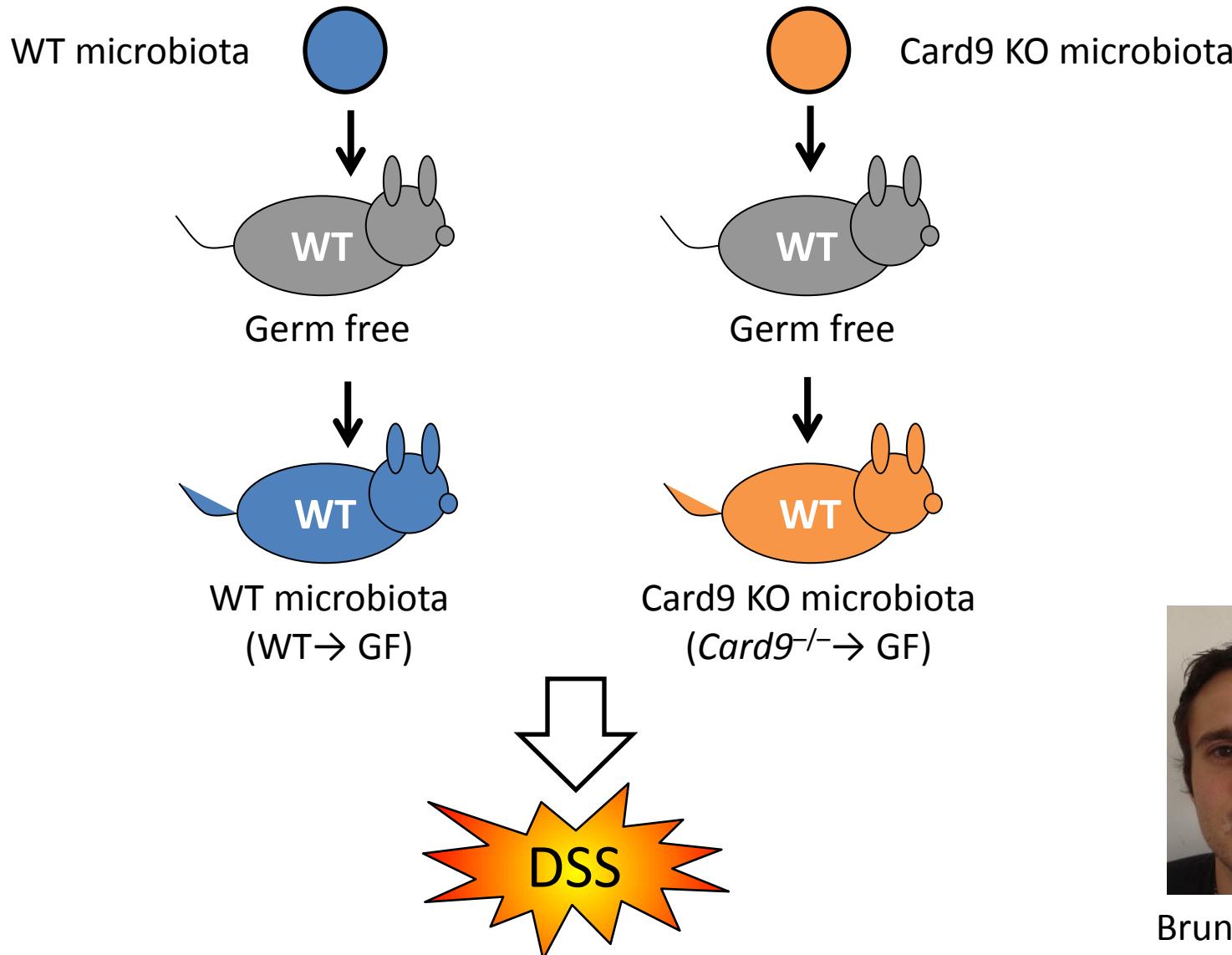


# Card9 KO mice have an altered bacterial and fungal microbiota



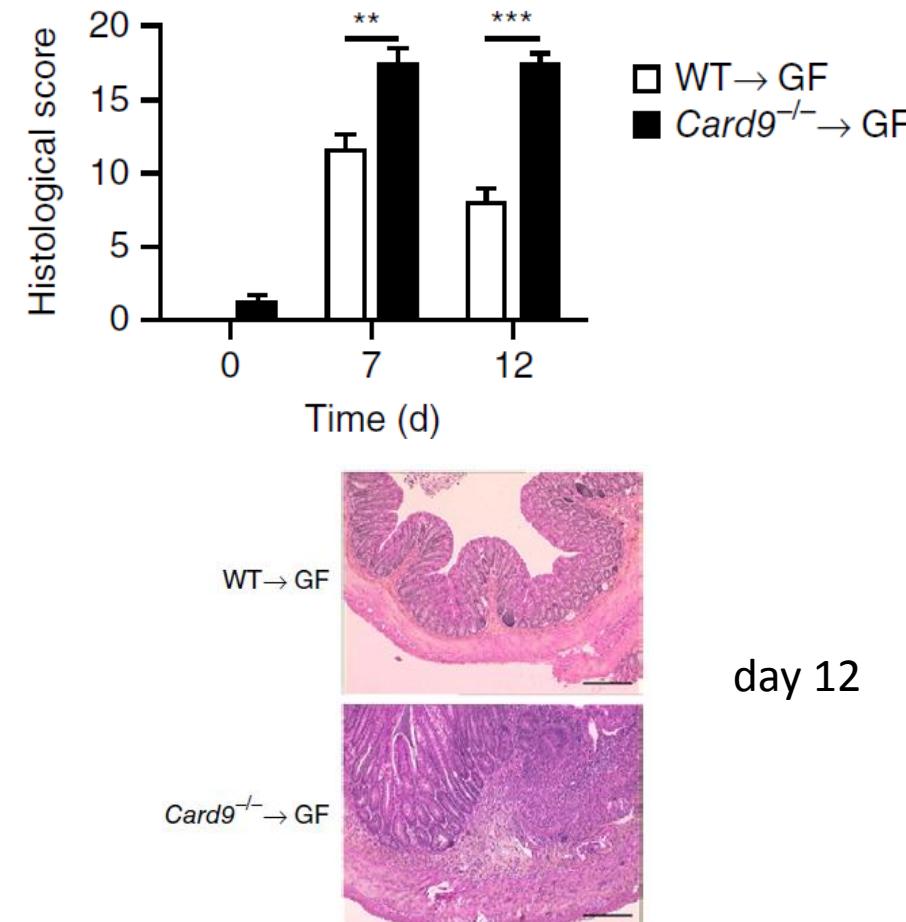
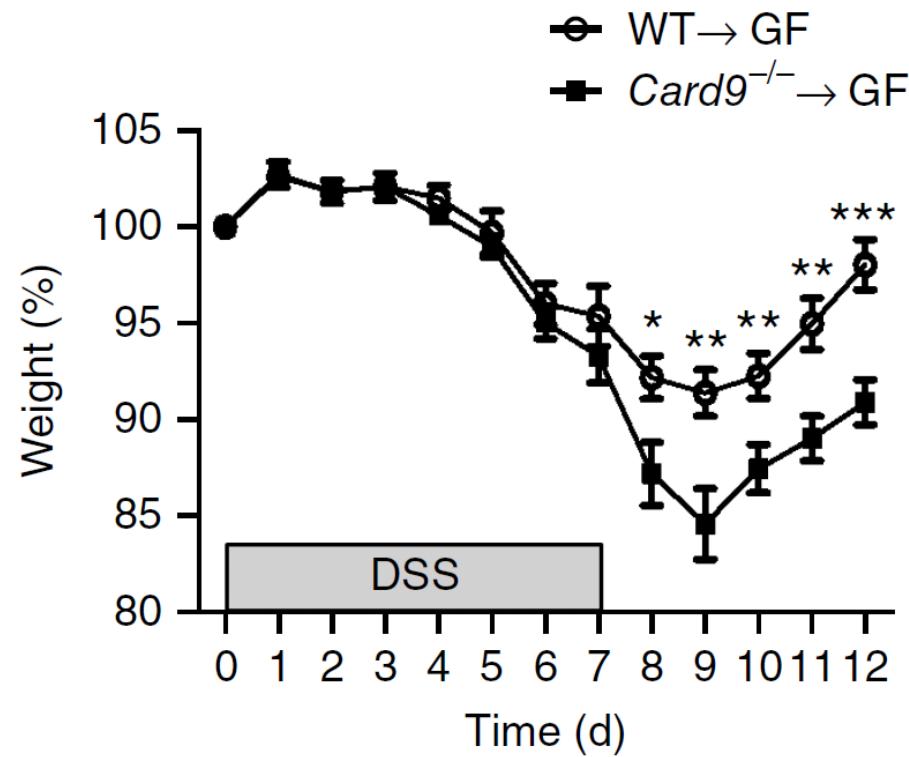
Does the dysbiotic microbiota play a role in inflammation by itself?

# Role of the microbiota in Card9 KO-mice susceptibility to inflammation



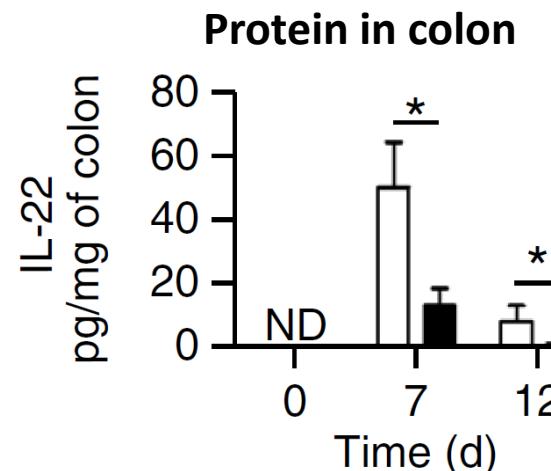
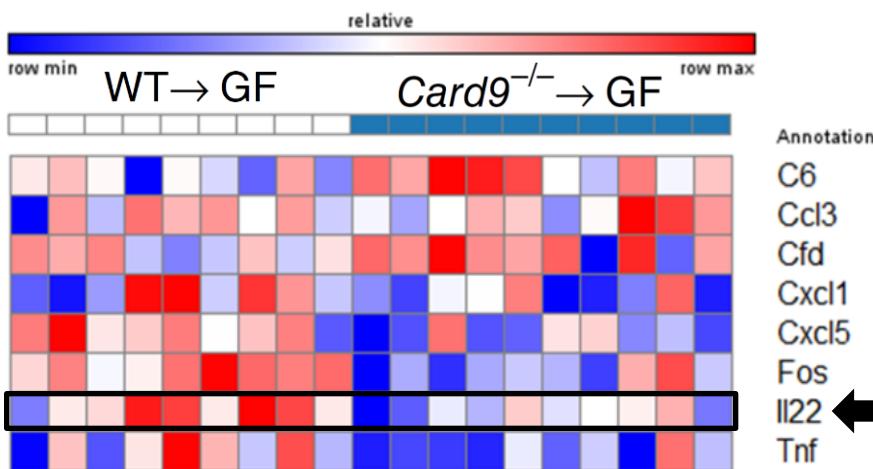
Bruno Lamas

# The microbiota from *Card9*<sup>-/-</sup> mice exerts proinflammatory effects

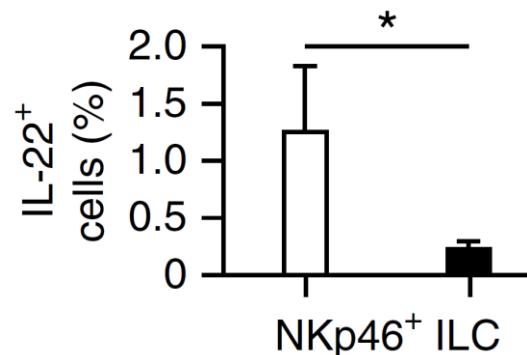
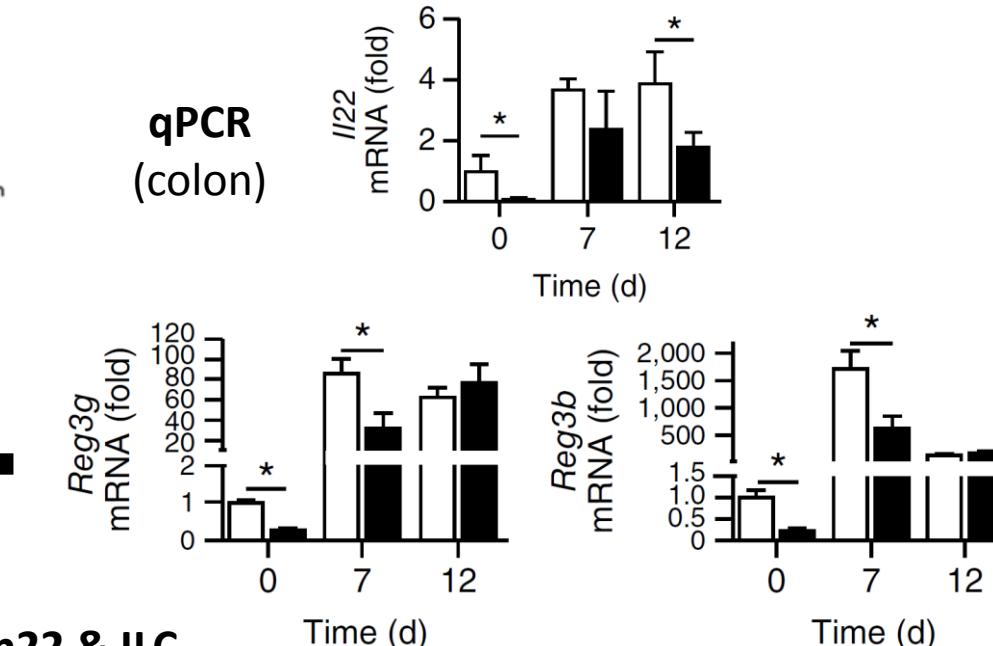
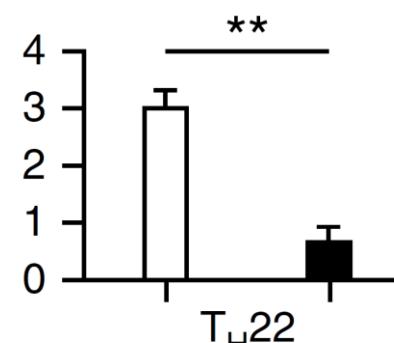


# The IL22 pathway is impaired *Card9*<sup>-/-</sup> → GF mice

Transcriptomic (Nanostring)

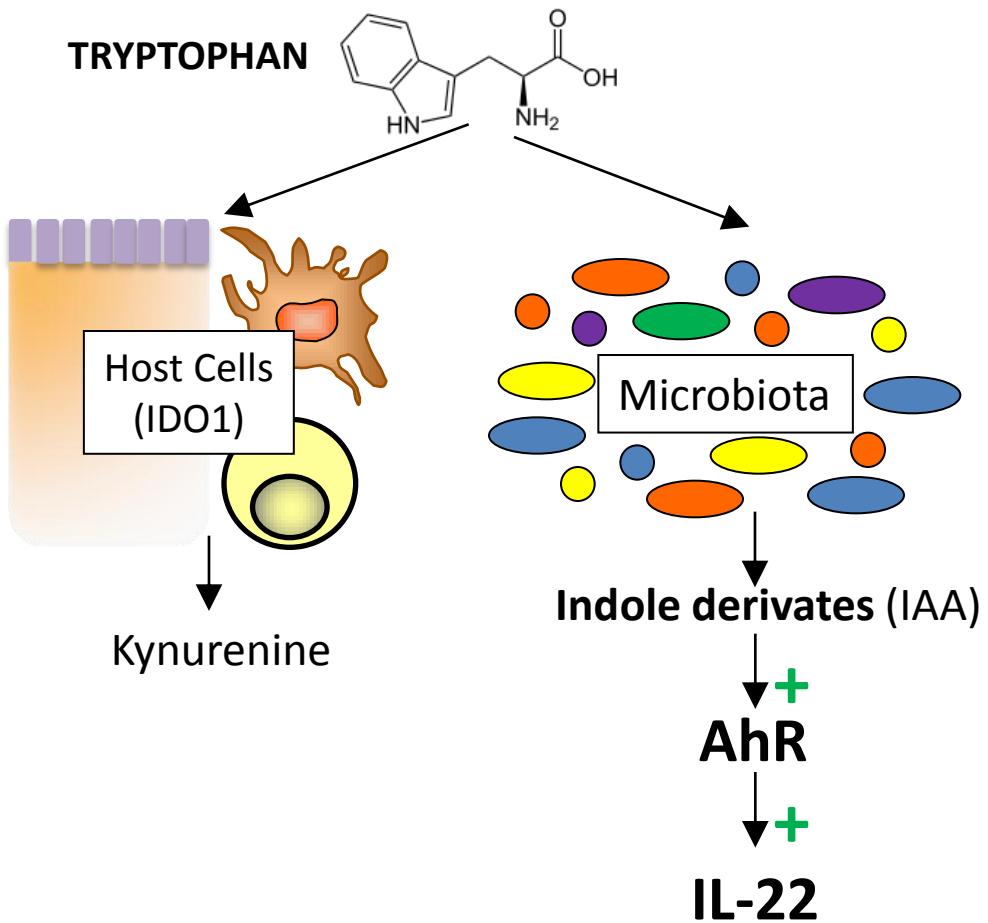


Th22 & ILC

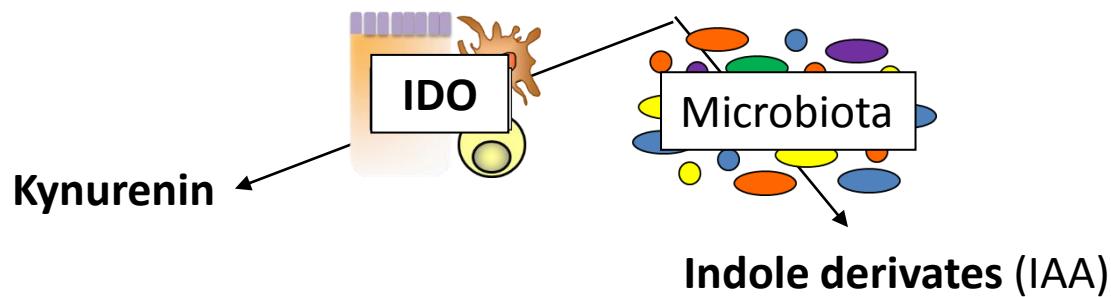
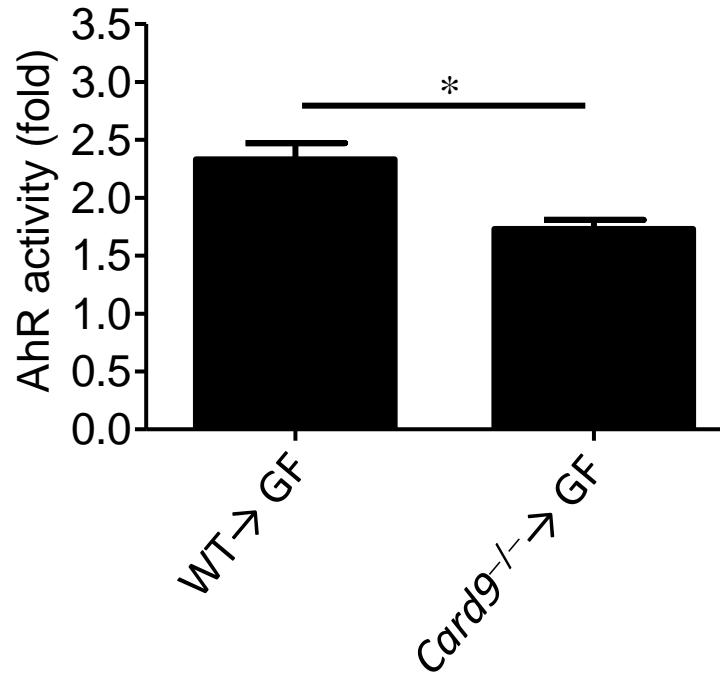
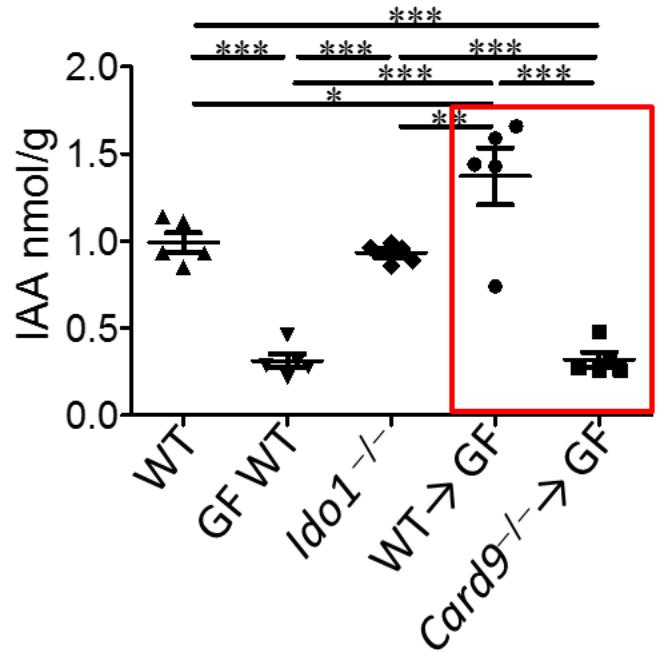


# Transfer of the IL22 defect through microbiota

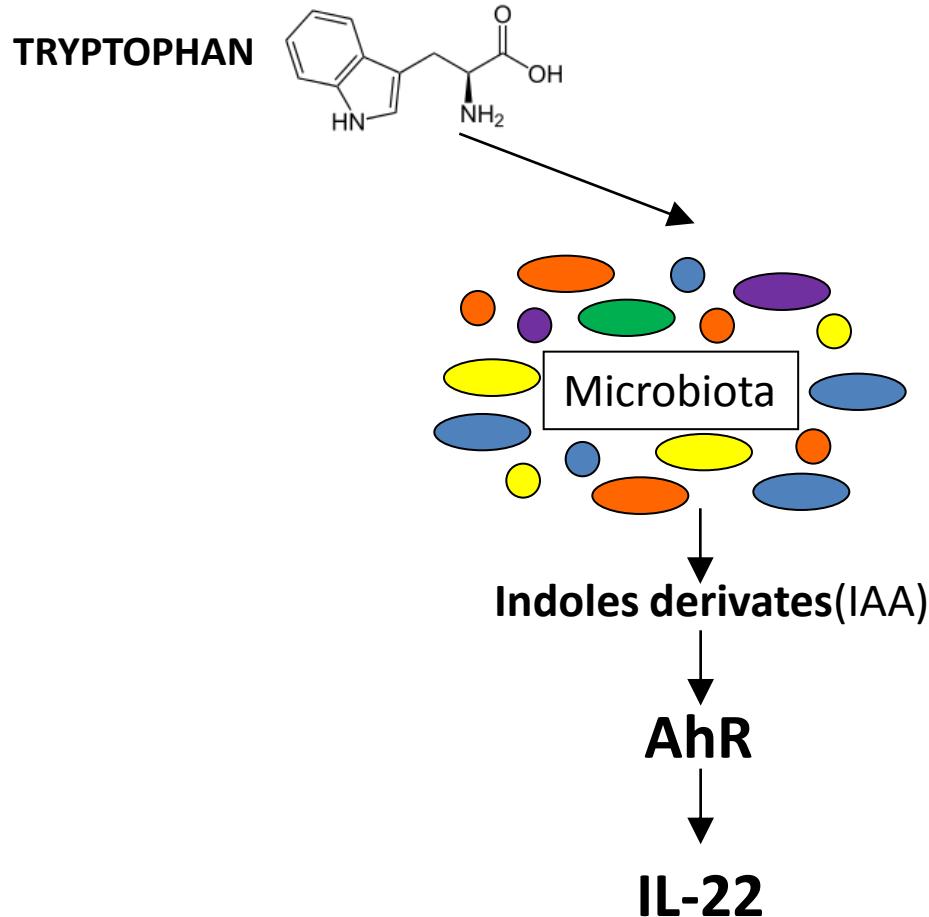
hypothesis = impaired function of the microbiota



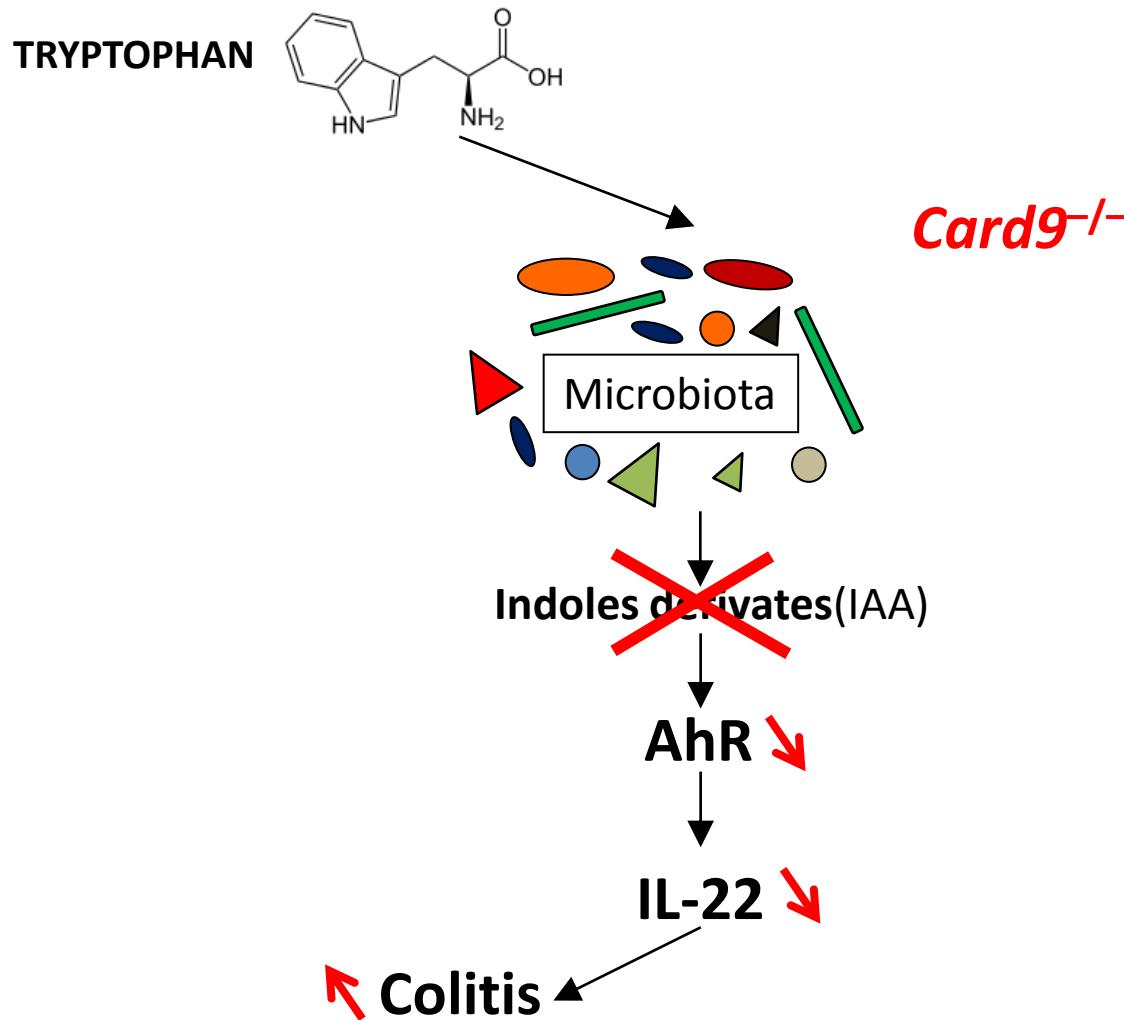
# *Card9<sup>-/-</sup>* → GF mice exhibits impaired tryptophan metabolism



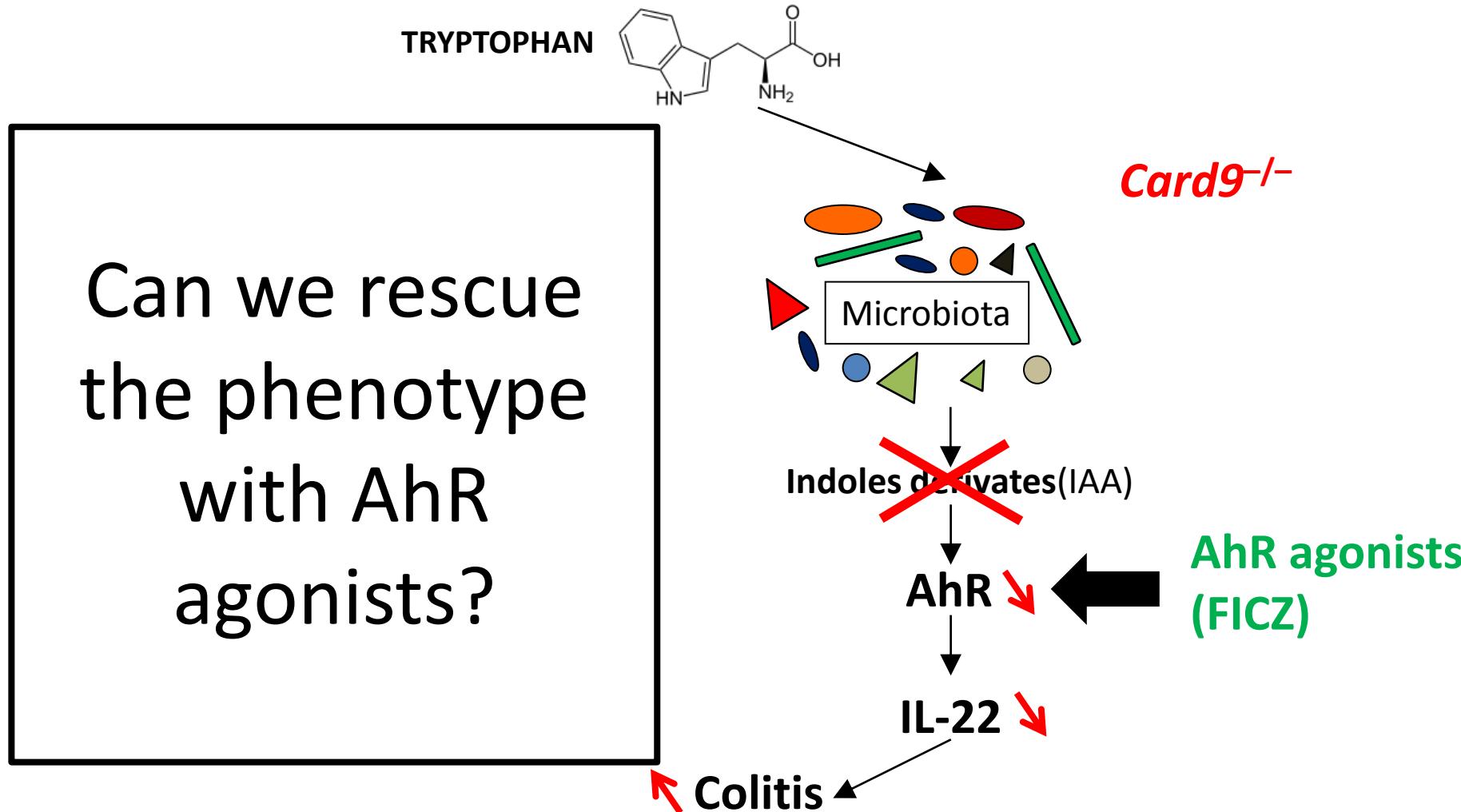
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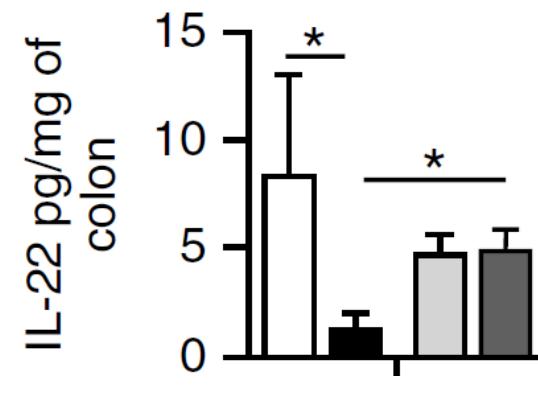
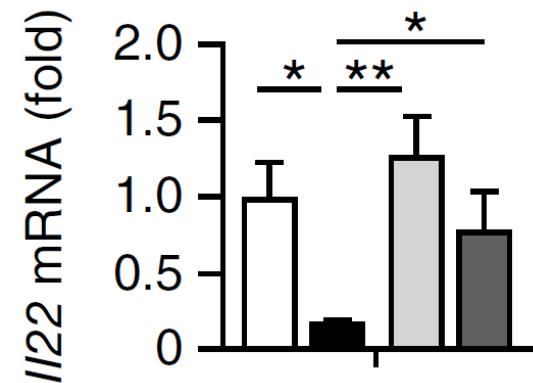
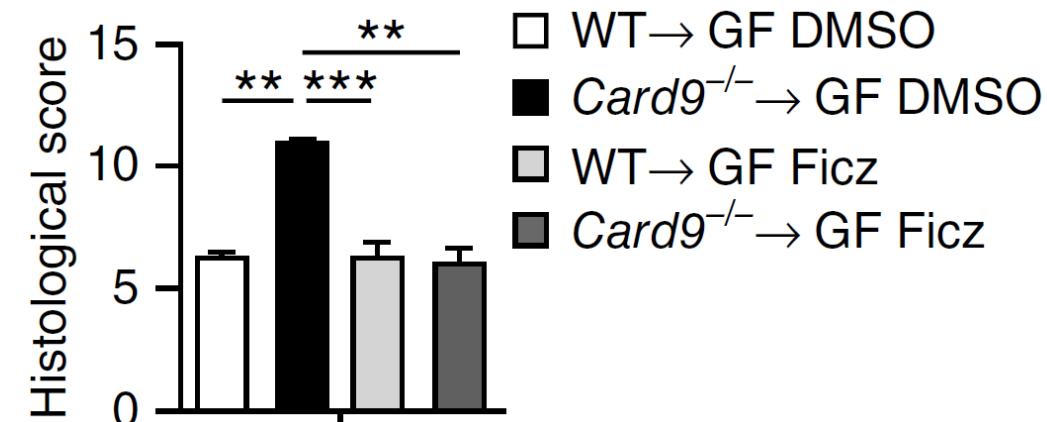
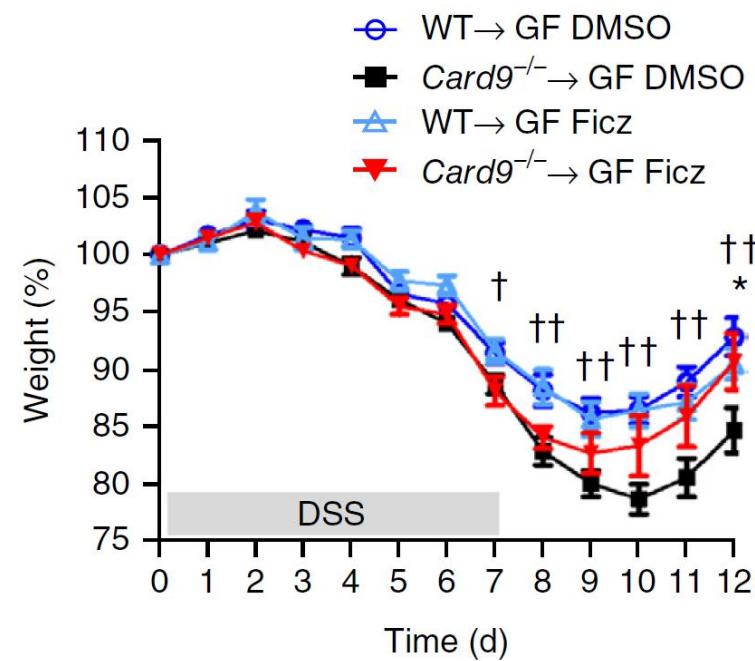
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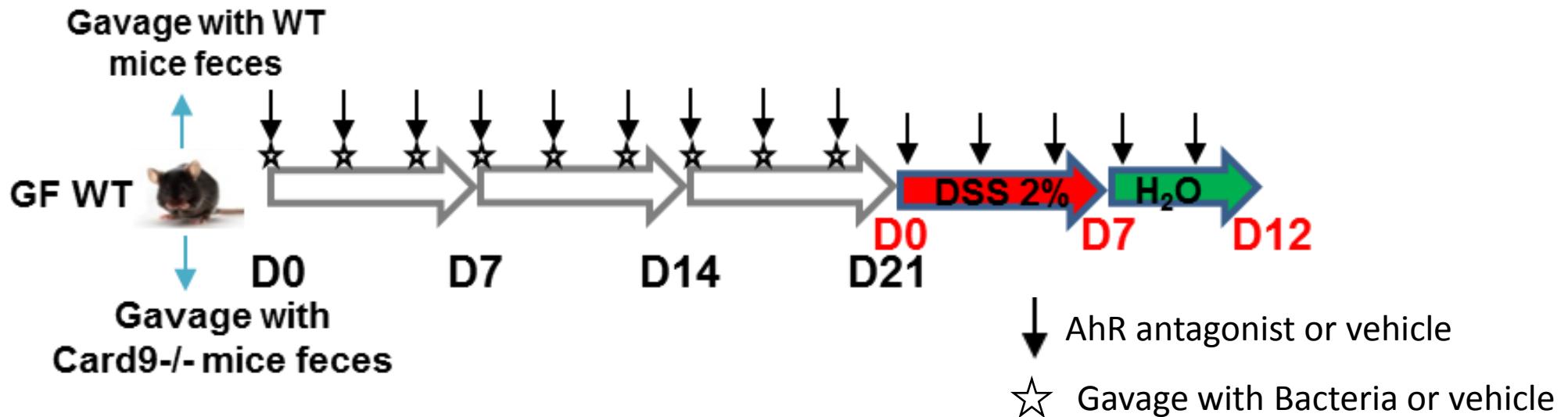
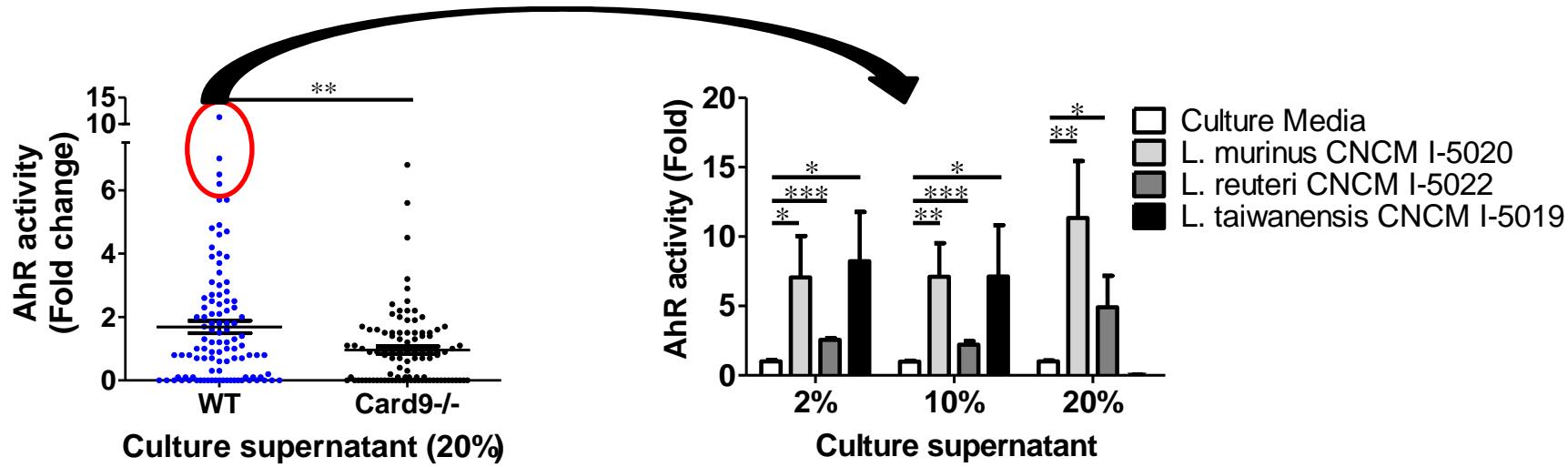
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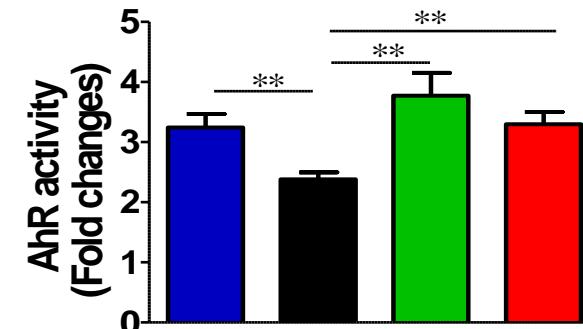
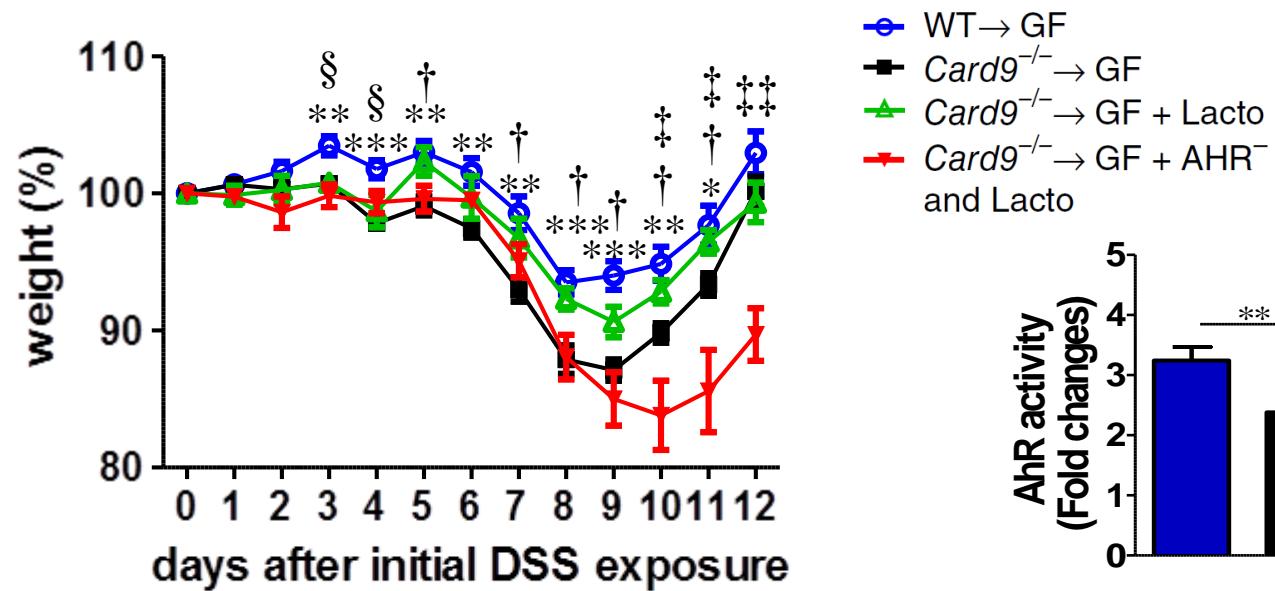
# AhR agonist rescue the phenotype in *Card9*<sup>-/-</sup> → GF mice



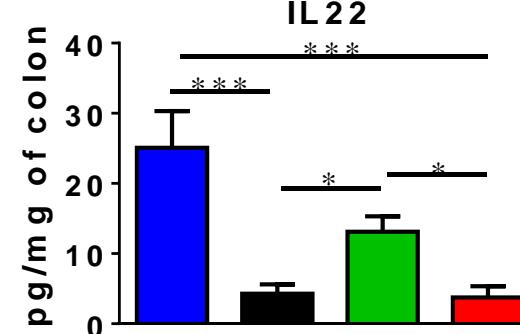
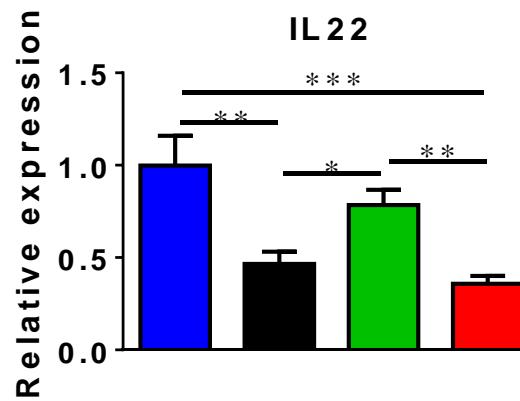
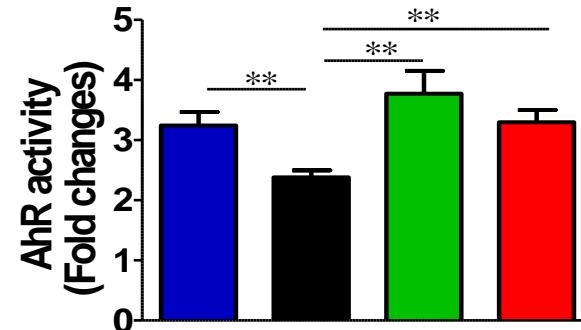
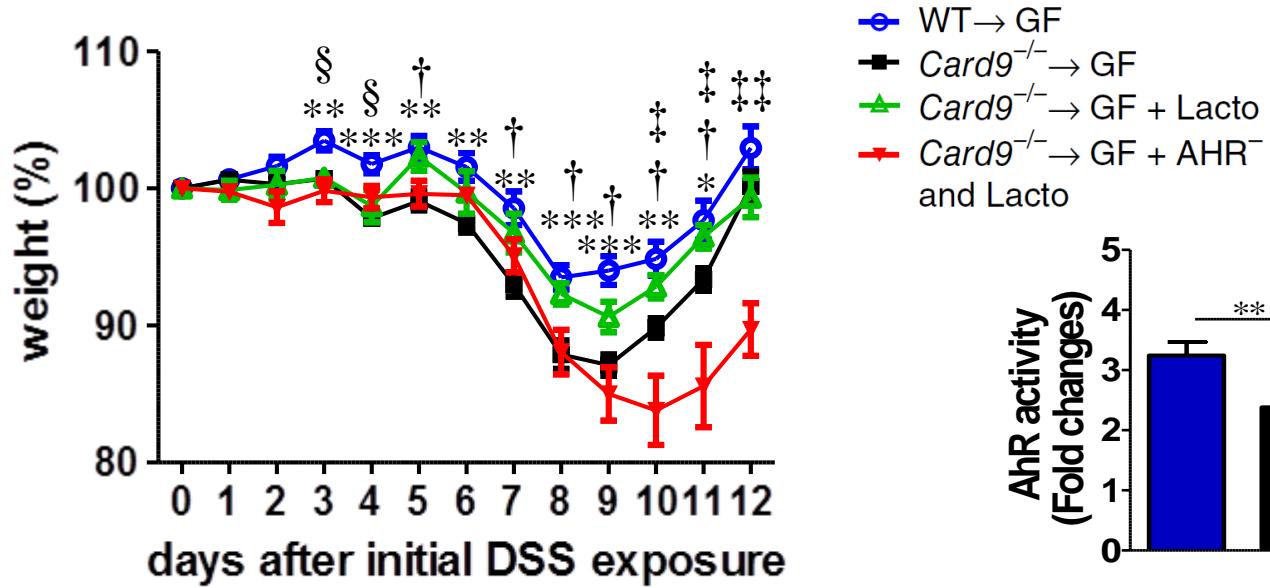
# AhR ligand-producing bacteria rescue the phenotype in *Card9<sup>-/-</sup>* → GF mice



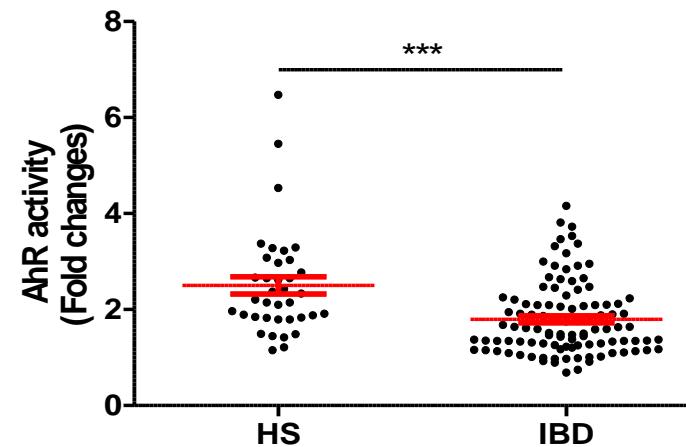
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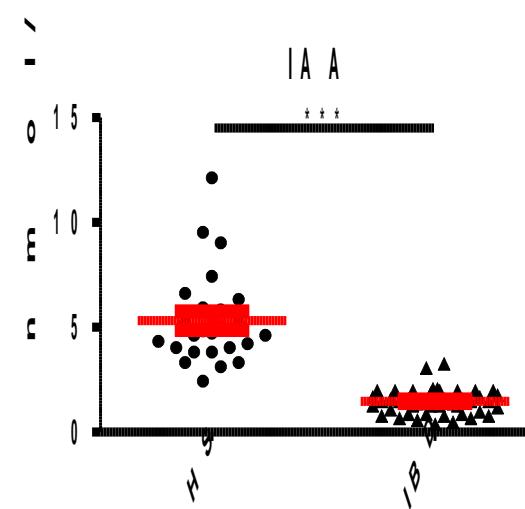
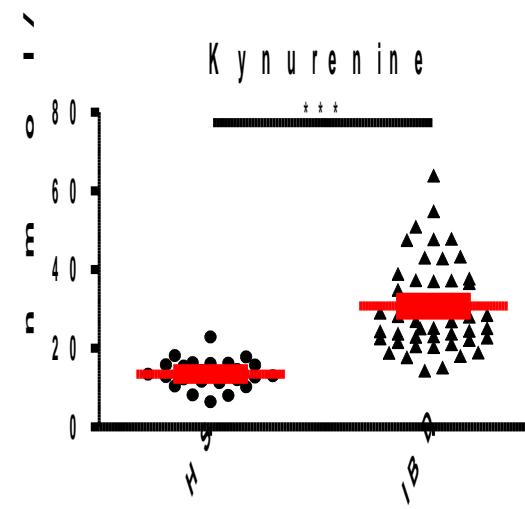
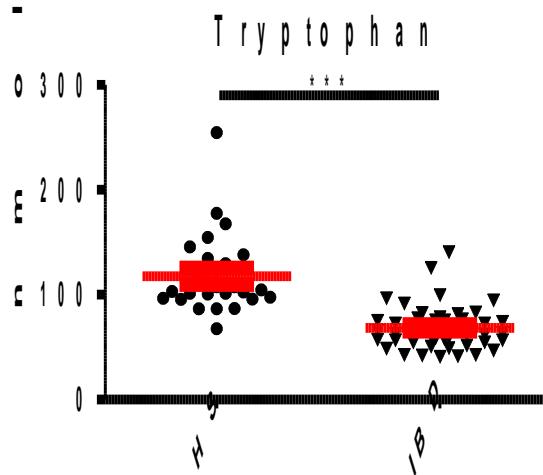
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# Relevance in Human?



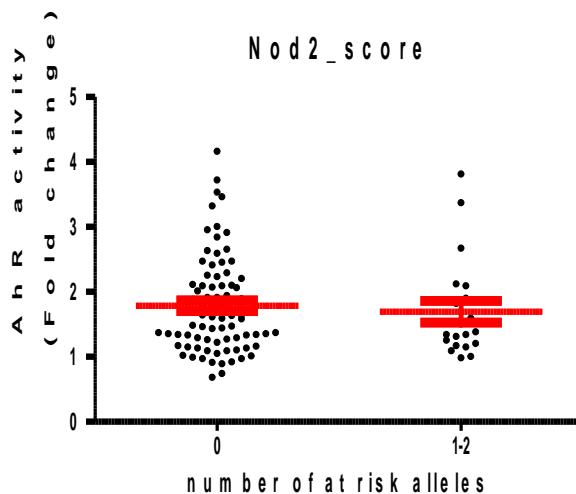
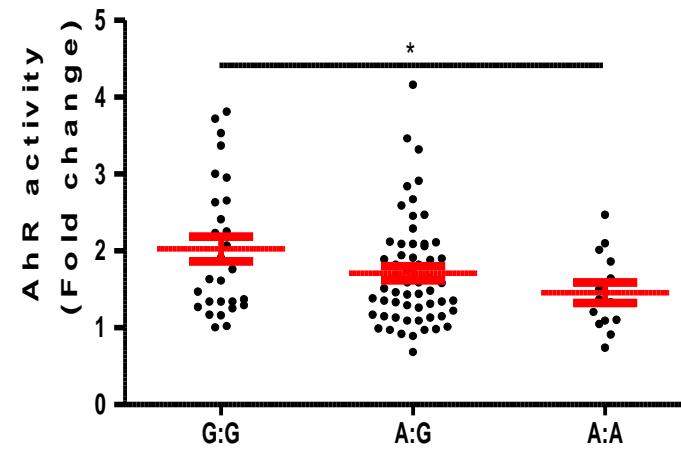
IBD: n = 112  
HS : n = 37



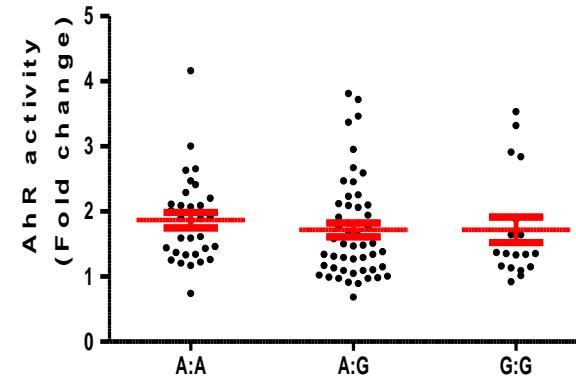
# Relevance in Human?

## Card9 SNP associated with IBD

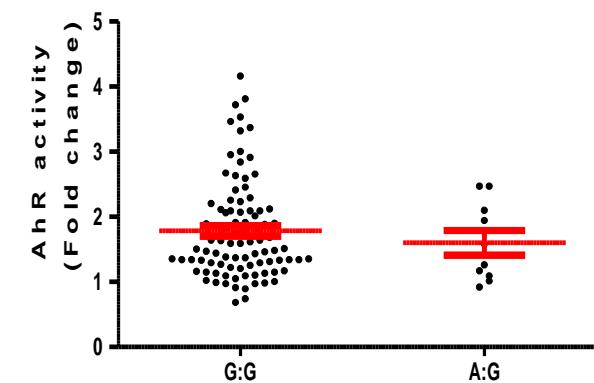
rs10781499



ATG16L1\_rs12994997

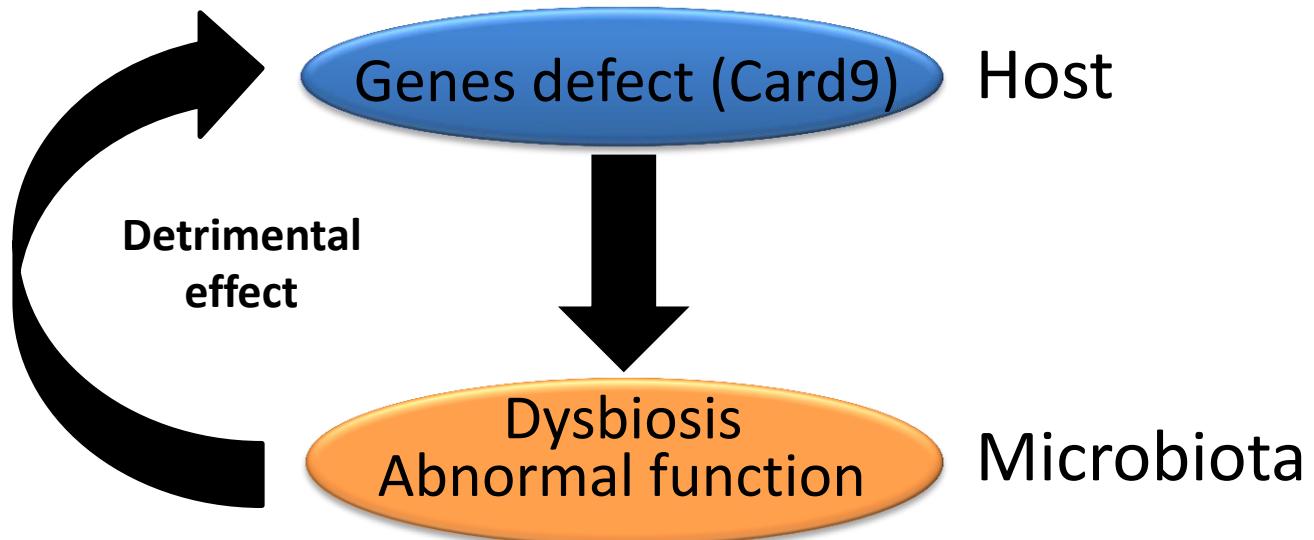


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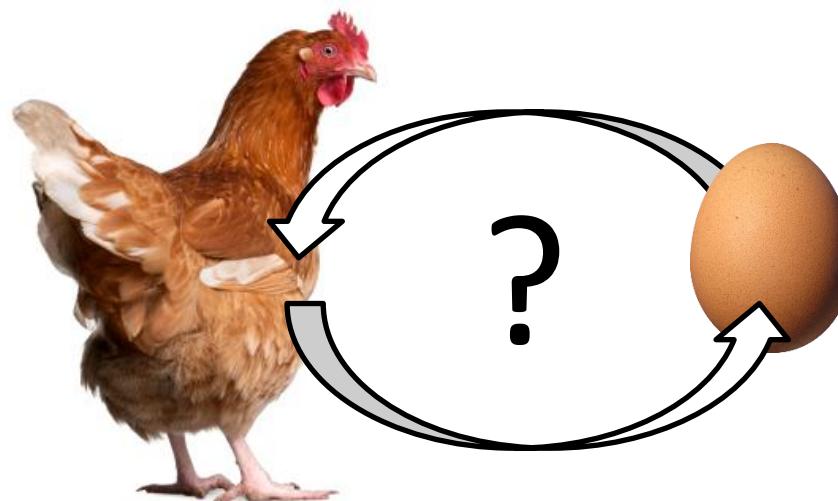
# Conclusion

- Gut microbiota has a central role in IBD pathogenesis
- It is under the influence of both environmental and genetic factors



# Conclusion

- Gut microbiota in IBD pathogenesis:  
chicken or egg?



**Both!**

P Seksik  
D Rainteau  
JP Grill  
L Brot



**B Lamas**  
V Leducq  
S Jegou  
C McQuitty



Saint Antoine Hospital  
**L Beaugerie** I Nion-Larmurier  
J Cosnes A Bourrier  
P Seksik C Landman

## Sokol's lab



### Collaborators

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JM Launay Unit 970 Inserm  
RJ Xavier Broad /Harvard / MGH



**P Langella**  
L Bermudes  
JM Chatel  
M Thomas  
F Chain

M Lavie-Richard  
ML Michel  
C Bridonneau  
G Da Costa  
J Natividad  
B Sovran  
J Planchais  
A Demoget  
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