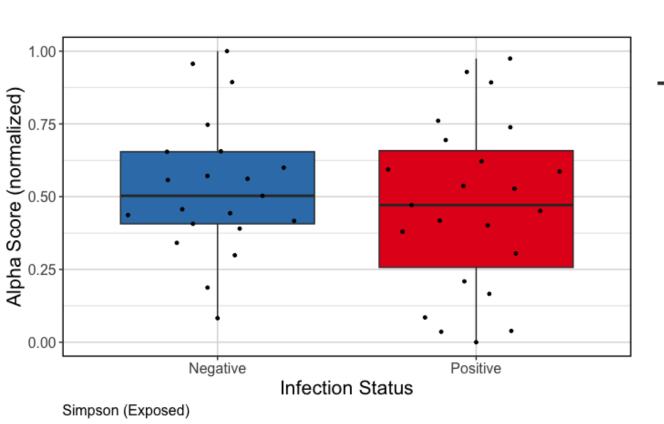
### Microbiome Analysis: Infection

Michael Sieler, December 19, 2022

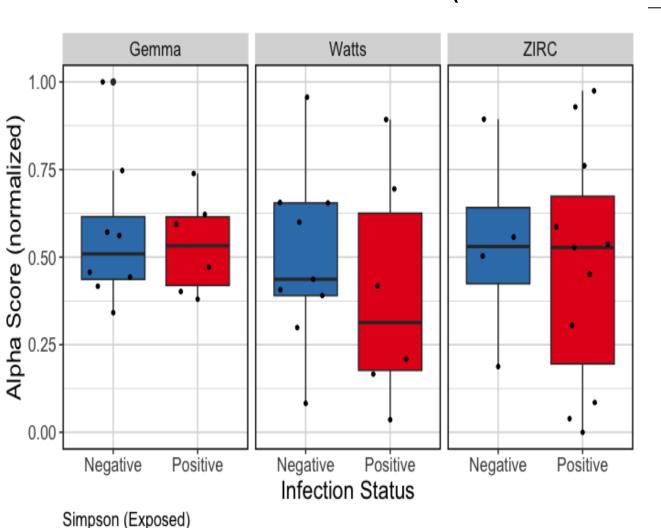
#### Alpha diversity

# We do not see a statistical difference in alpha diversity and presence of infection (males and females)



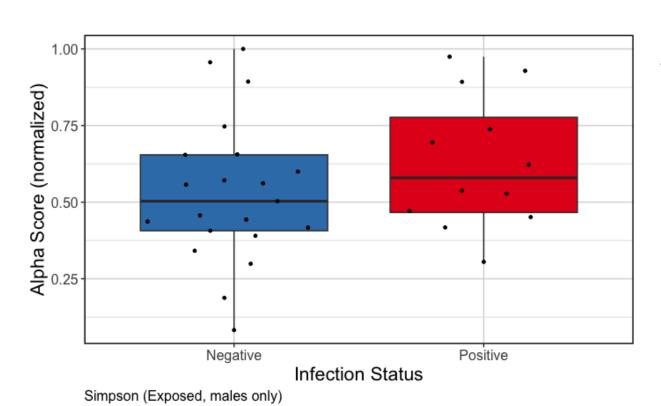
ANOVA( glm(Alpha.Score ~ Infection), family = quasibinomial))							
metric	term	statistic	df	p.value sig			
Observed	Infection.Status	0.042	1	0.838			
Shannon	Infection.Status	0.167	1	0.683			
Simpson	Infection.Status	0.582	1	0.446			

### We do not see a statistical difference in alpha diversity and presence of infection depending on diet (males and females)



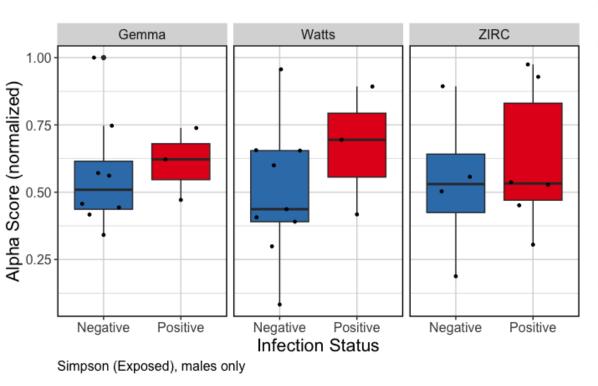
ANOVA( glm(Alpha.Score ~ Diet*Infection), family = quasibinomial))							
metric	term	statistic	df	p.value	sig		
Observed	Infection.Status	0.733	1	0.392			
	Diet	29.258	2	<0.001	*		
	Infection.Status:Diet	0.628	2	0.730			
Shannon	Infection.Status	0.593	1	0.441			
	Diet	14.156	2	<0.001	•		
	Infection.Status:Diet	0.044	2	0.978			
Simpson	Infection.Status	0.541	1	0.462			
	Diet	0.861	2	0.650			
	Infection.Status:Diet	0.091	2	0.956			

# We do not see a statistical difference in alpha diversity and presence of infection (males only)



ANOVA( glm(Alpha.Score ~ Infection), family = quasibinomial) ), males only							
metric	term	statistic	df	p.value si			
Observed	Infection.Status	0.588	1	0.443			
Shannon	Infection.Status	1.116	1	0.291			
Simpson	Infection.Status	1.421	1	0.233			

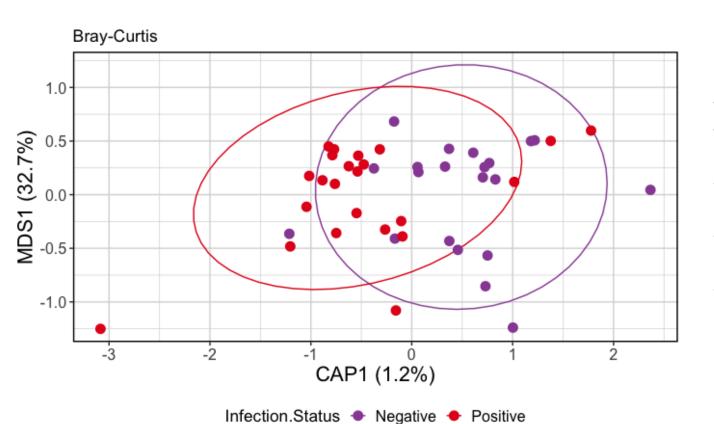
### We do not see a statistical difference in alpha diversity and presence of infection depending on diet (males only)



metric	term	statistic	df	p.value	sig
Observed	Infection.Status	0.118	1	0.732	
	Diet	23.626	2	< 0.001	*
	Infection.Status:Diet	1.196	2	0.550	
	Infection.Status	0.645	1	0.422	
Shannon	Diet	10.491	2	0.005	*
	Infection.Status:Diet	0.906	2	0.636	
	Infection.Status	1.154	1	0.283	
Simpson	Diet	0.131	2	0.936	
	Infection.Status:Diet	0.314	2	0.855	

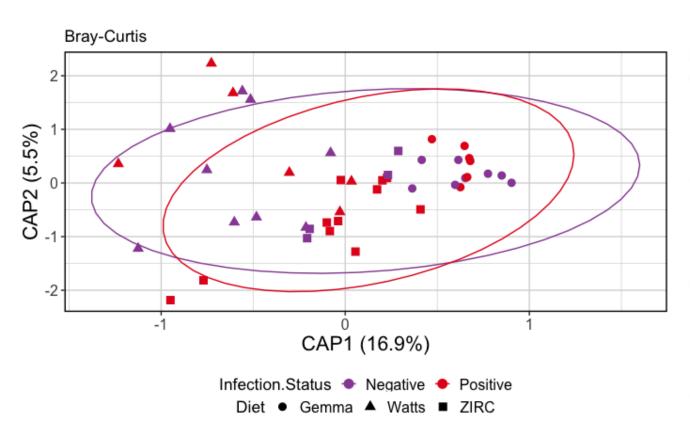
#### Beta diversity

## We do not see a statistical difference in beta diversity and presence of infection (males and females)



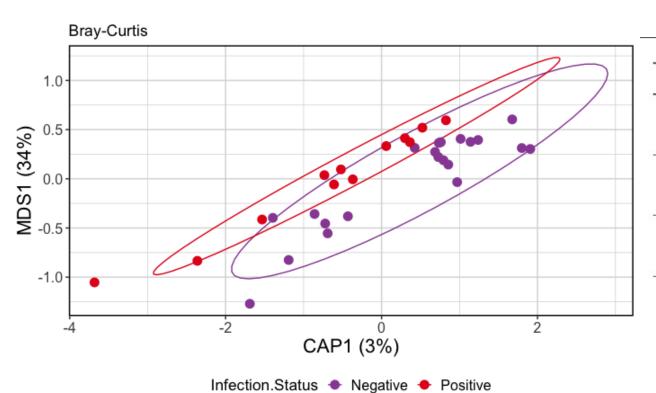
Distance	-based redundancy an	alysis (dbl	RDA) ordination	. Beta.Score	~ Infection	
metric	term	Df	SumOfSqs	statistic	p.value si	g
Bray-Curtis	Infection.Status	1.00	0.02915815	0.492	0.878	
	Residual	42.00	2.48963116			
Canberra	Infection.Status	1.00	0.24985811	1.059	0.338	
	Residual	42.00	9.91405427			
Sørensen	Infection.Status	1.00	0.09436585	1.081	0.345	
	Residual	42.00	3.66472024			

## We do not see a statistical difference in beta diversity and presence of infection depending on diet (males and females)



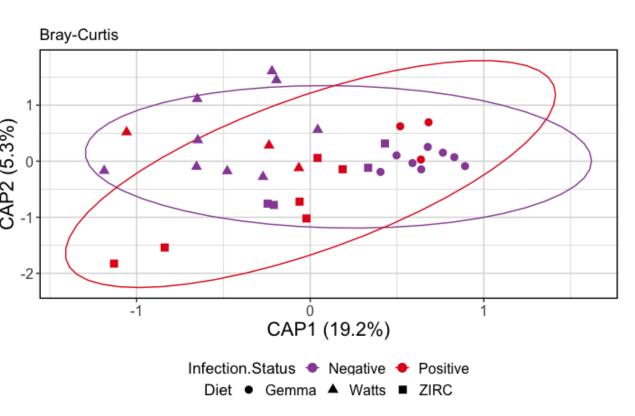
metric	term	Df	SumOfSqs	statistic	p.value	sig
	Diet	2.00	0.54487892	5.578	0.001	*
Denote Occation	Infection.Status:Diet	2.00	0.08875981	0.909	0.529	
Bray-Curtis	Infection.Status	1.00	0.02915815	0.597	0.760	
	Residual	38.00	1.85599243			
	Diet	2.00	1.75371316	4.296	0.001	*
	Infection.Status	1.00	0.24985811	1.224	0.173	
Canberra	Infection.Status:Diet	2.00	0.40351911	0.988	0.481	
	Residual	38.00	7.75682200			
Sørensen	Diet	2.00	0.84140066	5.992	0.001	*
	Infection.Status	1.00	0.09436585	1.344	0.166	
	Infection.Status:Diet	2.00	0.15530811	1.106	0.307	
	Residual	38.00	2.66801146			

## We do not see a statistical difference in alpha diversity and presence of infection (males only)



term	Df	SumOfSqs	statistic	p.value	sig
Infection.Status	1.00	0.05996175	0.972	0.405	
Residual	31.00	1.91188446			
Infection.Status	1.00	0.26875943	1.098	0.271	
Residual	31.00	7.58963052			
Infection.Status	1.00	0.12295920	1.349	0.161	
Residual	31.00	2.82654440			
	Infection.Status Residual Infection.Status Residual Infection.Status	Infection.Status 1.00 Residual 31.00 Infection.Status 1.00 Residual 31.00 Infection.Status 1.00	Infection.Status         1.00         0.05996175           Residual         31.00         1.91188446           Infection.Status         1.00         0.26875943           Residual         31.00         7.58963052           Infection.Status         1.00         0.12295920	Infection.Status         1.00         0.05996175         0.972           Residual         31.00         1.91188446           Infection.Status         1.00         0.26875943         1.098           Residual         31.00         7.58963052           Infection.Status         1.00         0.12295920         1.349	Infection.Status         1.00         0.05996175         0.972         0.405           Residual         31.00         1.91188446           Infection.Status         1.00         0.26875943         1.098         0.271           Residual         31.00         7.58963052           Infection.Status         1.00         0.12295920         1.349         0.161

## We do not see a statistical difference in alpha diversity and presence of infection depending on diet (males only)



Distance	-based redundancy analysis	(dbRDA	) ordination. Bet	a.Score ~ In	fection*Di	et
metric	term	Df	SumOfSqs	statistic	p.value	sig
	Diet	2.00	0.44583713	4.307	0.001	*
Brow Custin	Infection.Status	1.00	0.05996175	1.158	0.266	
Bray-Curtis	Infection.Status:Diet	2.00	0.06845416	0.661	0.800	
	Residual	27.00	1.39759316			
Canberra	Diet	2.00	1.42589844	3.316	0.001	*
	Infection.Status	1.00	0.26875943	1.250	0.139	
	Infection.Status:Diet	2.00	0.35943130	0.836	0.813	
	Residual	27.00	5.80430079			
	Diet	2.00	0.68456706	4.595	0.001	*
Sørensen	Infection.Status	1.00	0.12295920	1.651	0.061	
	Infection.Status:Diet	2.00	0.13056042	0.876	0.639	
	Residual	27.00	2.01141692			