		[Kullillococcus] gilavus [01594]
/er	•	
obLiv	•	] mnsom
Micro is	•	[Clostridium] clostridioforme/bolteae [03442] -
XY/N Cardi	•	Clostridiales sp. [13012] -
	•	[Clostridium] glycyrrhizinilyticum [03672] -
	•	Firmicutes sp.
•	•	Flavonifractor plautii [02971] -
	•	Veillonella atypica [01941]-
	•	Flavonifractor plautii [05238] -
	:	Streptococcus sp. [00283]-
	•	Clostridiales sp. [12768] -
	•	Clostridiales sp. [13140] -
	•	Proteobacteria sp. [00095] -
	•	Veillonella parvula [01938] -
	•	Clostridiales sp. [12473] -
	8	Eggerthella lenta [00719] -
		Veillonella dispar [01940] -
	•	Rothia mucilaginosa [05265] -
	•	Clostridium sp. CAG:299 [07619]-
•		Roseburia sp. 40_7 [12352]-
•		Clostridiales sp. [13168]-
•		Ruminococcaceae sp. [12239]-
*		Clostridiales sp. [13863]-
•		Clostridiales sp. [12631]-
•		Odoribacter splanchnicus [03697]-
•		Faecalibacterium prausnitzii [06109] -
•		Clostridiales sp. [12284] -
•		Clostridiales sp. [12563] -
8		Clostridiales sp. [12463]-
•		Clostridiales sp. [12334] -
•		Ruminococcus bicirculans [02792] -
•		Clostridium sp. [12417] -
•		Clostridiales sp. [12252] -
•		Faecalibacterium prausnitzii [06108] -
•		Oscillibacter sp. 57_20 [12610] -
•		[Eubacterium] eligens [02376] -
*		Clostridiales sp. [12635]-
•		Oscillibacter sp. [04664]-
•		Clostridiales sp. [12356]
•		Simpson diversity -
•		Species richness -
•		Shannon diversity -
0.4 -	0.0 <b>-</b> -0.2 <b>-</b>	
correlation	Pearson	
oortolorioo	00000	