

Gut microbiome: family\_Veillonellaceae

1e-04  
5e-05  
0e+00  
-5e-05

Correlation: 1

FDR adjusted P value: 0.017

2016-01-12

2016-01-13

2016-01-15

2016-01-17

2016-02-24

Exposome (chemical): Styrene oxide

