

Gut microbiome: family_Streptococcaceae

Correlation: -1

FDR adjusted P value: 0.017

0.00025

2016-01-15

2016-01-17

0.00000

2016-01-13

2016-01-12

-0.00025

2016-02-24

-0.00050

9

10

11

12

13

Exposome (chemical): Bergapten

