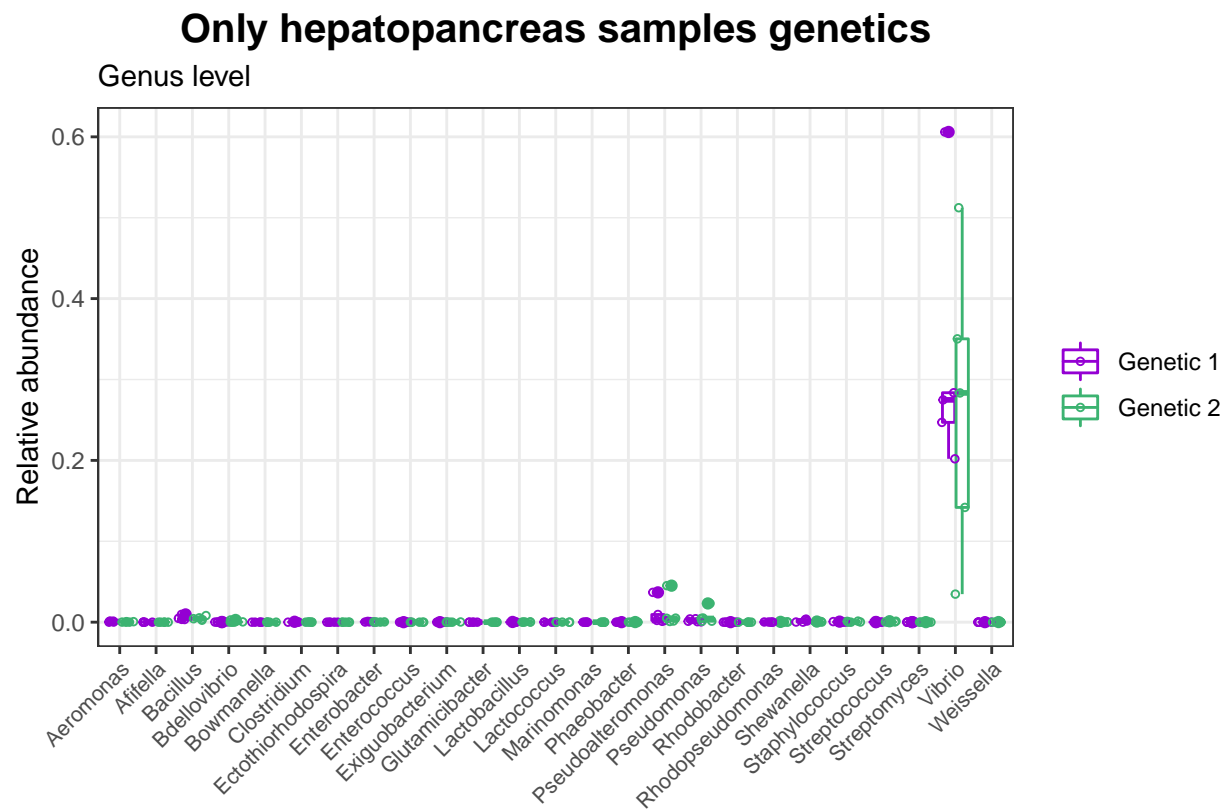


Probiotics boxplots

15.08.2021

Genus level

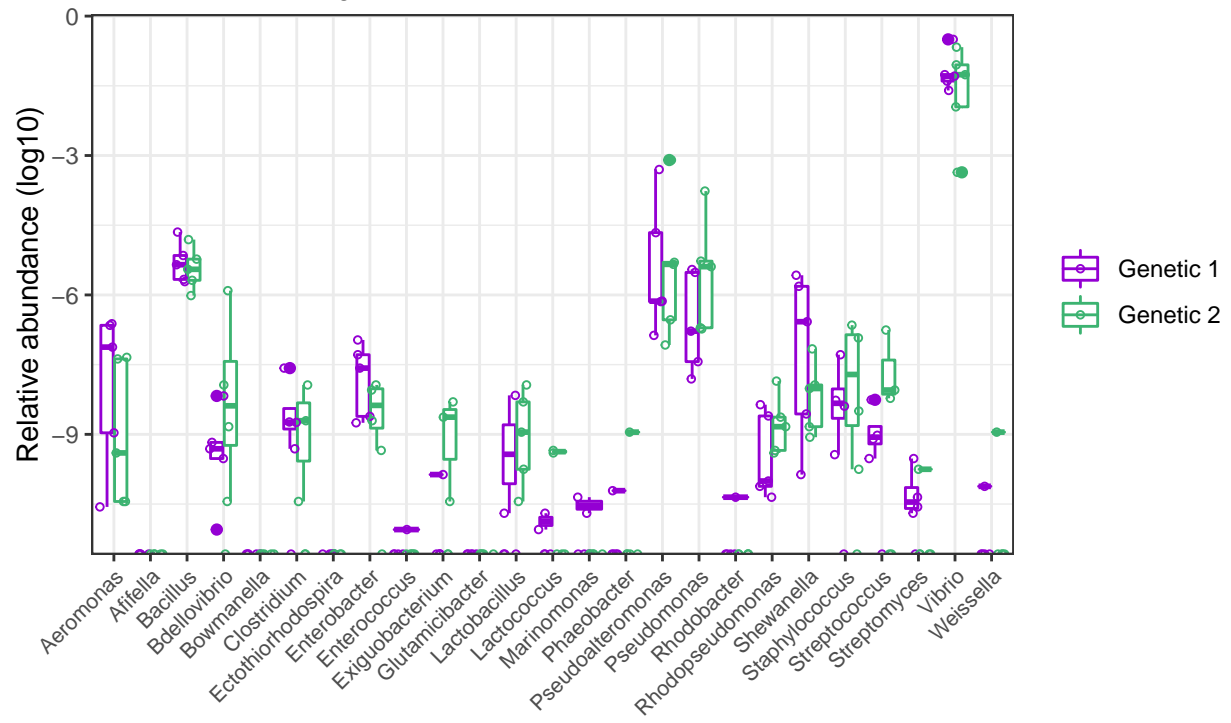
Only Hepatopancreas samples



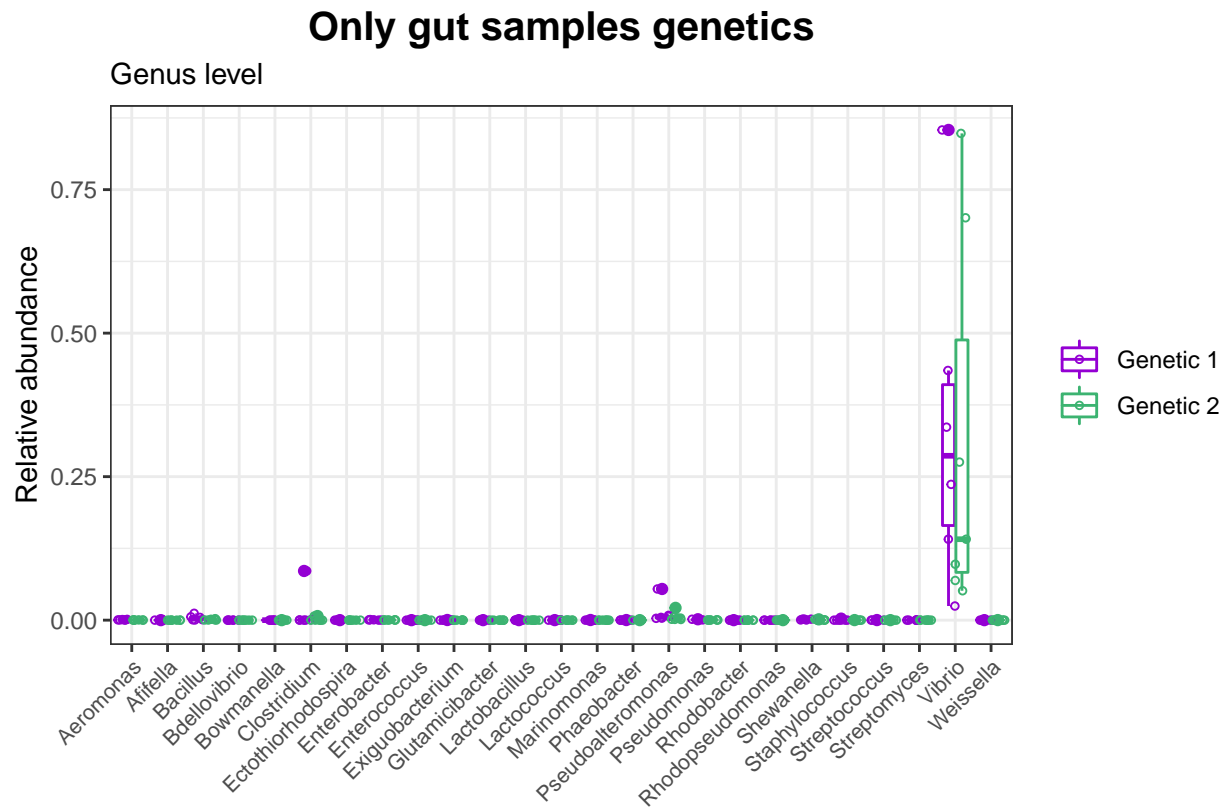
Warning: Removed 112 rows containing non-finite values (stat_boxplot).

Only hepatopancreas samples genetics

Genus level with log10 transformation



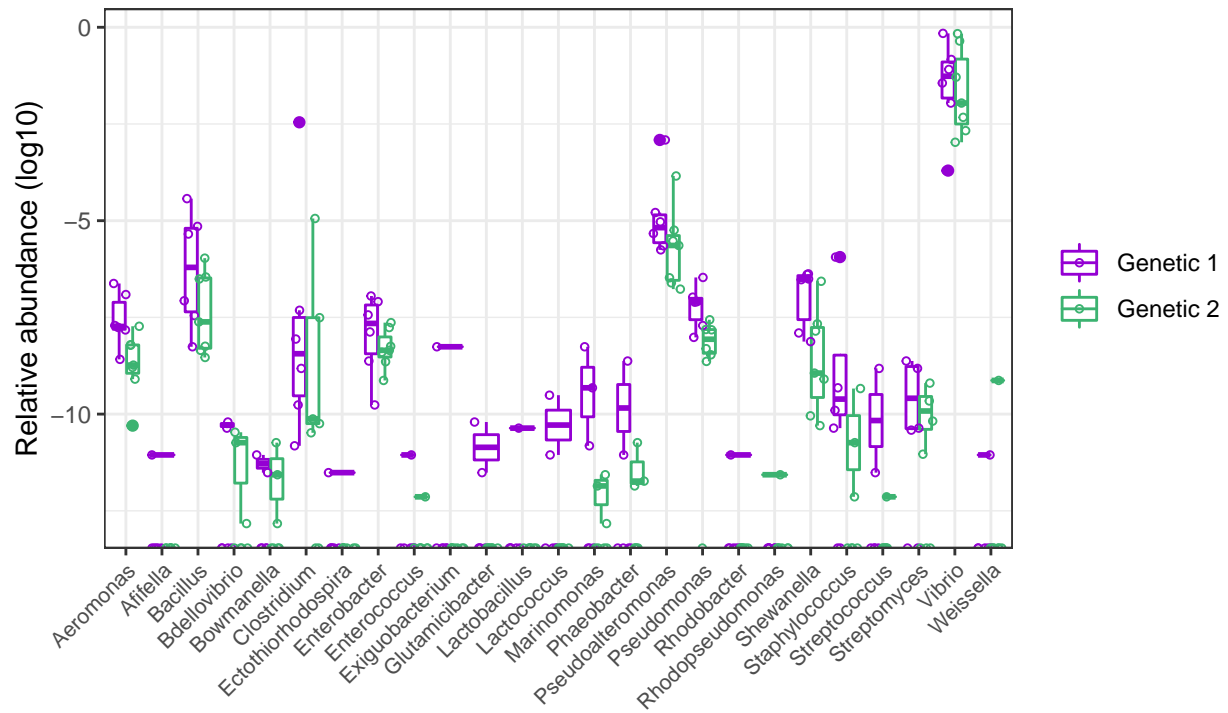
Only Gut samples Genetics



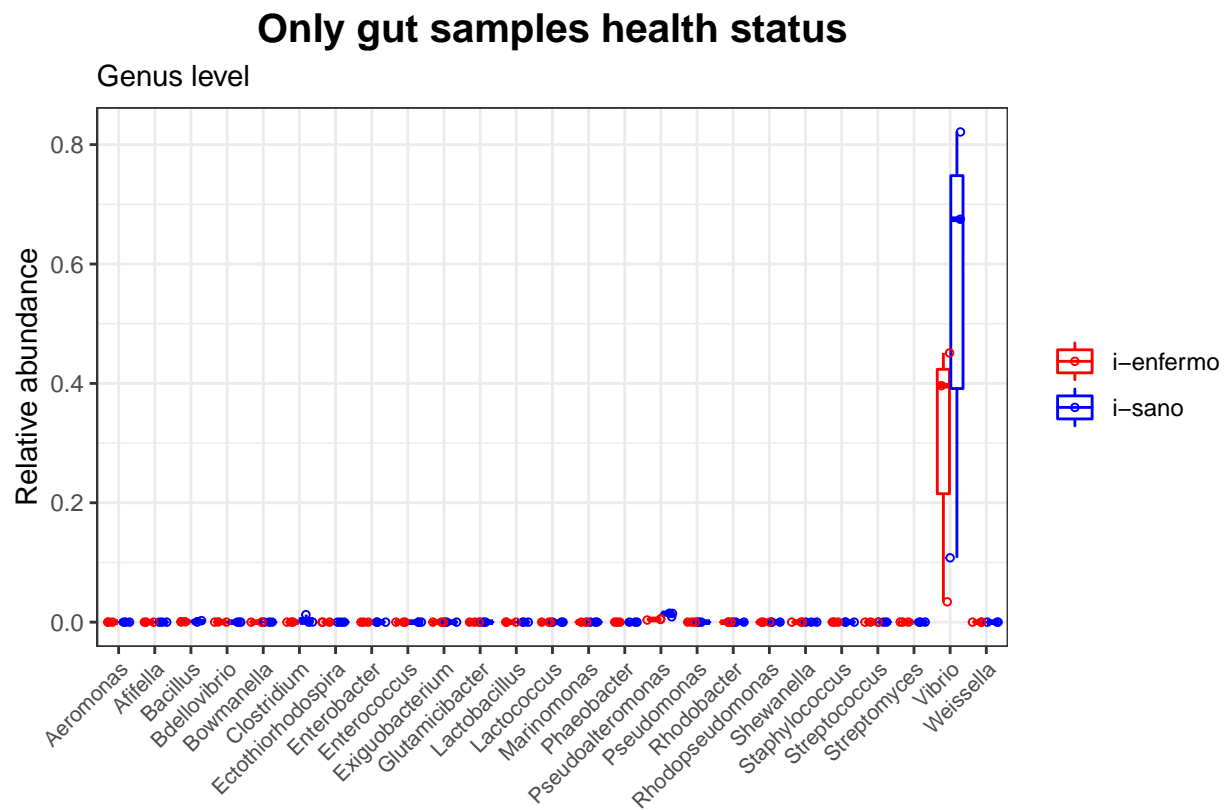
Warning: Removed 171 rows containing non-finite values (stat_boxplot).

Only gut samples genetics

Genus level with log10 transformation



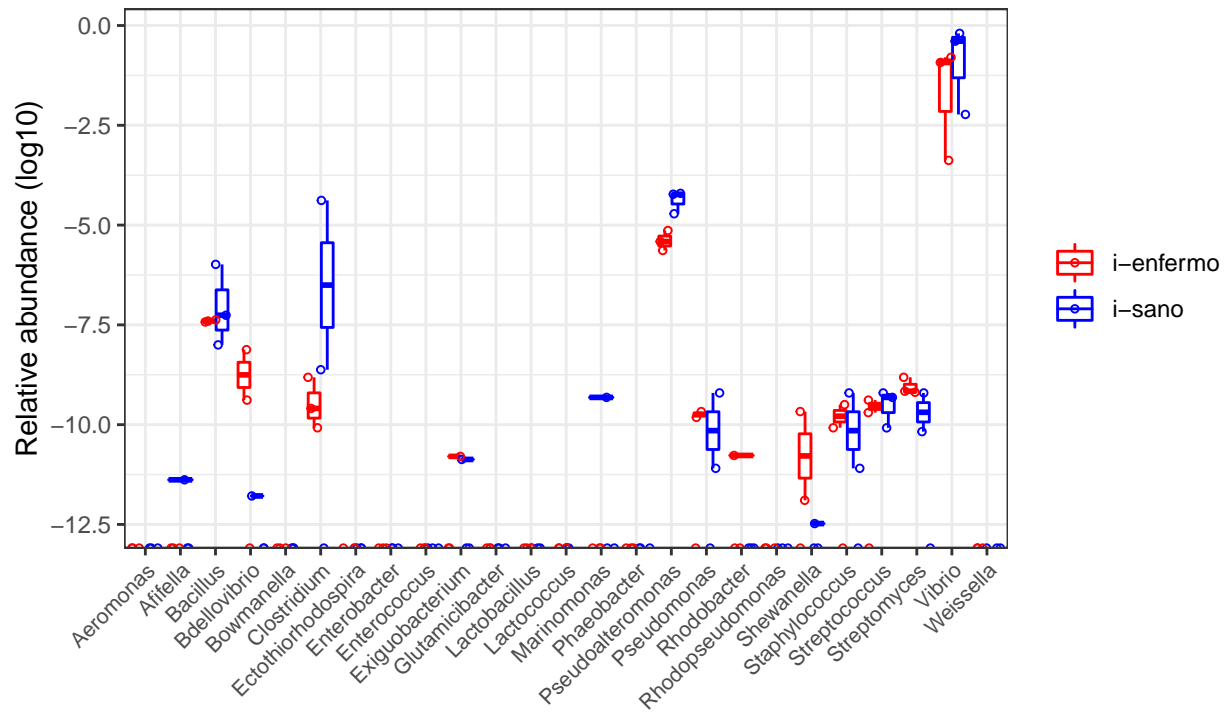
Only Gut samples Health status



Warning: Removed 98 rows containing non-finite values (stat_boxplot).

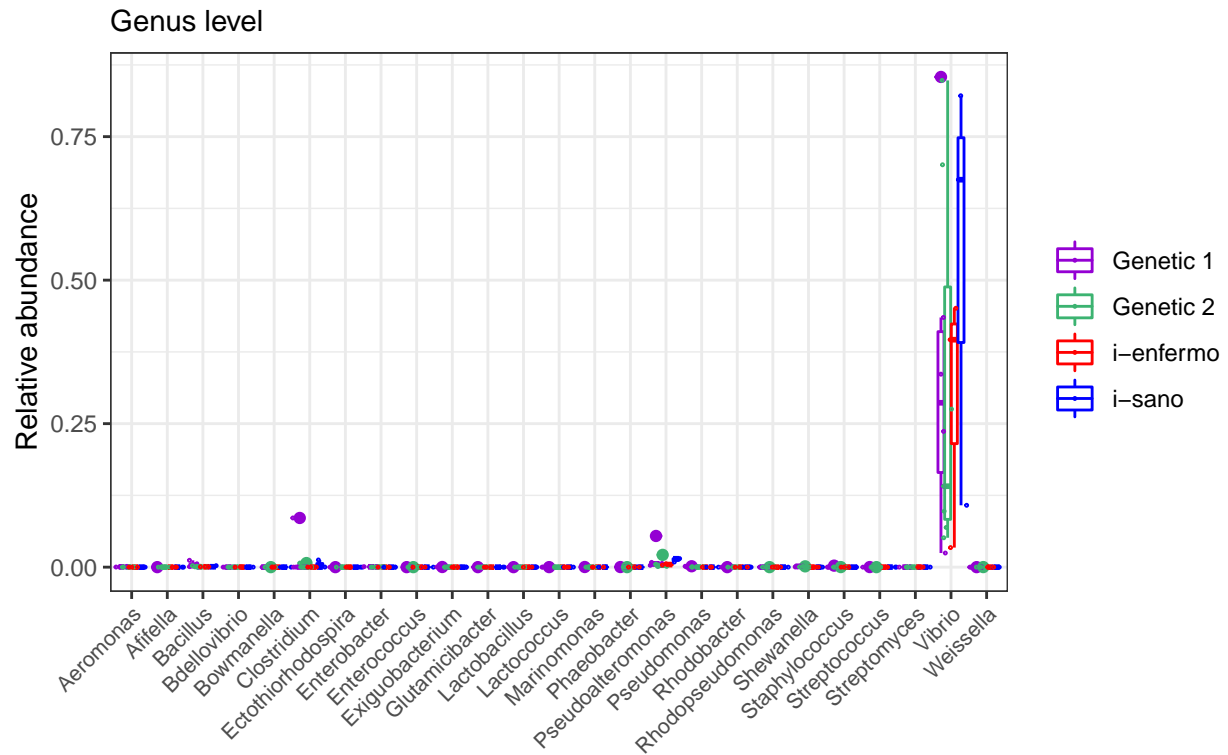
Only gut samples health status

Genus level with log10 transformation



All gut samples

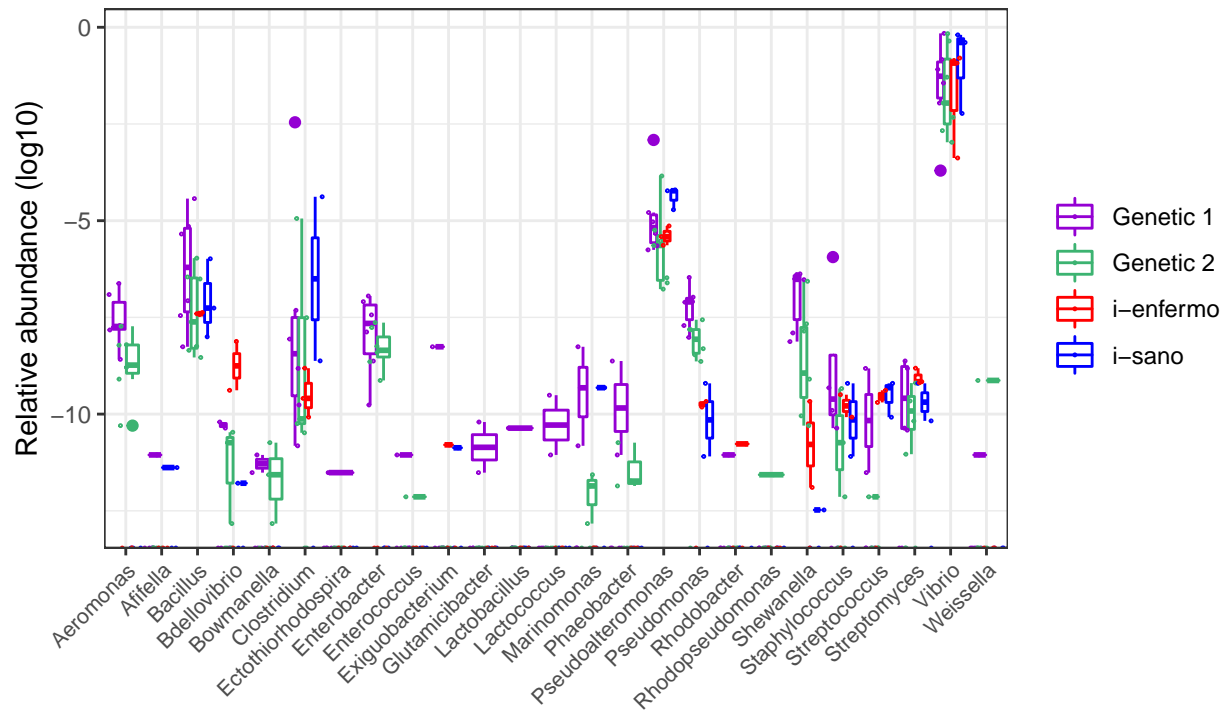
All gut samples



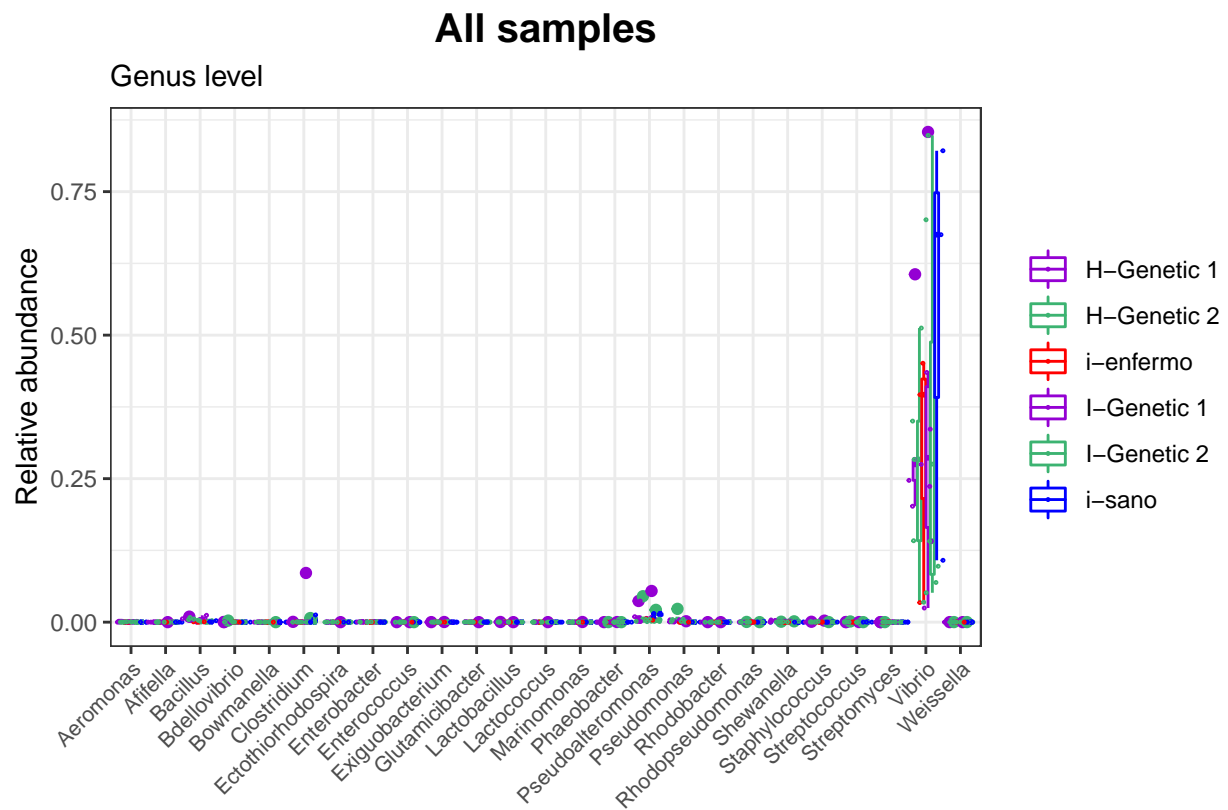
```
## Warning: Removed 269 rows containing non-finite values (stat_boxplot).
```

All gut samples

Genus level with log10 transformation



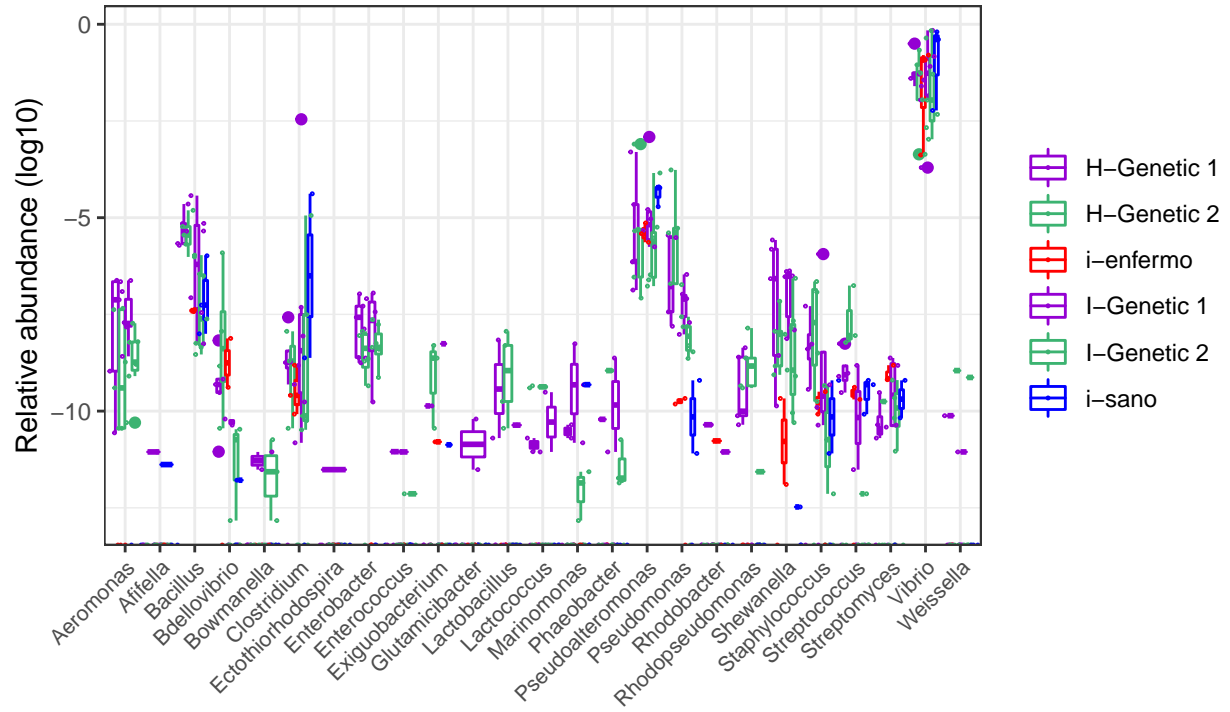
All samples



```
## Warning: Removed 381 rows containing non-finite values (stat_boxplot).
```

All samples

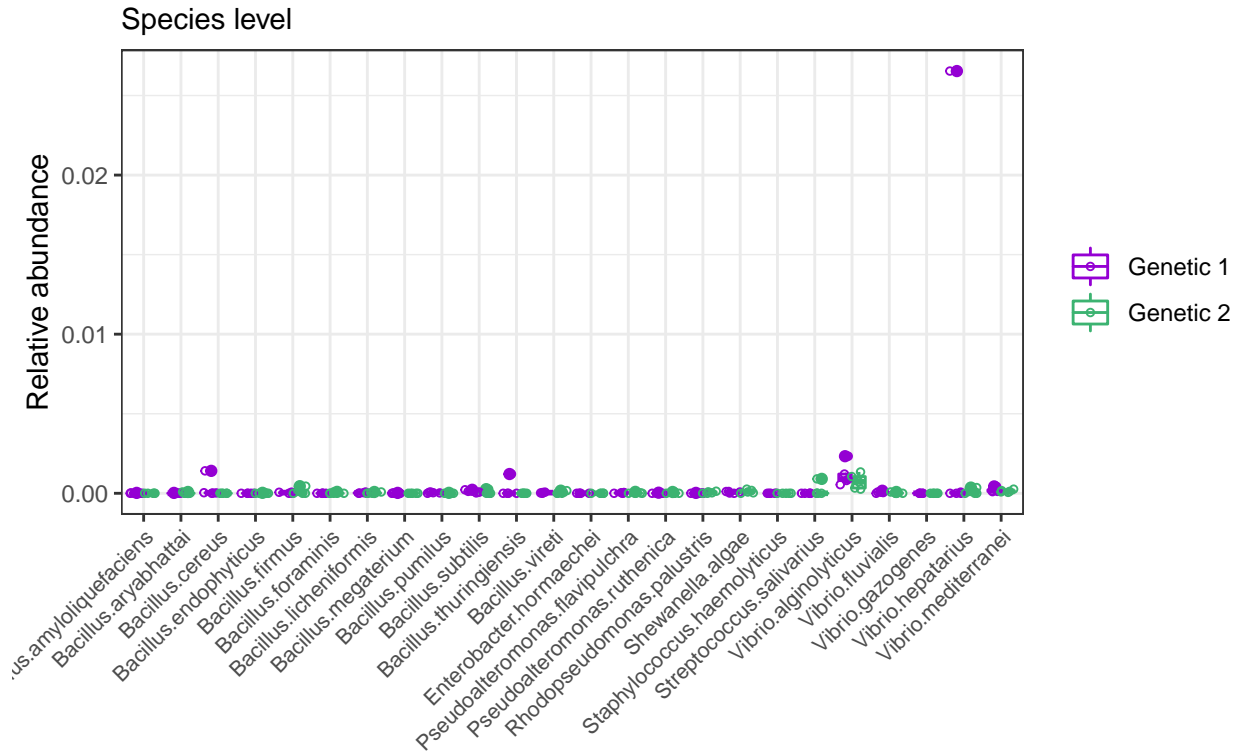
Genus level with log10 transformation



Species level

Only Hepatopancreas samples

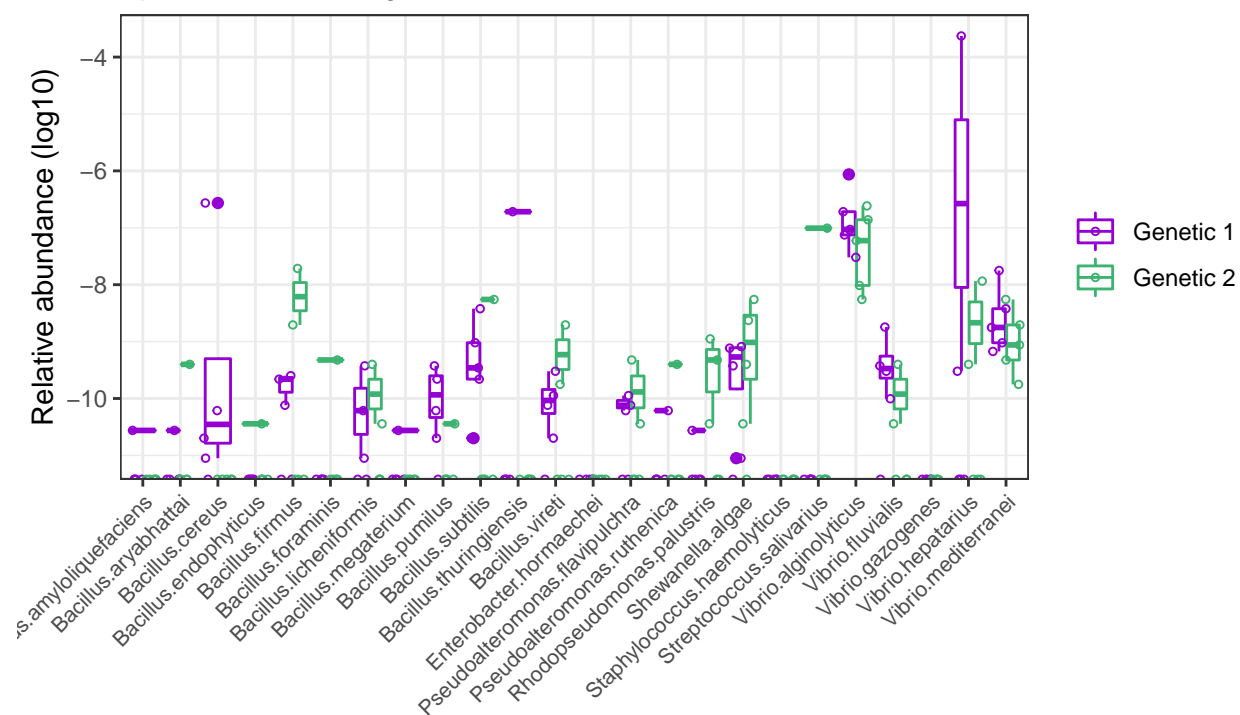
Only hepatopancreas samples genetics



Warning: Removed 152 rows containing non-finite values (stat_boxplot).

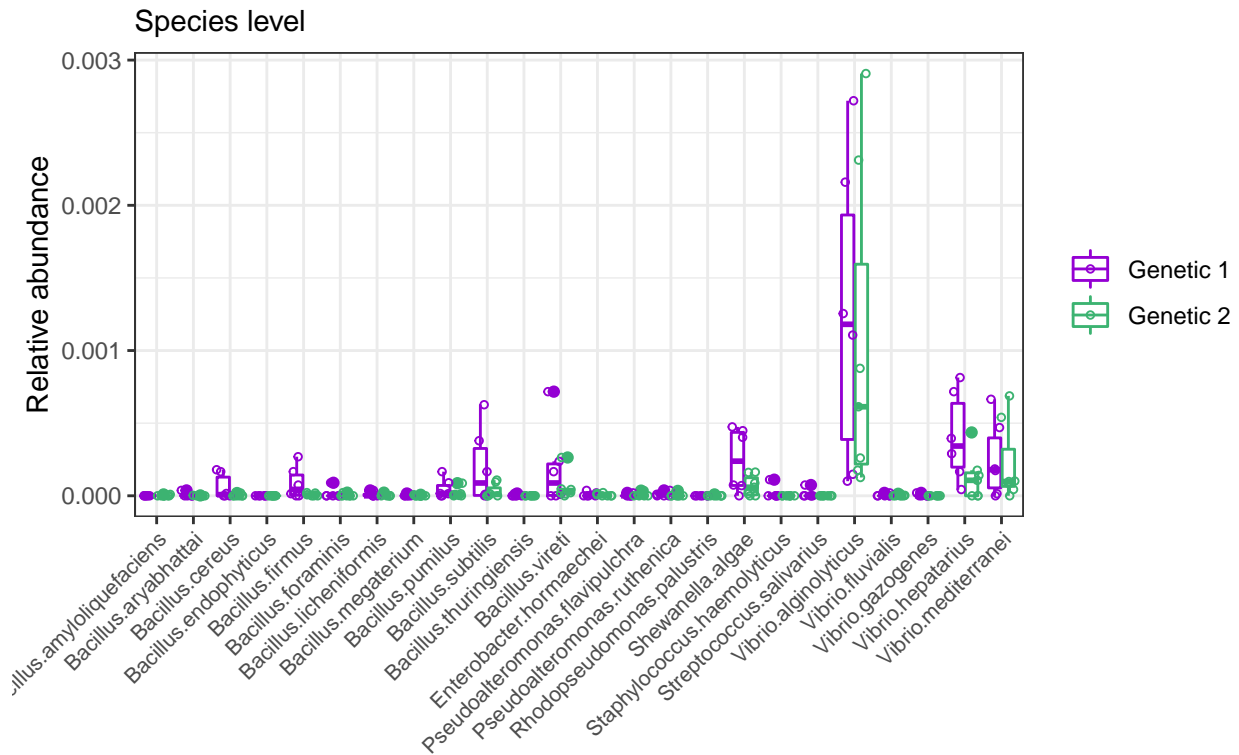
Only hepatopancreas samples genetics

Species level with log10 transformation



Only Gut samples Genetics

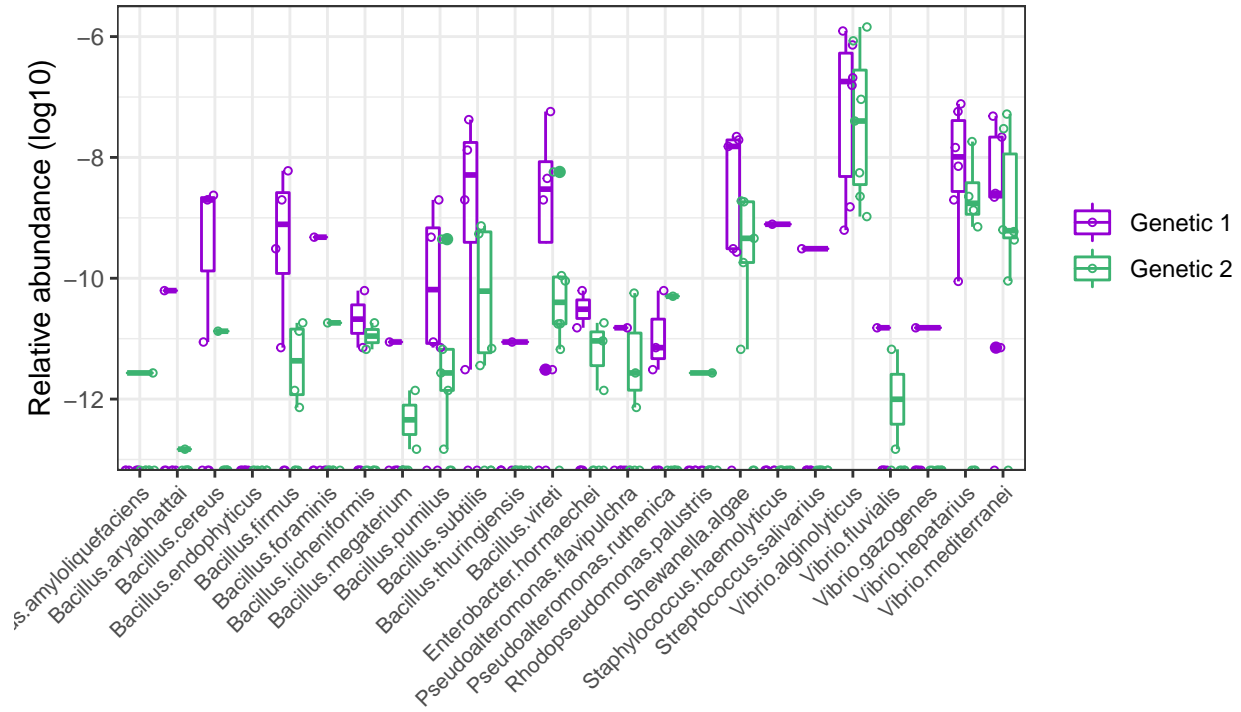
Only gut samples genetics



```
## Warning: Removed 196 rows containing non-finite values (stat_boxplot).
```

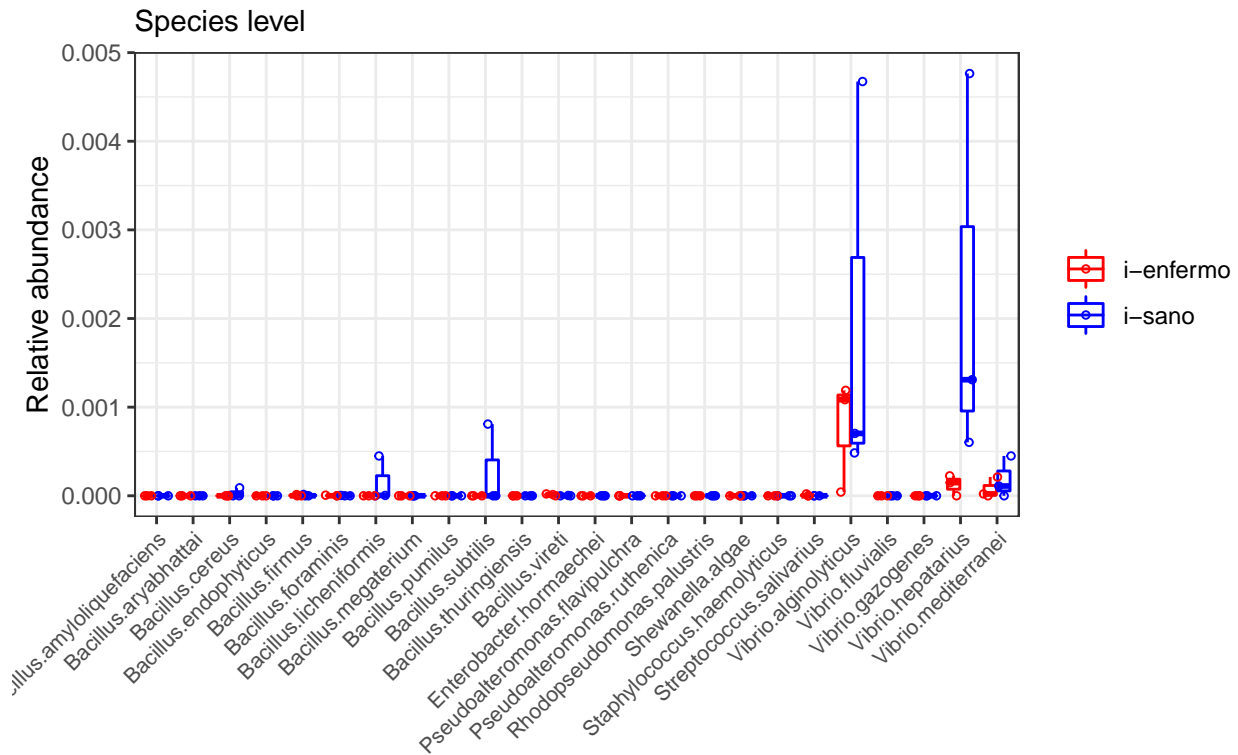
Only gut samples genetics

Species level with log10 transformation



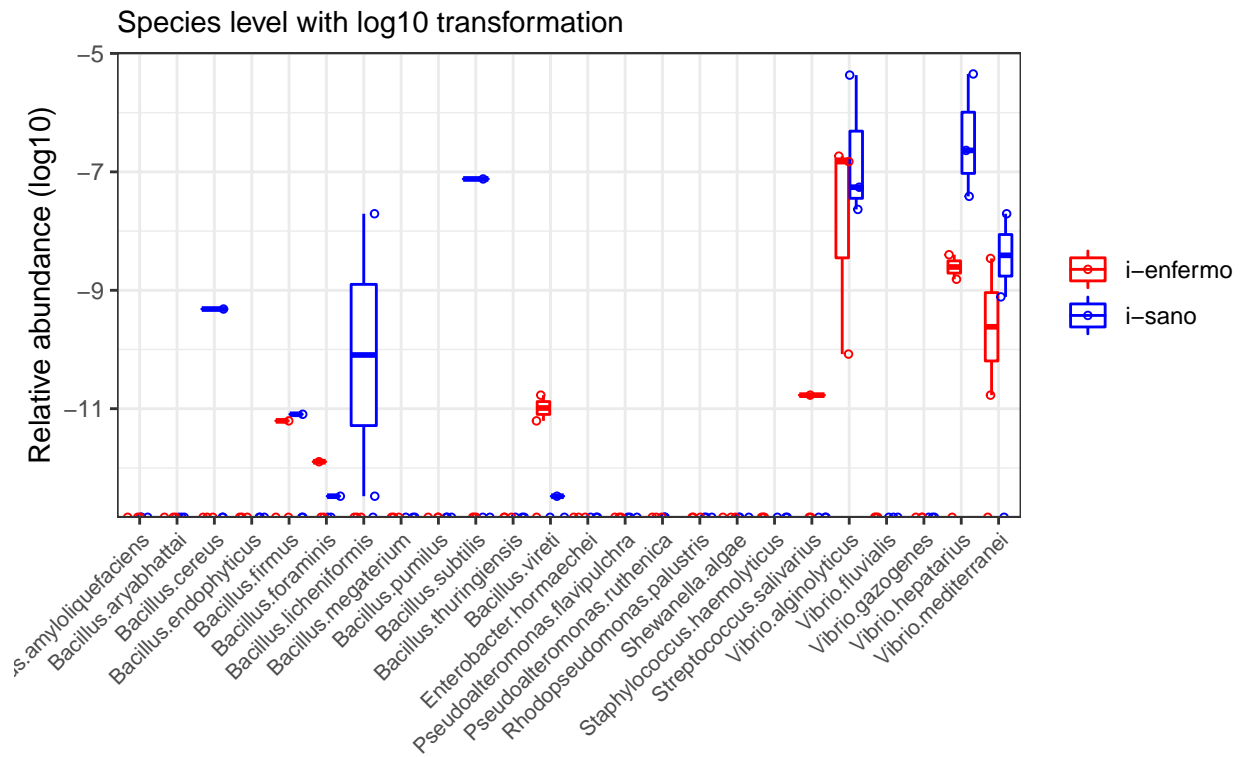
Only Gut samples Health status

Only gut samples health status

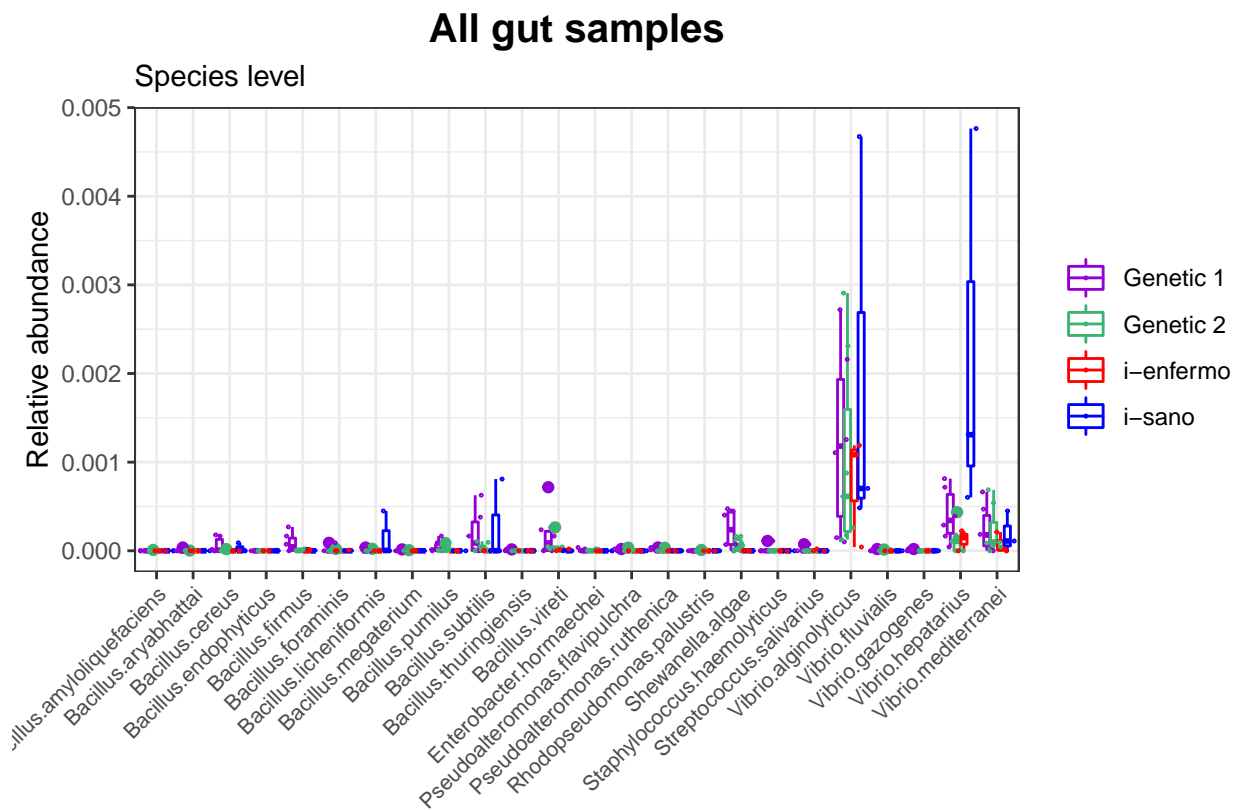


```
## Warning: Removed 117 rows containing non-finite values (stat_boxplot).
```

Only gut samples health status

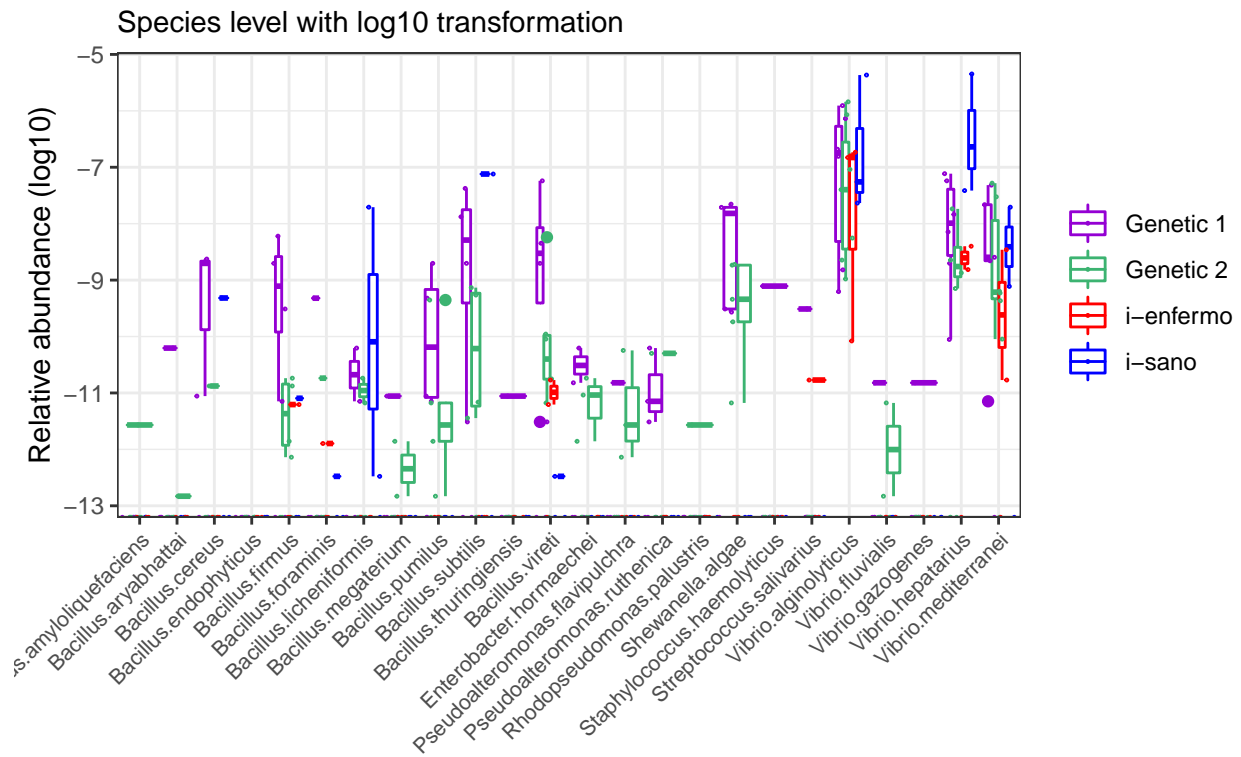


All gut samples

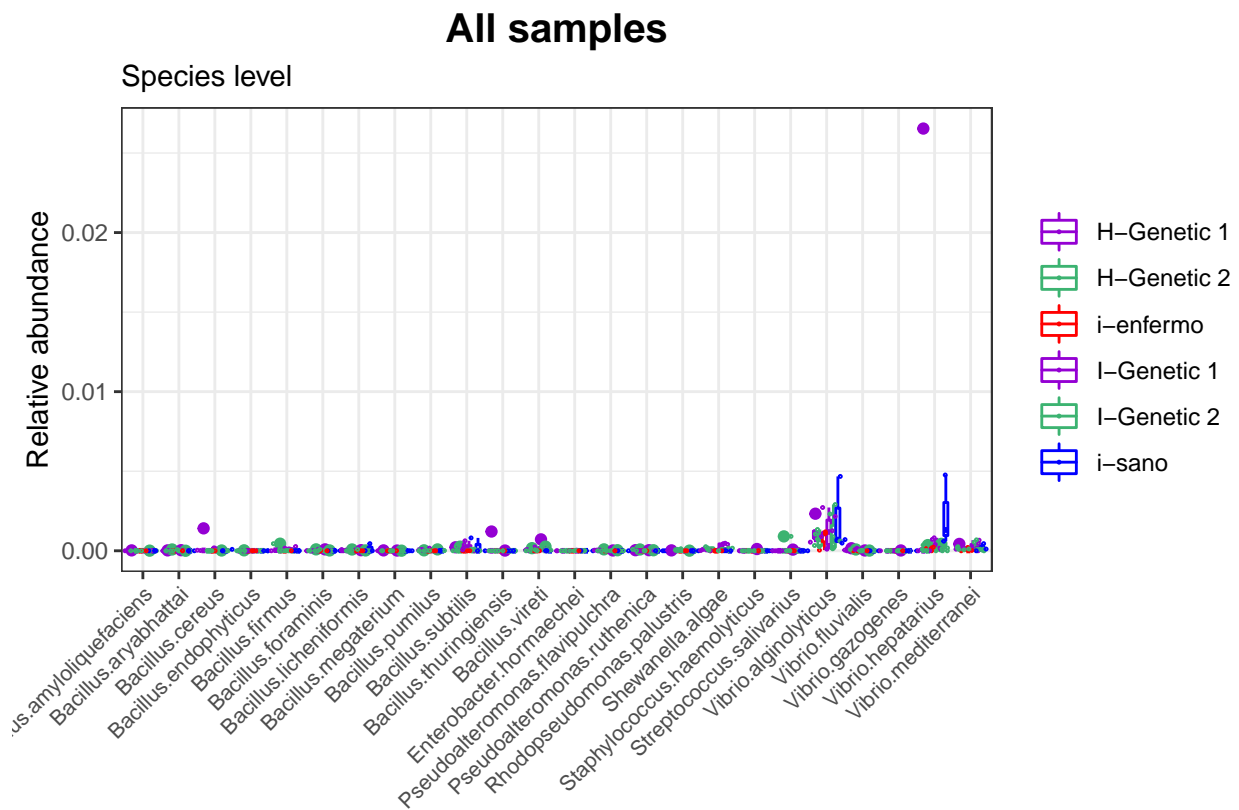


Warning: Removed 313 rows containing non-finite values (stat_boxplot).

All gut samples



All samples



Warning: Removed 465 rows containing non-finite values (stat_boxplot).

All samples

Species level with log10 transformation

