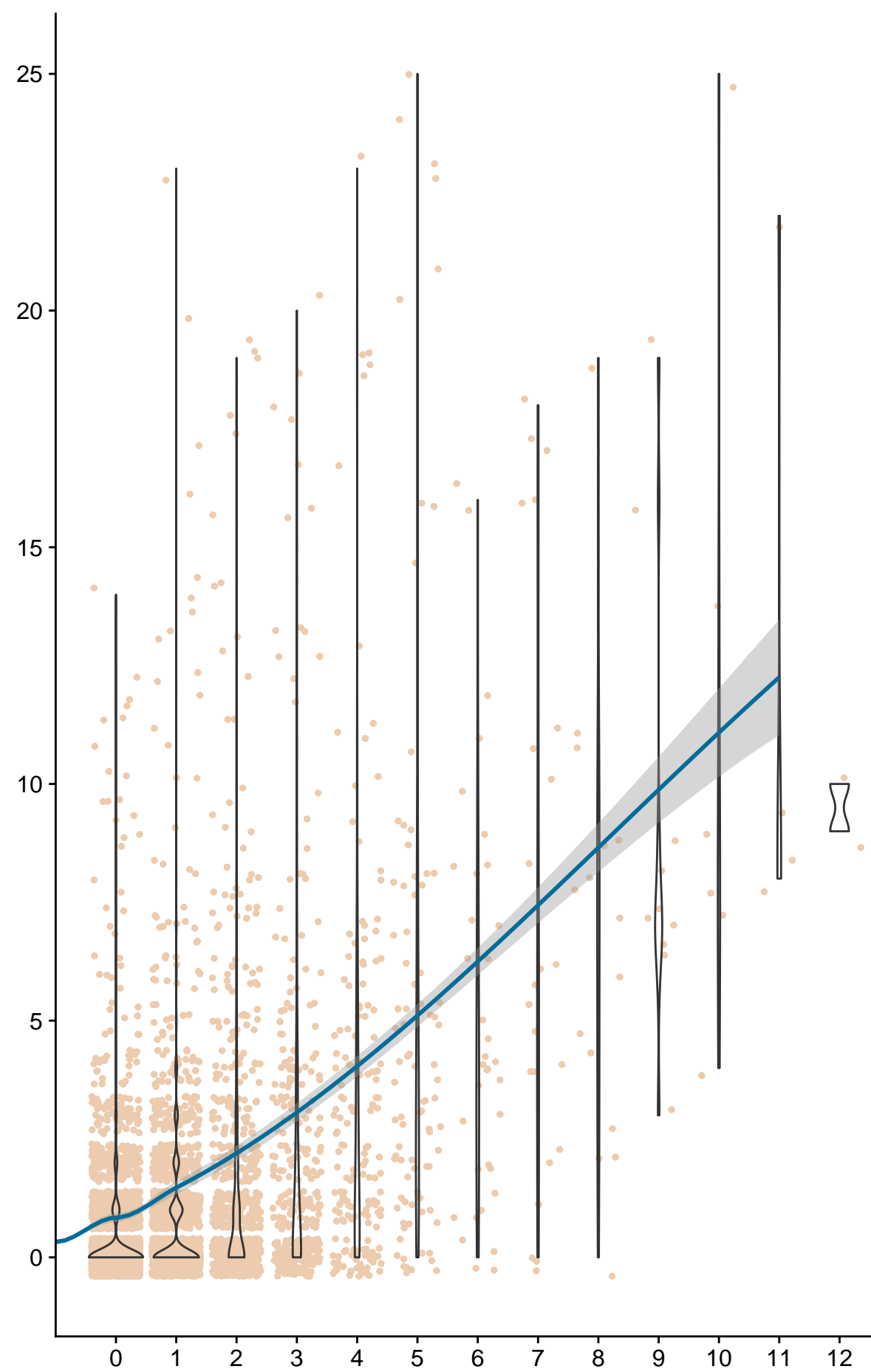
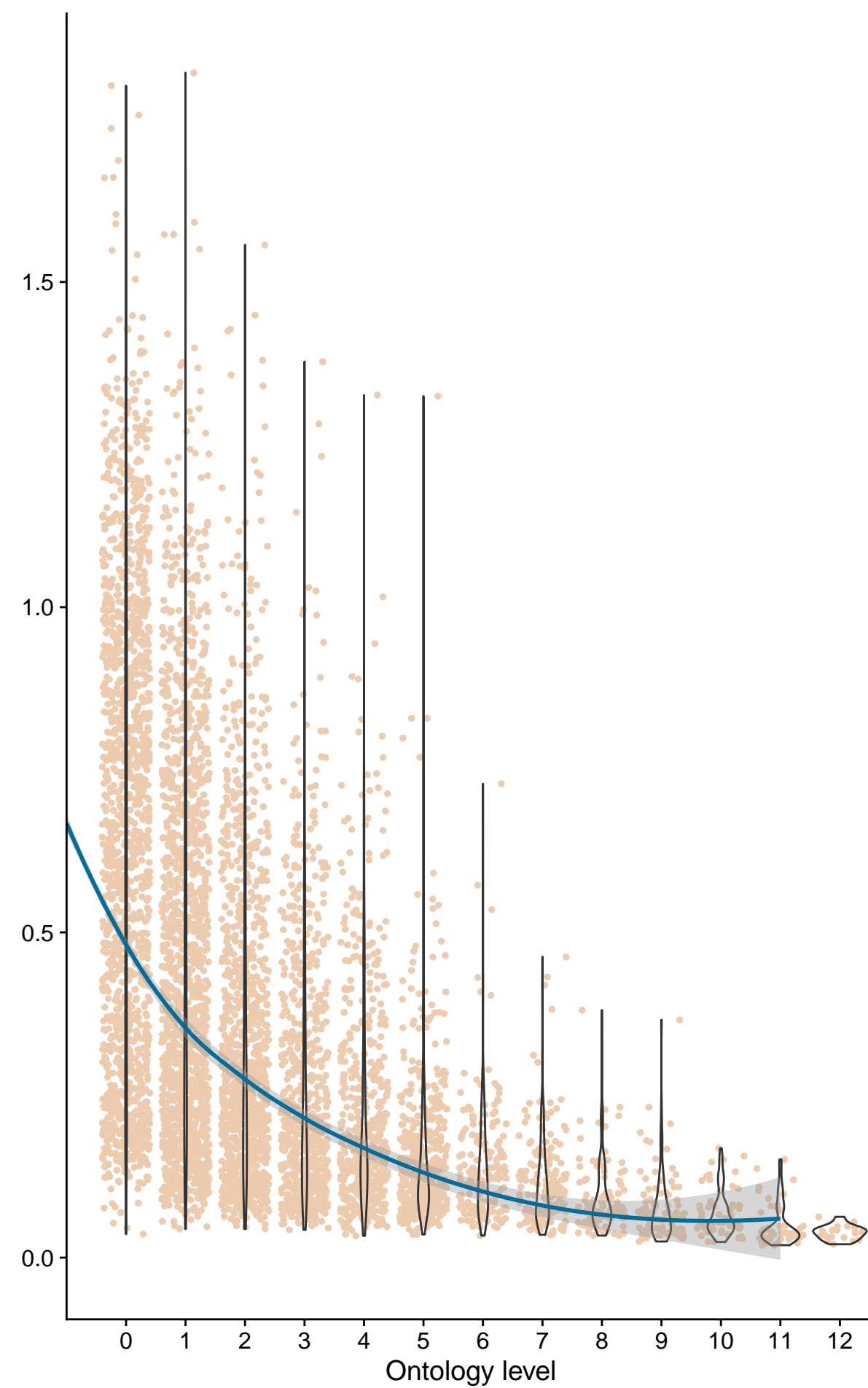


**A**

Number of associated cell types

**B**

log10(fold enrichment)

**C**

Number of genes

