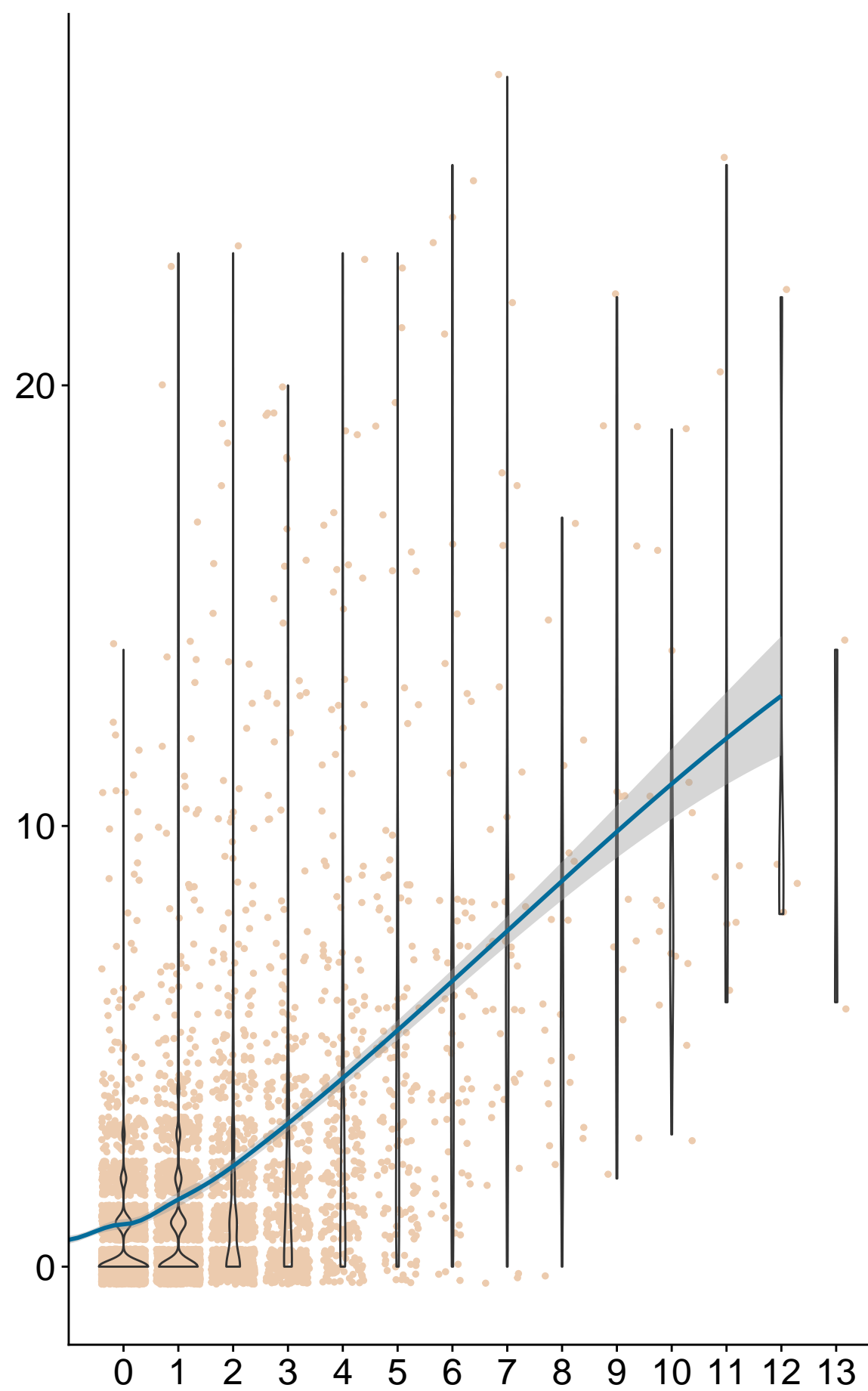
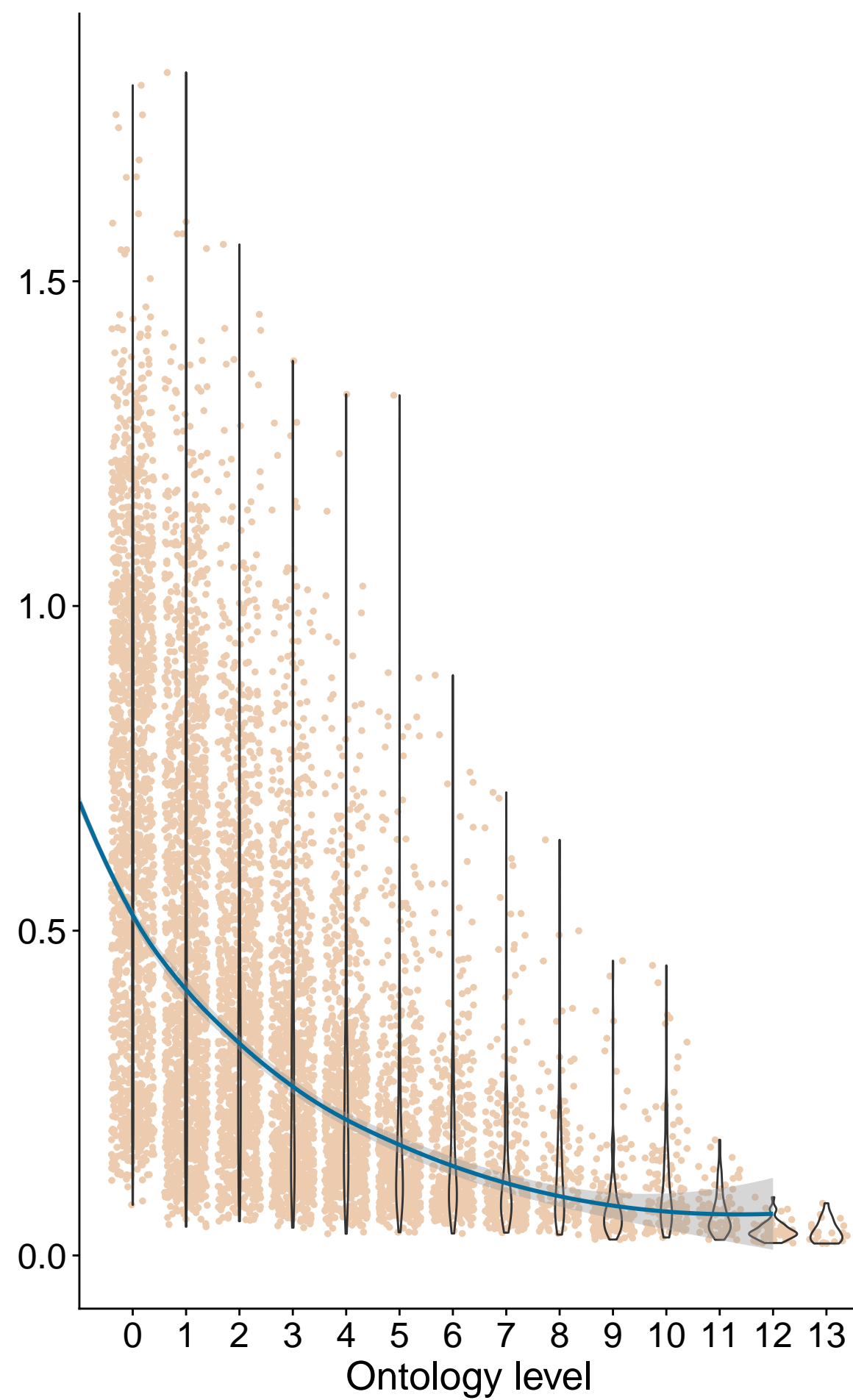


**A**

Number of associated cell types

**B** $\log_{10}(\text{fold enrichment})$ **C**

Number of associated genes

