

A

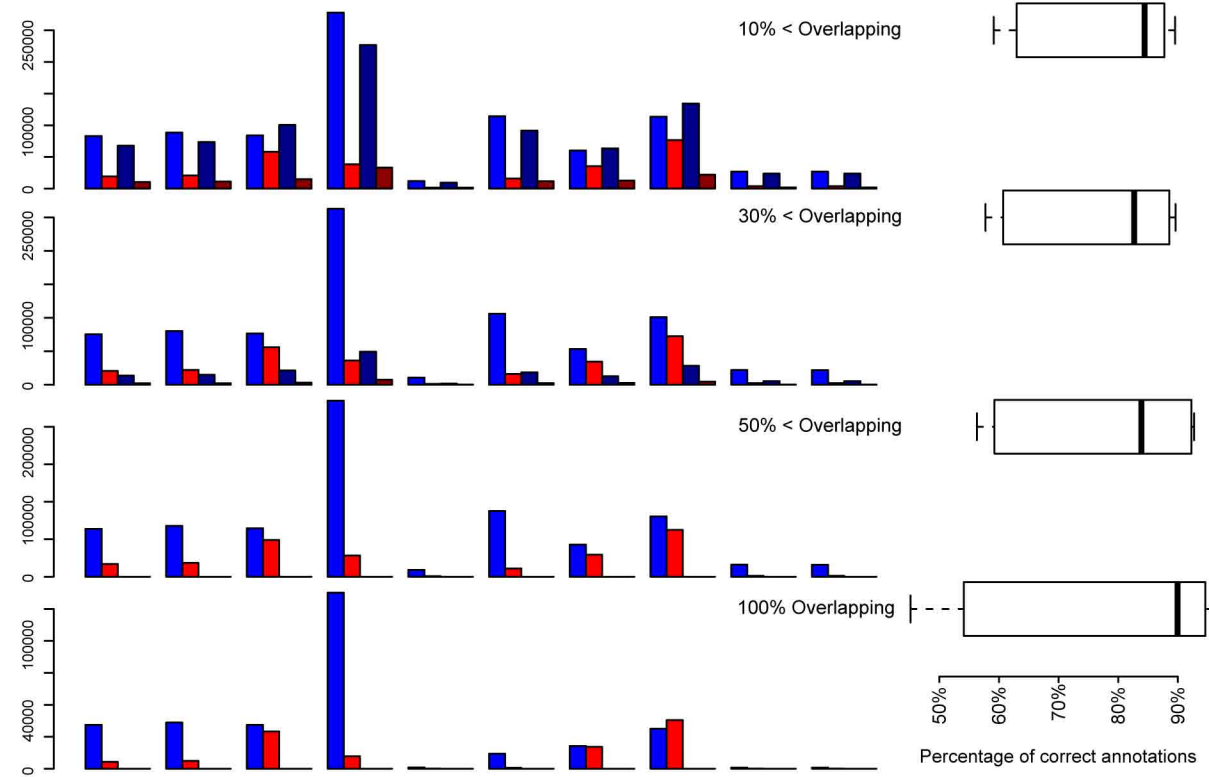
Functional annotation in the viral-bacterial metagenome

Percentage of correct annoations above
overlapping threshold

Number of reads

Assembler C 02 C 05 C meta Genovo M 100/98 M 60/95 N 100/98 N 60/95 Velvet Meta_Velvet

Percentage of correct annotations



B

Functional annotation in the viral metagenome

Percentage of correct annoations above
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Percentage of correct annotations

