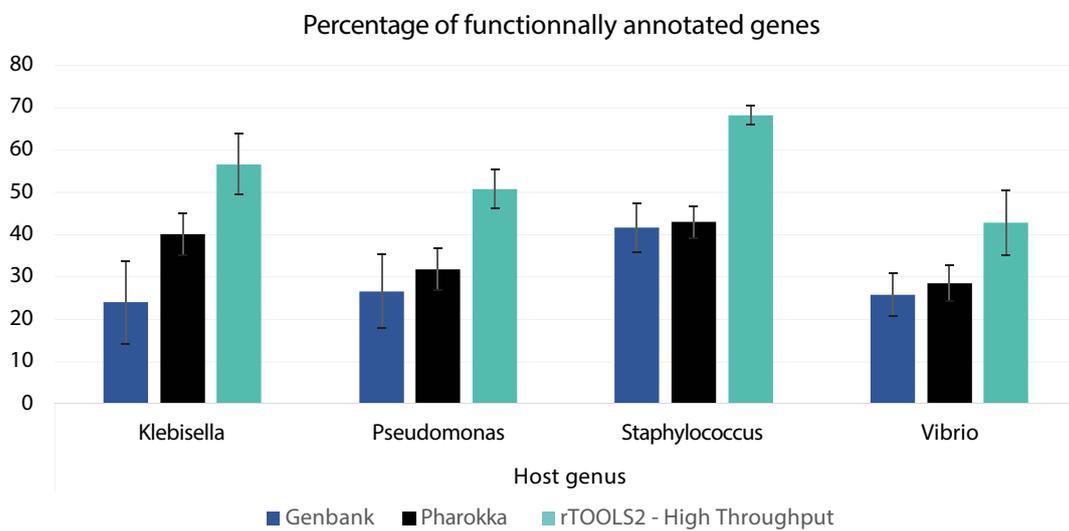
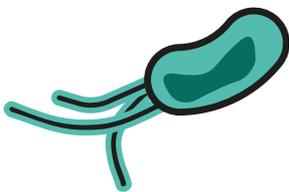


R I M E
B I O I N F O R M A T I C S

Is phage genome annotation **too cryptic** ?
time-consuming ?
yielding little results ?

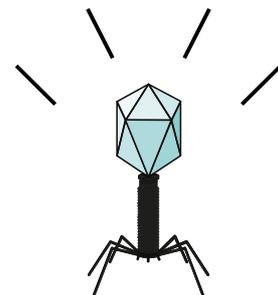


Boost your phage research with rTOOLS2



Production strain selection

- Justifiable, reproducible results
- Compliant with US and EU regulations



Phage characterization

- x2 more annotated genes
- Compliant with best publication practices

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