

USING BAT GENOMES TO STUDY THE EXTRAORDINARY FEATURES OF BATS

OUR MISSION:

Bat1K is a global initiative that aims to sequence the genomes of the >1400 species of living bats.

These genomes will all be open access and publicly available.

Our consortium aims to generate genomic tools to uncover genetic mechanisms behind the unique traits and extraordinary adaptations of bats.

BAT1K LEADERSHIP

FOUNDING DIRECTORS

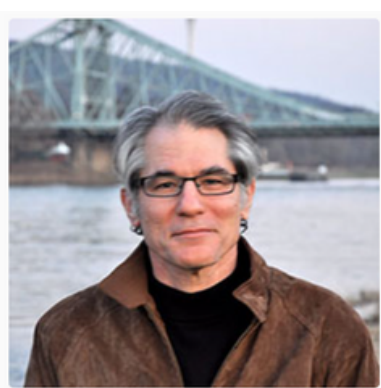


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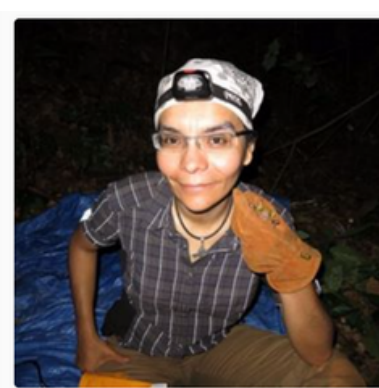


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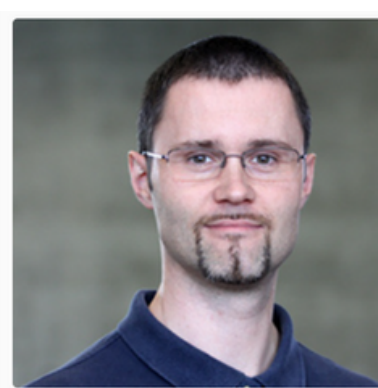
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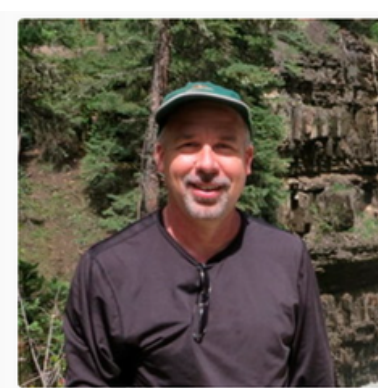
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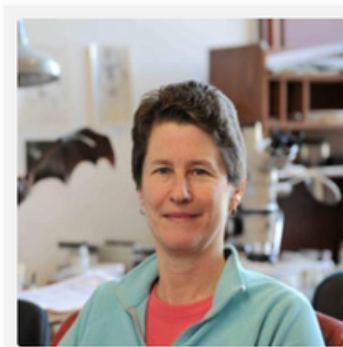


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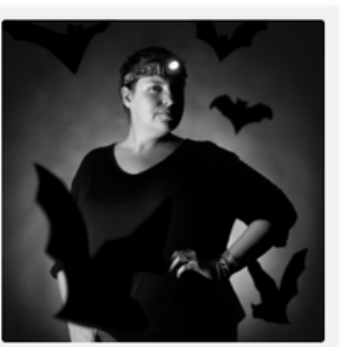
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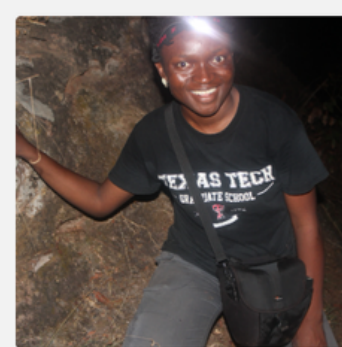
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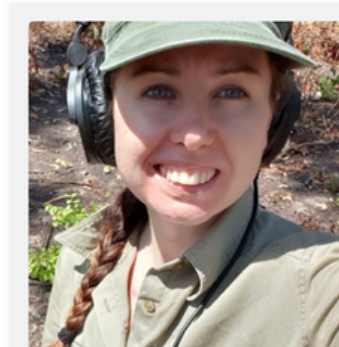
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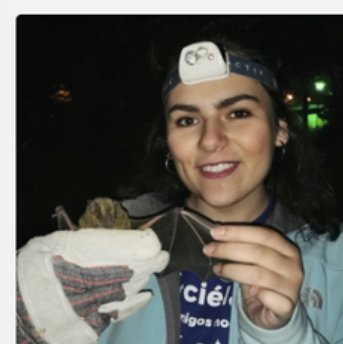
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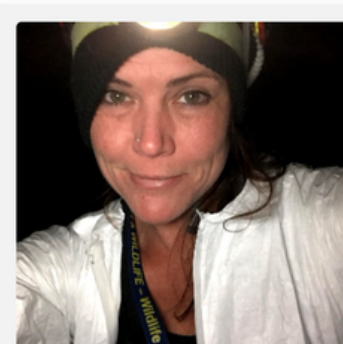
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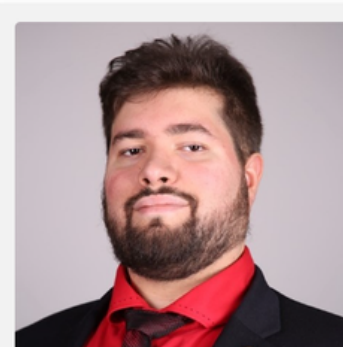
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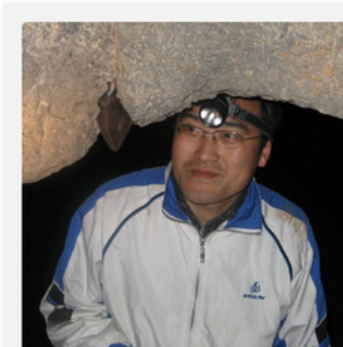
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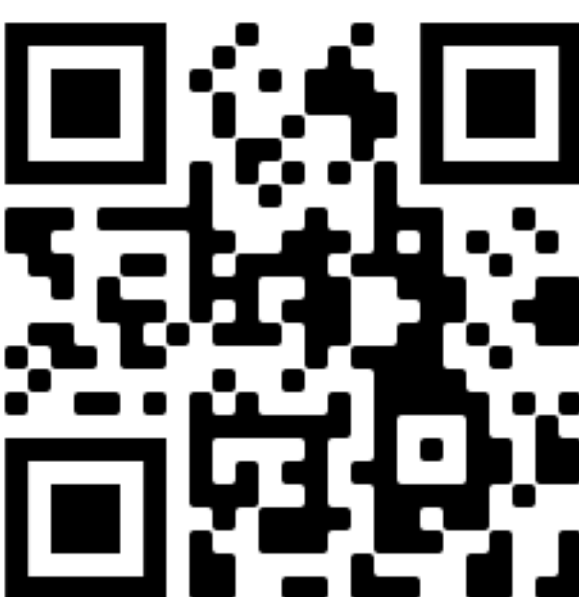
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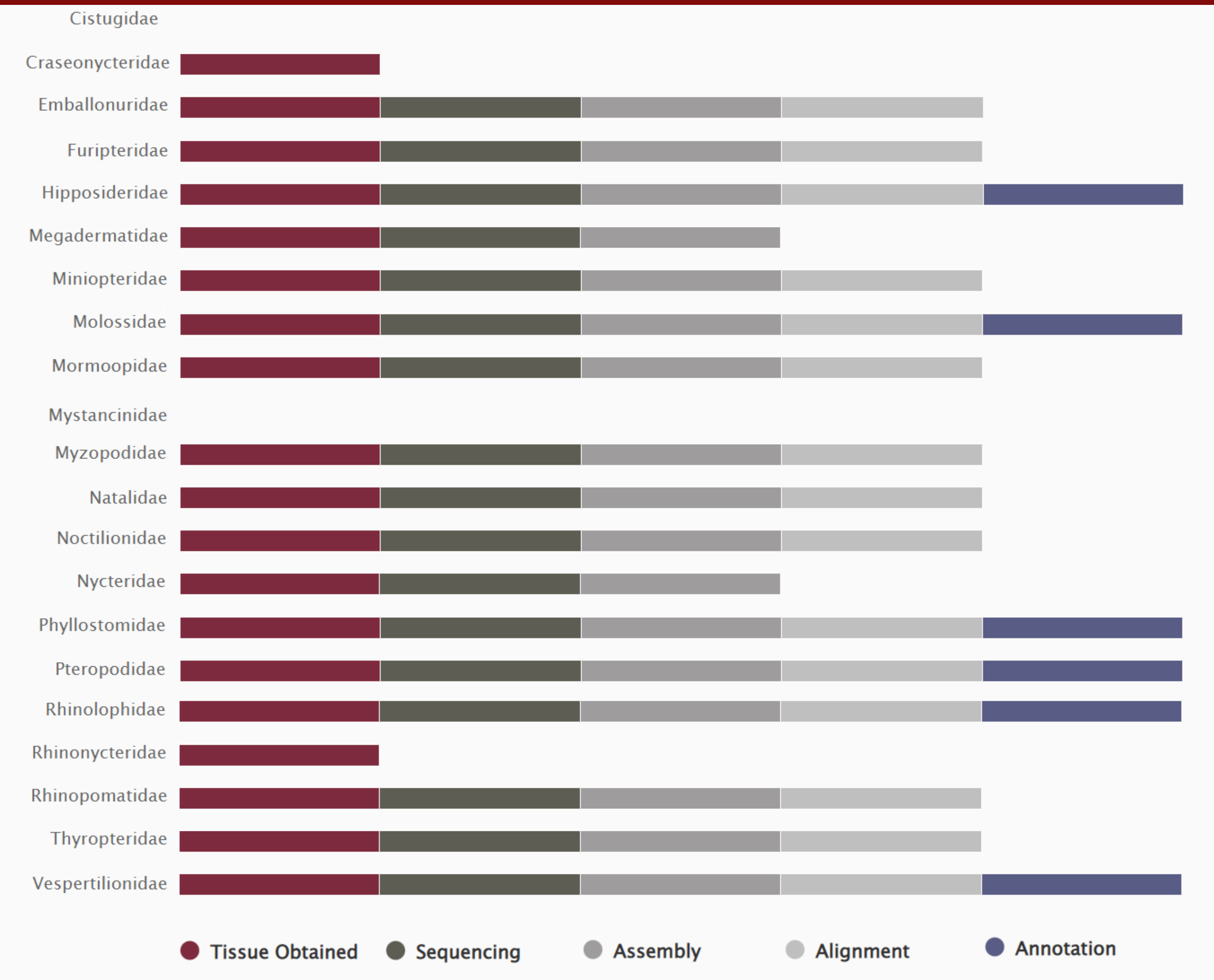


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BAT1K GENOME PHASE I SEQUENCING PROGRESS



GET ACCESS TO BAT1K GENOMES!

The 6 published reference quality bat genomes (Jebb et al, 2020) can be accessed via the original publication in Nature:

<https://www.nature.com/articles/s41586-020-2486-3>

They are hosted on, and can be downladed from the Senckenberg Genome Browser at:

<https://genome.senckenberg.de>

Further information about these genomes, species and genome databases and sequencing progress can be found on our website:

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