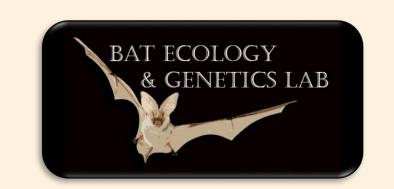
How Rare? Limits of Detection of a Genetic Assay for Species Identification from Guano

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Introduction

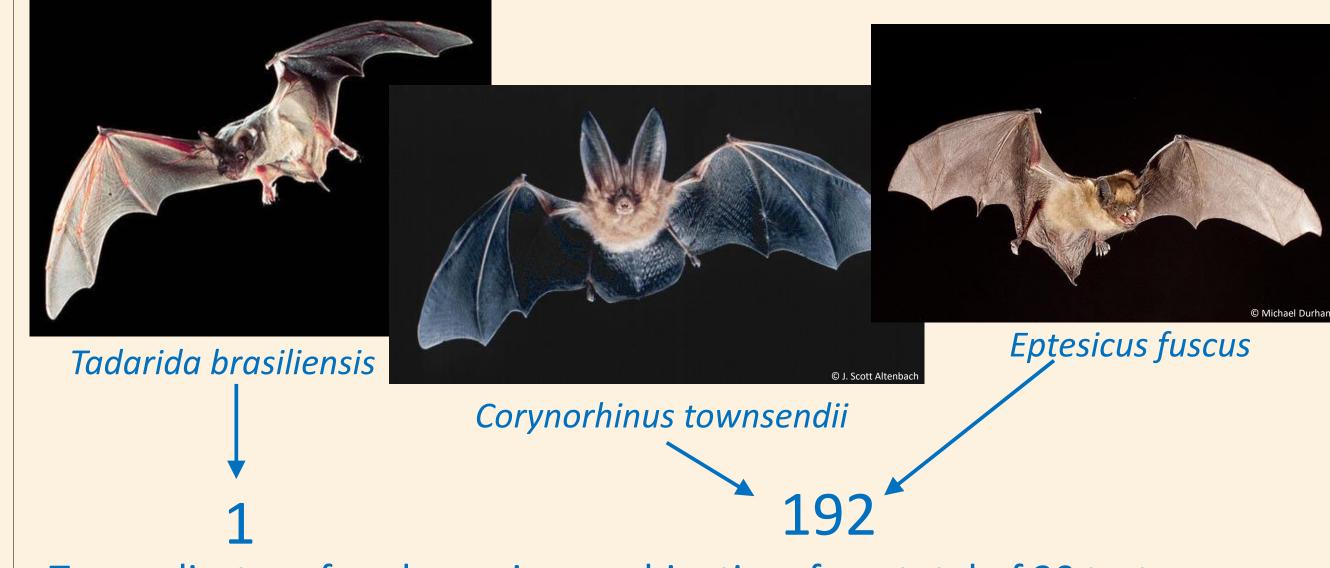
Species from Feces is a DNA mini-barcode assay that uses next-generation amplicon sequencing to identify all the bat species that contributed to a pooled guano sample (Walker *et al.* 2016). A pooled sample consists of about 200 fecal pellets collected from a roost. We wished to determine the capability of the assay in:

- 1. Detecting a rare bat
- 2. Detecting bat species in degraded guano of unknown age
- 3. Large-scale application across a landscape



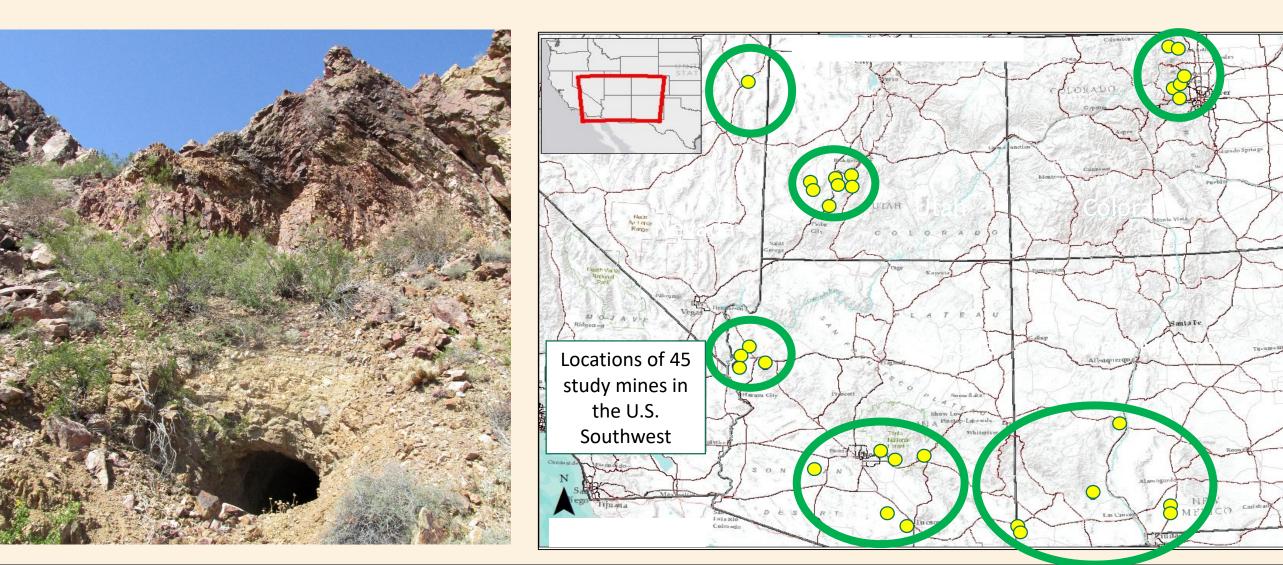
Methods

1. Mock community: Controlled lab study to assess ability to detect a "rare" bat in a pooled sample. Equivalent of 1 fecal pellet of a "rare" species combined with 192 pellets of "common" species (96 of each).



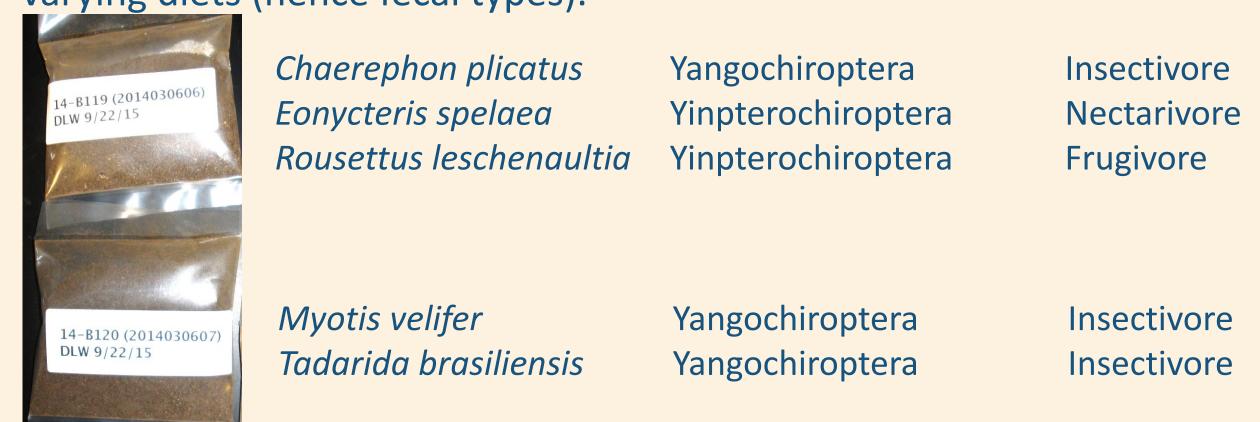
Ten replicates of each species combination, for a total of 30 tests.

- 2. Degraded samples: Single DNA extractions from two guano fertilizers of unknown age, source, or species composition.
- **3.** Large-scale application: Roost survey via a single pooled guano sample collected from each of 45 mines in 5 states in the U.S. southwest

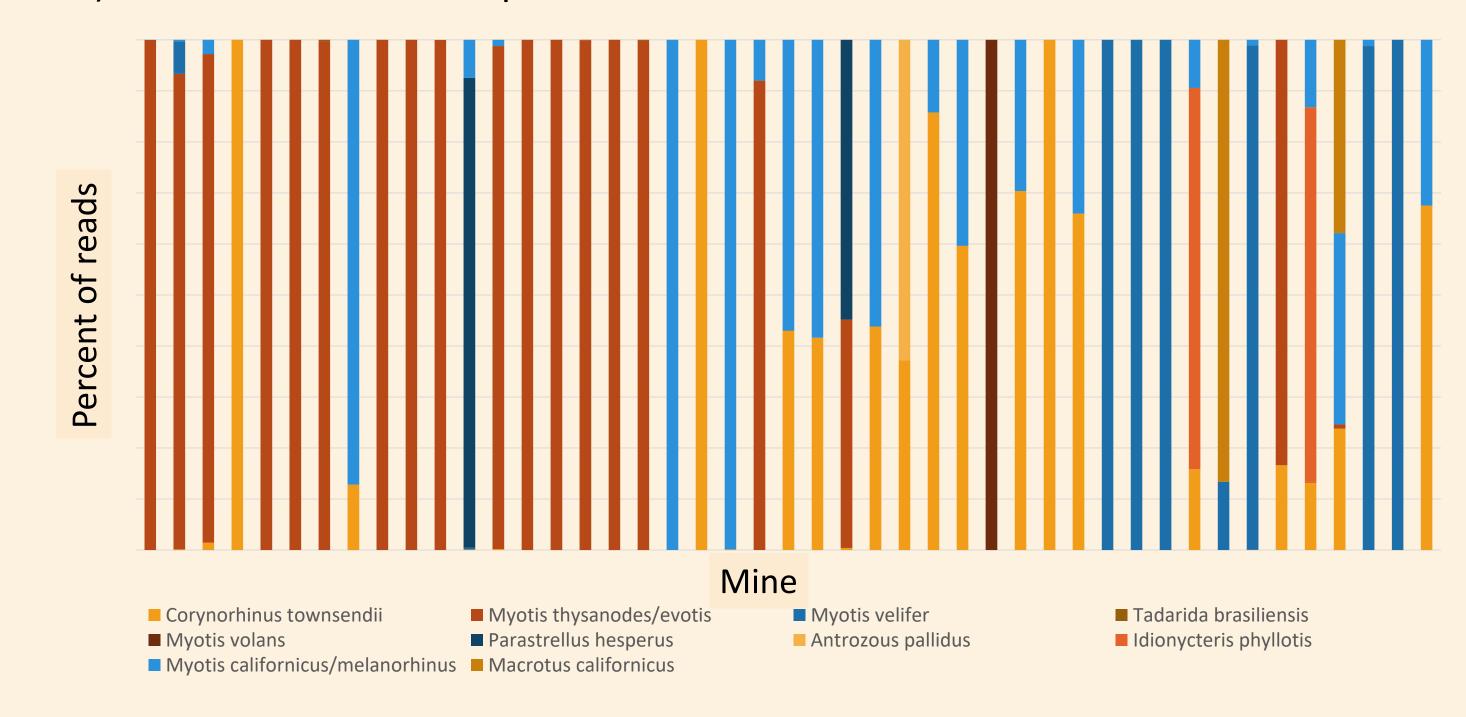


Results

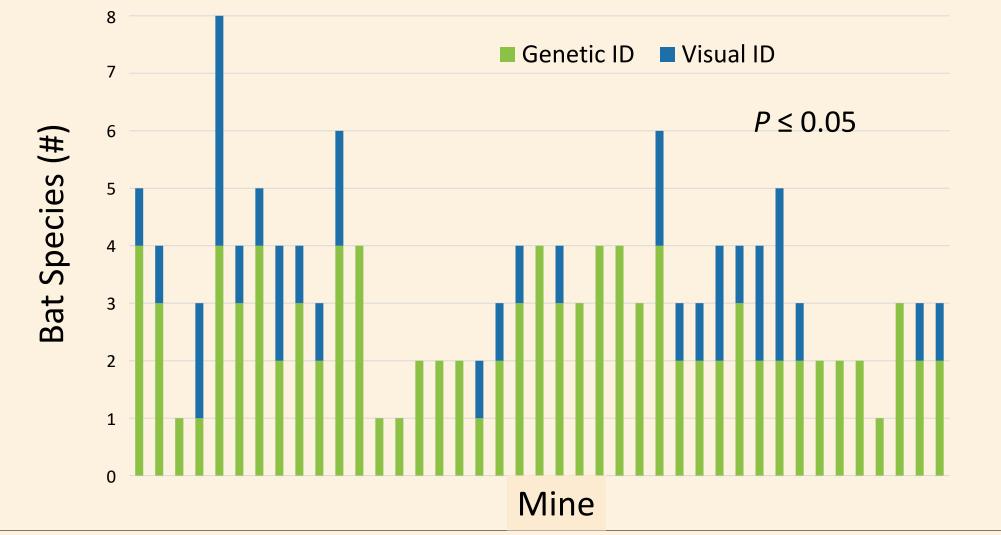
- 1. Regardless of which species of "rare" bat (at 1:192) was used, we always detected it.
- 2. In guano fertilizer, we identified species from both bat suborders and with varying diets (hence fecal types).



- 3. Roost survey:
- a) Between 1 and 4 bat species were detected at each mine.



b) Bat species were genetically identified in 100% of mines, and visually identified in 58% of mines.



Discussion

- 1. Our mock community experiment illustrated that even if a single fecal pellet in a pooled guano sample (typically ~200) is from a different species than the others, it will be detected by the Species from Feces assay.
- 2. The assay was sensitive with degraded DNA (guano fertilizer of unknown age).
- 3. Guano sampling for genetic identification of species was an effective means to survey roosts across a landscape, and provided more species information at lower cost than visual surveys.

Acknowledgements

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References: Walker, F.M., C.H.D. Williamson, D.E. Sanchez, C.J. Sobek, & C.L. Chambers. 2016. Species from Feces: Order-wide identification of Chiroptera from guano and other non-invasive genetic samples. *PLoS ONE* 11(9): e0162342.

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