

"We may see comprehensive barcoding projects within five

robots, offers a rapid way to label and identify new samples.

cause lunar eclipse 13 May 2003

years," says Hebert, "and it's feasible to imagine a database of all animal life on the planet in 20 years." Such a scheme would cost about US\$2 billion, he says.

Powerful code

The barcode is a sequence of about 650 DNA letters, found in a gene in mitochondrial DNA. Mitochondria - cellular powerhouses with their own genomes - are good for genetic identification because there are many copies in each cell, and their DNA evolves relatively quickly, creating differences between species.

By comparing more than 2,000 animal species, Hebert's team found that the sequence could tell nearly all of them apart. The only group that proved tricky was jellyfish and sea anemones, which seem to evolve too slowly for DNA differences between species to be a badge of identity.

Miller and his colleagues are including DNA barcodes in their descriptions of newly discovered moth species in Papua New Guinea. As well as describing new species, they have used the markers to match up the males, females and caterpillars of known species. The animals' different sexes and life stages look completely different.

Barcoding could also help to identify pest species, or to monitor commercial fishing, says Miller. But it won't replace the traditional methods of identifying and classifying species by their appearance, he says: "It's one more tool in the box, but we will always need backup."

References

 Hebert, P. D. N., Ratsingham, S. & Dewaard, J. R.Barcoding animal life: cytochrome c oxidase subunit 1 divergences among closely related species.. *Proceedings of the Royal Society B* (2003). |Article|

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