

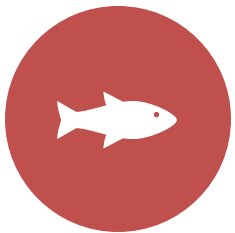


Using eDNA to Monitor Marine Biodiversity



Applied
Genomics

Why Monitor Marine Biodiversity?



Oceans support
complex food webs



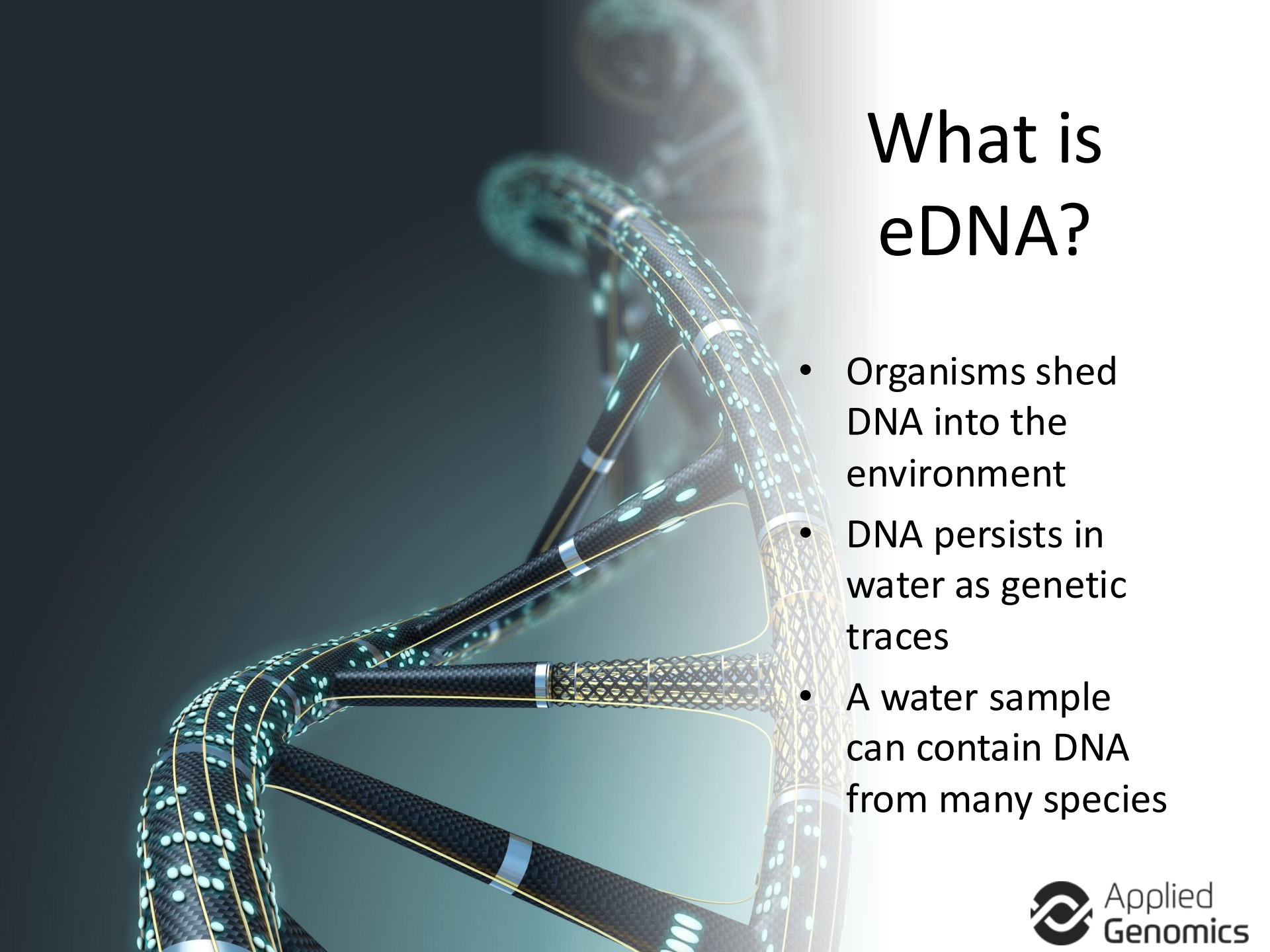
Many species are
hard to observe
directly



Traditional surveys
are expensive and
limited



Key question: How
do we know what
lives there?



What is eDNA?

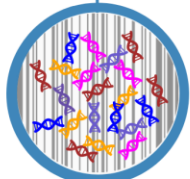
- Organisms shed DNA into the environment
- DNA persists in water as genetic traces
- A water sample can contain DNA from many species



Environmental sample
collection



Environmental DNA
isolation



Enrichment of species
barcoding genes



High throughput DNA
sequencing



Bioinformatics for
species identification

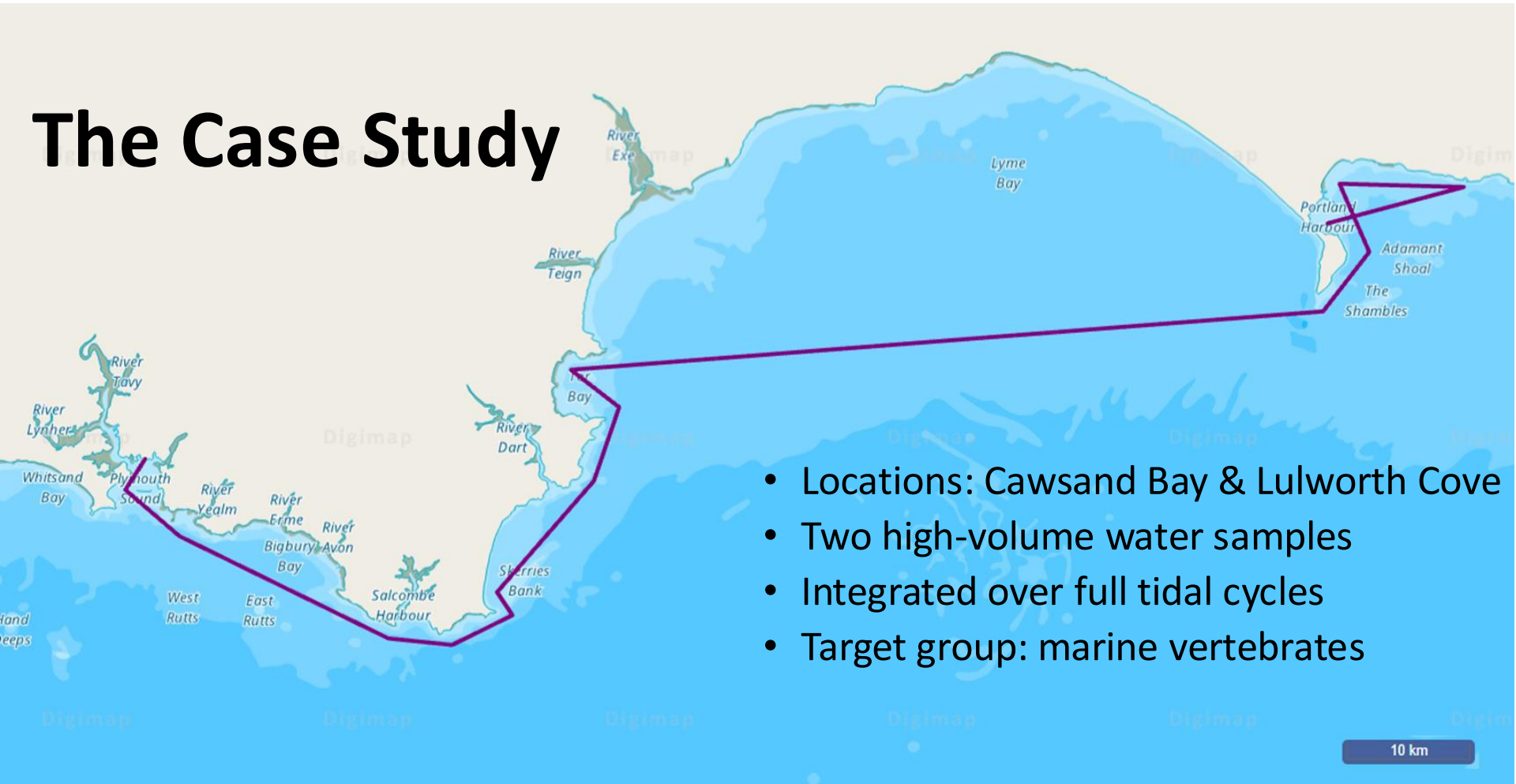


Ecological analysis

How Does eDNA Metabarcoding Work?

1. Collect water samples
2. Extract DNA
3. Amplify barcode genes
4. Sequence DNA
5. Match sequences to
reference databases
6. Analyse patterns

The Case Study



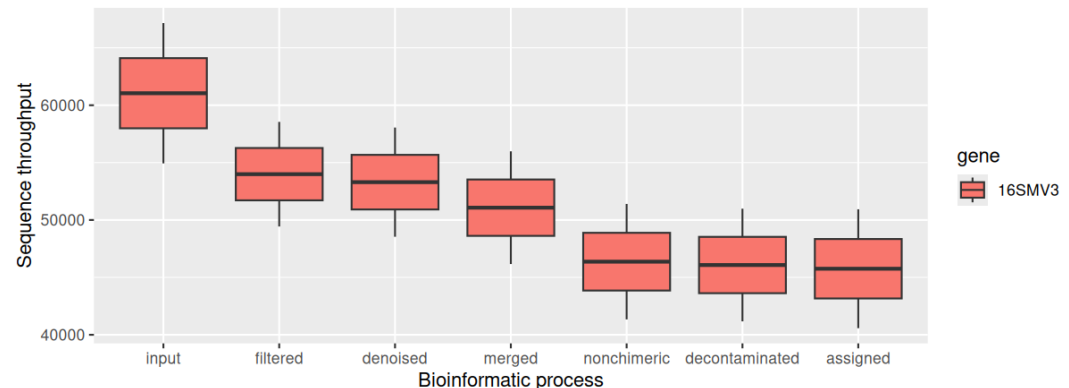
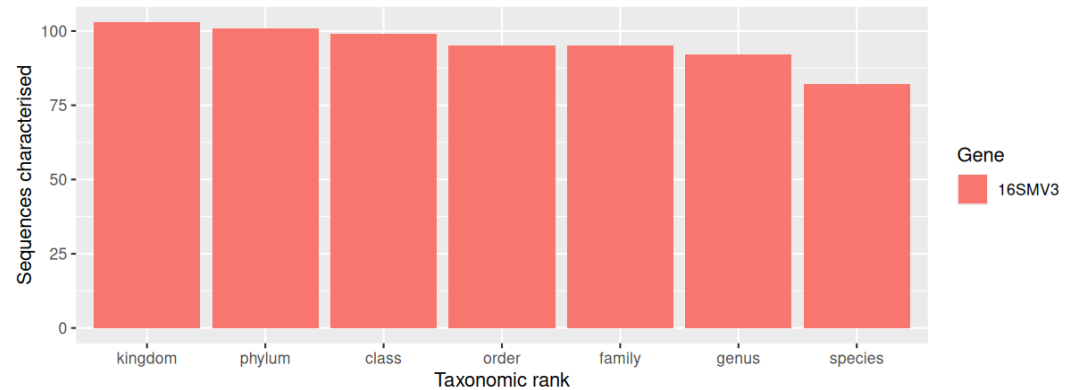
- Locations: Cawsand Bay & Lulworth Cove
- Two high-volume water samples
- Integrated over full tidal cycles
- Target group: marine vertebrates

Data Outputs

From just **2 water samples**, this project produced:

- Hundreds of thousands of DNA sequences
- Hundreds of unique DNA variants (ASVs)
- Dozens of vertebrate taxa detected
- Multiple genetic haplotypes within species
- Multiple quality-control filtering steps

Modern biodiversity monitoring is a data-science problem, not just a fieldwork problem.



Key Species Detected

- European conger eel (*Conger conger*)
- European sardine (*Sardina pilchardus*)
- European hake (*Merluccius merluccius*)
- Sand smelt (*Atherina presbyter*)
- Wrasse (*Labrus* spp.)

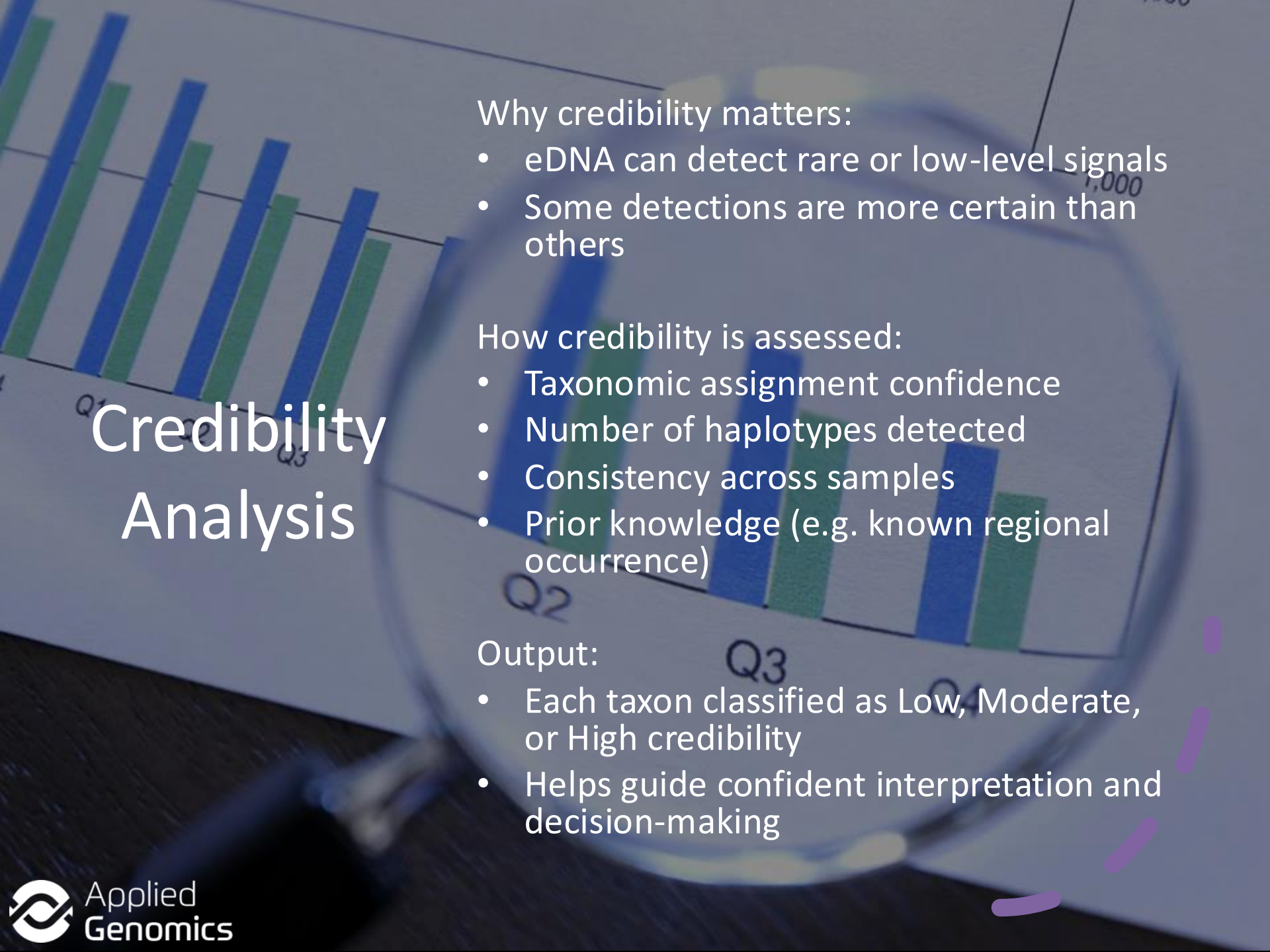
Why These Species Matter

- Important predators and food-web species
- Commercially valuable species
- Indicators of healthy marine ecosystems



What Are Haplotypes?

- Genetic variants within a species
- Multiple haplotypes suggest genetic diversity
- Not a count of individuals



Credibility Analysis

Why credibility matters:

- eDNA can detect rare or low-level signals
- Some detections are more certain than others

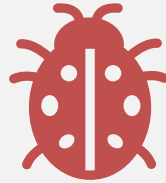
How credibility is assessed:

- Taxonomic assignment confidence
- Number of haplotypes detected
- Consistency across samples
- Prior knowledge (e.g. known regional occurrence)

Output:

- Each taxon classified as Low, Moderate, or High credibility
- Helps guide confident interpretation and decision-making

Strengths of eDNA Monitoring



Non-invasive



Detects elusive species



Broad taxonomic coverage



Efficient and scalable



Limitations to Understand

- Presence/absence, not abundance
- DNA can move with water
- Reference databases are incomplete

Why This Matters for the Future

- Conservation monitoring
- Fisheries management
- Long-term ecosystem assessment
- Growing field for future scientists