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# Meeting Energy Industry Biodiversity Challenges through the Environmental Genomics Joint Industry Program

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Image courtesy of IOGP



# Energy Industry Biodiversity Monitoring

## Environmental Baselines



Images courtesy of Chevron

## Requirements

- Detecting species presence and distribution at new project sites
- Data for Environmental Impact Assessment (EIA) and Approvals
- Baseline data for the unlikely event of an oil spill

## Opportunities for eDNA use

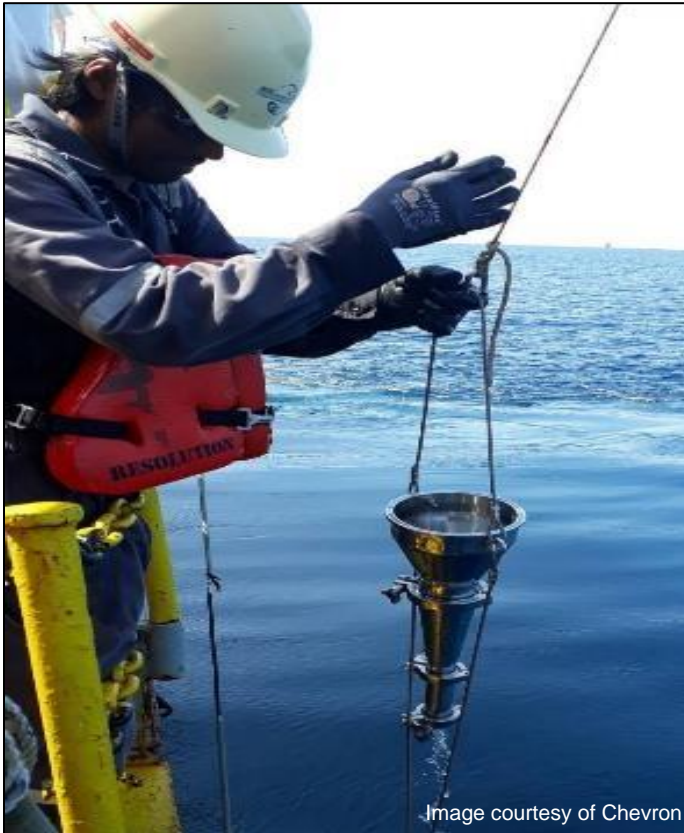
- Large reduction in field time (costs, safety risk)
- eDNA provides holistic data on entire assemblages

## Challenges for eDNA use

- Many regulatory agencies do not yet prescribe or accept eDNA for baselines
- eDNA still viewed by industry as an additional method/cost, not a replacement
- New geographies = limited reference sequences

# Energy Industry Biodiversity Monitoring

## Impact Assessments and Compliance Monitoring



### Requirements

- Assess impacts from construction of projects (EIA)
- Provide routine data for compliance monitoring and reporting

### Opportunities for eDNA use

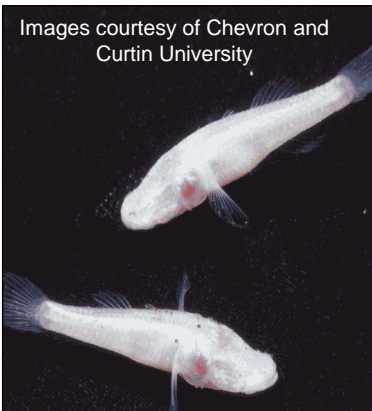
- Increased power to detect changes in biological communities

### Challenges for eDNA use

- Data needs to stand up to regulatory and legal scrutiny
- Lack of standard methods for assessing and reporting error in eDNA data
- Limited validation of spatio-temporal variation in eDNA
- Limited measures of abundance or condition

# Energy Industry Biodiversity Monitoring

## Species of Interest or Concern



Images courtesy of Chevron and Curtin University

## Requirements

- Monitoring and management of impacts to rare, threatened or endangered species or species that are difficult to monitor (e.g. cryptic, subterranean)
- Detection of invasive species

## Opportunities for eDNA use

- Improved detection rates (e.g. low numbers, juveniles, rare, cryptic species)
- Powerful when combined with geospatial data to map distributions
- Can be deployed by non-specialist staff

## Challenges for eDNA use

- Understanding levels of false positives and negatives
- Turnaround times (critical for invasive species)
- Detection of species in arid environments

# Energy Industry Biodiversity Monitoring

## Decommissioning, Remediation & Restoration



Images courtesy of Chevron and Curtin University

## Requirements

- Demonstrating biodiversity benefits and risks associated with decommissioning
- Assessing recovery of soil communities impacted by contamination
- Assessing site restoration success and driving factors

## Opportunities for eDNA use

- Holistic data on assemblages can provide additional insights into biodiversity value, ecosystem condition, factors driving success

## Challenges for eDNA use

- Applications relatively new and evolving
- Uptake by industry and acceptance by regulators a work in progress

# Energy Industry Biodiversity Monitoring

## Nature Reporting, Nature-Positive, Nature-based Solutions



### Requirements

- Nature-related financial disclosures (e.g. EU CSRD, TNFD)
- Demonstrating Nature-Positive Impacts
- Demonstrating Biodiversity Co-benefits of Nature-based Solutions (NbS)

### Opportunities for eDNA use

- Ability to cost-effectively scale biodiversity monitoring
- Can provides data on target species as well as holistic data on entire assemblages

### Challenges for eDNA use

- Integrating eDNA with other data sources and comparing data through time
- Agreement on metrics and indicators of ecosystem condition provided by eDNA

Corporate  
Sustainability  
Reporting  
Directive



# Addressing eDNA Challenges through the Environmental Genomics Joint Industry Program

## Goals:

- Mutualize R&D efforts to accelerate the use of environmental genomics for environmental management applications in the energy industry
- Develop environmental genomics guidance applicable to all phases of energy industry activities, study objectives and environments



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# Development of Energy Industry eDNA Guidance

## Objectives:

- Allow standardization of approaches and reproducibility of results
- Leverage existing guidance and publications (do not re-invent the wheel)
- Facilitate regulatory input to guidance to improve alignment on expectations
- Ensure guidance is evergreen and regularly updated as the field evolves

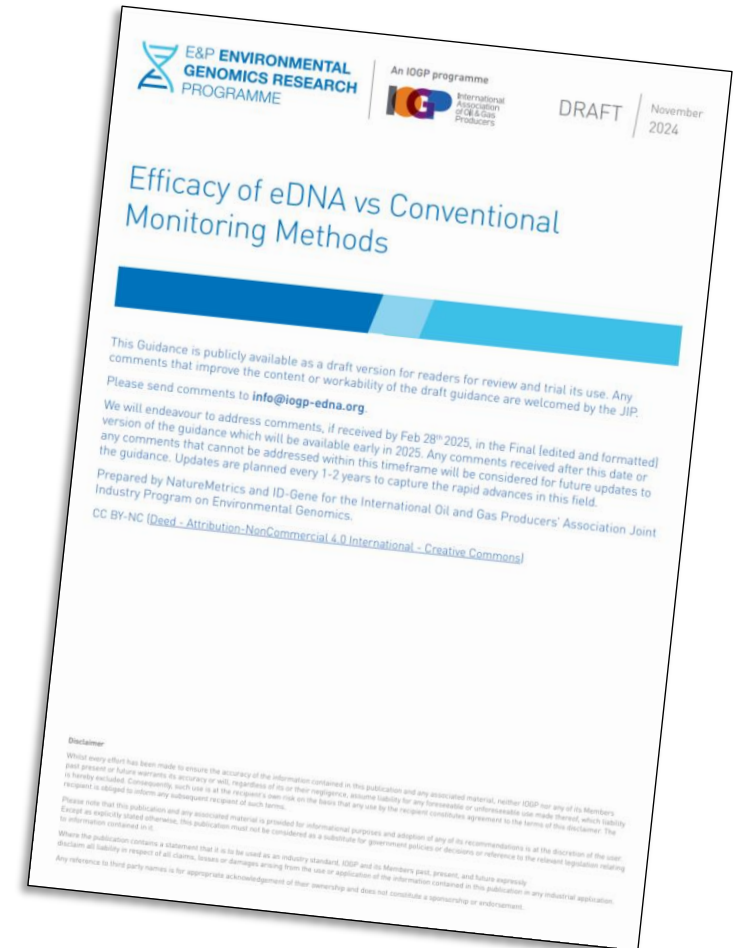
## Four comprehensive guidance chapters drafted (400+ pages):

Chapter 1: *Efficacy of eDNA versus conventional monitoring methods*

Chapter 2: *Sampling guidelines and considerations for industry applications*

Chapter 3: *Laboratory analysis guidelines*

Chapter 4: *Bioinformatics, data analysis and interpretation guidelines*



# Guidance Chapter 1: Efficacy of Genomic versus Conventional Monitoring Methods

- Compares the efficacy of environmental genomic versus conventional monitoring methods for a range of industry objectives, taxa and habitats, through a meta-analysis of published literature
- Presents decision trees that guide environmental practitioners and contractors on when, where and how to apply genomic methods
- Summarizes the current state of readiness of genomics for a range industry applications (evergreen, to be updated regularly)



# Guidance Chapter 1: Efficacy of Genomic versus Conventional Monitoring Methods

Habitat	Target community	eDNA methods	Conventional methods	Recommended use	Phase of operation
Marine	<b>Benthic sediment</b>	Sediment sample; Macrofauna sample	Macrofauna sample; Benthic imagery; Active acoustics	CA with further validation	B; E&P; OSR; R
	<b>Benthic hard substrate</b>	Surface scrape; Surface swabs; Surface suction	Surface scrape; Benthic imagery; Active acoustics	CA in combination	B; E&P; OSR; D; R
	<b>Fish</b>	Water filter	Netting; Benthic and aerial imagery; Active acoustics	Stand-alone CA & SD	B; E&P; OSR; D; R
	<b>Broad vertebrates</b>	Water filter	Visual observations; Passive acoustics; Aerial imagery	CA & SD in combination	B; OSR; R
	<b>Zooplankton &amp; Phytoplankton</b>	Water filter; Bulk net sample	Water sampling; Netting; Active acoustics; Aerial imagery	CA in combination, and SD stand-alone	B; OSR; D
Freshwater	<b>Fish</b>	Water filter	Netting; Electrofishing	CA in combination and SD stand-alone	B; E&P; OSR; D; R
	<b>Aquatic Invertebrates</b>	Water filter; Bulk sampling	Kicknet; Traps	CA with further validation and SD stand-alone	B; E&P; OSR; R
	<b>Phytobenthos</b>	Biofilm sampling	Biofilm sampling;	CA with further validation	B; E&P; OSR; R
Terrestrial	<b>Broad vertebrates</b>	Water filter; Air DNA; Surface swabs; iDNA	Visual and auditory; Camera traps; Vertebrate traps; Artificial cover	CA and SD in combination	B; OSR; D; R
	<b>Invertebrates</b>	Invertebrate traps; Air DNA; Surface swabs	Invertebrate traps	CA in combination	B; OSR; D; R
	<b>Soil</b>	Soil sampling	Soil sampling	CA with further validation	B; E&P; OSR; R

**Key:** Recommended use: CA = Community analysis; SD = Species detection

Phase of operation: B = Baseline; E&P = Exploratory and production; OSR = Oil spill and remediation; D = Decommissioning; R = Restoration

# Guidance Chapter 2: Development of eDNA Sampling Standards and Guidelines

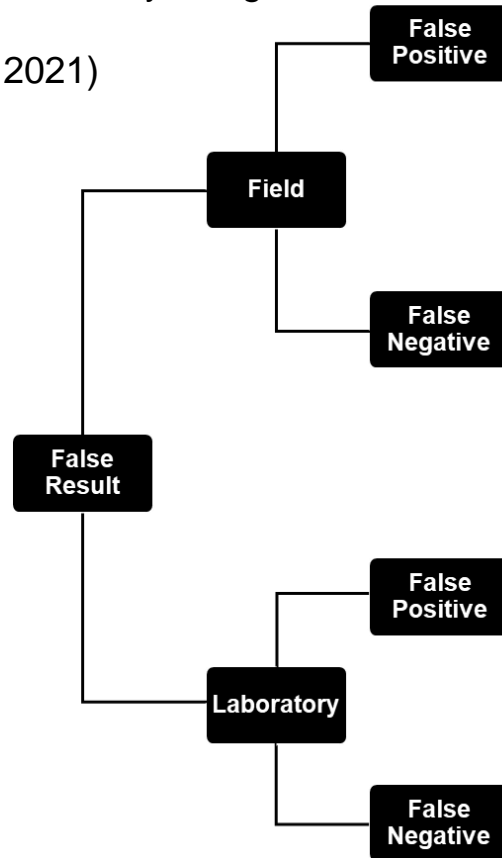
- Presents critical considerations for industry practitioners and contractors when designing genomics sampling programs
- Provides recommendations that improve robustness of sampling, representation of taxa and reproducibility of results
- Identifies critical knowledge gaps that need to be addressed to improve confidence in sampling and data obtained



# Guidance Chapter 2: Development of eDNA Sampling Standards and Guidelines

**Factors influencing a false result** introduced during eDNA field sampling or laboratory analysis, with the potential for mitigation based on study design

(modified from Burian et al., 2021)



Error Description	Mitigation Potential
<ul style="list-style-type: none"> <li>Contamination during sampling (accidental eDNA across site transfer)</li> <li>*Sampling independent contamination (e.g., by birds, predator feces, ballast water or other human activities)</li> <li>*Vertical transport (e.g., downstream eDNA along river networks influenced by hydrology)</li> <li>*eDNA persistence (e.g., historical eDNA introduced to sample)</li> </ul>	<ul style="list-style-type: none"> <li>Very High</li> <li>Intermediate</li> <li>Intermediate</li> <li>High</li> </ul>
<ul style="list-style-type: none"> <li>Inappropriate sampling method (wrong habitats within sites or insufficient media quantities are sampled)</li> <li>Environmental heterogeneity (of environment and populations)</li> <li>eDNA persistence (high microbial activity leads to rapid eDNA degradation and low concentrations)</li> <li>Physiological factors (e.g., seasonality in activity patterns)</li> <li>Environmental drivers (e.g., rain events can flush out eDNA)</li> </ul>	<ul style="list-style-type: none"> <li>Very High</li> <li>Intermediate</li> <li>Intermediate</li> <li>High</li> <li>Intermediate</li> </ul>
<ul style="list-style-type: none"> <li>Technical contamination (contamination across samples due to inappropriate lab procedures or mislabeling of samples)</li> <li>Insufficient specificity (non-target eDNA is amplified)</li> </ul>	<ul style="list-style-type: none"> <li>Very High</li> <li>High</li> </ul>
<ul style="list-style-type: none"> <li>Inhibition (amplification of target DNA is inhibited by co-extracted compounds)</li> <li>Inadequate extraction protocol (low extracted DNA concentrations)</li> <li>Inadequate sample preservation (if extraction is delayed)</li> <li>Low sensitivity (low amplification efficiency, high limit of detection)</li> <li>Insufficient specificity (assays fails to detect certain haplotypes)</li> </ul>	<ul style="list-style-type: none"> <li>Intermediate</li> <li>Very High</li> <li>Very High</li> <li>Very High</li> <li>High</li> </ul>

# Guidance Chapter 3: eDNA Laboratory Analysis Guidelines

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- Recommends minimum laboratory analysis requirements that allow industry practitioners to confidently rely on genomics data
- Provides recommendations that are relevant to a range of study objectives, sample substrates, environments and taxa
- Identifies critical knowledge gaps that currently introduce uncertainty or bias during laboratory analysis



Image courtesy of eDNATec

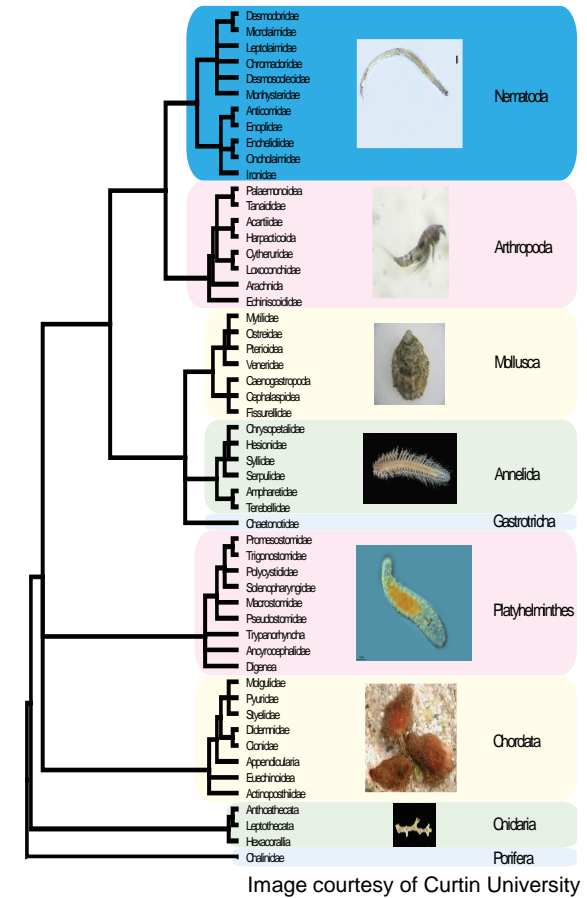
# Guidance Chapter 3: eDNA Laboratory Analysis Standards

## Guidance on meta-barcode primer selection

Gene region	Target Taxa	Example Marker	Mean amplicon length (bp) including primers	Taxonomic Resolution
12S	Fish	MiFishU (Miya et al., 2015) F: NNNNNNGTCGGTAAAACCTCGTGCCAGC R: NNNNNNCATAGTGGGGTATCTAATCCCAGTTTG	234	Medium
	Vertebrates	12SV05 (Riaz et al., 2011) F: TTAGATACCCCACTATGC R: TAGAACAGGCTCCTCTAG	133	Low
16S	Bacteria	515F–806R (V4 region) (Apprill et al., 2015; Parada et al., 2016) F: GTGYCAGCMGCCGCGGTAA R: GGACTACNVGGGTWTCTAAT	390	Medium
	Mammals	16Smam (P. Taylor, 1996) F: CGGTTGGGGTGACCTCGGA R: GCTGTTATCCCTAGGGTAACT	135	Low
18S	Eukaryotes	18SV4M1 (V4 region) (Stat et al., 2017) F: GCAGTAAAAAGCTCGTAG R: TCCAAGAATTRCACCTCT	346	Low
COI	Eukaryotes	F230 (Gibson et al., 2015) F: GGTCAACAAATCATAAAGATATTGG R: CTTATRTTRTTTATNCGNGGRAANGC	281.5	High
	Metazoans, Macroalgae	Leray XT (Wangensteen et al., 2018) F: GGWACWRGWTGRACWITITAYCCYCC R: TAIACYTCIGGRTGICCRAARAAYCA	368	High
	Insects	BF3BR2 (Roger et al., 2022) F: CCHGAYATRGCHTTYCCHCG R: TCDGGRTGNCCRAARAAYCA	458	High

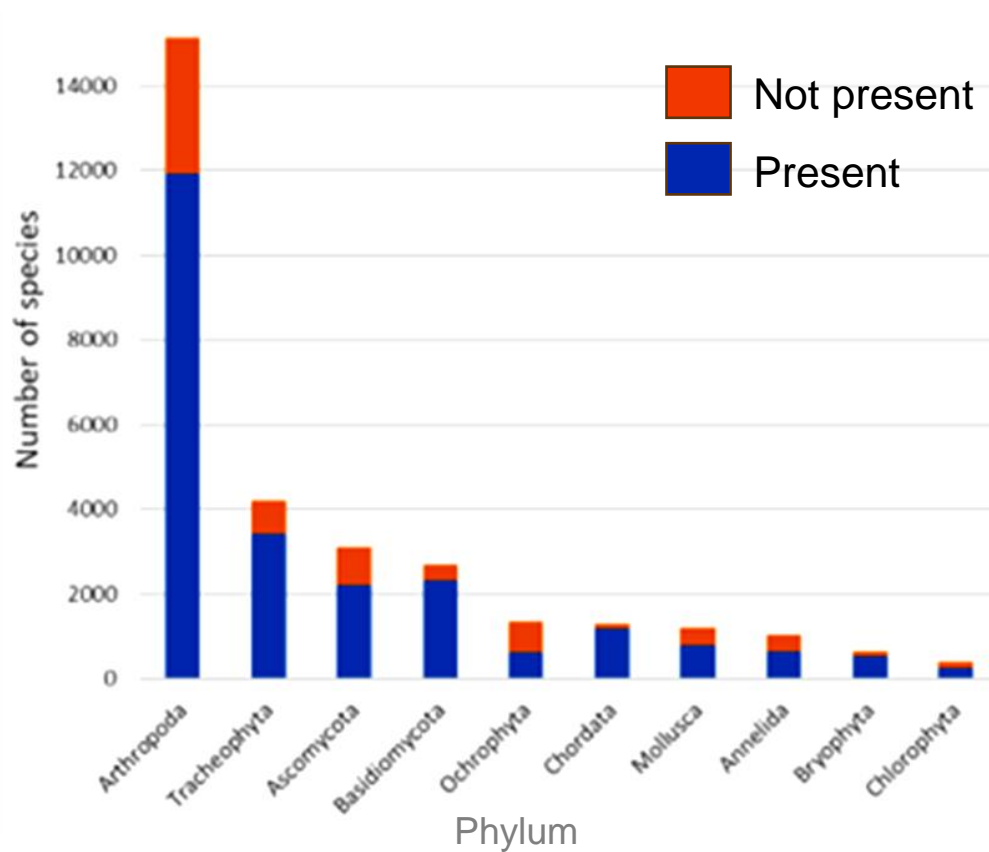
# Guidance Chapter 4: Bioinformatics, Data Analysis and Interpretation

- Provides recommended workflows and methods for bioinformatics and data analysis
- Includes interviews with thought leaders on controversial or rapidly-evolving topics
- Improves ease of interpretation, reproducibility of results and greater confidence in using genomics data in management decisions
- Includes status of sequence reference databases for many key regions where the energy industry operates

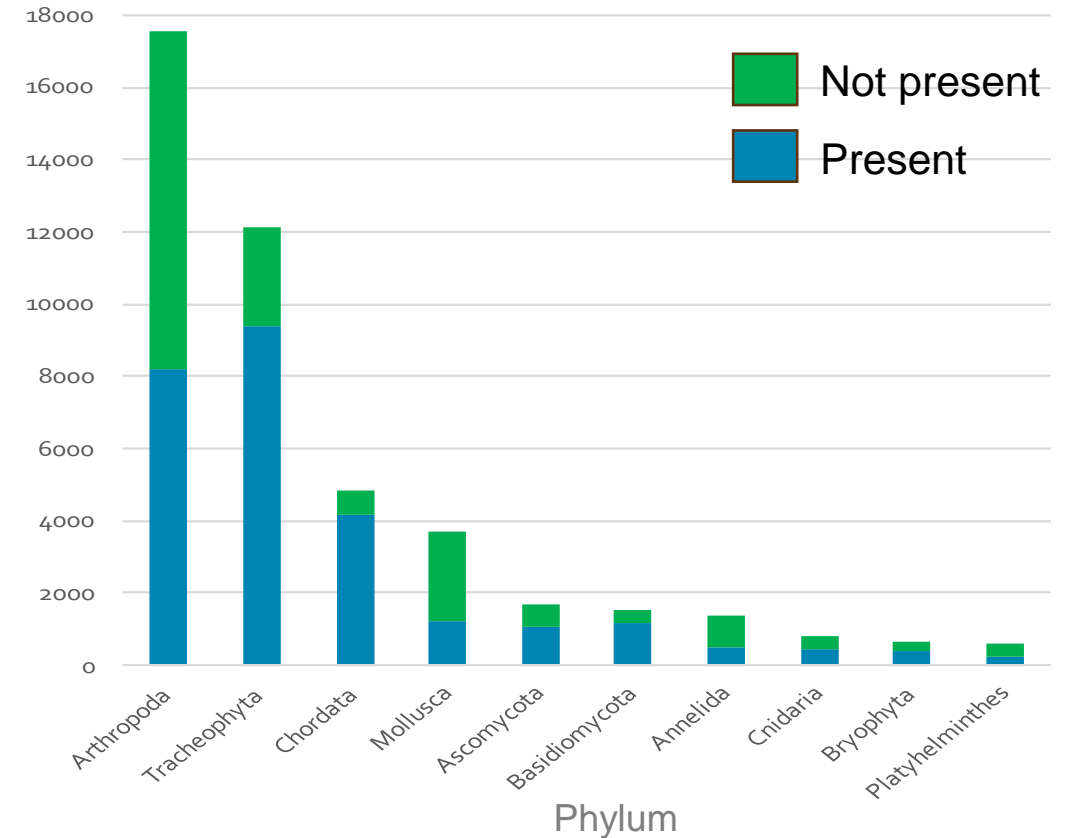


# Guidance Chapter 4: Bioinformatics, Data Analysis and Interpretation

Sequence Database Completeness:  
North Sea



Sequence Database Completeness:  
Gulf of Mexico



Data based on number of recorded species for region from GBIF versus Genbank nucleotide database (mid-2023); Figures courtesy of eDNAtec

# We need your help!

- **Draft guidance** publicly available on JIP website:  
<https://www.iogp-edna.org/publications/>



- Please review and use guidance and provide comments to:  
[info@iogp-edna.org](mailto:info@iogp-edna.org)
- Comments will be addressed Q1 2025 revision



Chapter 1 - Efficacy of eDNA vs Conventional Monitoring Methods



Chapter 2 - eDNA Sampling Standards and Guidelines



Chapter 3 - Laboratory Analysis Guidelines and Best Practices for Environmental Genomics Applications Relevant to the Energy Sector



Chapter 4 - Industry Guidance on Bioinformatics Analysis Standards and Guidelines for eDNA Data relevant to O&G

# JIP Phase 2 Priorities (2025-2028)

1. **Nature Reporting:** Framework on how to incorporate eDNA methods
2. **Nature-based Solutions (NbS):** Framework on how to incorporate eDNA methods
3. **Estimating and reporting levels of error in eDNA data:** Standardized approach
4. **Extracting greater value from eDNA data:** Guidance and case studies (e.g. using occupancy modelling, relative reads, co-occurrence networks)
5. **eDNA sampling technology development:** Maintain currency of knowledge within industry and outline needs to technology developers
6. **Industry sequence data:** Facilitate sharing for other uses such as conservation or validation of geospatial/AI models
7. **eDNA fluency:** Build fluency among industry, regulators and government agencies
8. **Guidance:** Update industry guidance to capture the rapid evolution of the field

Priorities 1-3: work commenced; outputs due mid-2025  
Priorities 4-8: work to commence later in 2025



Image courtesy of Chevron



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