**Data Compliance to FAIR Principles**

**Findable**

**F1. (Meta)data are assigned a globally unique and persistent identifier**

All of the data columns and individual data points as well as sampling locations are assigned a unique identifier. The data is available at a unique URL on GitHub.

**F2. Data are described with rich metadata**

As many of the Darwin Core Standard columns were included and filled in as was possible.

**F3. Metadata clearly and explicitly include the identifier of the data they describe**

All the necessary metadata is contained within the data file itself.

**F4. (Meta)data are registered or indexed in a searchable resource**

It is not at the moment of publication.

**Accessible**

**A1. (Meta)data are retrievable by their identifier using a standardised communications protocol**

The data file can be retrieved from GitHub from a unique URL using command-line software or within GUI programs that allow download from a URL.

**A1.1 The protocol is open, free, and universally implementable**

The protocol that can be used for retrieval is HTTP.

**A1.2 The protocol allows for an authentication and authorisation procedure, where necessary**

Files on GitHub can be restricted to be accessible only to authorised individuals or groups.

**A2. Metadata are accessible, even when the data are no longer available**

As most of the metadata is within the file, this is not the case.

**Interoperable**

**I1. (Meta)data use a formal, accessible, shared, and broadly applicable language for knowledge representation.**

The file format used for the data is XLSX, which is not as accessible as RDF, OWL or JSON-LD, but is both human-readable and machine-readable by many programs.

**I2. (Meta)data use vocabularies that follow FAIR principles**

Standard language and formats (such as the Darwin Core Standard and the GBIF taxonomy) are used where possible.

**I3. (Meta)data include qualified references to other (meta)data**

Where data from other sources was used, it will have been cited appropriately.

**Reusable**

**R1. (Meta)data are richly described with a plurality of accurate and relevant attributes**

Relevant sampling information is described in the dataset.

**R1.1. (Meta)data are released with a clear and accessible data usage license**

The data has been released with an MIT license.

**R1.2. (Meta)data are associated with detailed provenance**

Some detail on the context of the data has been provided, but not citation data.

**R1.3. (Meta)data meet domain-relevant community standards**

The Darwin Core Standard and GBIF taxonomy data was used while creating the dataset.