

This is a published notice on the Find a Tender service: <https://www.find-tender.service.gov.uk/Notice/007937-2022>

Not applicable

## **CEFAS22-11 Next Generation Sequencing Services Framework**

CEFAS

F14: Notice for changes or additional information

Notice identifier: 2022/S 000-007937

Procurement identifier (OCID): ocds-h6vhtk-031eee

Published 23 March 2022, 6:07pm

### **Section I: Contracting authority/entity**

#### **I.1) Name and addresses**

CEFAS

Pakefield Road

Lowestoft

SW1P 3JR

#### **Contact**

Emma Roberts

#### **Email**

[procure@cefas.co.uk](mailto:procure@cefas.co.uk)

#### **Telephone**

+44 2072385921

#### **Country**

United Kingdom

**NUTS code**

UK - United Kingdom

**Internet address(es)**

Main address

[www.cefas.o.uk](http://www.cefas.o.uk)

---

## Section II: Object

### II.1) Scope of the procurement

#### II.1.1) Title

CEFAS22-11 Next Generation Sequencing Services Framework

#### II.1.2) Main CPV code

- 71621000 - Technical analysis or consultancy services

#### II.1.3) Type of contract

Services

#### II.1.4) Short description

o help meet increasing needs, Cefas have a requirement to establish a new framework agreement for the provision of Next Generation Sequencing (NGS) services to include the approaches detailed below, and any proposed solution/awarded Provider must be able to fulfil the following requirements (as a minimum):

1. The service provided must include the receipt of sample(s) from Cefas, Library Preparation, Quality Control and sequencing, and delivery of data back to Cefas.
2. The service provided must be able to offer, depending on the requirements of the project, a range of NGS technical options including but not restricted to high throughput short and long read sequencing (including Illumina MiSeq, HiSeq, and NovaSeq platforms, PacBio, and Oxford Nanopore Technologies) for a diverse range of sequencing requirements, including but not restricted to amplicon sequencing (metabarcoding and multiplexed protocols), whole/partial genome sequencing, mitogenomics, genome skimming, metagenomics and metatranscriptomics, RNASeq, and hybrid approaches.

---

## Section VI. Complementary information

### VI.6) Original notice reference

Notice number: [2022/S 000-006229](#)

---

## Section VII. Changes

### VII.1.2) Text to be corrected in the original notice

Section number

1V.2.2

Instead of

Date

4 April 2022

Local time

12:00pm

Read

Date

29 April 2022

Local time

12:00pm