

## **QUASIMEME**

# Quality assurance of information for marine environmental monitoring

## **Certificate of Analysis**



**ASP shellfish toxins** 

REFERENCE MATERIAL
BT7 sample 72





#### Certificate of Analysis BT7 72

#### **General Information**

In this report an overview is given of analytical data for this sample collected in our proficiency testing program. The consensus values are calculated using a robust statistical model. With this NDA model, the mean and standard deviation are calculated using all reported data when at least 4 results are left after removal of reported 'lower than' (<) and 0 (= zero) values. No outliers are removed.

This report is divided into two sections: Consensus Values and Indicative Values. The division is made on the reliability of the data. Consensus Values are based on at least 8 results and a maximum relative uncertainty of 6.25%. Indicative Values are based on a maximum relative uncertainty of 35% and a minimum of 4 and maximum of 7 results, or a relative uncertainty greater than 6.25% when there are at least 8 results.

For each determinand the following parameters are given: mean, standard deviation, coefficient of variation, number of results, median, MAD (Median of Absolute Deviation), the uncertainty of the mean (consensus or indicative) value and the relative uncertainty.

The results of each determinand is expressed on a wet weight basis.

#### Sample information

QUASIMEME reference materials cover a range of natural Shellfish toxins species from contaminated waters from the North Sea and/or Mediterranean.

This BT7 sample 72 of Blue mussels from CEFAS, United Kingdom is prepared for the QUASIMEME proficiency programs. The results on which the values in this report are based were taken from the periods given in the following table.

Year.Round	Program	Sample
		Round Id
2024.2	BT7	QST361BT





### Consensus Values BT7

Method: Toxins(SF) - BT7

Element Domoic+EpiDomoic Unit Std.Dev. CV % Rel.Uncert. % Mean Ν Median MAD Uncertainty mg/kg 6.23 1.36 21.8 38 6.19 0.780 0.275 4.42