

## DOME – Disease in biota

### Description of DOME output fields

<b>DOME output field</b>	<b>Description</b>
MPROG	Monitoring programme (cf. <a href="#">MPROG</a> ). Multiple programmes are possible separated by “~” (ascii 126)
PURPM	Purpose of monitoring (cf. <a href="#">PURPM</a> ). Multiple purposes possible separated by “~” (ascii 126)
Country	Country that has reported the measurement
RLABO	Reporting institute (cf. <a href="#">RLABO</a> )
STATN	Station name as reported
MYEAR	Monitoring year. Note that MYEAR may not always be identical to the year in date of sampling
DATE	Date of sampling (DD/MM/YYYY)
Latitude	Latitude of measurement (decimal degrees)
Longitude	Longitude of measurement (decimal degrees)
Species	Species/taxon name of examined specimen
SEXCO	Sex of specimen (cf. <a href="#">SEXCO</a> )
NOINP	Number of individuals in sub-sample
MATRX	Matrix being analyzed (cf. <a href="#">MATRX</a> )
NODIS	Number of diseases looked for during a fish disease survey
PARGROUP	Parameter group of measured parameter (cf. <a href="#">PARGROUP</a> )
PARAM	Measured parameter (cf. <a href="#">PARAM</a> )
BASIS	Basis of determination (cf. <a href="#">BASIS</a> )
QFLAG	Qualifier flag for measured value, i.e. “<” (cf. <a href="#">QFLAG</a> ). Multiple flags possible separated by “~” (ascii 126)
Value	Value measured
MUNIT	Measurement unit
VFLAG	Validity flag, i.e. “S” suspect value (cf. <a href="#">VFLAG</a> ). Multiple flags possible separated by “~” (ascii 126)
DETLI	Detection limit value
LMQNT	Limit of quantification
UNCRT	Uncertainty value
METCU	Method of calculating uncertainty (cf. <a href="#">METCU</a> )
ALABO	Analytical laboratory (cf. <a href="#">RLABO</a> )
REFSK	Reference source or key (cf. <a href="#">REFSK</a> )
METST	Method of storage (cf. <a href="#">METST</a> )
METPT	Method of pretreatment (cf. <a href="#">METPT</a> )
METPS	Method of purification/separation (cf. <a href="#">METPS</a> )

METFP	Method of chemical fixation/preservation (cf. <a href="#">METFP</a> )
METCX	Method of chemical extraction (cf. <a href="#">METCX</a> ) multiple flags possible (separate multiple entries with “~” (ascii 126))
METOA	Method of analysis (cf. <a href="#">METOA</a> )
FORML	Formula used in calculation (cf. <a href="#">FORML</a> )
Test organism	<b>Not used in this data type</b>
SMTYP	Sampler type (cf. <a href="#">SMTYP</a> )
SUBNO	Sub-sample identification (can refer to an individual specimen or an aggregate pool of same species)
BULKID	If an individual (or parts thereof) has been analyzed in one or more bulks, the sub-sample (SUBNO) identification(s) of the bulk(s) can be indicated as a BULKID. BULKID refer to a SUBNO within the same sample (see “tblSampleID”). Multiple SUBNO’s possible separated by “~” (ascii 126)
FINFL	Factors potentially influencing guideline compliance and interpretation of data (cf. <a href="#">FINFL</a> ) Multiple factors possible separated by “~” (ascii 126)
DCFLGs	Data conversion flags from ICES Data Centre (cf. <a href="#">DCFLG</a> ). Multiple flags possible separated by “~” (ascii 126)
tblAnalysisID	Reference to analytical method (integer). Can be used to trace additional fields in ICES data systems
tblParamID	Reference to the measurement (integer). Unique for the data table
tblBioID	Reference to the sub-sample (integer). Unique for a sub-sample
tblSampleID	Reference to the sample (integer). Unique for a sample