

## BIOTRAITS TEMPLATE DESCRIPTION

Fields	Description	Variable codes
<b>OriginalID</b>	Unique ID that separates each thermal response curve (as comes in the original dataset)	
<b>FinalID</b>	Unique ID that separates each thermal response curve (to be assigned manually in the global dataset)	
<b>OriginalTraitName</b>	Trait name as it found in source (e.g. "Chlorophyll a - specific photosynthesis rate" or "max photosynthesis rate" or "net co2 consumption rate")	
<b>OriginalTraitDef</b>	Definition of OriginalTraitName	
<b>StandardisedTraitName</b>	Trait name for comparison: "net photosynthesis rate" / "gross photosynthesis rate" / "respiration rate"	
<b>StandardisedTraitDef</b>	Definition of the StandardisedTraitName	
<b>OriginalTraitValue</b>	Value of the measured trait as found in source	
<b>OriginalTraitUnit</b>	Units of OriginalTraitValue as found in source	
<b>OriginalErrorPos</b>	Positive bound of the error around OriginalTraitValue as found in source	
<b>OriginalErrorNeg</b>	Negative bound of the error around OriginalTraitValue as found in source	
<b>OriginalErrorUnit</b>	Units of ErrorPos and ErrorNeg as found in source	<i>SD</i> = standard deviation ; <i>SE</i> = standard error; <i>95% CI</i> = 95% confidence interval; <i>interquartile range</i> ; <i>range</i>
<b>StandardisedTraitValue</b>	Value of the measured trait after unit standardisation (SI)	
<b>StandardisedTraitUnit</b>	Units of StandardisedTraitValue (SI)	
<b>StandardisedErrorPos</b>	Standardised positive bound of the error around StandardisedTraitValue	
<b>StandardisedErrorNeg</b>	Standardised negative bound of the error around StandardisedTraitValue	
<b>StandardisedErrorUnit</b>	Units of StandardisedErrorPos and StandardisedErrorNeg	
<b>Replicates</b>	Number of replicates used to calculate ErrorPos and ErrorNeg	

<b>Habitat</b>	Habitat Type of Consumer	<i>Terrestrial / freshwater / marine</i>
<b>Labfield</b>	Experimental location	<i>Laboratory / field</i>
<b>ArenaValue</b>	Size of the experimental area (e.g. volume of chemostat)	
<b>ArenaUnit</b>	Units of ArenaValue (e.g. liter, square meter)	
<b>ArenaValueSI</b>	SI value of size of arena where trait performance was measured (ArenaValue in SI units)	
<b>ArenaUnitSI</b>	SI unit of size of arena where trait performance was measured.	<i>cubic meter; square meter; meter</i> (when only length of arena was stated)
<b>AmbientTemp</b>	Ambient temperature during experiment (i.e., field or experimental arena)	If NA then see ConTemp or ResTemp
<b>AmbientTempMethod</b>	Method used to measure ambient temperature	<i>Direct / indirect</i>
<b>AmbientTempUnit</b>	Units of AmbientTemp	
<b>AmbientLight</b>	Ambient light conditions during experiment	
<b>AmbientLightUnit</b>	Units of AmbientLight	
<b>SecondStressor</b>	Second stressor during experiment (nutrients, CO2, UV, salinity, etc)	
<b>SecondStressorDef</b>	Description of SecondStressor	
<b>SecondStressorValue</b>	Value of SecondStressor	
<b>SecondStressorUnit</b>	Units of SecondStressorValue	
<b>EquilibTimeValue</b>	How much time was given for equilibration of the measured rate to a given temperature	
<b>EquilibTimeUnit</b>	Units of EquilibTimeValue	
<b>ObsTimeValue</b>	How long the measure was taken for at each fixed temperature (e.g. how long was photosynthesis rate measured at a given temperature)	
<b>ObsTimeUnit</b>	Units of ObsTimeValue	
<b>ObsTimeValueSI</b>	ObsTimeValue in SI units	
<b>ObsTimeUnitSI</b>	Units of ObsTimeValueSI	- <i>second</i> ; - <i>prey caught</i> (time taken for ObsTimeValueSI number of prey caught)
<b>ObsTimeNotes</b>	Notes for ObsTimeValue (i.e., time over which experiment was run for measurement of trait performance).	
<b>ResRepValue</b>	How often the resource was replaced	

<b>ResRepUnit</b>	Units of ResRepValue	
<b>ResRepValueSI</b>	SI value for how often resources were replaced over observation time (same as ResRepValue using SI)	
<b>ResRepUnitSI</b>	SI units for ResRepValueSI	<ul style="list-style-type: none"> <li>- <i>not replaced</i>;</li> <li>- <i>second</i>;</li> <li>- <i>to satiation</i> (resources replaced sufficiently frequently so that consumer always had access to resources)</li> </ul>
<b>Climate</b>	Type of climate where the experiment was run (i.e., temperate, tropical, etc)	
<b>Location</b>	Named geographical location	
<b>LocationType</b>	Type of location:	<ul style="list-style-type: none"> <li>- <i>Organism</i> = where the study organism was collected/cultured from</li> <li>- <i>Experiment</i> = where the experiment was conducted (if Location of Organism is not stated)</li> <li>- <i>Author</i> = affiliation of corresponding author (if neither Location of Organism nor Experiment is stated)</li> </ul>
<b>LocationDate</b>	Date when organism was collected (not experiment conducted)	
<b>CoordinateType</b>	Coordinate Source (Article / Gazetteer)	
<b>Latitude</b>	Of Location	
<b>Longitude</b>	Of Location	
<b>TaxaPresent</b>	Whether one or two taxa are part of the trait measurement and definition. If only a single organism is involved (e.g., metabolic rate, heart rate), it is always listed as a trait for a consumer.	<ul style="list-style-type: none"> <li>- <i>consumer</i> = trait involves a single organism;</li> <li>- <i>consumer-resource</i> = trait involves two organisms</li> </ul>
<b>TraitMotivation</b>	Type of motivation of the trait (e.g., positive, negative, voluntary, neutral)	
<b>TraitOrg</b>	Level of biological organization of the trait:	<ul style="list-style-type: none"> <li>- <i>internal</i> = processes internal to the organism;</li> <li>- <i>individual</i> = processes at the level of individual organisms that include mechanical interactions with the external environment;</li> <li>- <i>population</i> = processes for a group of conspecific individuals;</li> <li>- <i>interaction</i> = processes involving interaction between two or more species</li> </ul>

<b>TraitRelevantTaxa</b>	Taxa (i.e., Consumer or Resource) for which the trait was measured	<i>consumer; resource</i>
<b>Consumer</b>	Binomial name of the consumer or lowest taxonomic identity (e.g. binomial name including sub-species or race name)	
<b>ConCommon</b>	Common name of Consumer	
<b>ConWholePart</b>	Specify here if the thermal response corresponds to a whole organism (a tree	
<b>ConWholePartType</b>	Type of consumer	<i>whole; part; tissue</i>
<b>ConIDLevel</b>	Minimum taxonomic level to which the consumer was identified (e.g. Kingdo, Class, Species, etc)	
<b>ConKingdom</b>	Kingdom of Consumer	
<b>ConPhylum</b>	Phylum of Consumer	
<b>ConClass</b>	Class of Consumer	
<b>ConOrder</b>	Order of Consumer	
<b>ConFamily</b>	Family of Consumer	
<b>ConGenus</b>	Genus of Consumer	
<b>ConSpecies</b>	Species of Consumer	
<b>ConTrophic</b>	Broad trophic group of consumer, as determined by published literature and expert opinion.	<ul style="list-style-type: none"> <li>- <i>carnivore</i>;</li> <li>- <i>detritivore</i>;</li> <li>- <i>herbivore</i>;</li> <li>- <i>omnivore</i>;</li> <li>- <i>producer</i>;</li> <li>- <i>self</i> = energy self-supplied (e.g., pupae, egg)</li> </ul>
<b>ConTrophicGroup_citation</b>	Source of information where ConTrophic was taken.	
<b>ConType</b>	Type of consumer.	<ul style="list-style-type: none"> <li>- <i>alive</i> = organism alive when trait performance measured;</li> <li>- <i>dead</i> = organism dead when trait performance measured;</li> <li>- <i>artificial</i> = 'organism' simulated by a physical stimulus (e.g., predator model, prodding, gravity)</li> </ul>
<b>ConStage</b>	Life stage of consumer, and sex in parenthesis when available (e.g. Egg, adult, larvae, etc)	
<b>ConThermy</b>	Thermy of consumer.	<i>ectotherm; endotherm</i>

<b>ConTemp</b>	Body temperature of Consumer (if this is missing, Ambient temperature) during the experiment	
<b>ConTempUnit</b>	Units of ConTemp	
<b>ConTempMethod</b>	Method of determining ConTemp can be:	<ul style="list-style-type: none"> <li>- <i>Direct</i> = measured directly from within or on the organism</li> <li>- <i>Inferred (ambient)</i> = estimated from known ambient temperature</li> <li>- <i>Inferred (endotherm)</i> = body temperature relatively constant and estimated from published literature</li> </ul>
<b>ConLabGrowthTemp</b>	Temperature at which Consumer was grown in the lab, prior to the start of the experiment (this may be different from ConAccTemp)	
<b>ConLabGrowthTempUnit</b>	Units of ConLabGrowthTemp	
<b>ConLabGrowthDur</b>	Duration for which Consumer was exposed to ConLabGrowthTemp	
<b>ConLabGrowthDurUnit</b>	Units of ConLabGrowthTempDur	
<b>ConAcc</b>	Broad details of how consumer was acclimated (i.e., prior to placement at test temperature), such as if it was captive bred or wild caught.	<ul style="list-style-type: none"> <li>- <i>field collected</i>;</li> <li>- <i>field collected (at acclimation temperature)</i> = see ConAccTemp;</li> <li>- <i>field collected (at test temperature)</i> = see AmbientTemp or ConTemp;</li> <li>- <i>captive bred</i>;</li> <li>- <i>captive bred (at acclimation temperature)</i> = see ConAccTemp;</li> <li>- <i>captive bred (at test temperature)</i> = see AmbientTemp or ConTemp</li> </ul>
<b>ConAccTemp</b>	Temperature (°C) at which consumer was kept prior to placement at test temperature (acclimated). Average temperature used when a temperature range reported in the original study. This is distinct from ConLabGrowthTemp.	
<b>ConAccTempNotes</b>	Notes for ConAccTemp, such as temperature range.	

<b>ConAccTime</b>	Time consumer spent at ConAccTemp. Every attempt was made to obtain a value, and where necessary text descriptions of time were converted to second (e.g., “few” or “several” = 3, “couple” = 2).	
<b>ConAccTimeNotes</b>	Notes for ConAccTimeNotes (e.g., entire life)	
<b>ConAccTimeUnit</b>	Units of ConAccTime.	<i>seconds</i>
<b>ConOrigTemp</b>	Temperature (°C) at which consumer was originally adapted. Average temperature given when a temperature range reported in the original study.	
<b>ConOrigTempNotes</b>	Notes for ConOrigTemp, such as temperature range.	
<b>ConOrigTime</b>	Time consumer spent at ConOrigTemp. Every attempt was made to obtain a value, and where necessary text descriptions of time were converted to second (e.g., “few” or “several” = 3, “couple” = 2).	
<b>ConOrigTimeNotes</b>	Notes for ConOrigTempTime (e.g., entire life)	
<b>ConOrigTimeUnit</b>	Units of ConOrigTime.	<i>seconds</i>
<b>ConSize</b>	Size of Consumer	
<b>ConSizeUnit</b>	Units of ConSize	
<b>ConSizeType</b>	What ConSize is referring to (e.g. "Ash-free dry weight")	
<b>ConSizeSI</b>	Size of Consumer in SI units	
<b>ConSizeUnitSI</b>	Units of SizeSI	
<b>ConDenValue</b>	Value of Consumer density (e.g. 10 individuals per square meter)	
<b>ConDenUnit</b>	Units of ConDenValue (e.g. individual m <sup>-2</sup> )	
<b>ConDenTypeSI</b>	Type of SI units of ConDenValueSI	<ul style="list-style-type: none"> <li>- <i>individual</i>;</li> <li>- <i>kilogram (dry body mass)</i>;</li> <li>- <i>kilogram (wet body mass)</i>;</li> <li>- <i>liter</i>;</li> <li>- <i>to satiation</i> = resource density above what consumer could fully consume (relevant for ResDenTypeSI, see below)</li> </ul>
<b>ConDenValueSI</b>	Same as ConDenValue in SI units	

<b>ConDenUnitSI</b>	Units for ConDenValueSI	<ul style="list-style-type: none"> <li>- <i>arena</i>;</li> <li>- <i>square meter</i>;</li> <li>- <i>cubic meter</i>;</li> <li>- <i>meter</i> = when only length of arena was stated (see ArenaUnitSI)</li> </ul>
<b>ConMassValueSI</b>	SI value of consumer mass as obtained from original source or estimated from other published literature (see Dell et al. (2011) for further details).	
<b>ConMassUnitSI</b>	SI unit of ConMassValueSI.	<ul style="list-style-type: none"> <li>- <i>kilogram (wet body mass)</i> = wet mass of entire body of consumer;</li> <li>- <i>kilogram (wet tissue mass)</i> = wet mass of tissue of consumer (e.g., excluding shell for gastropods)</li> </ul>
<b>ConTimeTestTemp</b>	Time consumer spent at test temperature (see AmbientTemp, ConTemp, or ResTemp) prior to start of the experiment over which trait performance was measured (i.e., ObsTime is zero).	
<b>ConTimeTestTempNotes</b>	Notes for ConTimeTestTemp, such as range of time.	
<b>ConTimeTestTempUnit</b>	Units of ConTimeTestTemp.	
<b>ConTimeTestTempSI</b>	Same as ConTimeTestTemp using SI units	
<b>ConTimeTestTempUnitSI</b>	Units of ConTimeTestTempSI	<i>seconds</i>
<b>Resource</b>	Binomial name of the resource or lowest taxonomic identity (e.g. binomial name including sub-species or race name), Or if the resource is a non-living resource, its name (e.g. "Co2" "N2")	
<b>ResCommon</b>	Common name of Resource	
<b>ResIDLevel</b>	Minimum taxonomic level to which the resource was identified (e.g. Kingdo, Class, Species, etc)	
<b>ResKingdom</b>	Kingdom of Resource	
<b>ResPhylum</b>	Phylum of Resource	
<b>ResClass</b>	Class of Resource	
<b>ResOrder</b>	Order of Resource	
<b>ResFamily</b>	Family of Resource	
<b>ResGenus</b>	Genus of Resource	

<b>ResSpecies</b>	Species of Resource	
<b>ResTrophic</b>	Broad trophic group of resource, as determined by published literature and expert opinion.	<ul style="list-style-type: none"> <li>- <i>carnivore</i>;</li> <li>- <i>detritivore</i>;</li> <li>- <i>herbivore</i>;</li> <li>- <i>omnivore</i>;</li> <li>- <i>producer</i>;</li> <li>- <i>self</i> = energy self-supplied (e.g., pupae, egg)</li> </ul>
<b>ResTrophicGroup_citation</b>	Source of information where ResTrophic was taken.	
<b>ResType</b>	Type of resource.	<ul style="list-style-type: none"> <li>- <i>alive</i> = organism alive when trait performance measured;</li> <li>- <i>dead</i> = organism dead when trait performance measured;</li> <li>- <i>artificial</i> = 'organism' simulated by a physical stimulus (e.g., predator model, prodding, gravity)</li> </ul>
<b>ResStage</b>	Life stage of resource, and sex in parenthesis when available (e.g. Egg, adult, larvae, etc)	
<b>ResThermy</b>	Thermy of resource.	<i>ectotherm; endotherm</i>
<b>ResTemp</b>	Body temperature of Ronsumer (if this is missing, Ambient temperature) during the experiment	
<b>ResTempUnit</b>	Units of ResTemp	
<b>ResTempMethod</b>	Method of determining ResTemp can be:	<ul style="list-style-type: none"> <li>- <i>Direct</i> = measured directly from within or on the organism</li> <li>- <i>Inferred (ambient)</i> = estimated from known ambient temperature</li> <li>- <i>Inferred (endotherm)</i> = body temperature relatively constant and estimated from published literature</li> </ul>
<b>ResLabGrowthTemp</b>	Temperature at which Resourechr was grown in the lab, prior to the start of the experiment (this may be different from ResAccTemp)	
<b>ResLabGrowthTempUnit</b>	Units of ResLabGrowthTemp	
<b>ResLabGrowthDur</b>	Duration for which Resource was exposed to ResLabGrowthTemp	



<b>ResLabGrowthDurUnit</b>	Units of ResLabGrowthTempDur	
<b>ResAcc</b>	Broad details of how resource was acclimated (i.e., prior to placement at test temperature), such as if it was captive bred or wild caught.	<ul style="list-style-type: none"> <li>- <i>field collected</i>;</li> <li>- <i>field collected (at acclimation temperature)</i> = see ResAccTemp;</li> <li>- <i>field collected (at test temperature)</i> = see AmbientTemp or ResTemp;</li> <li>- <i>captive bred</i>;</li> <li>- <i>captive bred (at acclimation temperature)</i> = see ResAccTemp;</li> <li>- <i>captive bred (at test temperature)</i> = see AmbientTemp or ResTemp</li> </ul>
<b>ResAccTemp</b>	Temperature (°C) at which resource was kept prior to placement at test temperature (acclimated). Average temperature used when a temperature range reported in the original study. This is distinct from ResLabGrowthTemp.	
<b>ResAccTempNotes</b>	Notes for ResAccTemp, such as temperature range.	
<b>ResAccTime</b>	Time resource spent at ResAccTemp. Every attempt was made to obtain a value, and where necessary text descriptions of time were converted to second (e.g., “few” or “several” = 3, “couple” = 2).	
<b>ResAccTimeNotes</b>	Notes for ResOrigTempTime (e.g., entire life)	
<b>ResAccTimeUnit</b>	Units of ResAccTemp.	seconds
<b>ResOrigTemp</b>	Temperature (°C) at which resource was originally adapted. Average temperature given when a temperature range reported in the original study.	
<b>ResOrigTempNotes</b>	Notes for ResnOrigTemp, such as temperature range.	
<b>ResOrigTime</b>	Temperature (°C) at which resource was at before lab growth. Average temperature given when a temperature range reported in the original study.	
<b>ResOrigTimeNotes</b>	Notes for ResOrigTempTime (e.g., entire life)	
<b>ResOrigTimeUnit</b>	Units of ResOrigTime.	seconds
<b>ResSize</b>	Size of resource	
<b>ResSizeUnit</b>	Units of ResSize	
<b>ResSizeType</b>	What ResSize is referring to (e.g. "Ash-free dry weight")	

<b>ResSizeSI</b>	Size of resource in SI units	
<b>ResSizeUnitSI</b>	Units of SizeSI	
<b>ResDenValue</b>	Value of resource density (e.g. 10 individuals per square meter)	
<b>ResDenUnit</b>	Units of ResDenValue (e.g. individual m <sup>-2</sup> )	
<b>ResDenTypeSI</b>	Type of SI units of ResDenValueSI	<ul style="list-style-type: none"> <li>- <i>individual</i>;</li> <li>- <i>kilogram (dry body mass)</i>;</li> <li>- <i>kilogram (wet body mass)</i>;</li> <li>- <i>liter</i>;</li> <li>- <i>to satiation</i> = resource density above what resource could fully Resume (relevant for ResDenTypeSI, see below)</li> </ul>
<b>ResDenValueSI</b>	Same as ResDenValue in SI units	
<b>ResDenUnitSI</b>	Units for ResDenValueSI	<i>arena; square meter; cubic meter; meter</i> = when only length of arena was stated (see ArenaUnitSI)
<b>ResMassValueSI</b>	SI value of resource mass as obtained from original source or estimated from other published literature (see Dell et al. (2011) for further details).	
<b>ResMassUnitSI</b>	SI unit of ResMassValueSI.	<ul style="list-style-type: none"> <li>- <i>kilogram (wet body mass)</i> = wet mass of entire body of resource;</li> <li>- <i>kilogram (wet tissue mass)</i> = wet mass of tissue of resource (e.g., excluding shell for gastropods)</li> </ul>
<b>ResTimeTestTemp</b>	Time resource spent at test temperature (see AmbientTemp, ResTemp, or ResTemp) prior to start of the experiment over which trait performance was measured (i.e., ObsTime is zero).	
<b>ResTimeTestTempNotes</b>	Notes for ResTimeTestTemp, such as range of time.	
<b>ResTimeTestTempUnit</b>	Units of ResTimeTestTemp.	
<b>ResTimeTestTempSI</b>	Same as ResTimeTestTemp using SI units	
<b>ResTimeTestTempUnitSI</b>	Units of ResTimeTestTempSI	<i>seconds</i>
<b>PhysicalProcess</b>	First classification of physical process (ej. Foraging Time, Angular Velocity, Metabolic Efficiency)	
<b>PhysicalProcess_1</b>	Second level of physical process classification (ej. Interaction, Life history, Reproduction)	

<b>PhysicalProcess_2</b>	Third level of physical process classification (ej. Population density, Life history, body movement)	
<b>CitationID</b>	Full citation of source (typically a published paper)	
<b>Citation</b>	DOI number for each citation	
<b>DOI</b>	Is the data published or not	<i>True/False</i>
<b>Published</b>	Figure or table from Citation from which data were extracted	
<b>FigureTable</b>	Clarification notes of data-logger (e.g. are the different thermal responses collected from a particular study)	
<b>Notes</b>	For treatments not captured by this data sheet?)	
<b>SubmittedBy</b>	The person who input the data into the datasheet	
<b>ContributorEmail</b>	E-mail address of the person who input the data into the datasheet	
<b>ManualCheck</b>	<i>The entry has been manually checked or not</i>	<i>Yes/No</i>