

Bioschemas

Structured data markup for life sciences with Schema.org

Roberto Preste



Data infrastructure for Europe's life-science research:



Data



Interoperability



Tools



Compute



Training



Marine metagenomics



Crop and forest plants



Human data



Rare diseases





Research and information sharing

Information is exposed in the Internet through web pages

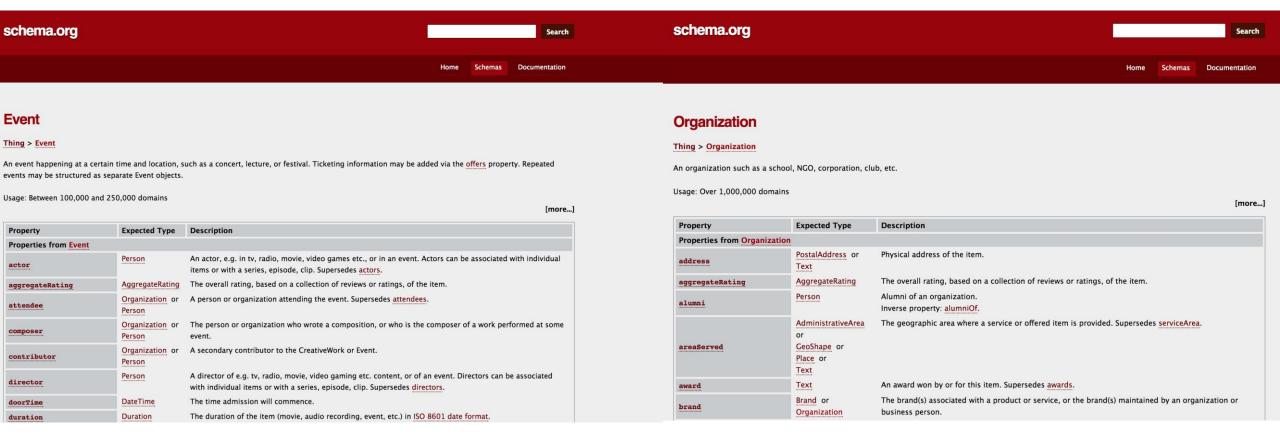


Great for common users

Bad for automated and programmatical data collection



Schema.org

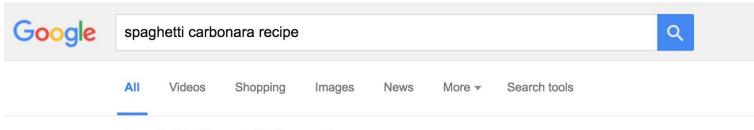


Community and collaborative initiative to integrate structured data markup on web pages, email messages and on the Internet in general, without altering the presentation layer



Schema.org

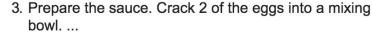
Allows search engines to easily access websites' content and provide more useful results based on their underlying metadata

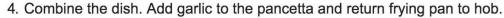


About 2,460,000 results (0.42 seconds)

Method

- 1. Cook the spaghetti. Bring a large, deep pan of salted water to the boil. ...
- 2. Fry the pancetta. Meanwhile, heat the oil in a frying pan.





Learn to make spaghetti carbonara | BBC Good Food www.bbcgoodfood.com/recipes/2876/learn-to-make-spaghetti-carbonara

About this result . Feedback

Ultimate spaghetti carbonara | BBC Good Food

www.bbcgoodfood.com/recipes/1052/ultimate-spaghetti-carbonara •

*** Rating: 4.6 - 126 votes - 35 mins - 655 cal

Discover how to make the perfect **Spaghetti carbonara** - with a little help from Gennaro Contaldo ... While the spaghetti is **cooking**, fry the pancetta with the garlic.

Learn to make spaghetti ... • The ultimate makeover • Courgette carbonara

Learn to make spaghetti carbonara | BBC Good Food



www.bbcgoodfood.com/recipes/2876/learn-to-make-spaghetti-carbonara •

*** Rating: 4.2 - 28 votes - 20 mins - 734 cal

Method. Cook the spaghetti. Bring a large, deep pan of salted water to the boil. Fry the



Adopted by > 10 million websites



...but still not this popular in the life science community!



Bioschemas





Home

Groups

Specifications

Community

- Biosoftware description using bio.tools and schema.org (NETTAB 2016 Workshop). October 25-26, 2016, Rome, Italy
- Bioschemas hands-on meeting. November 8-9, 2016, Harpenden, Hertfordshire, UK

Bioschemas aims to improve data interoperability in life sciences. It does this by encouraging people in life science to use schema.org markup, so that their websites and services contain consistently structured information. This structured information then makes it easier to discover, collate and analyse distributed data. The main outcome of Bioschemas is a collection of specifications that provide guidelines to facilitate a more consistent adoption of schema.org markup within the life sciences. Bioschemas operates as an open community initiative. You are welcome to join a community group.

Bioschemas aims to apply the Schema.org markup in life science Better discoverability and usability of structured data



Bioschemas: organization

Open community initiative:

















Organized in groups:

Group	Email	Email archive	Tasks/Issues	Shared Folder
Event	events@bioschemas.org	P	P	P
Training Material	training-material@bioschemas.org	P	P	P
Organization	organization@bioschemas.org	P	P	P
Person	person@bioschemas.org	P	P	P
Standard	standard@bioschemas.org	P	P	P
Data	data@bioschemas.org	P	P	P
Tool	tool@bioschemas.org	P	P	P
Data Repositories	datarepositories@bioschemas.org	P	P	P
Community	community@bioschemas.org	P	P	P
Commons	commons@bioschemas.org	P	P	P



Content types



Generic content types

E.g.: training materials, datasets

Biological content types

E.g.: pathways, proteins

Can be used in many disciplines, besides life science

Are domain-specific

In collaboration with

health-lifesci.schema.org

to integrate and extend

Schema.org types and properties



- Introduction
- Data model
- Content guidelines
- Cardinality and controlled vocabularies
- New properties
- Examples

Overview of the problem, goal of the specification, technologies used to implement the solution

Problem statement

Conferences, workshops, meetings and events in general play a very important role in knowledge sharing and acquisition of new skills. Though there are several technological solutions for event sharing, events in life sciences are still not described in a consistent manner that would make them easy to discover, exchange or compare.

Because there is no single standard to follow, the dissemination, discovery and aggregation of events is not effective. As a result, the advertisement of an event in third party websites normally requires manual curation to shape the content to each provider's requirements.

Proposed solution

Rationale

In the development of the Life Science Events standard we have considered the following design goals.

Consensus

Many organisations and repositories providing events already exist. It is important this standard takes into account their experience and contribution.

Adoption

Many organisations already have a website or system providing information about events. They will not be willing to change their methods unless there is a clear benefit and a low barrier for adoption.

Reuse

There are existing formats and technologies suitable to represent at least some information about events. This specification will avoid reinvention and seek to extend existing standards.



- Introduction
- Data model
- Content guidelines
- Cardinality and controlled vocabularies
- New properties
- Examples

Describes how to elaborate and implement more specific guidelines into a Schema.org skeleton

Data model

The data model proposed involves:

- 1. Adopting the Schema.org Event type, and extending it with additional properties. Schema.org already has a way of describing events, through its Event type. In this document we suggest using this type to describe life science events, but we also suggest new properties for this type, so that event descriptions can be more accurate and useful in life sciences. If the community agrees, these additional properties will be put forward for adoption to Schema.org.
- 2. Adopting a standard way of using the Schema.org Event type. Many properties in the Schema.org Event type are loosely defined, and we propose guidelines on how to use them so that they are more specific and consistent. These guidelines include concepts not supported by Schema.org, such as cardinality, controlled vocabularies and content guidelines (minimum, optional and recommended fields). For example, we suggest the use of a controlled vocabulary based on the EDAM ontology for the 'topics' property. These recommendations will not be part of the Schema.org Event type, but are proposed as best practices in using that type in life science.



- Introduction
- Data model
- Content guidelines
- Cardinality and controlled vocabularies
- New properties
- Examples

Defines a set of Minimum fields that are mandatory, Recommended fields for optimal discovery and integration, and Optional fields to enhance the user experience

Bioschemas

Event type definition

Data fields

Legend:

CN: Cardinality (one, many)

CG: Content Guideline (M: minimum; R: recommended; O: optional)

CV: Controlled Vocabulary

Property	Expected Type	Description		CG	cv
Existing properties in schema.org/Event					
aggregateRating	AggregateRati ng	The overall rating, based on a collection of reviews or ratings, of the item.			
attendee	Organization or Person	A person or organization attending the event. Supersedes attendees.			
doorTime	DateTime	The time admission will commence.			
duration	Duration	The duration of the item (movie, audio recording, event, etc.) in ISO 8601 date format.			
endDate	Date	The end date and time of the item (in ISO 8601 date format).		М	
eventStatus	EventStatusTy pe	An eventStatus of an event represents its status; particularly useful when an event is cancelled or rescheduled.			x
offers	Offer	An offer to provide this item—for example, an offer to sell a product, rent the DVD of a movie, or give away tickets to an event. Can use eligibleCustomerType and eligibleDuration properties to express any special offers.	One	0	
organizer	Organization or Person	An organizer of an Event.		R	
performer	Organization or Person	A performer at the event—for example, a presenter, musician, musical group or actor. Supersedesperformers.		0	

- Introduction
- Data model
- Content guidelines
- Cardinality and controlled vocabularies
- New properties
- Examples

Define the number of times a property can be instantiated and specific terms that need to be used

Event type definition

Data fields

Legend:

CN: Cardinality (one, many)

CG: Content Guideline (M: minimum; R: recommended; O: optional)

CV: Controlled Vocabulary

	ocabalal y				
Property	Expected Type	Description		CG	cv
Existing propertie	Existing properties in schema.org/Event				
aggregateRating	AggregateRati ng	The overall rating, based on a collection of reviews or ratings, of the item.			
attendee	Organization or Person	A person or organization attending the event. Supersedes attendees.			
doorTime	DateTime	The time admission will commence.			
duration	Duration	The duration of the item (movie, audio recording, event, etc.) in ISO 8601 date format.			
endDate	Date	The end date and time of the item (in ISO 8601 date format).		М	
eventStatus	EventStatusTy pe	An eventStatus of an event represents its status; particularly useful when an event is cancelled or rescheduled.			х
offers	Offer	An offer to provide this item—for example, an offer to sell a product, rent the DVD of a movie, or give away tickets to an event. Can use eligibleCustomerType and eligibleDuration properties to express any special offers.	One	0	
organizer	Organization or Person	An organizer of an Event.	Many	R	
performer	Organization or Person	A performer at the event—for example, a presenter, musician, musical group or actor. Supersedesperformers	Many	0	



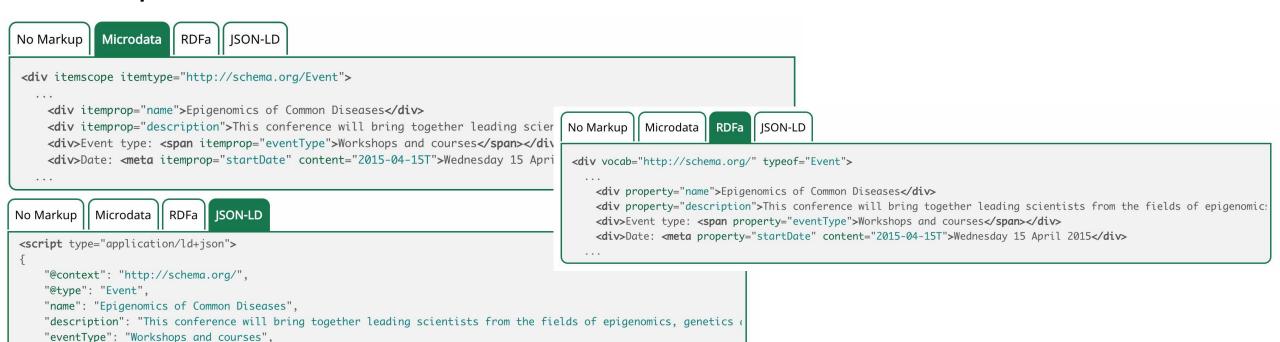
- Introduction
- Data model
- Content guidelines
- Cardinality and controlled vocabularies
- New properties
- Examples

New properties can be suggested in order to provide more accurate and useful type description in life science

Proposed new properties for the Event type						
eventId	Text	Unique Id for the event.		0		
prerequisite	Text	A list of prerequisites to be able to attend the event.		R		
accreditation	Organization or Text	Type of accreditation or organisation that accredits the event.		R		
eligibility	Text	Defines the type of eligibility to attend this event e.g first come first served.		R	x	
capacity	Integer	Available number of spaces	One	R		
contact	Organization or Person	Main point of contact that can be contacted for general queries. This would be an event organiser or an administrator.	Many	М		
attachment	URL	Any files or related websites which give more information about this event. e.g. flyers, third party sites handling tickets.		0		



- Introduction
- Data model
- Content guidelines
- Cardinality and controlled vocabularies
- New properties
- Examples



Microdata RDFa

Epigenomics of Common Diseases

Event type: Workshops and courses

Date: Wednesday 15 April 2015

JSON-LD

This conference will bring together leading scientists from the fields of epigenomics, genetics and bioinformatics.

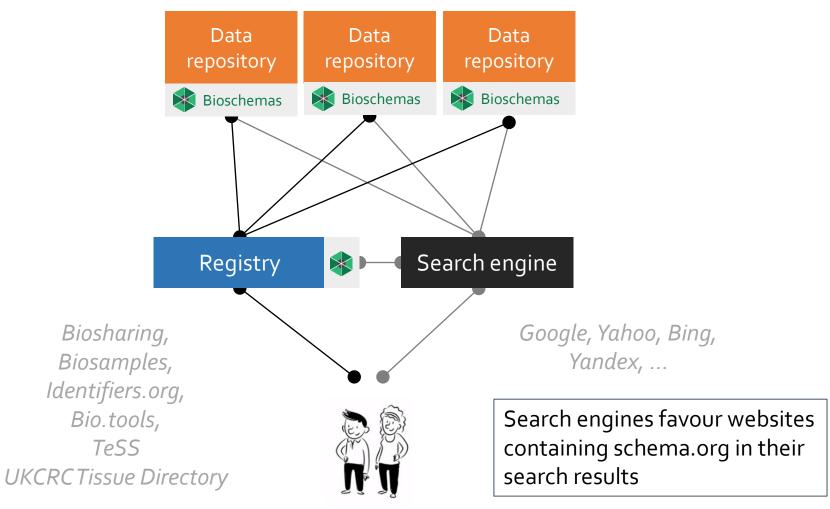
No Markup



Bioschemas: applications

Improving search results via Bioschemas

PDBe, UniProt, Pfam, ...





Bioschemas: applications

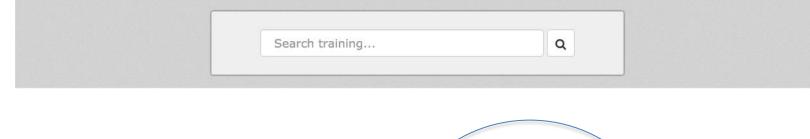
TeSS: ELIXIR's Training Portal

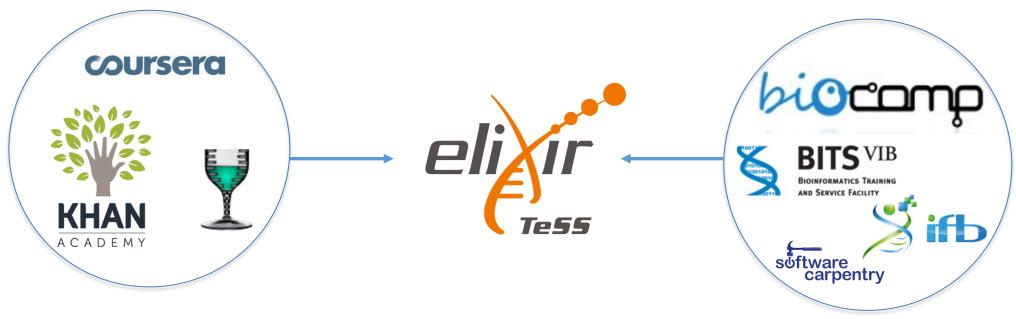
https://tess.elixir-uk.org



Welcome to TeSS: ELIXIR's Training Portal

Browsing, discovering and organising life sciences training resources, aggregated from ELIXIR nodes and 3rd-party providers.





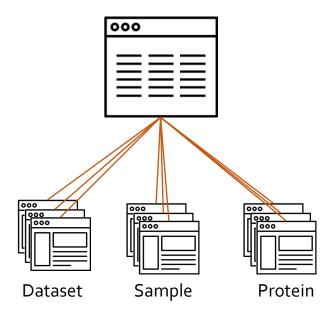


Bioschemas: validation

Discovery, Registration and Validation

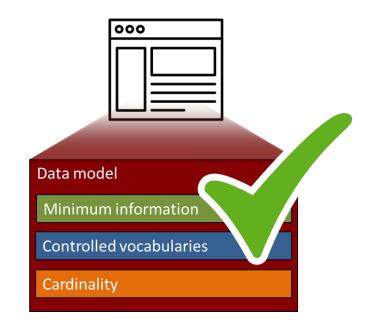
DISCOVERY

Provide a simple registry where providers or consumers can describe of sites providing content in schema.org compliant with Bioschemas



VALIDATION

Provide a GUI to validate Bioschemas compliant websites and Validate data repositories adopting Bioschemas





Bioschemas: validation



Home Registry Test Website Submit Website

Test And Submit A Website

With the Submit tool you can test a specific URL for its compliance with the Bioschemas specifications and submit it to the Registry for public consultation. You car choose to test all the children links for that page (links with the same root URL) as well as scraping your URL for one specific Bioschemas type (Event, Organization Person, Training Material).

PerspectiveDrawings http://d.lib.ncsu.edu/collections/catalog/mc00383-001-ff0004-001001_0004 Bioschemas type: CreativeWork (9 entries) Guideline Minimum property Recomm. property

Website URL:

,

Website Name:

If you choose not to provide a name for the URL you are testing, the title field content from the tested website will be used as its nam

Validate children links too

Scrape a specific Bioschemas type: All Types

Mail notification when the process is done example@domain.com

Validate!

Minimum and Recommended Properties

Show properties description

Compliance -	Property ▼	Type ▼	Guideline ▼	CV→	Cardinality 🕶
93%	<u>url</u>	URL	Recommended	No	Many
100%	<u>name</u>	Text	Minimum	No	One
0%	<u>license</u>	URL or CreativeWork	Recommended	Yes	One
47%	<u>keywords</u>	Text	Recommended	No	Many
13%	<u>genre</u>	Text or URL	Minimum	Yes	Many
40%	description	Text	Recommended	No	One
7%	dateModified	Date or DateTime	Minimum	No	One
0%	contributor	Person or Organization	Recommended	No	Many
0%	author	Person or Organization	Recommended	No	Many
0%	audience	Audience	Minimum	Yes	Many
7%	about	Thing	Minimum	No	One
0%	level	Text	Recommended	Yes	One



Upcoming activities

Implementation Study Objectives

Life sciences
Content Types

Schema.org content types for life science Data

- Data repository, Dataset
- Samples, Protein annotations
- Phenotype annotations



Discovery and validation

- Publication of metadata
- Automated integration of metadata in specialised registries

Discovery and validation of Bioschemas entries

Community
Support and
Promotion

Support community and adoption
Alignment of technical activities
Working group within ELIXIR
Collaboration between ELIXIR and BD2K
Test benefits and issues

Hands-on workshops Small delivery team



Upcoming activities

• **Bioschemas AGM**, 8th-9th November, Rothamsted (UK) Details on https://goo.gl/hu7uYK

More content types for life science in development:

Tools

Data repository

Dataset

Sample

Phenotype

Protein annotations



Acknowledgements

Bioschemas community

Group chairs

- Community
 - Premysl Velek
- Event
 - Martin Cook
- Training materials
 - Aleksandra Nenadic & Gabriella Rustici

- Organization
 - Richard Holland & Rafael C Jimenez
- Person
 - Niall Beard
- Standard
 - A Gonzalez-Beltran & P McQuilton

Organization representatives

- ELIXIR
 - Premysl Velek
- Pistoia Alliance
 - Richard Holland
- GOBLET

Terry Atwood

Bioschemas

- BBMRI
 - Michaela Mayrhofer
- TeSS
 - Niall Beard
- BioSharing
 - SA Sansone, A Gonzalez-Beltran, P McQuilton, P Rocca-Serra
- NIH BD2K bioCADDIE

SA Sansone, A Gonzalez-Beltran, Jeff Grethe

Contributors

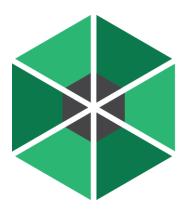
- Aleksandra Nenadic
- Adam Hospital
- Gabriella Rustici
- Carlos Horro
- Martin Cook
- Niall Beard
- Rafael C Jimenez
- Andy Jenkinson
- Manuel Corpas
- Roberto Preste
- Richard Holland
- Alejandra Gonzalez-Beltran
- Andrew Lonie
- Carole Goble
- Peter McQuilton
- Premysl Velek
- Ian Dunlop
- Jef Grethe

- Milo Thurston
- Niklas Blomberg
- Yasset Perez-Riverol
- Sarala Wimalaratne
- Nick Juty
- Jose Luis Ambite
- Brane Leskošek
- Celia van Gelder
- Christa Janko
- Christine Staiger
- Dan Brickley
- Daniel Faria
- Dmitry Repchevsky
- Daniel Sobral
- Daniel Vaughan
- Ian Fore
- Frederik Coppens
- Josep Gelpi
- ChuQiao Gong

- Hedi Peterson
- Hervé Ménager
- Nina Hrtonova
- Isabelle Perseil
- Jaap Heringa
- Jon Ison
- John Hancock
- Simon Jupp
- John D. Van Horn
- Ivana Krenkova
- Laura Furlong
- Morris Swertz
- Mateusz Kuzak
- Mario Alberich
- Mark Thompson
- Maria Martin
- Mikael Borg
- Montserrat Gonzàlez
- Norman Morrison

- Nùria Queralt-Rosinach
- Olivier Sallou
- Robert Pergl
- Pedro Fernandes
- Pierre Larmande
- Rob Finn
- Renzo Kottmann
- Rodrigo Lopez
- Sameer Velankar
- Sara Light
- Carol Shreffler
- Silvano Squizzato
- Susanna Sansone
- Tony Burdett
- Terry Atwood
- Cath Brooksbank
- Luc Deltombe
- Michaela Mayrhofer
- Philippe Rocca-Serra





Bioschemas

Thank you!

all@bioschemas.org

http://bioschemas.org

