# Machine-Actionability and Interoperability

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#### Collective Motion of Humans in Mosh and Circle Pits at Heavy Metal Concerts

Jesse L. Silverberg,\* Matthew Bierbaum, James P. Sethna, and Itai Cohen

Department of Physics and Laboratory of Atomic and Solid State Physics, Cornell University, Ithaca, New York 14853, USA (Received 13 February 2013; published 29 May 2013)

Human collective behavior can vary from calm to panicked depending on social context. Using videos publicly available online, we study the highly energized collective motion of attendees at heavy metal concerts. We find these extreme social gatherings generate similarly extreme behaviors: a disordered gaslike state called a *mosh pit* and an ordered vortexlike state called a *circle pit*. Both phenomena are reproduced in flocking simulations demonstrating that human collective behavior is consistent with the predictions of simplified models.

DOI: 10.1103/PhysRevLett.110.228701

PACS numbers: 89.65.Ef, 47.32.-y, 87.15.Zg, 87.23.Cc

Human collective behaviors vary considerably with social context. For example, lane formation in pedestrian traffic [1], jamming during escape panic [2], and Mexican waves at sporting events [3] are emergent phenomena that have been observed in specific social settings. Here, we study large crowds  $(10^2-10^5 \text{ attendees})$  of people under the extreme conditions typically found at heavy metal concerts [4]. Often resulting in injuries [5], the collective mood is influenced by the combination of loud (130 dB [6]), fast

Fig. 2, the decay length was  $0.39 \pm 0.03$  m, which is approximately human shoulder width. Taken together, these findings offer strong support for the analogy between mosh pits and gases. As a further check, we examined the 2D speed distribution. Previous observations of human pedestrian traffic and escape panic led us to expect a broad distribution not well described by simple analytic expressions [2,13]. However, the measured speed distribution in mosh pits was well fit by the 2D Maxwell-Boltzmann

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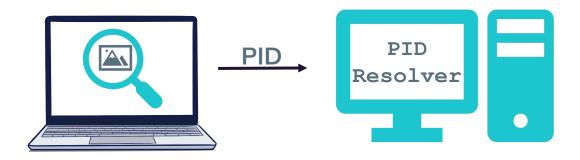
Fig. 2, the decay length w approximately human shou these findings offer strong su mosh pits and gases. As a fu 2D speed distribution. Prev pedestrian traffic and escape distribution not well desc



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# **Client – PID Resolver Interaction**





# PID Record Request



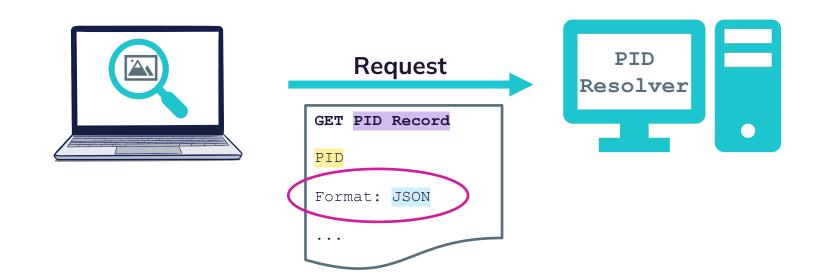


Request	PID Resolver	
GET PID Record		
PID		
Format: JSON		

I 6

# PID Record Request





# PID Record Request





# <text>



# **PID Record Schemas**



```
Crossref Metadata Input Schema v. 5.3.1
```

This is the latest iteration of the Crossref Metadata Input Schema, used to register metadata records with Crossref. This schema was originally developed for Crossref by Inera (http://www.in

As of version 4.5.0 all changes are noted in the release notes for the schema repository in GitLab: https://gitlab.com/crossref/schema/-/releases

```
Prior updates are noted as comments within previous schema versions. -->
 <!-- include common schema -->
 <xsd:include schemaLocation="common5.3.1.xsd"/>
 <!-- imported schema -->
 <xsd:import namespace="http://www.ncbi.nlm.nih.gov/JATS1" schemaLocation="JATS-journalpublishing1-3d2-mathml3.xsd"/>
 <xsd:import namespace="http://www.w3.org/1998/Math/MathML" schemaLocation="http://www.w3.org/Math/XMLSchema/mathml3.xsd"/>
 <xsd:import namespace="http://www.crossref.org/fundref.xsd" schemaLocation="fundref.xsd"/>
 <xsd:import namespace="http://www.crossref.org/clinicaltrials.xsd" schemaLocation="clinicaltrials.xsd"/>
 <xsd:import namespace="http://www.crossref.org/AccessIndicators.xsd" schemaLocation="AccessIndicators.xsd"/>
 <xsd:import namespace="http://www.crossref.org/relations.xsd" schemaLocation="relations.xsd"/>
 <!-- *** Head elements (used when processing submission XML) -->
w<xsd:element name="doi batch">
 ▼<xsd:annotation>
    <xsd:documentation>Top level element for a metadata record submission. This element indicates the start and end of the XML file. The version number is fixed to the version of the schema.
  </xsd:annotation>
 w<xsd:complexType>
  ▼<xsd:sequence>
      <xsd:element ref="head"/>
      <xsd:element ref="body"/>
    </xsd:sequence>
    <xsd:attribute name="version" type="xsd:string" fixed="5.3.1"/>
  </xsd:complexType>
 </xsd:element>
v<xsd:element name="head">
 ▼<xsd:annotation>
    <xsd:documentation>Container for information related to the DOI batch submission. This element uniquely identifies the batch deposit to Crossref and contains information that will be user
    processing. </xsd:documentation>
  </xsd:annotation>
 w<xsd:complexType>
  ▼<xsd:sequence>
      <xsd:element ref="doi batch id"/>
      <xsd:element ref="timestamp"/>
      <xsd:element ref="depositor"/>
      <xsd:element ref="registrant"/>
    </xsd:sequence>
  </xsd:complexType>
 </xsd:element>
                                                                                                                                                                                           9
v<xsd:element name="body">
  - duad . annat at i ant
```

# **PID Record Schemas**

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<1-- include common schema -->
<xsd:include schemaLocation="common5.3.1.xsd"/>
<1-- imported schemaLocation="common5.3.1.xsd"/>
<1-- imported schema -->
<xsd:import namespace="http://www.ncbi.nlm.nih.gov/JATS1" schemaLocation="JATS-journalpublishing1-3d2-mathm13.xsd"/>
<xsd:import namespace="http://www.crossref.org/fundref.xsd" schemaLocation="fundref.xsd"/>
<xsd:import namespace="http://www.crossref.org/clinicaltrials.xsd" schemaLocation="clinicaltrials.xsd"/>
<xsd:import namespace="http://www.crossref.org/clinicaltrials.xsd" schemaLocation="clinicaltrials.xsd"/>
<xsd:import namespace="http://www.crossref.org/AccessIndicators.xsd" schemaLocation="clinicaltrials.xsd"/>
<<!-- \*\*\* Head elements (used when processing submission XML) -->
<xsd:element name="doi\_batch">
</->
</->

<xsd:documentation>Top level element for a metadata record submission. This element indicates the start and end of the XML file. The version number is fixed to the version of the schema.



<xsd:element n
v<xsd:annotati
<xsd:docume;
processing.
</xsd:annotat
v<xsd:complex1
v<xsd:eleme;
<xsd:eleme;
<xsd:eleme;
</xsd:eleme;
</xsd:seque;
</xsd:seque;
</xsd:complex2
</pre>

### PID Record Schemas

are also called

## **Kernel Information Profiles**

entifies the batch deposit to Crossref and contains information that will be used

# One Last Thing about PIDs... PIDs in Schemas, Vocabularies and Ontologies







#### 🖸 http://purl.obolibrary.org/obo/MI\_1159 🍺 Copy

-configurational pre-organization

A molecule whose binding or catalytic properties at one site are altered by allosteric post-translational modification or binding of an allosteric effector at a distinct site. An allosteric molecule is identified by referring to its participant id.

Search MI		Searc		
Exact match 🔲 Include obsolete terms 🗹 Include imported	d terms			
Tree Graph	_	Class Information created_by		
Lattribute name (139)	Show counts	orchard		
interaction attribute name (38)	Show obsolete terms			
-3d-r-factors	Show obsolete terms	creation_date		
-3d-resolution	Show all siblings	2012-06-07T12:55:09Z		
-affected interaction		has_obo_namespace		
agonist		PSI-MI		
-allosteric change in dynamics		In authorit		
-allosteric change in structure		in_subset		
-allosteric effector		PSI-MI_slim		
allosteric k-type response				
allosteric molecule		Class Relations		
allosteric post-translational modification				
allosteric v-type response		Subclass of		
allostery		<ul> <li>biological role</li> </ul>		
altered physicochemical compatibility		interaction attribute name		
-antagonist -author-confidence				
<ul> <li>autnor-confidence</li> <li>binding site hiding</li> </ul>		cooperative interaction		
caution				
comment				
complex-properties				
composite binding site formation				





*	en 🗸	JSON

#### Purl 🗹 http://purl.obolibrary.org/obo/MI\_1159 🍺 Copy A molecule whose binding or catalytic properties at one site are altered by allosteric post-translational modification or binding of an allosteric effector at a distinct site. An allosteric molecule is identified by referring to its participant id. () Search Search MI... Exact match Include obsolete terms ▼ Class Information < Graph T- Tree molecular interaction (1,454) created\_by Show counts attribute name (139) orchard interaction attribute name (38) Show obsolete terms creation\_date 3d-r-factors 3d-resolution 2012-06-07T12:55:09Z Show all siblings 3d-structure has\_obo\_namespace affected interaction PSI-MI agonist allosteric change in dynamics in\_subset -allosteric change in structure PSI-MI\_slim allosteric effector allosteric k-type response allosteric molecule Class Relations -allosteric post-translational modification allosteric v-type response Subclass of allostery biological role altered physicochemical compatibility • interaction attribute name antagonist author-confidence cooperative interaction binding site hiding caution comment complex-properties composite binding site formation configurational pre-organization





# **Persistent URLs** (**PURLs**) are a form of **PIDs**. A PURL Resolver redirects HTTP requests to the current URL of a resource.

Class Information

created\_by orchard

reation\_date 2012-06-07T12:55:09Z

has\_obo\_namespace PSI-MI

<mark>in\_subset</mark> PSI-MI slim

#### Class Relations

#### Subclass of

- biological role
- interaction attribute name
- cooperative interaction

attribute name (139)
 ▲ interaction attribute name (38)
 ▲ 3d-resolution
 3d-structure
 affected interaction
 agonist
 allosteric change in dynamics
 allosteric change in structure
 allosteric v-type response
 allosteric v-type response
 allostery
 altered physicochemical compatibility
 antagonist
 author-confidence
 binding site hiding
 caution
 complex-properties
 composite binding site formation

Show counts

Show obsolete tern

Show all siblings





Help

About

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👪 en 🗸	JSON

#### Purl 🗹 http://purl.obolibrary.org/obo/MI\_1159 📗 Copy A molecule whose binding or catalytic properties at one site are altered by allosteric post-translational modification or binding of an allosteric effector at a distinct site. An allosteric molecule is identified by referring to its participant id. () Search Search MI... Exact match Include obsolete terms ▼ Class Information < Graph T- Tree molecular interaction (1,454) created\_by Show counts attribute name (139) orchard interaction attribute name (38) Show obsolete terms creation\_date 3d-r-factors 3d-resolution 2012-06-07T12:55:09Z Show all siblings 3d-structure has\_obo\_namespace affected interaction PSI-MI agonist allosteric change in dynamics in\_subset -allosteric change in structure PSI-MI\_slim allosteric effector allosteric k-type response allosteric molecule Class Relations -allosteric post-translational modification allosteric v-type response Subclass of allostery biological role altered physicochemical compatibility • interaction attribute name antagonist author-confidence cooperative interaction binding site hiding caution comment complex-properties composite binding site formation configurational pre-organization



Home	Intro	Statistics	SPARQL	Ontobeep	Annotator	Tutorial	FAQs	References	Links	Contact	Acknowledge	News
	Molecular Interactions Controlled Vocabulary											
Keywords: Search terms												
Class: allost	eric m	olecule										
Term IRI:	http://	purl.obolib	rary.org/ob	o/MI_1159								
								steric post-trans [database_cro			or binding of an 18706817]	allosteric
nnotations												
<ul> <li>has_obo_namespace: PSI-MI</li> <li>http://www.geneontology.org/formats/obolnOwl#created_by: orchard</li> <li>http://www.geneontology.org/formats/obolnOwl#creation_date: 2012-06-07T12:55:09Z</li> <li>http://www.geneontology.org/formats/obolnOwl#id: MI:1159</li> <li>in_subset: http://purl.obolibrary.org/obo/mi#PSI-MI_slim</li> </ul>												
Class Hierarchy												
Thing + molecular interaction												

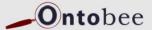
- + attribute name + interaction attribute name
  - figure legend
  - comment
  - function
  - <u>url</u>
  - disease
  - caution
  - pathway
  - author-confidence
  - inhibition
  - stimulant
  - agonist

  - more... allosteric molecule

Superclasses & Asserted Axioms

- interaction attribute name
- biological role
- <u>cooperative interaction</u>

Show RDF Source



Home	Intro	Statistics	SPARQL	Ontobeep	Annotator	Tutorial	FAQs	References	Links	Contact	Acknowledge	News	
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#### **Molecular Interactions Controlled Vocabulary**

Keywords:

Search terms

Class: allosteric molecule

#### Term IRI: http://purl.obolibrary.org/obo/MI\_1159

Definition: A molecule whose binding or catalytic properties at one site are altered by allosteric post-translational modification or binding of an allosteric effector at a distinct site. An allosteric molecule is identified by referring to its participant id. [database cross reference: PMID:18706817]

#### Annotations

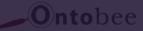
- has\_obo\_namespace: PSI-MI
- http://www.geneontology.org/formats/obolnOwl#created\_by: orchard
- http://www.geneontology.org/formats/obolnOwl#creation\_date: 2012-06-07T12:55:09Z
- http://www.geneontology.org/formats/obolnOwl#id: MI:1159
- in\_subset: http://purl.obolibrary.org/obo/mi#PSI-MI\_slim

#### **Class Hierarchy**

Thing
+ molecular interaction
+ attribute name
+ interaction attribute name
- figure legend
- comment
- function
- url
- disease
- caution
- <u>pathway</u>
- author-confidence
- inhibition
- <u>stimulant</u>
- <u>agonist</u>
more
- <u>allosteric molecule</u>
Superclasses & Asserted Axioms
ouperclasses a Asserted Axionis

- interaction attribute name
- biological role
- <u>cooperative interaction</u>

#### Show RDF Source



	Molecular Interactions Controlled Vocabulary											
Key	eywords: Search terms											
Clas	s: allost	eric mo	olecule									
	Term IRI: http://purl.obolibrary.org/obo/MI_1159 Definition: A molecule whose binding or catalytic properties at one site are altered by allosteric post-translational modification or binding of an allosteric effector at a distinct site. An allosteric molecule is identified by referring to its participant id. [database_cross_reference: PMID:18706817] Annotations											
<ul> <li>has_obo_namespace: PSI-MI</li> <li>http://www.geneontology.org/formats/obolnOwi#created_by: orchard</li> <li>http://www.geneontology.org/formats/obolnOwi#creation_date: 2012-06-07T12:55:09Z</li> <li>http://www.geneontology.org/formats/obolnOwi#id: MI:1159</li> <li>in_subset: http://purl.obolibrary.org/obo/mi#PSI-MI_slim</li> </ul>												

#### **Class Hierarchy**



**Internationalized Resource Identifier (IRI)** is extending the URI, allowing for a greater set of characters.





🗹 http://purl.obolibrary.org/obo/MI\_1159 🍺 Copy

A molecule whose binding or catalytic properties at one site are altered by allosteric post-translational modification or binding of an allosteric effector at a distinct site. An allosteric molecule is identified by referring to its participant id.

Search MI		Search
Exact match 📋 Include obsolete terms 🗹 Include imported terms		
Tree Graph		▼ Class Information
Putative self ML_0918	Legend Relationship Color Visibility Extended nodes (*) subClassOf Select/Deselect all List of extended nodes (*): allosteric molecule (MI_1159) biological role (MI_0500)	created_by orchard creation_date 2012-06-07T12:55:09Z has_obo_namespace PSI-MI in_subset PSI-MI_slim Class Relations
MI_0898 G B B C B C B		Subclass of • biological role • interaction attribute name • cooperative interaction
Create clusters Open all clusters Auto rearrange ( Hierarchical layout		

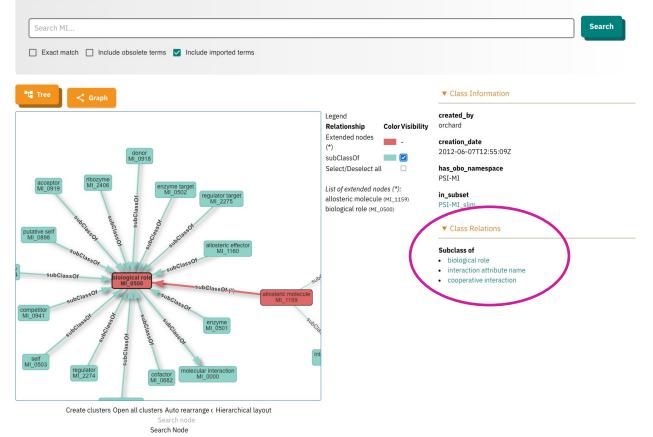
Search node Search Node





🖸 http://purl.obolibrary.org/obo/MI\_1159 🍺 Copy

A molecule whose binding or catalytic properties at one site are altered by allosteric post-translational modification or binding of an allosteric effector at a distinct site. An allosteric molecule is identified by referring to its participant id.

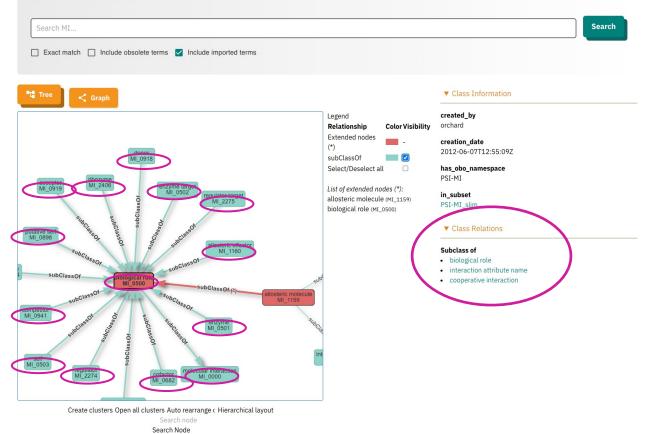






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A molecule whose binding or catalytic properties at one site are altered by allosteric post-translational modification or binding of an allosteric effector at a distinct site. An allosteric molecule is identified by referring to its participant id.



en 🗸 🛛 🛛



# The study of meaning, reference, or truth is called **Semantics**.

Illosteric effector at a distinct site. An allosteric molecule is

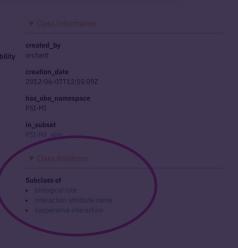
earch

📔 Exact match 🔄 Include obsolete terms 🛛 Include imported terms 🚽



Formally describing **concepts**, **relationships** between (meta)data entities, and **categories**, encodes semantics with the data.

This technology is known as the **Semantic Web.** 



Create clusters Open all clusters Auto rearrange c Hierarchical layout Search node The Semantic Web is an extension of the current Web in which information is given well-defined meaning, better enabling computers and people to work in cooperation.

Berners-Lee et al. (2001), The Semantic Web, Scientific American

image: http://images.wired.it/wp-content/uploads/2014/10/141449444

# Semantic Web Technologies











<!-- http://purl.obolibrary.org/obo/IAO\_0100001 -->

<!-- http://purl.obolibrary.org/obo/NCIT\_C42610 -->

<!-- http://purl.obolibrary.org/obo/NCIT\_C93495 -->

<owl:AnnotationProperty rdf:about="http://purl.obolibrary.org/obo/NCIT\_C03405">
 <obo:IA0\_0000115>The non-unique initials or abbreviated name used for identification. [Definition Source: NCI]</obo:IA0\_0000115>
 </rdfs:label xml:lang="en">acronym=</rdfs:label>
 </rdfs:label xml:label</rd>

<rdfs1subProperty0f rdf1resource="http://www.w3.org/2004/02/skos/core#altLabel"/>
</owl:AnnotationProperty>

<!-- http://purl.obolibrary.org/obo/R0\_0001900 -->

<owl:AnnotationProperty rdf:about="http://purl.obolibrary.org/obo/R0\_0001900"/>

<!-- http://purl.obolibrary.org/obo/R0\_0002259 -->

</owl:AnnotationProperty>

# "But this is another story and

shall be told another time.'

Michael Ende, The Neverending Story (1979)

# Questions?

## DISCLAIMER

#### This slide deck is part of the Lesson

#### <u>Fundamentals of Scientific Metadata:</u> <u>Why Context Matters</u>

published on The Carpentries Incubator.

#### Please cite this presentation as:

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image: https://c.pxhere.com/photos/35/f5/coffee\_notebook\_wooden\_backgr ound\_orange\_work\_table\_office-1222115.jpg!d