eHealth

Life in Scales and Data Sources

André Santanchè Laboratory of Information Systems – LIS Institute of Computing – UNICAMP August 2019



Life in Scales



Building Life

Francis Crick

- British molecular biologist, biophysicist, and neuroscientist
- Co-discoverer of the structure of the DNA molecule in 1953
- Nobel Prize in Physiology or Medicine

(Wikipedia, 2018)



Photo: Marc Lieberman - Siegel RM, Callaway EM: Francis Crick's Legacy for Neuroscience: Between the α and the Ω . PLoS Biol 2/12/2004: e419. https://dx.doi.org/10.1371/journal.pbio.0020419

Central Dogma of Molecular Biology

(Crick, 1970)

"The central dogma of molecular biology deals with the detailed residue-by-residue **transfer of sequential information**. It states that such information cannot be transferred back from protein to either protein or nucleic acid."

RNA

Phenotype

- Sets of organism observable characteristics
- Expression of organism's genotype interacting with the environment

Genotype to Phenotype



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Genotype to Phenotype







Genotype



Complexity: From Genes to Phenotypes



EQS eye decreased size cornea opaque lens fused_to corneal epithelium anterior chamber decreased size eye decreased size cornea opaque iris decreased size or absent retina malformed lens opaque anterior chamber decreased size

eye decreased size lens decreased size retina malformed

eye absent

(Mungall, 2009)



By Opabinia regalis - Self created from PDB entry 1TIM using the freely available visualization and analysis package VMD, CC BY-SA 3.0, https://commons.wikimedia.org/w/index.php?curid=1068554

TNF and its Receptor



Protein–Protein Interactions in Virus–Host Systems (Brito, A. F., & Pinney, 2017)

"To study virus-host protein interactions, knowledge about viral and host protein architectures and repertoires, their particular evolutionary mechanisms, and information on relevant sources of biological data is essential."

"From a biomedical perspective, blocking such interactions is the main mechanism underlying antiviral therapies."

Data Sources

Protein Data Base

http://www.rcsb.org/



PDB Data Fue										
ATOM	1	N	ILE	в	16	18,871	65.715	12.731	1.00	21.86
ATOM	2	CA	ILE	Ξ	16	19.782	64.969	13.587	1.00	21.86
ATOM	3	C	ILE	Ξ	16	21.173	64.987	12.945	1.00	21.86
ATOM	4	0	ILE	Е	16	21.316	64.450	11.815	1.00	21.86
ATOM	5	CB	ILE	Ξ	16	19.336	63.476	13.649	1.00	21.86
ATOM	6	CG1	ILE	Ξ	16	17.903	63.230	14.154	1.00	18.04
ATOM	7	CG2	ILE	Ε	16	20.336	62.527	14.373	1.00	18.04
ATOM	8	CD1	ILE	Ε	16	17.785	63.415	15.666	1.00	18.04
ATOM	9	N	VAL	Ε	17	22.160	65.538	13.640	1.00	21.83
ATOM	10	CA	VAL	Е	17	23.595	65.525	13.234	1.00	21.82
ATOM	11	C	VAL	Ε	17	24.317	64.322	13.799	1.00	21.82
ATOM	12	0	VAL	Ξ	17	24.087	63.910	14.971	1.00	21.82
ATOM	13	CB	VAL	E	17	24.328	66.745	13.770	1.00	21.82
ATOM	14	CG1	VAL	Ε	17	25.764	66.765	13.241	1.00	9.61
ATOM	15	CG2	VAL	E	17	23.598	68.073	13.459	1.00	9.61
ATOM	16	N	GLY	Ε	18	24.967	63.607	12.952	1.00	20.38
ATOM	17	CA	GLY	E	18	25.730	62.514	13.495	1.00	20.38
ATOM	18	С	GLY	Ε	18	24.949	61.242	13.653	1.00	20.38
ATOM	19	0	GLY	Ε	18	25.511	60.285	14.234	1.00	20.38
ATOM	20	N	GLY	Ε	19	23.764	61.201	13.121	1.00	18.65

(Wiltgen et al., 2007)



GO - Molecular function¹

- cytokine activity Source: BHF-UCL -
- identical protein binding Source: BHF-UCL +
- protease binding Source: BHF-UCL +
- transcription regulatory region DNA binding Source: UniProtKB -
- tumor necrosis factor receptor binding Source: BHF-UCL +

View the complete GO annotation on QuickGO ...

GO - Biological process¹

- activation of cysteine-type endopeptidase activity involved in apoptotic process Source: UniProtKB -
- activation of MAPK activity
 Source: BHF-UCL +
- activation of MAPKKK activity Source: BHF-UCL -

TNF - UniProt

https://www.uniprot.org/

Drugs Pair Analysis



DrugBank

https://www.drugbank.ca/



ORUGBANK

The DrugBank database is a unique bioinformatics and cheminformatics resource that combines detailed drug data with comprehensive drug target information.

The latest release of DrugBank (version 5.0.11, released 2017-12-20) contains 11,002 drug entries including 2,504 approved small molecule drugs, 943 approved biotech (protein/peptide) drugs, 109 nutraceuticals and over 5,110 experimental drugs. Additionally, 4,910 non-redundant protein (i.e. drug

PDB & DrugBank

- Search by Drugs & Drug Targets
 - <u>http://www.rcsb.org/pages/search_features#search_drugs</u>

U.S. Food & Drug Administration Adverse Event Reporting Systems (AERs)

- Drug Approvals and Databases
 - <u>https://www.fda.gov/Drugs/Informat</u> <u>ionOnDrugs/</u>



U.S. Food & Drug Administration Adverse Event Reporting Systems (AERs)

- FAERS FDA Adverse Event Reporting Systems
 - <u>https://www.fda.gov/Drugs/Informat</u> <u>ionOnDrugs/ucm135151.htm</u>



Note: ir you need neip accessing information in dimerent file formats, see instructions for Downloading Viewers and Players. Language Assistance Available: Español | 繁體中文 | Tiếng Việt | 한국어 | Tagalog | Русский | Laguage Assistance Available: Español | 繁體中文 | Tiếng Việt | 한국어 | Tagalog | Русский | Je [Jenglish

Online Mendelian Inheritance in Man (OMIM)

"[...] catalog of human genes and genetic disorders and traits, with a particular focus on the gene-phenotype relationship." (Wikipedia, 2018)

Mendelian Trait

"Mendelian trait is one that is controlled by a single locus in an inheritance pattern. In such cases, a mutation in a single gene can cause a disease that is inherited according to Mendel's laws. Examples include sickle-cell anemia, Tay-Sachs disease, cystic fibrosis and xeroderma pigmentosa." (Wikipedia, 2018)

HIV-1 Human Interaction Database

https://www.ncbi.nlm.nih.gov/genome/viruses/retroviruses/hiv-1/interactions/browse/

S NCBI Resources 🕑 How To 🛇							Sign in to NC
Retroviruses				Search			
HIV-1 Human Interaction Database	Browse	About	Help	Publications	Releases		
HIV-1 Interactions: Browse an — Filters HIV protein ALL	v All	ein interact	tion on types	~	Replication	interaction ction types	; v
		20 -61000		Do	ownload	View NCBI	Gene records
	Items 1	- 20 of 1806	9 << Fi	rst < Prev F	Page 1	of 904	Next> Last>>
HIV-1		1	nteractio	n	Human	genes	Details
Asp		t	pinds		HLA-A		more

binds

enhances

HLA-B

IFNG

more.

more

Asp

Asp



Estimates of Cells in Human Body

• Reference man

- 70 kilograms
- 20-30 years old
- 1.7 metres tall
- 30 trillion human cells
- 39 trillion bacteria

(Sender et al., 2016)



NIH Human Microbiome Project

https://hmpdacc.org/

NIH Human Microbiome Project



Characterization of the microbiomes of healthy human subjects at five major body sites, using 16S and metagenomic shotgun sequencing.



Characterization of microbiome and human host from three cohorts of microbiomeassociated conditions, using multiple 'omics technologies.

Enter HMP1

Enter iHMP



- Initial Phase (2008)
- 300 healthy individuals
- Sites on the human body
 - nasal passages
 - oral cavity
 - o Skin
 - gastrointestinal tract
 - urogenital tract
- 16S rRNA sequencing
- Metagenomic whole genome shotgun (wgs) sequencing
- Over 14.23 terabytes of data

Model Organisms

Model Organisms

- "Most of our knowledge about the basic properties of metabolism, growth, and division in living cells is a result of studies on species described as 'model organisms'".
- These species include:
 - bacterium Escherichia coli
 - bakers' yeast (Saccharomyces cerevisiae),
 - the fruit fly (Drosophila melanogaster)
 - the nematode worm (Caenorhabditis elegans)
 - the mouse (Mus musculus)
 - the thale cress (Arabidopsis thaliana)

(Oliver et al., 2016)

Model Organism Databases (MOD)

 "Model organism databases (MODs) host the genomic and functional information produced by organism-specific research projects and provide query and visualization tools to access these data"

(Oliver et al., 2016)

PortEco

http://www.porteco.org/



۷	VHAT CAN I DO?
	Find data for a particular gene
	Search
	Enter a list of genes and test for enriched functions
	separate IDs by newline
	Search
	Find and analyze datasets

Currently Unavailable

PortEco is a next-generation data resource for the bacterial model organism, Escherichia coli [Read more...]

PortEco Resources

- Pathway/Genome Databases for 130 E. coli genomes are available at BioCyc.org including curated databases for *E. coli* B Rel 606 and *E. coli* W3110
- EcoliHouse provides a publicly queryable MySQL database warehouse for E. coli data
- Community features including colleague search, event calendar, job postings
- E. coli systems models at BioModels
- PortEco data downloads and database access
- PortEco also supports manual curation of Gene Ontology terms from published papers at EcoCyc

PortEco News and Events

PortEco needs letters of support

PortEco blog > PortEco: by jimhu (4 years ago.): Dear Colleagues, Since taking on the project in 20...[Read more...]

New version of the MC16EE seguence at Conhenk

EcoCyc https://ecocyc.org/



LOGIN | Why Login? | Create New Account

Gene Search

Ouick Search

5

6

Sites - Search - Genome - Metabolism - Analysis - SmartTables - Help -

EcoCyc E. coli Database

EcoCyc is a scientific database for the bacterium *Escherichia coli* K-12 MG1655. The EcoCyc project performs literature-based curation of the entire genome, and of transcriptional regulation, transporters, and metabolic pathways.

New to EcoCyc? Take the guided tour of the EcoCyc.org Web site, watch our free online instructional videos, or read our 2017 article: "EcoCyc: reflecting new knowledge about *Escherichia coli* K-12"

EcoCyc User Guide >>

REACTION × Bar XY Heat REACTION umma X Bar XY Her REACTION Bar XY Heat -3.5 ilvh 0 1 2 3 ilvN **IIvB** ilvG ilvG ilvM 0 signal transduction pathways

Searching Escherichia coli K-12 substr. MG1655 (EcoCyc) change organism database

Enter a gene, protein, metabolite or pathway ...

Gene Expression Data Analysis

Multiple tools are available in this website for analysis of gene expression data.

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Learn More


Extending Worms Life

https://www.npr.org/2015/05/22/408027400/how-do-you-make-an-elderly-worm-feel-young-again



A journey through fascinating ideas, astonishing inventions, and new ways to think and create. Based on riveting TEDTalks from the world's most remarkable minds.



How Do You Make An Elderly Worm Feel Young Again?

May 22, 2015 - 8:24 AM ET Heard on TED Radio Hour





A C. elegans mutant that lives twice as long as wild type

 "WE have found that mutations in the gene daf-2 can cause fertile, active, adult Caenorhabditis elegans hermaphrodites to live more than twice as long as wild type." (Kenyon et al., 1993)

Caenorhabditis elegans



WormBase

https://www.wormbase.org



efr-3

fubp-31

WBGene00016311

WBGene00007534

C32D5.3

View More >

Zebrafish - ZFIN

https://zfin.org/



Mouse - MGI

http://www.informatics.jax.org/







Fly - FlyBase



Rat - Rat Genome Database (RGD)

https://rgd.mcw.edu/

	Help Publications Poster Archive FT	P Download REST API C	iting RGD Contact Us Sign In	
	Enter Search Term	Q	Advanced Search (OLGA)	
~ Rod		Gene Editing Rat Resource Center		
DME DATA ANALYSIS & VISUALIZATION DI	SEASES PHENOTYPES & MODELS GENETI	C MODELS PATHWAY	COMMUNITY	
earch RGD Grant Resources Citing RGD About	Us Contact Us			
Not statisticate angular Market Angular Map positions, functions a more	nd Strains		Find us on Facebook	
Map positions, functions a more Map positions, functions a more Function Gene Ontology, Phenotype Pathway	nd Strains Search Strains e, Diseases Genes, QTL & Strains related to Disease		Phenotypes & Traits linked to the genome Phenotypes & Models Phenotype data, Assays, tusbandry and more	

Yeast - Saccharomyces Genome Database (SGD)

https://www.yeastgenome.org/





About SGD

The *Saccharomyces* Genome Database (SGD) provides comprehensive integrated biological information for the budding yeast *Saccharomyces cerevisiae* along with search and analysis tools to explore these data, enabling the discovery of functional relationships between sequence and gene products in fungi and higher organisms.

Try this?

Meetings

31st VHYC Yeast Conference ☑ April 16 to April 17, 2018 -Leuven, Belgium New & Noteworthy

In Memoriam: André Goffeau - April 12, 2018 It was with great sadness that we learned that André Goffeau, renowned yeast researcher and Professor at the Université Catholique de Louvain in Belgium, passed away on April 2, 2018, Brof, Goffeau

Tweets by @yeastgenome

@yeastgenome Full set of curated #yeast protein (and other) complexes, aka the yeast #complexome, now available at the @complexportal at the @complexbill.aback it out:

0

Fungal Pathogen Genomics 2

Alliance of Genome Resources

http://www.alliancegenome.org/







WormMine

http://intermine.wormbase.org/tools/wormmine/



Digital Patient

in the second

annanannannan

"a technological framework that, once fully developed, will make it possible to create a computer representation of the health status of each citizen that is descriptive and interpretive, integrative and predictive." Discipulus Consortium (2013)

eVip Electronic Virtual Patients https://virtualpatients.eu/

Referatory electronic

Co-funded by the European Commission

Search...

HOMEPAGE ABOUT EVIP » R

RESOURCES » NEWS »

CONTACT

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Referatory

These eViP virtual patients are available to use, by clicking on the URL link. In some cases this will immediately open the case in its original virtual patient player, in other cases you will be guided to a form, where you will be able to register for free and then open the case. There are several different type of players so don't be surprised if the different VPs open in very different player systems.

In addition to the URL link, the VPs are all available as a content package, which conforms to the virtual patient ANSI-accredited standard. To use these packages, you will need to import them into a suitable virtual patient player i.e. a player which is able to accept VP standard-compliant content. This will require the support of a suitably-experienced learning technologist.

Search For:

search

Displaying 340 virtual patients

Title	Keywords	Language	Institution	License	Content Package	URL
Catherine Miller	Meningitis, Bacterial Meningitis, Sepsis	English	St George's, University of London	(CC) BY-NC-SA	٢	Link
Anna-Lena Olofsson	Failure to thrive	English	St George's, University of London	(CC) BY-NC-SA	O	Link
John M	Idiopathic thrombocytopenic purpura, Bruises, Immunoglobulin	English	St George's, University of London	(CC) BY-NO-SA	U	Link
Florian	Prematurity, Respiratory distress syndrome,	English	St George's, University of London	(CC) BY-NC-SA	C	Link

Semantic Web

"... the idea of having data on the web defined and linked in a way that it can be used by machines not just for display purposes, but for automation, integration and reuse of data across various applications."



W3C Semantic Web Activity Group, 2011

Linked Data



Infobox

Wikipedia







DBPedia - English

4.58 million things

• 4.22 million classified in a consistent ontology

- 1,445,000 persons
- 735,000 places (478,000 populated)
- 411,000 creative works
 - 123,000 music albums; 87,000 films; 19,000 video games
- 241,000 organizations

DBPedia - International

- 125 languages
- 38.3 million things
- 23.8 million interlinked with English

Linked Data 05/2007 FOAF Revyu Musicbrainz DBpedia Geo-**RDF Book** names Mashup DBtune World Factbook Jamendo DBLP US Project Census Guten-Data As of May 2007 berg

Source: http://lod-cloud.net/

Datasets published following Linked Data 'format': 05/2007



Datasets published following Linked Data 'format': **11/2007**

Linked Data



Source: http://lod-cloud.net

Datasets published following Linked Data 'format': 2008





Datasets published following Linked Data 'format': 2010







Linked Data 03/2019

1,239 datasets

16,147 links

https://lod-cloud.net/



Medical Subject Headings (MeSH) https://www.nlm.nih.gov/mesh/meshhome.html

- "National Library of Medicine's controlled vocabulary thesaurus."
- used by the MEDLINE/PubMed article database
- 28,000 descriptors
- 90,000 entry terms

Hierarchical Tree Myocardial Infarction

- MeSH Browser
 - https://meshb.nlm.nih.gov/search
- Myocardial Infarction
 - C14.280.647.500
 - C stands for Diseases
 - C14 Cardiovascular Diseases
 - C14.280 Heart Diseases
 - C14.2280.647 Myocardial Ischemia

https://en.m.wikipedia.org/wiki/Medical_Subject_Headings

Cardiovascular Diseases [C14] Heart Diseases [C14.280] Myocardial Ischemia [C14.280.647] Acute Coronary Syndrome [C14.280.647.124] Angina Pectoris [C14.280.647.187] Coronary Disease [C14.280.647.250] Kounis Syndrome [C14.280.647.375] Myocardial Infarction [C14.280.647.500] Anterior Wall Myocardial Infarction [C14.280.647.500.093] Inferior Wall Myocardial Infarction [C14.280.647.500.187] Non-ST Elevated Myocardial Infarction [C14.280.647.500.469] Shock, Cardiogenic [C14.280.647.500.750] ST Elevation Myocardial Infarction [C14.280.647.500.875] Myocardial Reperfusion Injury [C14.280.647.625]

PubMed

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PubMed

PubMed comprises more than 29 million citations for biomedical literature from MEDLINE, life science journals, and online books. Citations may include links to full-text content from PubMed Central and publisher web sites.

Using PubMed

PubMed Quick Start Guide

Full Text Articles

PubMed FAQs

Dealer and Transition

PubMed Tools

PubMed Mobile

Single Citation Matcher

Batch Citation Matcher

01:-----

More Resources

MeSH Database

Journals in NCBI Databases

Clinical Trials

- I MATTAL - - / A POIN

PubMed MeSH Search

-

PubMed Search Builder

"Myocardial Infarction" [Mesh] AND "Chest Pain"[Mesh]

Add to search builder

AND ~

Search PubMed

PubMed MeSH Search

	"Myocardial Infarction"[Mesh] AND "Chest Pain"[Mesh] - PubMed - NCBI - Mozilla Firefox	_ 0	(
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S NCBI Resources	⊠ How To ⊠ Sign in to	NCBI	
Publiced.gov US National Library of Medicine National Institutes of Health	MeSH V "Myocardial Infarction"[Mesh] AND "Chest Pain"[Mesh] Search Create RSS Create alert Advanced	Help	
Article types Clinical Trial Review Customize	Format: Summary - Sort by: Most Recent - Per page: 20 - Send to - Filters: Manage Filters Search results Sort by:		
Text availability Abstract Free full text Full text Publication dates 5 years 10 years Custom range	Items: 1 to 20 of 14939 Effect of Prehospital Blood Draws on Length of Stay for Chest Pain Patients in the Emergency Effect of Prehospital Blood Draws on Length of Stay for Chest Pain Patients in the Emergency Results by year 1. Department: A Pilot Study. DuCharme B, Macci Bires A, Montanye E, Khan M, DuCharme S, Linse M, Carlson JN. Crit Care Nurs Q. 2019 Apr/Jun;42(2):208-214. doi: 10.1097/CNQ.0000000000257. Results by year PMID: 30807348 Similar articles Image: Comparison of Chest Pain Patients in the Emergency Page I of 747 Next > Last >>	nt	
Species Humans Other Animals <u>Clear all</u> <u>Show additional filters</u>	 Targeted metabolomic analysis of plasma metabolites in patients with coronary heart disease in southern China. Zhong Z, Liu J, Zhang Q, Zhong W, Li B, Li C, Liu Z, Yang M, Zhao P. Medicine (Baltimore). 2019 Feb:98(7):e14309. doi: 10.1097/MD.00000000014309. PMID: 30762730 Free PMC Article Spontaneous Coronary Artery Dissection Masquerading as Coronary Artery Stenosis in a Young Patient. Rawala MS, Naqvi STS, Yasin M, Rizvi SB. Am J Case Rep. 2019 Feb 6;20159-162. doi: 10.12659/AJCR.913522. PMID: 30723187 Free PMC Article 	tratity in rt. 2018] ic. 2018] e id. 2018] se more	
Gene Ontology (GO)

http://www.geneontology.org/

Gene Ontology Consortium Home

Documentation - Downloads -

About - Contact us

Enrichment analysis

Y	'our gene IDs here
ł	
H	Homo sapiens

Help Powered by PANTHER

Statistics



Gene Ontology Consortium

Search GO data

Search for terms and gene products...

Search

Ontology

Filter classes

Download ontology

Gene Ontology: the framework for the model of biology. The GO defines concepts/classes used to describe gene function, and relationships between these concepts. It classifies functions along three aspects:

molecular function molecular activities of gene products cellular component

Annotations

Tools -

Annotations

Download annotations (standard files)

Filter and download (customizable files <100k lines)

GO annotations: the model of biology. Annotations are statements describing the functions of specific genes, using concepts in the Gene Ontology. The simplest and most common annotation links one gene to one function, e.g. FZD4 + Wnt signaling pathway. Each statement is based on a specified piece of evidence. more The mission of the GO Consortium is to develop an up-to-date, comprehensive, computational model of biological systems, from the molecular level to larger pathways, cellular and organismlevel systems, more

Search documentation



What is the Gene Ontology?

- An introduction to the Gene
 Ontology
- What are annotations?
- Enrichment analysis
- Downloads



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Biomedical Ontology

https://www.bioontology.org/



BioPortal

http://bioportal.bioontology.org/

BioPortal			
		BioPortal Statistics	
Welcome to BioP comprehensive repositor	ortal, the world's most y of biomedical ontolog	Ontologies	766
Search for a class	Find an ontology	Classes	9,238,120
Enter a class, e.g. Melanoma	Start typing ontology name, th		
Advanced Search	Browse Ontologies	Resources Indexed	48
Ontology Visits (March 2019)	BioPortal Statistics		
CPT	Ontologies	Indexed Records	39,537,360
MEDDRA SNOMEDCT RXNORM	Classes		
NDDF	Resources Indexed	Direct Annotations	95,468,433,792
		Direct Plus Expanded Annotations	144,789,582,932



-

Human Phenotype Ontology http://human-phenotype-ontology.github.io/

Machine Learning

Traditional Programming



(Domingos, 2017)



uuunnan anuuuuuu



Iris Data Set Download: Data Folder, Data Set Description

Abstract: Famous database; from Fisher, 1936



Search

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att:

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Data Set Charac	teristics:	Multivariate	Number of Instances:	150	Area:	Life
Attribute Charac	teristics:	Real	Number of Attributes:	4	Date Donated	1988-07-01
Associated Task	s:	Classification	Missing Values?	No	Number of Web Hits:	2512440

Source:

Creator:

R.A. Fisher

Donor:

Michael Marshall (MARSHALL%PLU '@' io.arc.nasa.gov)

https://archive.ics.uci.edu/ml/datasets/iris

Modeling Healthcare

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NII	C National Institute for
	L Health and Care Excellence

Find NICE guidance

NICE	NICE
Pathways	Guidance

Evidence

services

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Search NICE...

Improving health and social care through evidence-based guidance

Browse guidance by area: Conditions and diseases Health protection Lifestyle and wellbeing Population groups Service delivery, organisation and staffing Settings

	About us	Put guidance into practice	Find journals and databases	Financial planning	Get involved
https://www.nice.org.uk/					

Q

NICE

https://pathways.nice.org.uk/pathways/bacterial-meningitis-and-meningococcal-septicaemia-in-under-16s



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NICE

https://pathways.nice.org.uk/pathways/heart-rhythm-conditions/heart-rhythm-conditions-overview#content=view-node:nodes-ventricular-arrhythmias



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AHRQ Search

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The AHRQ National Gu through AHRQ will 20 of 21 results for	uideline Clearinghe I no longer be avai	ouse (NBC, guideline.gov) Web alte will not be available after July 16, 2018 because f riable to support the NGC as of that date. For additional information, read our full anno	ederal funding uncement
Cardiology			
NARROW RESULTS	Clear All	1 2	Next >
Meets 2013 Inclusion Criteria (Includes NEATS Assessment (1) U.Sbased Organizations (14) Addresses Multiple Chronic Cr	11)) • • S inditions (4)	SGRT av Relevance Date SHOW 20150 100 Compare Summaries O × Clinical Algorithm × Diagnosis × Cardiology	
Publication Date	r	GUIDELINE SUMMARY NGC 010428 2014 MAY	
From: 2008 - To: 2017 -	h t	Management of acute atrial fibrillation and atrial flutter in non-pregnant hospitalized adults. University of Michigan Health System	Compare
Apply			
Age Age Aged, 80 and over (17) Aged (65 to 79 years) (17) Middle Age (45 to 64 years) (17) Middle Age (45 to 64 years) (17)	ristics	■ CONSTRUCT SUBMARY NOCESSAL 1998 Nov (100080704.JBB 19) 2014 AHA/ACC guideline for the management of patients with valvular heart disease: a report of the American College of Cardiology/American Heart Association Task Force on Practice Guidelines. American College of Cardiology Foundation: American Heart Association	Compane
Adult (19 to 44 years) (17) Show More Gender Female (21) Male (21)	B c z	CODELINE SUMMARY INCC 001997 2013 JR. Myocardial Infarction with ST-segment elevation. The acute management of myocardial Infarction with ST-segment elevation. National Guidence Centre	Compare
Clinical Specialty	- r	GUIDELINE SUMMARY NGC 010473 2006 (REVISED 2014 JUN)	
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Dermatology (1) Show More	F	CUIDELINE SUMMARY NGC 010600 2008 SEP (SEVISED 2014 JUL)	Compare
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Cardiac arrhythmias in coronary heart disease

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Cardiac arrhythmiais a group of conditions in which the heartbeat is irregular. Symptoms include: palpitations or a pause between heartbeats; light-headedness; shortness of breath; and chest pain.

Remit and target users

This guideline provides evidence based recommendations for the management of cardiac arrest and the arrhythmias associated with acute coronary syndromes, chronic coronary heart disease and cardiac surgery. It excludes arrhythmias not associated with coronary heart disease such as supraventricular tachycardias associated with accessory pathways or dual atrioventricular (AV) nodal physiology, arrhythmias caused by inherited ion channel disorders (eg long QT syndrome, Brugada syndrome) and arrhythmias associated with non-ischaemic cardiomyopathies.

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Management of invasive meningococcal disease in children & young people

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The most common clinical manifestation of invasive meningococcal disease is meningitis, but up to 20% of patients will develop meningococcal septicaemia, associated with the highest mortality. The majority of deaths continue to occur in the first 24 hours and the recorded case fatality rate varies between 2.6% and10% each year.

Remit and target users

This guideline makes recommendations on best practice in the recognition and management of meningococcal disease in children and young people up to 16 years of age. It addresses pre-hospital care, referral, diagnostic testing, disease management, follow-up care and rehabilitation and considers public health issues.

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http://www.sign.ac.uk/assets/grg102.pdf

- Akinsola, J. E. T., O, A., & J, A. (2017). A Framework for Web Based Detection of Journal Entries Frauds using Data Mining Algorithm. International Journal of Computer Trends and Technology, 51, 1–9. http://doi.org/10.14445/22312803/IJCTT-V51P101
- Alvarado, P. (2016). Building the Machine Learning Infrastructure. Retrieved March 12, 2018, from http://blogs.teradata.com/data-points/building-machine-learning-infrastructure-2/
- Brailsford, S. C. (2007). Advances and Challenges in Healthcare Simulation Modeling: Tutorial. In Proc. of the 39th Conf. on Winter Simulation (pp. 1436–1448). Washington D.C.: IEEE Press.
- Brito, A. F., & Pinney, J. W. (2017). Protein–Protein Interactions in Virus–Host Systems. Frontiers in Microbiology, 8, 1557.
- Brownlee, J. (2018). Supervised and Unsupervised Machine Learning Algorithms. Retrieved March 12, 2018, from

https://machinelearningmastery.com/supervised-and-unsupervised-machine-learning-algorithms/

- Cheng, F., Liu, C., Jiang, J., Lu, W., Li, W., Liu, G., ... Tang, Y. (2012). Prediction of Drug-Target Interactions and Drug Repositioning via Network-Based Inference. PLoS Computational Biology, 8(5), e1002503. http://doi.org/10.1371/journal.pcbi.1002503
- Clancy, S. & Brown, W. (2008) Translation: DNA to mRNA to Protein. Nature Education 1(1):101
- Combs, C. D., Sokolowski, J. A., & Banks, C. M. (Eds.). (2016). The Digital Patient : Advancing Healthcare, Research, and Education. New Jersey: John Wiley & Sons.
- Cook, S., Conrad, C., Fowlkes, A. L., & Mohebbi, M. H. (2011). Assessing Google Flu Trends Performance in the United States during the 2009 Influenza Virus A (H1N1) Pandemic. PLoS ONE, 6(8), e23610.
- Crick, F. (1970). Central Dogma of Molecular Biology. Nature, 227(5258), 561–563. https://doi.org/10.1038/227561a0
- Carlos Greg, Duik (2014) The Formation of Love. Retrieved from https://www.facebook.com/notes/facebook-data-science/the-formation-of-love/10152064609253859

- Holzinger, A. (2014). Biomedical Informatics : Discovering Knowledge in Big Data. Switzerland: Springer.
- Wiltgen, M., Holzinger, A., & Tilz, G. P. (2007). Interactive Analysis and Visualization of Macromolecular Interfaces between Proteins. In HCI and Usability for Medicine and Health Care (pp. 199–212). Berlin, Heidelberg: Springer Berlin Heidelberg. http://doi.org/10.1007/978-3-540-76805-0_17
- Jeong, H., Mason, S. P., Barabási, A.-L., & Oltvai, Z. N. (2001). Lethality and centrality in protein networks. Nature, 411(6833), 41–42. http://doi.org/10.1038/35075138
- KhanAcademy (2018). The genetic code. 5 March 2018. Retrieved from <u>https://www.khanacademy.org/science/biology/gene-expression-central-dogma/central-dogma-trans</u> <u>cription/a/the-genetic-code-discovery-and-properties</u>
- Lee, T.B.; Hendler, J. & Lassila, O. The Semantic Web. Scientific American, 2001, 284, 28-37
- Leise, F.; Fast, K.; Steckel, M. What Is A Controlled Vocabulary? Boxes and Arrows, Dezembro 2002, online: http://www.boxesandarrows.com/view/what_is_a_controlled_vocabulary_

- Miettinen, P. (2012). Decomposing Binary Matrices: Where Linear Algebra Meets Combinatorial Data Mining. Bristol: ECML-PKDD 2012. Retrieved from <u>https://people.mpi-inf.mpg.de/~pmiettin/bmf_tutorial/</u>
- Mungall, C. (2009) Integrating phenotype ontologies across multiple species. Caltech.
- Raja, K., Patrick, M., Elder, J. T., & Tsoi, L. C. (2017). Machine learning workflow to enhance predictions of Adverse Drug Reactions (ADRs) through drug-gene interactions: application to drugs for cutaneous diseases. Scientific Reports, 7(1), 3690. http://doi.org/10.1038/s41598-017-03914-3
- Shortliffe, E. H., & Cimino, J. J. (Eds.). (2014). Biomedical Informatics Computer Applications in Health Care and Biomedicine. London: Springer London.

- Spiegler, I., & Maayan, R. (1985). Storage and retrieval considerations of binary data bases. Information Processing & Management, 21(3), 233–254.
 https://doi.org/10.1016/0306-4573(85)90108-6
- Stratton, M. R. et al. (2009) Nature 458, 719-724 doi:10.1038/nature07943
- Studer, R., Benjamins, V. R., & Fensel, D. (1998). Knowledge engineering: Principles and methods. Data & Knowledge Engineering, 25(1-2), 161-197.
- Tatonetti, N. P., Ye, P. P., Daneshjou, R., & Altman, R. B. (2012). Data-Driven Prediction of Drug Effects and Interactions. Science Translational Medicine, 4(125), 125ra31-125ra31. https://doi.org/10.1126/scitranslmed.3003377
- O'Connor, Timothy and Wong, Hong Yu, (2012) Emergent Properties, The Stanford Encyclopedia of Philosophy (Spring 2012 Edition), Edward N. Zalta (ed.), URL = http://plato.stanford.edu/archives/spr2012/entries/properties-emergent/.

- van Riel, Raphael and Van Gulick, Robert. (2016) "Scientific Reduction", The Stanford Encyclopedia of Philosophy (Winter 2016 Edition), Edward N. Zalta (ed.), URL = https://plato.stanford.edu/archives/win2016/entries/scientific-reduction/.
- Wang, X., Gorlitsky, R., & Almeida, J. S. (2005). From XML to RDF: how semantic web technologies will change the design of "omic" standards. Nat Biotech, 23(9), 1099-1103.
- Welty, C., Lehmann, F., Gruninger, G., and Uschold, M. (1999) Ontology: Expert Systems All Over Again? Invited panel at AAAI-99: The National Conference on Artificial Intelligence. Austin, Texas
- Wiltgen, M., Holzinger, A., & Tilz, G. P. (2007). Interactive Analysis and Visualization of Macromolecular Interfaces between Proteins. In HCI and Usability for Medicine and Health Care (pp. 199–212). Berlin, Heidelberg: Springer Berlin Heidelberg. http://doi.org/10.1007/978-3-540-76805-0_17
- Zhou, X., Menche, J., Barabási, A.-L., & Sharma, A. (2014). Human symptoms–disease network. Nature Communications, 5(1), 4212. https://doi.org/10.1038/ncomms5212

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