

## Web of Science 最新情報のご紹介

トムソン・ロイター 学術情報ソリューション

宮田 亮

2013年7月10日(水) NIMS LIBRARY OPEN SEMINAR Web of Science活用ワークショップ



## Web of Scienceの特徴

- 各分野をリードする国際誌(コア・ジャーナル)を集めてい ます
- 各文献を引用関係でリンクし、ある文献が何に引用されてい るか、何を引用しているかがわかります。
- インパクトファクターのソースデータです。

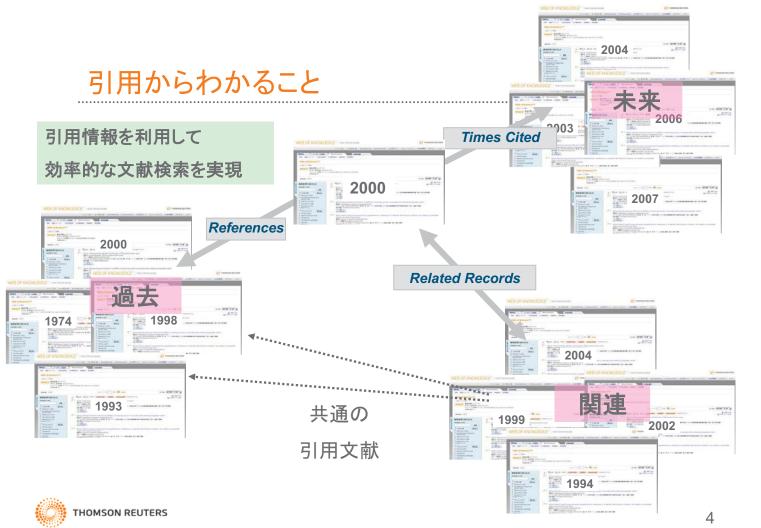


## Web of Scienceの基本 • Web of Knowledgeに搭載されている12種類の文献データベー スのひとつで、Web of Knowledgeの基幹コンテンツです

• 幅広い分野をカバーしています



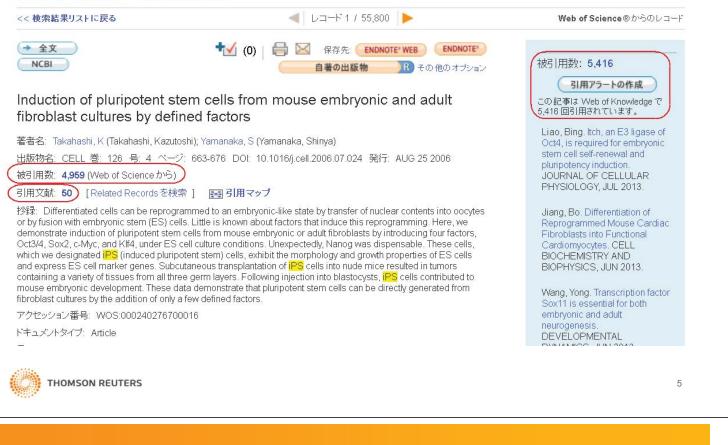






## Web of Scienceのレコード例

### Web of Science® now with books



## Web of Scienceの活用

- •研究者や学生の先行研究のための文献調査として
- •評価・分析用の統計データとして
  - Times Higher Education World University Rankings
  - CWTS Leiden Ranking
  - ・研究論文に着目した日本の大学ベンチマーキング2011
  - ベース(Web of Science)の連結によるデータ分析
  - •科学技術指標 一科学技術白書
  - •米国国立科学財団:NSF: Science and Engineering Indicators・・・など 国立大学の目指すべき方向





THOMSON REUTERS

## **RESEARCH OUTPUTS**



# **NEW CONTENT: BOOK CITATION INDEX**

7

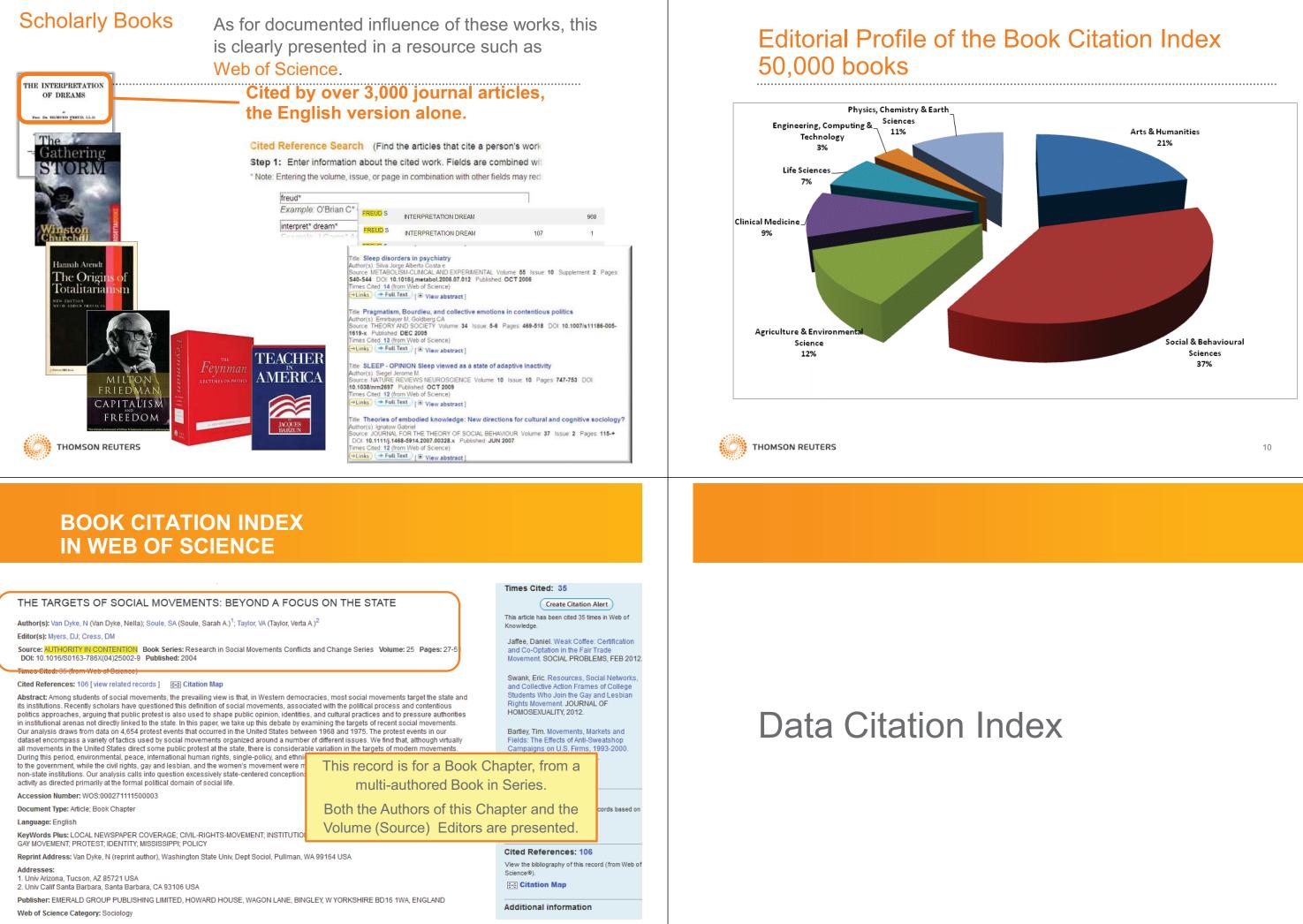
•科学研究費助成事業データベース(KAKEN)と論文データ











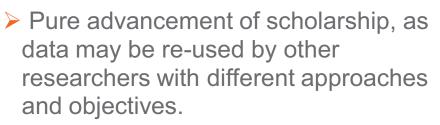


12

## **Data Citation Index**

Availability of datasets and various forms of research data has varied from field to field and region to region. However, the benefits of data sharing are obvious.





- > Verification of results.
- Promotion of one's work



### Why? Data Citation Index -2

- DISCONNECTED RESOURCES
  - Inability to discover relevant datasets.
  - Lack of standard attribution conventions
  - Difficulty in measuring data use and impact
  - Challenges in seeing the full funding picture
- HISTORICAL LACK OF INCENTIVE
- DATA CITATION INDEX
  - a single source of data discovery

http://wokinfo.com//products tools/multidisciplinary/dci/collaborative science/

10 K 15 K 20 K 25 K 30 K





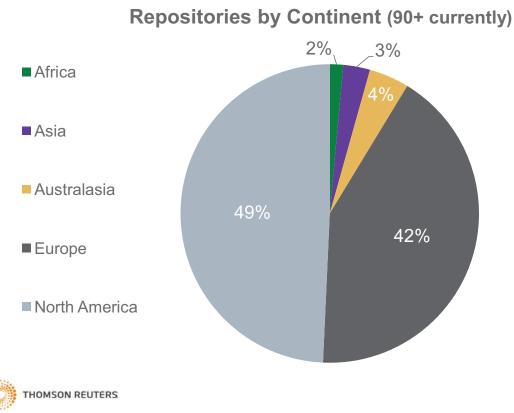
### Whv? Data Citation Index -1

- NSF: "Any data obtained with federal funds be accessible to the general public"
  - NSF was the first organization to mandate that all funding proposals include a two-page "data management plan"
  - Many funders and organizations followed: Economic and Social Research Council (ESRC), National Environment Research Council (NERC), European Commission FP7, Wellcome Trust (UK), Australian National Data Service (ANDS), Research Council (ARC).
- Benefits of Data Sharing
  - Advancing scholarship
  - Increasing transparency
  - Promoting work in new ways



- Curbing the double-funding tide

## **REGIONAL DISTRIBUTION**

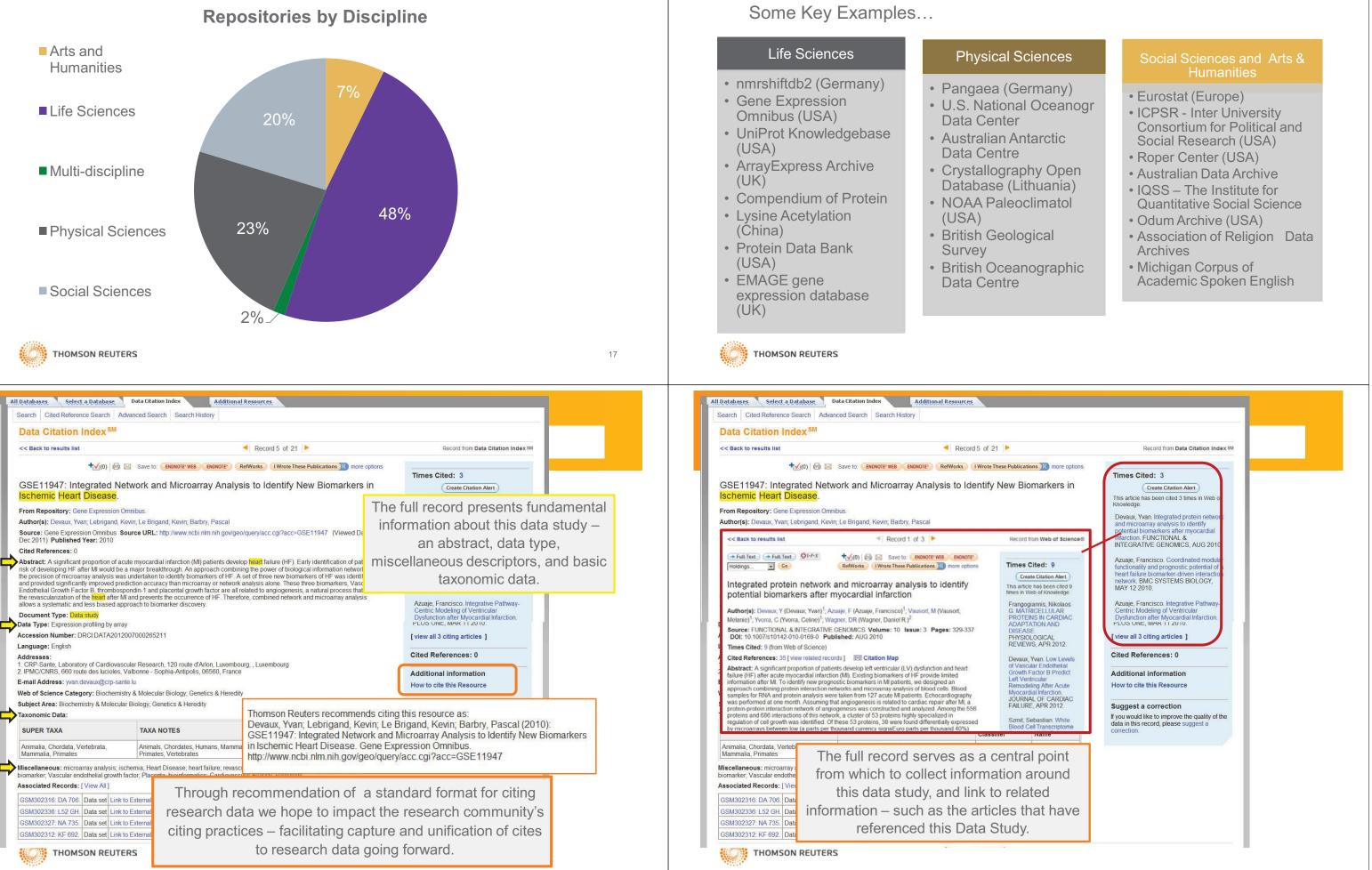




14

16

### DISCIPLINE COVERAGE BREAKDOWN

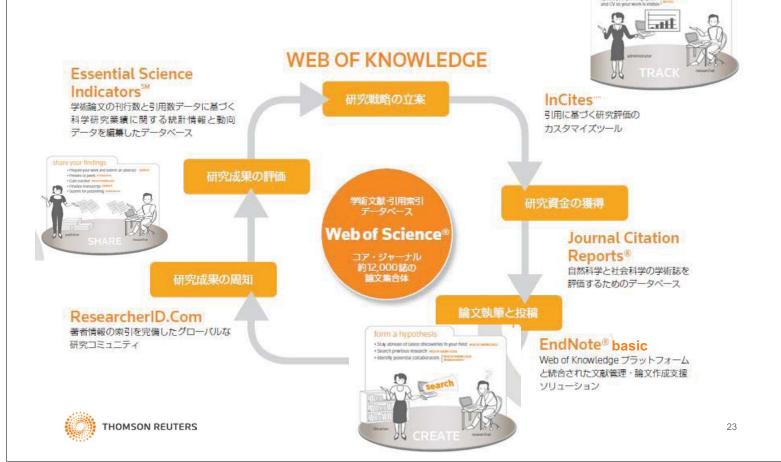


### THE REPOSITORIES AT LAUNCH

earch Cited Reference S	earch Advanced Search Search History						
				hove all th	ough – th	ne Data	
Data Citation Index <sup>SM</sup>				Above all though – the Data			
<< Back to results list			Cit	Citation Index is about getting		t getting	
ty(0) ⊖ ⊠ Save to: ENDNOTE' WEB ENDNOTE' RefWorks   Wrote These Put			otions US	ers to res	ers to research data itself.		
GSE11947: Integrated Network and Microarray Analysis to Identify New Ischemic Heart Disease.			Link	Link to the Data Set information			
From Repository: Gene Expression Omnibus.			within the repository.			torv.	
and the second							
and the second second second second second second	brigand, Kevin: Le Brigaad, Kovin, Barky, Fascal Immibile: <b>Source URL:</b> http://www.ncbi.nlm.nlh.gov/geo/query/acc.cgi?acc=GSE1 : 2010	1947 (Newed Date	: 12 pote infar	microarray analysis to ntial biomarkers after r ction. FUNCTIONAL & GRATIVE GENOMICS	nyocardial		
Abstract: A significant prop isk of developing HF after M e precision of microarray a ind p Endot	ortion of acute myocardial infarctum (MI) patients develop heart failure (HF). Early I would be a major breakthrough An approach combining the power of biological halysis was undertaken to identify biomarkers of HF. A set of three new biomarker information of the set of the	information networks	and func ti hear	aje, Francisco. Coordir tionality and prognostic t failure biomarker-driv rork. BMC SYSTEMS I 12 2010.	potential of a en interaction BIOLOGY,		
	Gene Expression Omnibus		n	aje, Francisco. Integrat tric Modeling of Ventric	cular		
ata HOME SEARCH SIT	MAP GEO Publications	FAQ MIAME	Email GEO	S UNE, MAR 11 2010.	al Infarction.		
CCes NCBI > GEO > Acce	ssion Display 🕐	Not log	ged in   Login 🛽	all 3 citing articles	í.		
and the second second second	er screen elements for information.			and clarig aracico			
ddre				d References: 0			
CRI IPM Scope: Self	Format: HTML 💽 Amount: Quick 💽 GEO accession: GSE11947	GO		tional information			
/eb c Series GSE119	47 Query DataSet	Platforms (1)	GPL1947 RNG-MR	C_HU25k_STRASBOU	RG		
ubje Status	Public on May 25, 2010	Samples (32)	GSM302309 L88 T	G			
axor Title	Integrated Network and Microarray Analysis to Identify New Biomar Heart Disease						
SUP Organism	Heart Disease Homo sapiens		GSM302311 L38 C	P			
Experiment type		Relations					
Anim Mam	A significant proportion of acute myocardial infarction (MI) patient failure (HF). Early identification of patients at risk of developing HF	t BioProject 105803					
lisce	be a major breakthrough. An approach combining the power of biolog networks and the precision of microarray analysis was underta						
oma	Download family Format				ormat		
ssod	significantly improved prediction accuracy than microarray or n	SOFT formatted family file(s) SOFT 2			OFT 🕐		
alone. These three biomarkers, Vascular Endothelial Grow SM thrombospondin-1 and placental growth factor are all related to		MINIML formatted family file(s) MINIML 2					
GSM	(s) TXT 😨						
3SM	a systematic and less biased approach to biomarker discovery.	Sup	plementary file	Size	Download	File type/resource	
GSM		GSE11947_RAW.ta	Welling and an and a second	222.2 Mb	(ftp)(http)(custom		
Overall design The 32 patients of this study were divided in 2 groups corresponding				supplementary file			
Overall design	quartiles of FE values. The first group are the patients with low E	Raw data provided	d as supplementary	file			

	Data Citation Index <sup>SM</sup>						
	<< Back to results list	Record 5 of 21					
	11.10	Save to: ENDNOTE' WEB ENDNOT		ote These Publication			
	GSE11947: Integrated Net Ischemic Heart Disease. From Repository: Gene Expression Orr	work and Microarray Ar	nalysis to Ident				
	Author(s): Devaux, Yvan; Lebrigand, Key	vin; Le Brigand, Kevin; Barbry, Pasc	al				
	Associated Records This: GSE11947; Integrated Network and Microarray Analysis to Identify New Biomarkers in I Heart Disease. Author(s): Devaux, Yvan; Lebrigand, Kevin; Le Brigand, Kevin; et al. Source: Gene Expression Omnibus Source URL: http://www.ncbi.nlm.nih.gov/igeo/query/acc.cgi?acc=GSE11947 Document Type: Data study						
	Results: 32	Mage 1 of 4 Go 1	Sort by: Year	Published - newest to			
	Refine Results	🔩 (0) 🖶 🖾 Save to	ENDNOTE" WEB ENDNOT	•			
	Refine Results Search within results for	RefWorks   I Wrote These Pu	ote These Publications R more options				
	Search within results for	T 1. Title: GSM302316: DA 70		997			
	T Web of Science Categories Refine	Author(s): Devaux, Yvan; Le	Sample GSM30231				
	BIOCHEMISTRY MOLECULAR BIOLOGY (32)	Source: Gene Expression Source URL: http://www.	Status	Public on May 2			
	GENETICS HEREDITY (32) more options / values	acc=GSM302316 Document Type: Data set [  • View abstract ]	Title Sample type	DA 706 RNA			
	V Subject Areas		Channel 1				
	BIOCHEMISTRY MOLECULAR BIOLOGY (32)	C 2. Title: GSM302336: L52 G Author(s): Devaux, Yvan; L		Ref			
	GENETICS HEREDITY (32) more options / values.	Source: Gene Expression	Organism	Homo sapiens			
	Document Types     Refine     DATA SET (32)	Source URL: http://www. acc=GSM302336 Document Type: Data set [ .View abstract ]	Characteristics	Labeling_referer Extraction_referen Sample_referen			
	► Authors	T 3. Title: GSM302327: NA 73		Subject_referen Tissue: blood			
	► Group Authors	Author(s): Devaux, Yvan; Li		Extraction_amo			
	► Editors	Source: Gene Expression Source URL: http://www.i					
		B6680 SM302327	Extracted molecule	total RNA			
	Animalia, Chordata, Vertebrata Mammalia, Primates	Animals, Chordates, Humans, N Primates, Vertebrates	Extraction protocol Label	Trizol Cy3, Cy5			
	1 <u></u>		Label protocol	Ambion			
	Miscellaneous: microarray analysis; isch		Laber protocor	Ambion			
	biomarker: Vascular endotherial growth fa	nor Placenta bioinformatics card	Channel 2				
	Associated Records: [View All]		Source name	DA 706			
	GSM302316: DA 706. Data set Link to	External Source	Organism	Homo sapiens			
	GSM302336: L52 GH. Data set Link to	Characteristics	Labeling_refere				
	GSM302327: NA 735. Data set Link to		Extraction_refe Sample_referen				
	GSM302327: NA 735. Data set Link to GSM302312: KF 692. Data set Link to			Subject_referen Tissue: Blood Extraction_amo			
				exclaction_allio			
	THOMSON REUTER		Extracted molecule				

トムソン・ロイターの研究サポート



Thank you!

