



基于Histcite的文献 的分析与筛选

华南农业大学图书馆信息部

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2021年11月23日



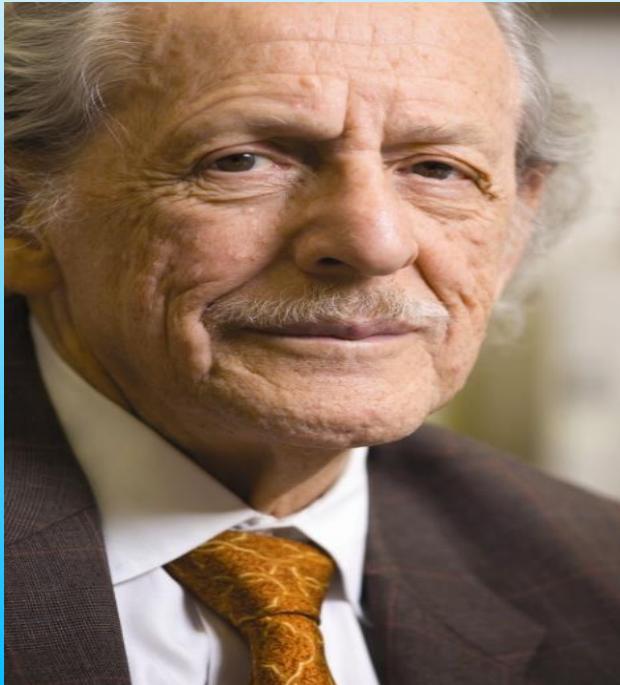
PART 01

引文索引与Web of science



一、引文索引

Dr. Garfield 1955年在《Science》发表的论文提出将引文索引作为一种新的文献检索与分类工具：将一篇文献作为检索起点从而跟踪一个Idea的发展过程及学科之间的交叉渗透的关系。



Dr. Eugene Garfield
Founder & Chairman Emeritus
ISI, Thomson Scientific

Citation Indexes for Science

A New Dimension in Documentation
through Association of Ideas

Eugene Garfield

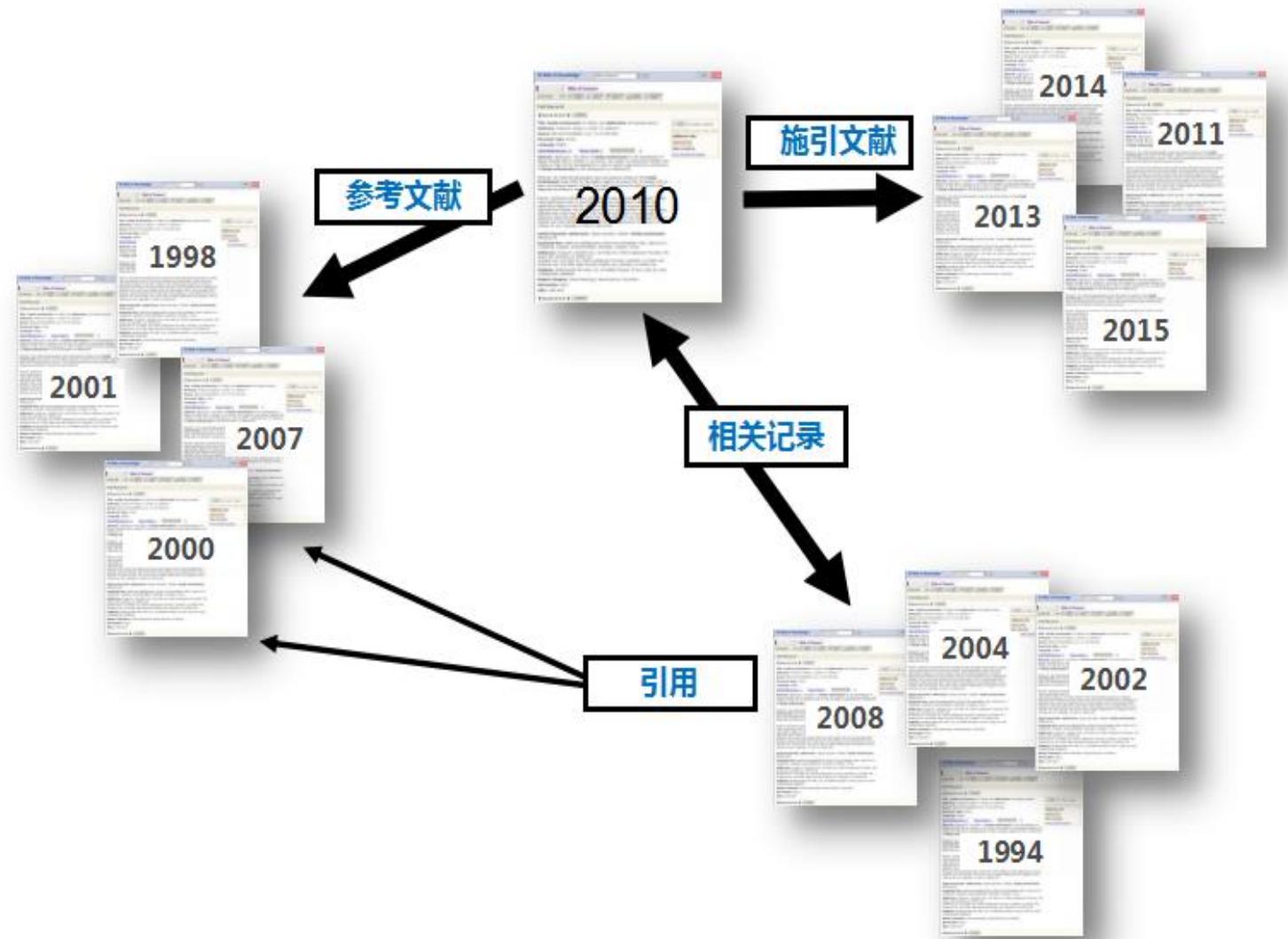
"The uncritical citation of disputed data by a writer, whether it be deliberate or not, is a serious matter. Of course, knowingly propagandizing unsubstantiated claims is particularly abhorrent, but just as many naive students may be swayed by unfounded assertions presented by a writer who is unaware of the criticisms. Buried in scholarly journals, critical notes are increasingly likely to be overlooked with the passage of time, while the studies to which they pertain, having been reported more widely, are discovered

approach to subject control of the literature of science. By virtue of its different construction, it tends to bring together material that would never be collated by the usual subject indexing. It is best described as an association-of-ideas index, and it gives the reader as much leeway as he requires. Suggestiveness through association-of-ideas is offered by conventional subject indexes but only within the limits of a particular subject heading.

If one considers the book as the macro unit of thought and the periodical article



二、引文索





引文索引系统打破了传统的学科分类界限，既能揭示某一学科的继承与发展关系，又能反映学科之间的交叉渗透的关系。



三、Web of science核心合集

Web of Science™核心合集是获取全球学术信息的重要数据库,由以下几个重要部分组成:

- Science Citation Index-Expanded™ (SCI-E, 科学引文索引) 1900年-
- Social Sciences Citation Index™ (SSCI, 社会科学引文索引) 1900年-
- Arts & Humanities Citation Index® (A&HCI, 艺术与人文引文索引) 1975年-
- Conference Proceedings Citation Index™ (CPCI, 会议论文引文索引) 1990年-
- Book Citation Index™ (BkCI, 图书引文索引) 截止至2017年收录了60,000多种图书, 共1,277,000多条记录, 同时每年增加10,000种新书 2005年-
- Current Chemical Reactions® 收录了1985年以来的最新化学反应 1985年-
- Index Chemicus® 收录了1993年以来的化学物质的事实型数据 1993年-
- Emerging Sources Citation Index (ESCI) 展示重要的新兴研究成果 2015年-

Web of Science™核心合集数据库收录了18,000多种世界权威的、高影响力的学术期刊,内容涵盖自然科学、工程技术、生物医学、社会科学、艺术与人文等领域,最早回溯至1900年。Web of Science™核心合集收录了论文中所引用的参考文献、并按照被引作者、出处和出版年代编制成独特的引文索引。



Clarivate

简体中文

产品

Web of Science™

检索

标记结果列表

历史

跟踪服务

登录

注册

探索跨学科内容

来自最值得您信赖的全球引文数据库

选择数据库: Web of Science 核心合集 引文索引: All

文献 作者 被引参考文献 化学结构

所有字段

示例: liver d

+ 添加行

+ 添加日期范围

高级检索

全选

Science Citation Index Expanded
(SCI-EXPANDED)--2002-至今

Social Sciences Citation Index
(SSCI)--2018-至今

Current Chemical Reactions
(CCR-EXPANDED)--1985-至今

Index Chemicus
(IC)--1993-至今

清除

检索



PART 02

Histcite的简介 1

2 数据下载 From WOS 核心合集

导入数据 To Histcite 3

4 作图分析 Make Graph





一、Histcite简介

Histcite=history of cite，意味引文历史，或者叫引文图谱分析软件。该软件是SCI的发明人加菲尔德开发，能够用图示的方式展示某一领域不同文献之间的关系。

优点：操作界面简洁，操作步骤简单，软件参数设置简单，容易掌握，上手快。

缺点：只适用于Windows系统，只支持WOS核心合集里面的数据。

Histcite有以下5个功能：

- 01 | 快速绘出一个领域的发展脉络
- 02 | 快速锁定该领域的重要文献
- 03 | 快速锁定该领域的重要科学家，机构
- 04 | 洞察该领域的最新进展
- 05 | 找出无指定关键词的重要文献



Histcite中的几个重要参数：

GCS: Global citation score总引用次数，WOS上看到的引用次数，表示某篇文章被整个WOS数据库中的文献所引用的次数，有些引用这篇参考文献的文章可能和你的研究方向毫无关系，但GCS还是会把这个数据记录下来。



LCS: Local citation score本地引用次数，某篇文献在当前数据集中被引用的次数。因为你导入Histcite的文章都是和你检索词有关的，可以认为这些文章是你的研究方向相关，因此如果某篇文献LCS值很高，意味着它肯定是你研究领域的重要文献。因此，LCS比GCS更重要，LCS高的极有可能是你领域内的开创性文章。



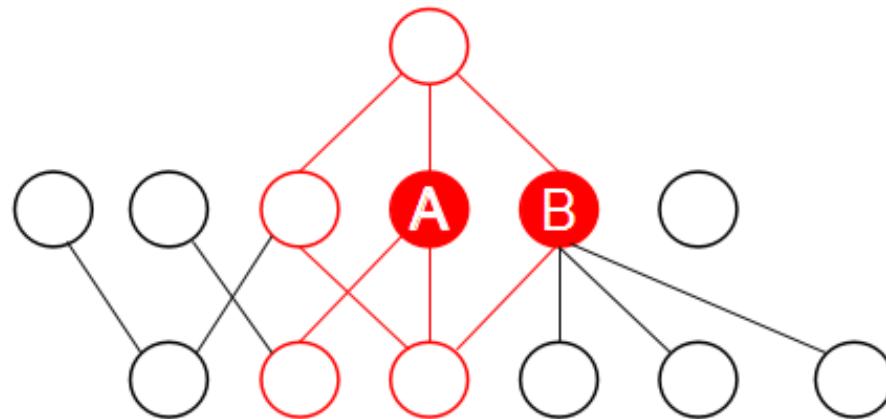
CR: Cited references 某篇文献在整个WOS数据中的参考文献数，这个值越高，说明这篇文章很可能 是综述文章，可根据该值的排序，快速定位综述。



LCR: Local Cited references 本地参考文献数，指某篇文献引用的所有参考文献中，有多少篇在当前数据集中。根据LCR值的排序，可以迅速定位近期关注该领域的重要文献。LCR高的极有可能是你领域内的综述文章。如果某篇文章的LCR为10，则表示有10篇该文章的参考文献在当前本地数据集中。



GCS&LCS

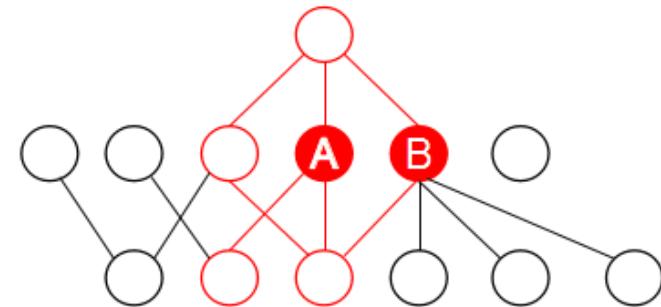


GCS: A文章发表之后被引用2次
B文章发表之后被引用4次

A、B文章的LCS值分别是 () 、 () 。

- A 3
- B 4
- C 2
- D 1

GCS&LCS



GCS: A文章发表之后被引用2次
B文章发表之后被引用4次

提交



Histcite如何判断文章的重要性：

不是看大家的意见，而是看这个领域同行专家的意见。

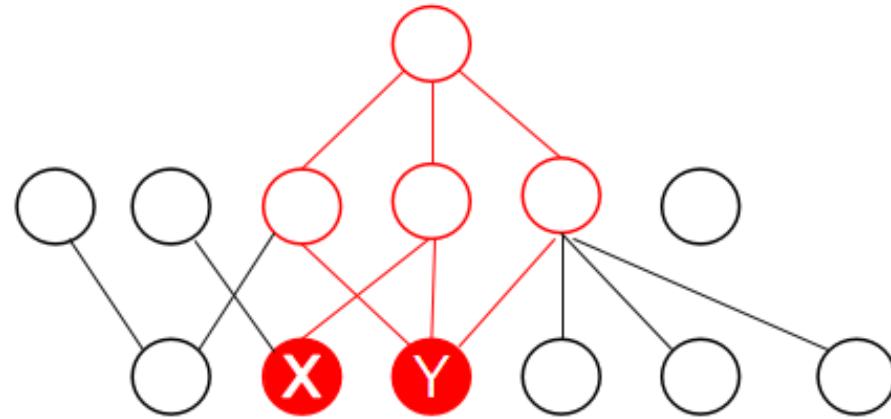


Histcite又如何判断最新发表文章的重要性：

主要看这篇新文章所引用的参考文献。



CR&LCR

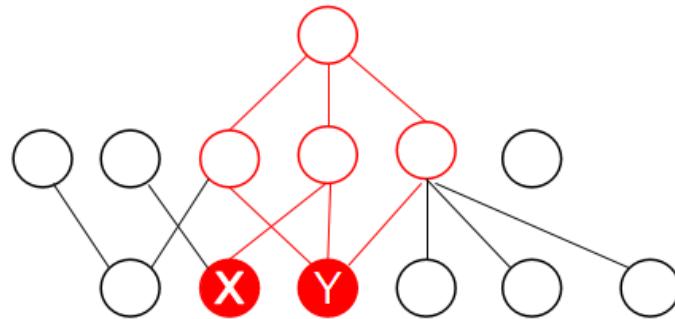


CR: X文章引用了2篇参考文献
Y文章引用了3篇参考文献

X、Y文章的LCR值分别是 () 、 () 。

- A 1
- B 2
- C 3
- D 4

CR&LCR



CR: X文章引用了2篇参考文献
Y文章引用了3篇参考文献

提交



LCS可以快速定位一个领域内的经典文献，LCR可快速找出最新文献中哪些是和自己研究方向最相关的文献。



二、数据下载 From WOS 核心 合集



第一步：在SCI数据库中进行检索

Clarivate

Web of Science™ 检索 标记结果列表 历史 跟踪服务 简体中文 产品 登录 注册

探索跨学科内容
来自最值得您信赖的全球引文数据库

选择数据库 Web of Science 核心合集 引文索引 Science Citation Index Expanded (SCI-EXPANDED)–2002–至今

文献 作者 被引参考文献 化学结构

主题 cold-pcr

+ 添加行 + 添加日期范围 高级检索 × 清除 检索



第二步：导出检索数据

The screenshot shows the Web of Science search results page for the query "cold-pcr (主题)". The results list 184 items from the Science Citation Index Expanded (SCI-Expanded). A modal dialog box is open, titled "将记录导出为纯文本文件" (Export results as plain text file). The dialog box contains the following settings:

- 记录选项: 记录: 1 至 184
- 一次不能超过 500 条记录 (highlighted with a red box)
- 记录内容: 全记录与引用的参考文献 (highlighted with a red box)
- 导出 (Export) and 取消 (Cancel) buttons

The background search results page shows the first item: "CR Sequencing for Early" by Wong, DK, et al. (Sep 2014). The results list includes various filters and a sidebar for related records.



HistCite Pro下载地址:

<https://pan.baidu.com/s/1hsIwJzQ>



File Analyses View Tools Help

Untitled Collection

List of All Records

HistCite™

HistCite™

Grand Totals: LCS 589, GCS 3494, CR 5228

Collection span: 2002 - 2021

Records: 183, Authors: 945, Journals: 92, Cited References: 3357, Words: 682

[Yearly output](#) | [Document Type](#) | [Language](#) | [Institution](#) | [Institution with Subdivision](#) | [Country](#)

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#	Date / Author / Journal	LCS	GCS	LCR	CR
1	144 Mauger F, How-Kit A, Tost J COLD-PCR Technologies in the Area of Personalized Medicine: Methodology and Applications MOLECULAR DIAGNOSIS & THERAPY. 2017 JUN; 21 (3): 269-283	7	16	33	81
2	59 Castellanos-Rizaldos E, Liu PF, Milbury CA, Guha M, Brisci A, et al. Temperature-Tolerant COLD-PCR Reduces Temperature Stringency and Enables Robust Mutation Enrichment CLINICAL CHEMISTRY. 2012 JUL; 58 (7): 1130-1138	0	25	17	30
3	69 Castellanos-Rizaldos E, Milbury CA, Makrigiorgos GM Enrichment of Mutations in Multiple DNA Sequences Using COLD-PCR in Emulsion PLOS ONE. 2012 DEC 6; 7 (12): Art. No. e51362	0	6	14	33
4	175 Mortazavipour MM, Shahbazi S, Mahdian R Detection of Paternal IVS-II-1 (G>A) (HBB: c.315+1G>A) Mutation in Cell-Free Fetal DNA Using COLD-PCR assay HEMOGLOBIN. 2020 MAY 3; 44 (3): 168-173	0	0	12	27
5	122 Castellanos-Rizaldos E, Richardson K, Lin R, Wu G, Makrigiorgos MG Single-Tube, Highly Parallel Mutation Enrichment in Cancer Gene Panels by Use of Temperature-Tolerant COLD-PCR CLINICAL CHEMISTRY. 2015 JAN; 61 (1): 267-277	0	7	11	38
6	34 Milbury CA, Li J, Liu PF, Makrigiorgos GM COLD-PCR: improving the sensitivity of molecular diagnostics assays EXPERT REVIEW OF MOLECULAR DIAGNOSTICS. 2011 MAR; 11 (2): 159-169	18	33	10	43
7	35 Milbury CA, Chen CC, Mamon H, Liu PF, Santagata S, et al. Multiplex Amplification Coupled with COLD-PCR and High Resolution Melting Enables Identification of Low-Abundance Mutations in Cancer Samples with Low DNA Content JOURNAL OF MOLECULAR DIAGNOSTICS. 2011 MAR; 13 (2): 220-232	12	25	10	62
8	53 Milbury CA, Correll M, Quackenbush J, Rubio R, Makrigiorgos GM COLD-PCR Enrichment of Rare Cancer Mutations prior to Targeted Amplicon Resequencing CLINICAL CHEMISTRY. 2012 MAR; 58 (3): 580-589	23	47	10	45
9	124 Castellanos-Rizaldos E, Paweletz C, Song C, Oxnard GR, Mamon H, et al. Enhanced Ratio of Signals Enables Digital Mutation Scanning for Rare Allele Detection JOURNAL OF MOLECULAR DIAGNOSTICS. 2015 MAY; 17 (3): 284-292	0	23	10	37
10	33 Milbury CA, Li J, Makrigiorgos GM Ice-COLD-PCR enables rapid amplification and robust enrichment for low-abundance unknown DNA mutations NUCLEIC ACIDS RESEARCH. 2011 JAN; 39 (1): Art. No. e2	0	79	9	24



2、重要化

#	Author	Recs	TLCS	TGCS
1	Makrigiorgos GM	30	330	1057
2	Richardson K	14	6	53
3	Wu G	13	0	11
4	Li J	12	269	741
5	Milbury CA	12	157	475
6	Castellanos-Rizaldos E	11	2	121
7	Eastlake P	11	6	46
8	Ferrari M	10	38	133
9	Tost J	10	53	257
10	Cremonesi L	9	37	122
11	Legendre B	9	0	3
12	Galbiati S	8	38	108
13	How-Kit A	8	19	210
14	Mamon H	8	101	411
15	Cubrich C	7	6	46
16	Brisci A	6	26	96
17	Lin R	6	0	8
18	Luthra R	6	38	114
19	Mancini I	6	43	147
20	Damin F	5	17	58
21	Daunay A	5	23	80
22	Guerrero JM	5	2	35
23	Liu PF	5	45	116
24	Macher HC	5	2	35

#	Author	Recs	TLCS	TGCS
1	Makrigiorgos GM	30	330	1057
2	Li J	12	269	741
3	Milbury CA	12	157	475
4	Wang LL	5	112	326
5	Mamon H	8	101	411
6	Berboco R	2	89	280
7	Kulke MH	2	89	321
8	Tost J	10	53	257
9	Liu PF	5	45	116
10	Mancini I	6	43	147
11	Pinzani P	5	40	132
12	Orlando C	3	39	103
13	Pratesi N	2	39	90
14	Santucci C	2	39	90
15	Simi L	4	39	128
16	Ferrari M	10	38	133
17	Galbiati S	8	38	108
18	Luthra R	6	38	114
19	Cremonesi L	9	37	122
20	Sestini R	3	32	65
21	Li C	1	30	44
22	Cianchi F	1	28	52
23	Valanzano R	1	28	52



3、核心期刊

File Analyses View Tools Untitled Collection Journal List (92)

Records: 183, Authors: 945, Journals: 92, Cited References: 3357, Words: 682
[Yearly output](#) | [Document Type](#) | [Language](#) | [Institution](#) | [Institution with Subdivision](#) | [Country](#)

#	Journal	Recs	TLCS	TGCS
1	JOURNAL OF MOLECULAR MEDICINE	15	153	519
2	CLINICAL CHEMISTRY	1	89	280
3	CANCER RESEARCH	7	75	173
4	PLOS ONE	16	45	157
5	HUMAN MUTATION	5	39	160
6	JOURNAL OF CLINICAL MICROBIOLOGY	1	35	53

File Analyses View Tools Help Untitled Collection List Totals: LCS 89, GCS 280, CR 27

List of 1 Records for Journal **NATURE MEDICINE**

Records: 183, Authors: 945, Journals: 92, Cited References: 3357, Words: 682
[Yearly output](#) | [Document Type](#) | [Language](#) | [Institution](#) | [Institution with Subdivision](#) | [Country](#)

#	Date / Author / Journal	LCS	GCS	LCR	CR
1	4 LI J, Wang LL, Mamon H, Kulke MH, Berbeco R, et al. Replacing PCR with COLD-PCR enriches variant DNA sequences and redefines the sensitivity of genetic testing <i>NATURE MEDICINE</i> . 2008 MAY; 14 (5): 579-584	89	280	0	27

14	NUCLEIC ACIDS RESEARCH	14	LEUKEMIA	1	6	43
15	ANALYTICAL AND BIOANALYTICAL CHEMISTRY	15	JOURNAL OF CLINICAL MICROBIOLOGY	2	5	17
16	APPLIED IMMUNOHISTOCHEMISTRY & MOLECULAR MORPHOLOGY	16	JOURNAL OF INVESTIGATIVE DERMATOLOGY	1	5	84
17	DIAGNOSTIC MOLECULAR PATHOLOGY	17	ANALYTICAL AND BIOANALYTICAL CHEMISTRY	2	3	36
18	EUROPEAN RESPIRATORY JOURNAL	18	CLINICAL CHEMISTRY AND LABORATORY MEDICINE	3	3	8
19	HUMAN PATHOLOGY	19	EXPERT OPINION ON BIOLOGICAL THERAPY	1	3	9
20	INTERNATIONAL JOURNAL OF CLINICAL MICROBIOLOGY	20	LAB ON A CHIP	2	3	368
21	JOURNAL OF CLINICAL MICROBIOLOGY	21	UROLOGIA INTERNATIONALIS	1	3	6
22	JOURNAL OF HEPATOLOGY	22	APPLIED IMMUNOHISTOCHEMISTRY & MOLECULAR MORPHOLOGY	2	2	3
23	JOURNAL OF THORACIC AND CARDIOVASCULAR SURGERY	23	CLINICAL BIOCHEMISTRY	1	2	5
24	LAB ON A CHIP	24	EUROPEAN RESPIRATORY JOURNAL	2	2	53
25	INTERNATIONAL JOURNAL OF CANCER	25	INTERNATIONAL JOURNAL OF CANCER	1	2	105



4、参考文献分析

排在前面的就是被特定领域学者反复引用的重要文献。后面带有+号的表示本地数据集中没有包含，这些往往就是被遗漏的重要文献。

File Analyses View Tools Help

Untitled Collection

Cited Reference List (3357) including 66 records, 32 on this page (Hide 32 records)

Records: 183, Authors: 945, Journals: 92 Cited References: 3357, Words: 682

Yearly output | Document Type | Language | Institution | Institution with Subdivision | Country

| < << < > >> > |

#	Author / Year / Journal	Recs
1	Li J, 2008, NAT MED, V14, P579, DOI 10.1038/nm1708	WoS 89
2	Milbury CA, 2011, NUCLEIC ACIDS RES, V39, DOI 10.1093/nar/gkq899	+ WoS 39
3	Li J, 2009, BIOCHEM SOC T, V37, P427, DOI 10.1042/BST0370427	WoS 35
4	Milbury CA, 2009, CLIN CHEM, V55, P2130, DOI 10.1373/clinchem.2009.131029	WoS 31
5	Li J, 2009, HUM MUTAT, V30, P1583, DOI 10.1002/humu.21112	WoS 30
6	Mancini I, 2010, J MOL DIAGN, V12, P705, DOI 10.2353/jmoldx.2010.100018	WoS 28
7	Milbury CA, 2009, CLIN CHEM, V55, P632, DOI 10.1373/clinchem.2008.113035	WoS 28
8	Zuo Z, 2009, MODERN PATHOL, V22, P1023, DOI 10.1038/modpathol.2009.59	WoS 26
9	Li J, 2009, CLIN CHEM, V55, P748, DOI 10.1373/clinchem.2008.113381	WoS 23
10	Milbury CA, 2012, CLIN CHEM, V58, P580, DOI 10.1373/clinchem.2011.176198	WoS 23
11	Galbiati S, 2011, CLIN CHEM, V57, P136, DOI 10.1373/clinchem.2010.155671	WoS 18
12	Kit AH, 2013, HUM MUTAT, V34, P1568, DOI 10.1002/humu.22427	WoS 18
13	Milbury CA, 2011, EXPERT REV MOL DIAGN, V11, P159, DOI [10.1586/erm.10.115, 10.1586/ERM.10.115]	WoS 18
14	Castellanos-Rizaldos E, 2012, CLIN CHEM, V58, P1130, DOI 10.1373/clinchem.2012.183095	+ WoS 17
15	Diehl F, 2008, NAT MED, V14, P985, DOI 10.1038/nm.1789	+ WoS 16
16	Ogino S, 2005, J MOL DIAGN, V7, P413, DOI 10.1016/S1525-1578(10)60571-5	+ WoS 16
17	Vogelstein B, 1999, P NATL ACAD SCI USA, V96, P9236, DOI 10.1073/pnas.96.16.9236	+ WoS 16
18	Kristensen LS, 2010, HUM MUTAT, V31, P1366, DOI 10.1002/humu.21358	WoS 15
19	Lo YMD, 1997, LANCET, V350, P485, DOI 10.1016/S0140-6736(97)02174-0	+ WoS 15
20	Song C, 2011, DIAGN MOL PATHOL, V20, P81, DOI 10.1097/PDM.0b013e3181fde92f	WoS 15
21	Pritchard CC, 2010, BMC CLIN PATHOL, V10, DOI 10.1186/1472-6890-10-6	+ WoS 14
22	Amado RG, 2008, J CLIN ONCOL, V26, P1626, DOI 10.1200/JCO.2007.14.7116	+ WoS 13
23	Delaney D, 2009, MODERN PATHOL, V22, P718, DOI 10.1038/modpathol.2009.32	WoS 12



5、关键词分析

辨析研究热点，拓展检索范围，调整检索策略，
了解交叉学科，激发研究新思路。

File Analyses View Tools Help

Untitled Collection

Word(i) List (682) Word count: 2380, All words count: 3183

Records: 183, Authors: 945, Journals: 92, Cited References: 3357, Words: 682

Yearly output | Document Type | Language | Institution | Institution with Subdivision | Country

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#	Word	Recs	TLCS	GCS
1	PCR	113	531	1741
2	COLD	99	470	1399
3	DETECTION	77	282	1691
4	MUTATIONS	81	236	1347
5	MUTATION	48	160	935
6	KRAS	43	144	954
7	CANCER	7	142	399
8	DNA	42	137	698
9	ANALYSIS	13	128	378
10	USING	10	127	450
11	TEMPERATU	38	122	889
12	ENRICHME	21	113	337
13	SENSITIVE	18	109	276
14	CELL	14	109	277
15	HIGH	27	91	390
16	ICE	2	89	280
17	PATIENTS	2	89	280
18	COLORECTA	2	89	280
19	DENATURAT	4	89	287
20	LOWER	2	89	280
21	SEQUENCIN	22	87	433
22	MELTING	17	86	374
23	SAMPLES	11	85	669
24	LOW	13	78	176



File Analyses Vi File Analyses View Tools Help

Untitled Coll

Publication Year

Records: 183, Authors:

Yearly output Document

Untitled Collection

Publication Year List (15: 2002 - 2021) Histogram

Records: 183, Authors: 945, Journals: 92, Cited References: 3357, Words: 682

[Yearly output](#) | [Document Type](#) | [Language](#) | [Institution](#) | [Institution with Subdivision](#) | [Country](#)

#	Publication
1	2002
2	2008
3	2009
4	2010
5	2011
6	2012
7	2013
8	2014
9	2015
10	2016
11	2017
12	2018
13	2019
14	2020
15	2021

#	Publication Year	Recs	TLCS	TGCS
1	2009	14	200	514
2	2008	3	89	280
3	2011	17	80	775
4	2012	24	60	343
5	2010	10	51	116
6	2013	19	30	442
7	2016	10	25	168
8	2014	32	24	302
9	2017	9	17	162
10	2015	9	5	132
11	2018	12	4	116
12	2019	12	4	77
13	2002	2	0	48
14	2020	7	0	19
15	2021	3	0	0

6、年份分



7、文献类型分析

File Analyses View Tools Help

Untitled Collection

Document Type List (6)

Records: 183, Authors: 945, Journals: 92, Cited References: 3357, Words: 682

[Yearly output](#) | [Document Type](#) | [Language](#) | [Institution](#) | [Institution with Subdivision](#) | [Country](#)

#	Document Type	Recs	TLCS	TGCS
1	Article	114	479	2913
2	Meeting Abstract	45	4	13
3	Review	12	57	395
4	Letter	6	29	99
5	Article; Proceedings Paper	3	1	13
6	Editorial Material	3	19	61



8、文献所用的语言类型分析

File Analyses View Tools Help

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Document Language List (1)

Records: 183, Authors: 945, Journals: 92, Cited References: 3357, Words: 682

[Yearly output](#) | [Document Type](#) **Language** [Institution](#) | [Institution with Subdivision](#) | [Country](#)

#	Language	Recs	TLCS	TGCS
1	English	183	589	3494



File Analyses View Tools Help

Untitled Collection

Institution List (301)

Records: 183, Authors: 945, Journals: 92, Cited References: 3357, Words: 61
Yearly output | Document Type | Language | **Institution** | Institution with Sub

|< << < > >> >|

#	Institution	Recs	TLC
1	Harvard Univ	24	31
2	Transgenomic Inc	14	
3	Univ Texas MD Anderson Canc Ctr	13	4
4	Dana Farber Canc Inst	9	6
5	Univ Vita Salute San Raffaele	9	3
6	CEA	8	4
7	Ist Sci San Raffaele	8	2
8	Univ Florence	8	4
9	INSERM	7	1
10	CNR	6	1
11	Fdn Jean Dausset CEPH	6	3
12	Diagnost & Ric San Raffaele SpA	5	2
13	Univ Paris 05	5	2
14	Univ Seville	5	
15	Hop St Louis	4	
16	Aarhus Univ Hosp	3	1
17	Brigham & Womens Hosp	3	1
18	Fujian Med Univ	3	
19	Harvard Med Sch	3	
20	Sun Yat Sen Univ	3	
21	Univ Calif San Diego	3	
22	Univ Paris 06	3	1
23	Univ Paris Diderot	3	2

File Analyses View Tools Help

Untitled Collection

Institution with Subdivision List (437)

Records: 183, Authors: 945, Journals: 92, Cited References: 3357, Words: 682

Yearly output | Document Type | Language | Institution | **Institution with Subdivision** | Country

|< << < > >> >|

#	Institution with Subdivision	Recs	TLCS	TGCS
1	Harvard Univ, Sch Med	22	277	923
2	Transgenomic Inc	13	6	54
3	Univ Vita Salute San Raffaele	9	38	133
4	Dana Farber Canc Inst	8	30	74
5	Harvard Univ, Brigham & Womens Hosp	7	52	125
6	Univ Texas MD Anderson Canc Ctr	7	1	37
7	CNR, Ist Chim Riconoscimento Mol	5	17	58
8	Diagnost & Ric San Raffaele SpA	5	25	98
9	CEA, Inst Genom	4	26	67
10	Fdn Jean Dausset CEPH, Lab Funct Genom	4	23	125
11	Ist Sci San Raffaele, Ctr Translat Genom & Bioinformat	4	3	47
12	Aarhus Univ Hosp, Dept Pathol	3	19	56
13	Brigl, Inst Genom	3	20	47
14	CNR	3	3	11
15	Fujian Med Univ, Affiliated Hosp 1	3	25	58
16	Ist Sci San Raffaele, Genom Unit Diag Human Pathol	3	2	10
17	Sun Yat Sen Univ, Affiliated Hosp 3	3	39	103
18	Univ Paris 05	3	5	111
19	Univ Seville, Univ Hosp Virgen del Rocio Seville	3	1	13
20	CEA Inst Genom, Ctr Natl Genotypage	2	7	137
21	Chinese Acad Sci, Beijing Inst Genom	2	2	10
22	Grp Hosp Pitie Salpetriere, AP HP	2	13	118
23	Harvard Med Sch, Brigham & Womens Hosp	2	0	41



File Analyses View Tools Help

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Country List (28)

Records: 183, Authors: 945, Journals: 92, Cited References: 3357, Words: 682
Yearly output | Document Type | Language | Institution | Institution with Subdivision

Country

#	Country	Recs	TLCS	TGCS
1	USA	74	382	1854
2	Italy	25	95	346
3	Peoples R China	24	34	177
4	France	19	66	801
5	Spain	14	6	241
6	UK	10	21	276
7	Japan	6	4	61
8	Germany	4	4	357
9	Netherlands	4	3	190
10	Canada	3	1	76
11	Czech Republic	3	0	106
12	Denmark	3	19	56
13	Iran	3	1	1
14	Singapore	3	4	29
15	India	2	0	27
16	Taiwan	2	0	23
17	Albania	1	0	5
18	Australia	1	0	17
19	Cyprus	1	0	3
20	Iraq	1	0	1
21	Nigeria	1	0	1
22	Norway	1	0	6
23	Poland	1	0	3

10、国家列表



其他参数

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List of All References
Custom

Records: 183, Authors: 92, Cited References: 3357, Words: 682

Yearly output | Document Type | Language | Institution | Institution with Subdivision | Country

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Grand Totals: LCS 589, LCSx 414, GCS 3494, OCS n/a, CR 5228, NA 1366
Means: LCS 3.22, LCSx 2.26, GCS 19.09, OCS n/a, CR 28.57, NA 7.46
Collection span: 2002 - 2021 (20 years)

#	Date / Author / Journal	LCS	LCS/t	LCSx	GCS	GCS/t	OCS	NA	LCR	CR	LCSb	LCSe	LCS(e/b)
1	144 Mauger F, How-Kit A, Tost J COLD-PCR Technologies in the Area of Personalized Medicine: Methodology and Applications MOLECULAR DIAGNOSIS & THERAPY. 2017 JUN; 21 (3): 269-283	7	1.40	6	16	3.20		3	33	81			
2	59 Castellanos-Rizaldos E, Liu PF, Milbury CA, Guha M, Brisci A, et al. Temperature-Tolerant COLD-PCR Reduces Temperature Stringency and Enables Robust Mutation Enrichment CLINICAL CHEMISTRY. 2012 JUL; 58 (7): 1130-1138	0	0.00	0	25	2.50		9	17	30	0	0	0
3	69 Castellanos-Rizaldos E, Milbury CA, Makrigiorgos GM Enrichment of Mutations in Multiple DNA Sequences Using COLD-PCR in Emulsion PLOS ONE. 2012 DEC 6; 7 (12): Art. No. e51362	0	0.00	0	6	0.60		3	14	33	0	0	0
4	175 Mortazavipour MM, Shahbazi S, Mahdian R Detection of Paternal IVS-II-1 (G>A) (HBB: c.315+1G>A) Mutation in Cell-Free Fetal DNA Using COLD-PCR assay HEMOGLOBIN. 2020 MAY 3; 44 (3): 168-173	0	0.00	0	0	0.00		3	12	27			
5	122 Castellanos-Rizaldos E, Richardson K, Lin R, Wu G, Makrigiorgos MG Single-Tube, Highly Parallel Mutation Enrichment in Cancer Gene Panels by Use of Temperature-Tolerant COLD-PCR CLINICAL CHEMISTRY. 2015 JAN; 61 (1): 267-277	0	0.00	0	7	1.00		5	11	38	0	0	0
6	34 Milbury CA, Li J, Liu PF, Makrigiorgos GM COLD-PCR: improving the sensitivity of molecular diagnostics assays EXPERT REVIEW OF MOLECULAR DIAGNOSTICS. 2011 MAR; 11 (2): 159-169	18	1.64	14	33	3.00		4	10	43	6	3	0.50
7	35 Milbury CA, Chen CC, Mamon H, Liu PF, Santagata S, et al. Multiplex Amplification Coupled with COLD-PCR and High Resolution Melting Enables Identification of Low-Abundance Mutations in Cancer Samples with Low DNA Content JOURNAL OF MOLECULAR DIAGNOSTICS. 2011 MAR; 13 (2): 220-232	12	1.09	7	25	2.27		6	10	62	6	1	0.17
8	53 Milbury CA, Correll M, Quackenbush J, Rubio R, Makrigiorgos GM COLD-PCR Enrichment of Rare Cancer Mutations prior to Targeted Amplicon Resequencing CLINICAL CHEMISTRY. 2012 MAR; 58 (3): 580-589	23	2.30	15	47	4.70		5	10	45	11	3	0.27
9	124 Castellanos-Rizaldos E, Paweletz C, Song C, Oxnard GR, Mamon H, et al. Enhanced Ratio of Signals Enables Digital Mutation Scanning for Rare Allele Detection JOURNAL OF MOLECULAR DIAGNOSTICS. 2015 MAY; 17 (3): 284-292	0	0.00	0	23	3.29		7	10	37	0	0	0
10	33 Milbury CA, Li J, Makrigiorgos GM Ice-COLD-PCR enables rapid amplification and robust enrichment for low-abundance unknown DNA mutations NUCLEIC ACIDS RESEARCH. 2011 JAN; 39 (1): Art. No. e2	0	0.00	0	79	7.18		3	9	24	0	0	0



LCS/t: 表示平均每年被引用多少次，LCS/t值越高，说明该文章每年都被大量引用，生命力强。

LCSx: 带x的表示去掉自引，LCSx值越高，说明该文章被同行认可度越高。

NA: 作者数，这篇文章有多少个作者。

LCSb: b表示begin;

LCSe: e表示end;

LCS (e/b) : LCSe和LCSb两个数的比值。LCSb表示文章发表头3年被引用的次数，LCSe表示文章最近3年被引用的次数，LCS (e/b) 越高表示该文章最近几年受关注程度越高。



四、引文编年图的绘制

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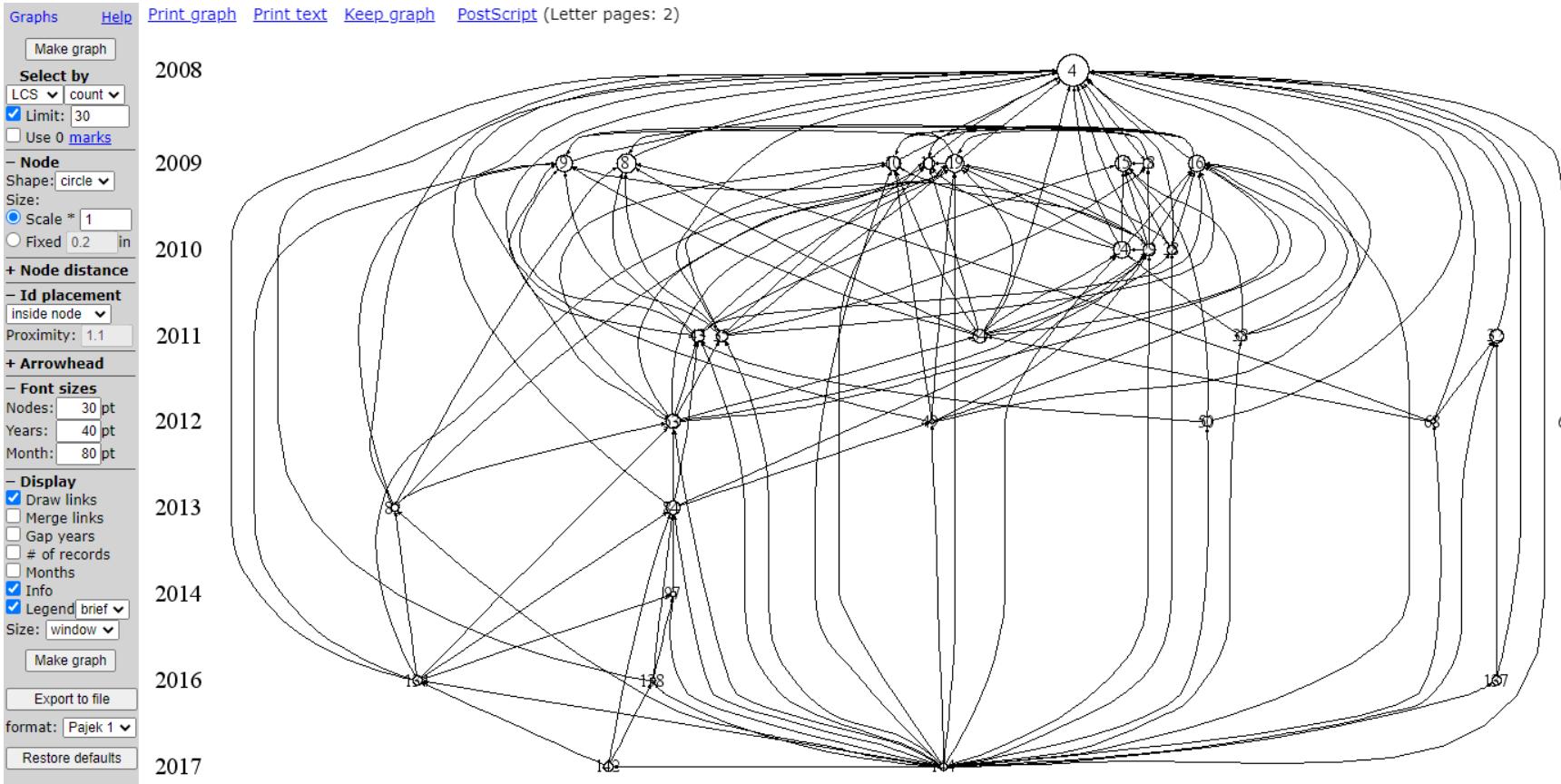
List of All Records

Records: 183, Authors: 193, References: 3357, Words: 682

Yearly output | Document | Institution | Institution with Subdivision | Country

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#	Date / Author / Journal
1	144 Mauger F. COLD-PCR Technologies in the Area of Personalized Medicine: Methodology and Applications. MOLECULAR DIAGNOSIS & THERAPY. 2017 JUN; 21 (3): 269-283
2	59 Castellanos-Rizaldos E, Liu PF, Milbury CA, Guha M, Brisci A, et al. Temperature-Tolerant COLD-PCR Reduces Temperature Stringency and Enables Robust Mutation Enrichment. CLINICAL CHEMISTRY. 2012 JUL; 58 (7): 1130-1138
3	69 Castellanos-Rizaldos E, Milbury CA, Makrigiorgos GM. Enrichment of Mutations in Multiple DNA Sequences Using COLD-PCR in Emulsion. PLOS ONE. 2012 DEC 6; 7 (12): Art. No. e51362
4	175 Mortazavipour MM, Shahbazi S, Mahdian R. Detection of Paternal IVS-II-1 (G>A) (HBc: c.315+1G>A) Mutation in Cell-Free Fetal DNA Using COLD-PCR assay. HEMOGLOBIN. 2020 MAY 3; 44 (3): 168-173
5	122 Castellanos-Rizaldos E, Richardson K, Lin R, Wu G, Makrigiorgos MG. Single-Tube, Highly Parallel Mutation Enrichment in Cancer Gene Panels by Use of Temperature-Tolerant COLD-PCR. CLINICAL CHEMISTRY. 2015 JAN; 61 (1): 267-277
6	34 Milbury CA, Li J, Liu PF, Makrigiorgos GM. COLD-PCR: improving the sensitivity of molecular diagnostics assays. EXPERT REVIEW OF MOLECULAR DIAGNOSTICS. 2011 MAR; 11 (2): 159-169
7	35 Milbury CA, Chen CC, Mamon H, Liu PF, Santagata S, et al. Multiplex Amplification Coupled with COLD-PCR and High Resolution Melting Enables Identification of Low-Abundance Mutations in Cancer Samples with Low DNA Content. JOURNAL OF MOLECULAR DIAGNOSTICS. 2011 MAR; 13 (2): 220-232
8	53 Milbury CA, Correll M, Quackenbush J, Rubio R, Makrigiorgos GM. COLD-PCR Enrichment of Rare Cancer Mutations prior to Targeted Amplicon Resequencing. CLINICAL CHEMISTRY. 2012 MAR; 58 (3): 580-589
9	124 Castellanos-Rizaldos E, Pawletz C, Song C, Oxnard GR, Mamon H, et al. Enhanced Ratio of Signals Enables Digital Mutation Scanning for Rare Allele Detection. JOURNAL OF MOLECULAR DIAGNOSTICS. 2015 MAY; 17 (3): 284-292
10	33 Milbury CA, Li J, Makrigiorgos GM. Ice-COLD-PCR enables rapid amplification and robust enrichment for low-abundance unknown DNA mutations. NUCLEIC ACIDS RESEARCH. 2011 JAN; 39 (1): Art. No. e2



图上有30个圆圈，每个圆圈表示一篇文献，中间的数字是这篇文献在数据集中的序号。圆圈越大表示被引次数越多，关注度越高。



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Records: 183, Authors: 945, Journals: 92, Cited References: 3357, Words: 68, Marks: 5

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OR

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Description:

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#	Date / Author / Journal	LCS	GCS	LCR	CR	CS	LCR	CR
1	4 Li J, Wang LL, Mamon H, Kulke MH, Berboco R, et al. Replacing PCR with COLD-PCR enriches variant DNA sequences and redefines the sensitivity of genetic testing NATURE MEDICINE. 2008 MAY; 14 (5): 579-584	89	280	0	27	80	0	27
2	8 Li J, Makrigiorgos GM. COLD-PCR: a new platform for highly improved mutation detection in cancer and genetic testing BIOCHEMICAL SOCIETY TRANSACTIONS. 2009 APR; 37: 427-432	35	53	1	15	53	1	15
3	19 Milbury CA, Li J, Makrigiorgos GM. COLD-PCR-Enhanced High-Resolution Melting Enables Rapid and Selective Identification of Low-Level Unknown Mutations CLINICAL CHEMISTRY. 2009 DEC; 55 (12): 2130-2143	31	60	4	33	60	4	33
4	16 Li J, Milbury CA, Li C, Makrigiorgos GM. Two-Round Coamplification at Lower Denaturation Temperature-PCR (COLD-PCR)-Based Sanger Sequencing Identifies a Novel Spectrum of Low-Level Mutations in Lung Adenocarcinoma HUMAN MUTATION. 2009 NOV; 30 (11): 1583-1590	30	44	5	41	44	5	41
5	9 Milbury CA, Li J, Makrigiorgos GM. PCR-Based Methods for the Enrichment of Minority Alleles and Mutations CLINICAL CHEMISTRY. 2009 APR; 55 (4): 632-640	28	130	1	41	30	1	41
6	24 Mancini I, Santucci C, Sestini R, Simi L, Pratesi N, et al. The Use of COLD-PCR and High-Resolution Melting Analysis Improves the Limit of Detection of KRAS and BRAF Mutations in Colorectal Cancer JOURNAL OF MOLECULAR DIAGNOSTICS. 2010 SEP; 12 (5): 705-711	28	52	5	40	52	5	40
7	15 Zuo Z, Chen SS, Chandra PK, Galbincea JM, Soape M, et al. Application of COLD-PCR for improved detection of KRAS mutations in clinical samples MODERN PATHOLOGY. 2009 AUG; 22 (8): 1023-1031	26	90	1	26	90	1	26
8	10 Li J, Wang LL, Janne PA, Makrigiorgos GM. Coamplification at Lower Denaturation Temperature-PCR Increases Mutation-Detection Selectivity of TaqMan-Based Real-Time PCR CLINICAL CHEMISTRY. 2009 APR; 55 (4): 748-756	23	46	1	35	46	1	35
	Coamplification at Lower Denaturation Temperature-PCR Increases Mutation-Detection Selectivity of TaqMan-Based Real-Time PCR EXPERT REVIEW OF MOLECULAR DIAGNOSTICS. 2011 MAR; 11 (2): 159-169							



2. 文献导出

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HTML presentation
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Cited References: 3357, Words: 682, Marks: 5
Document Type | Language | Institution | Institution with Subdivision | Country

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 Records citing selected records
 Records cited by selected records

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Description:
Tag Untag Remove All Tags

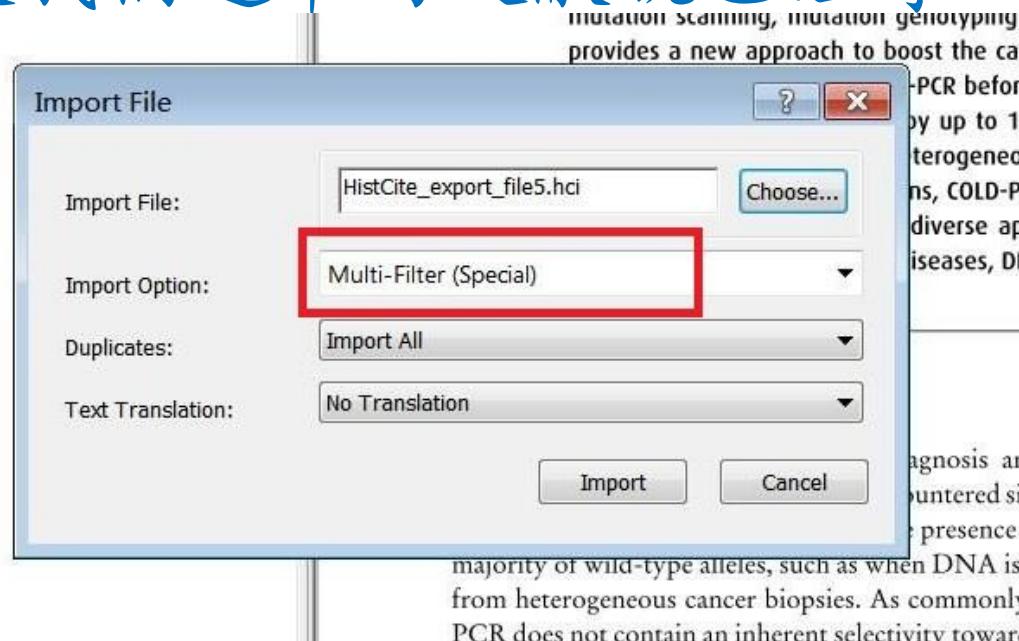
Date / Author / Journal LCS GCS LCR CR

#	Date / Author / Journal	LCS	GCS	LCR	CR
1	2008 4 Li J, Wang LL, Mamon H, Kulke MH, Berbeco R, et al. Replacing PCR with COLD-PCR enriches variant DNA sequences and redefines the sensitivity of genetic testing NATURE MEDICINE. 2008 MAY; 14 (5): 579-584	89	280	0	27
2	2009 8 Li J, Makrigiorgos GM COLD-PCR: a new platform for highly improved mutation detection in cancer and genetic testing BIOCHEMICAL SOCIETY TRANSACTIONS. 2009 APR; 37: 427-432	35	53	1	15
3	2009 9 Milbury CA, Li J, Makrigiorgos GM PCR-Based Methods for the Enrichment of Minority Alleles and Mutations CLINICAL CHEMISTRY. 2009 APR; 55 (4): 632-640	28	130	1	41
4	2009 16 Li J, Milbury CA, Li C, Makrigiorgos GM Two-Round Coamplification at Lower Denaturation Temperature-PCR (COLD-PCR)-Based Sanger Sequencing Identifies a Novel Spectrum of Low-Level Mutations in Lung Adenocarcinoma HUMAN MUTATION. 2009 NOV; 30 (11): 1583-1590	30	44	5	41
5	2009 19 Milbury CA, Li J, Makrigiorgos GM COLD-PCR-Enhanced High-Resolution Melting Enables Rapid and Selective Identification of Low-Level Unknown Mutations CLINICAL CHEMISTRY. 2009 DEC; 55 (12): 2130-2143	31	60	4	33



3. 导入Endnote

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Author	Year	Title	Rating	Journal	Last Updated	Reference Type
Li, J.; Makrigiorg...	2009	COLD-PCR: a new platform for highly improved...	• • • •	Biochemical So...	2021/11/2	Journal Article
Li, J.; Milbury, C. ...	2009	Two-Round Coamplification at Lower Denatura...	• • • •	Human Mutation	2021/11/2	Journal Article
Li, J.; Wang, L. L.; ...	2008	Replacing PCR with COLD-PCR enriches variant...	• • • •	Nature Medicine	2021/11/2	Journal Article
Milbury, C. A.; Li, ...	2009	COLD-PCR-Enhanced High-Resolution Melting ...	• • • •	Clinical Chemistry	2021/11/2	Journal Article
Milbury, C. A.; Li, ...	2009	PCR-Based Methods for the Enrichment of Min...	• • • •	Clinical Chemistry	2021/11/2	Journal Article

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Reference Preview Li-2009-COLD-PCR_a new pl

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COLD-PCR: a new platform for highly improved mutation detection in cancer and genetic testing

Jun Li and G. Héike Makrigiorgou*

*Correspondence: junli@scau.edu.cn (J. Li); heike.makrigiorgou@harvard.edu (G. H. Makrigiorgou)

Address: In Molecular Diagnostics and Medical Physics, Department of Radiation Oncology, Dana-Farber Cancer Institute and Brigham and Women's Hospital, Harvard Medical School, Boston, MA, USA.

Abstract

PCR is widely employed as the initial DNA amplification step for genetic testing and cancer biomarker detection, serving as a key link in the process of diagnosis. However, standard PCR is the ability to selectively amplify only a portion of genetic DNA in a sample. In this background, new detection assays are limited in their ability to identify subtle genetic changes that can have a profound impact on clinical decision-making and outcome that can serve as cancer biomarkers. We developed COLD-PCR (co-amplification at lower denaturation temperature) [Li, Wang, Stevens, Koller, Berber and Milbury, 2009] that can detect mutations in DNA samples with a high degree of sensitivity, even from mixtures of wild-type and mutation-containing sequences irrespective of the mutation type or position on the sequence. Consequently, COLD-PCR amplification from genomic DNA yields PCR products containing high-frequency variant alleles that can be detected. Since PCR constitutes a ubiquitous initial step for almost all genetic analyses, COLD-PCR has the potential to revolutionize the field of genetic testing. In addition, COLD-PCR is also a valuable tool for the detection of mutations in heterogeneous cancer samples that are usually missed by existing methods. For example, COLD-PCR can detect mutations in EGFR (epidermal growth factor receptor) mutations in heterogeneous cancer samples that were missed by all existing methods. For clinical applications, COLD-PCR has the potential to revolutionize the field of cancer diagnostics. COLD-PCR is expected to have diverse applications in the fields of biomarker identification and testing, genomic instability, retinoblastoma, DNA methylation testing and prenatal/diagnostic of fetal alleles in maternal blood.

Introduction

PCR plays a key role in molecular diagnosis and in the detection of mutations. Aversely, unexpected situations when variant DNA sequences exist in the presence of a large amount of wild-type DNA are often encountered. For example, a tumor may contain a mixture of wild-type and mutation-containing sequences, irrespective of where an obvious mutation lies (variant alleles); thus both variant and non-variant alleles are largely masked by the wild-type alleles. The problem of identifying and sequencing the variants in a PCR product falls into two main categories. Despite being reliable for sequencing point mutations, sequencing of PCR products that contain unknown low-frequency mutations using other sequencing powerful technologies is still problematic. The significance of identifying these mutations, however, is critical as several fields of medicine, including cancer, prenatal diagnosis and infectious diseases [1-3]. We recently developed COLD-PCR (co-amplification at lower denaturation temperature-PCR), a novel PCR-based technique that can selectively amplify only a portion of genetic DNA in a sample. In this technique, a single nucleotide polymorphism (SNP) in a double-stranded DNA sequence generates a small but predictable change to the 'melting' temperature of DNA (T_m) for that sequence [4,5]. Depending on the sequence context,

Principle of COLD-PCR

The principle of COLD-PCR is described in Figure 2. A single nucleotide variant anywhere along a double-stranded DNA sequence generates a small but predictable change to the 'melting' temperature of DNA (T_m) for that sequence [4,5]. Depending on the sequence context and

Figure 1 | Illustration of the concept

Figure 1 illustrates the concept of COLD-PCR. It shows a diagram of a double-stranded DNA molecule with a single nucleotide change (a SNP) highlighted. The DNA is being amplified by PCR, and the melting temperature (T_m) is indicated for both strands. The wild-type strand has a higher T_m than the variant strand. The COLD-PCR process is shown as a two-step PCR cycle where the temperature is lowered to a point where only the variant strand remains, allowing for its selective amplification.

PCR

Figure 1 shows a standard PCR cycle with denaturation, annealing, and extension steps. The COLD-PCR cycle is shown as a two-step PCR cycle where the temperature is lowered to a point where only the variant strand remains, allowing for its selective amplification.

COLD-PCR

Figure 1 shows a standard PCR cycle with denaturation, annealing, and extension steps. The COLD-PCR cycle is shown as a two-step PCR cycle where the temperature is lowered to a point where only the variant strand remains, allowing for its selective amplification.



4. 导入NoteExpress

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- 11.16
- 笔记
- 检索
- 组织
- 回收站

题录

年份	作者	标题
2009	Li, Jin; Makrigiorgos, G Mike	COLD-PCR: a new platform for highly improved mutation detection in cancer and genetic testing
2009	Milbury, Coren A; Li, Jin; Makrigiorgos, G Mike	COLD-PCR-Enhanced High-Resolution Melting Enables Rapid and Selective Identification of Low-Level Unknown Mutations
2009	Milbury, Coren A; Li, Jin; Makrigiorgos, G Mike	PCR-Based Methods for the Enrichment of Minority Alleles and Mutations
2008	Li, Jin; Wang, Lili; Mamon, Harvey; Kulke, Matthew H; Berbeco, Ross; Makrigiorgos, G Mike	Replacing PCR with COLD-PCR enriches variant DNA sequences and redefines the sensitivity of genetic testing
2009	Li, Jin; Milbury, Coren A; Li, Cheng; Makrigiorgos, G Mike	Two-Round Coamplification at Lower Denaturation Temperature-PCR (COLD-PCR)-Based Sanger Sequencing Identifies

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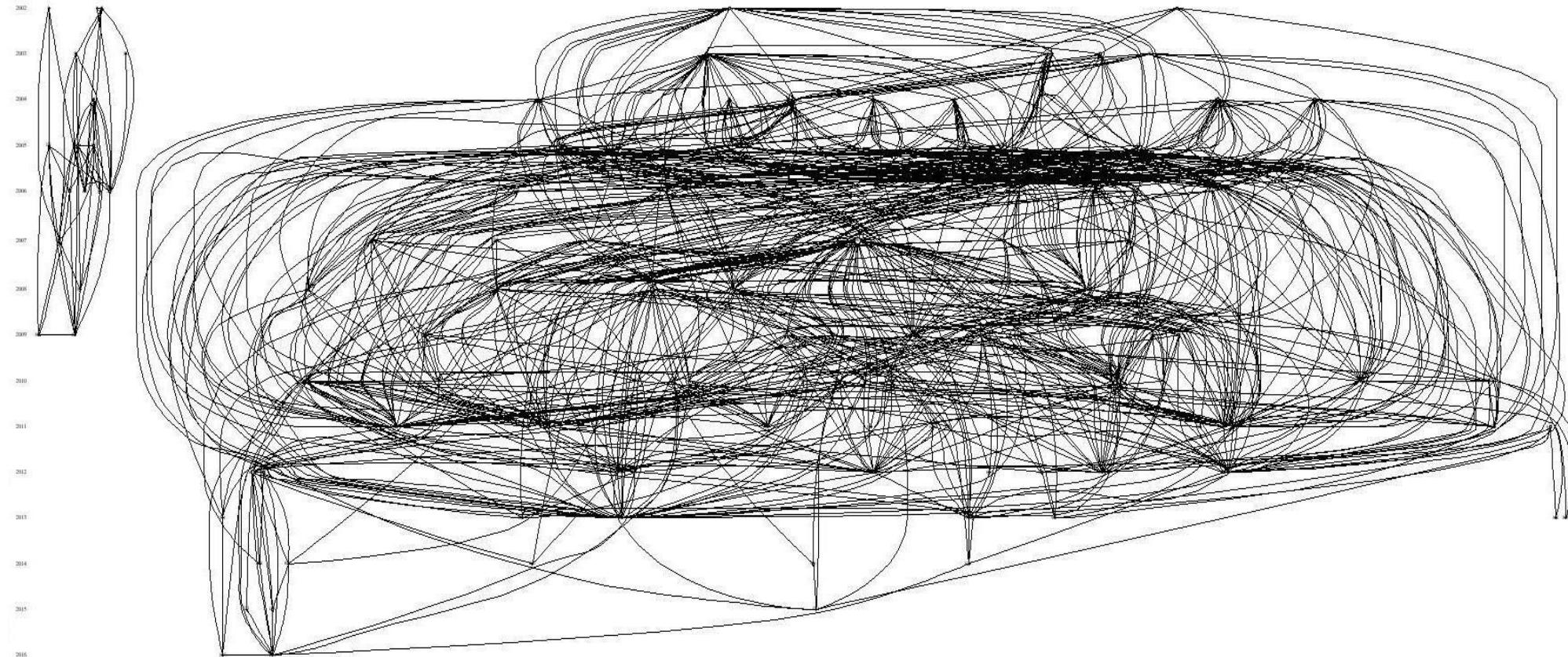


PART 03

Histcite作图分析结果示例



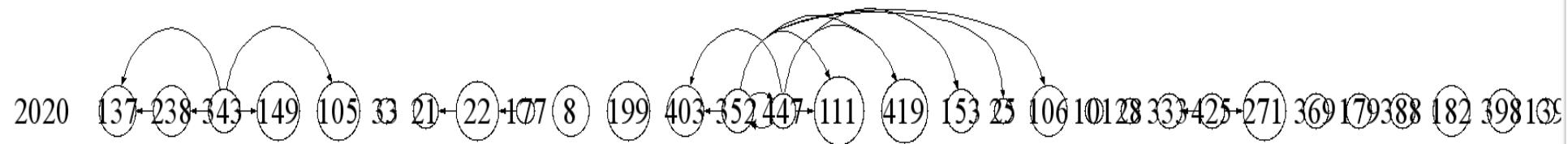
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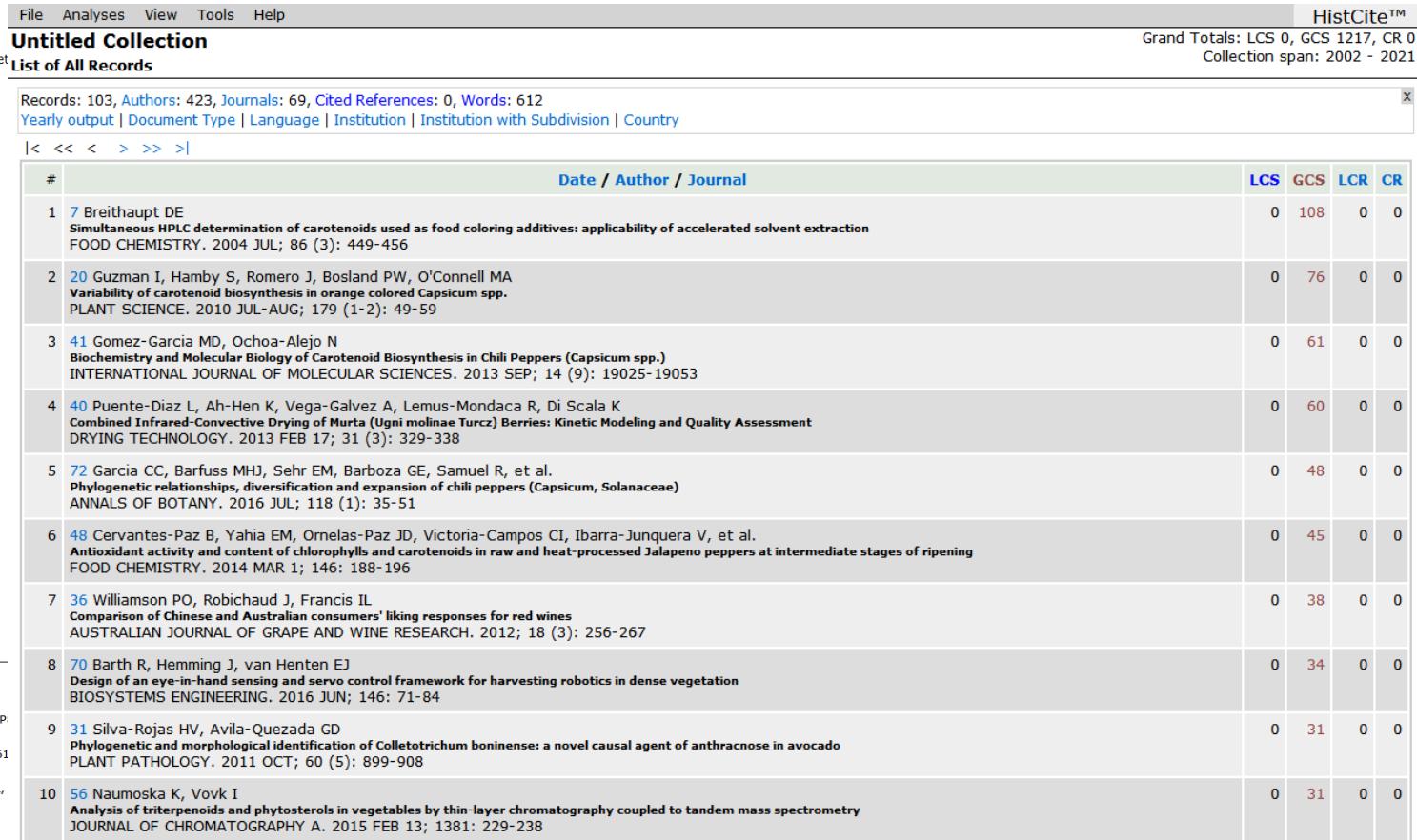
研究方向有两个分支



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注意：一些较新发表的文章因为年代近，因而被引用次数暂时还不多，没有入围到LCS前30，并不代表这个领域近几年不活跃了，没有人做这方面的研究了。



经典指标：

LCS是Local citation score的简写，即本地引用次数（本研究领域对该文献的关注程度）。与GCS相对应，GCS是总被引次数（所有的领域对该文献的关注程度）。LCS是某篇文章在当前数据库中被引用的次数。所以LCS一定是小于或等于GCS的。

如果一篇文章GCS很高，说明被全球科学家关注较多。但是如果一篇GCS很高，而LCS很小，说明这种关注主要来自与你不是同一领域的科学家。此时，这篇文章对你的参考意义可能不大。举个例子，有两篇文章P1(GCS:580,LCS:12) 和 P2(GCS:36,LCS:24)。第一篇文章GCS很高，LCS很低，说明关注这篇文章的绝大部分作者与你关注的方向不同。而第二篇文章GCS较低，但LCS比第一篇要高，即很多引用P2的文章都在当前数据库，也即与你的研究方向相关。所以，P1与P2相比，P2应该更贴近你的研究方向，参考价值更大。



LCR是Local cited references，是指某篇文章引用的所有参考文献中，有多少篇在当前数据库中。如果最近有两篇文章，P1 P2,都引用了30篇参考文献，其中P1引用的30篇文献中有20篇在当前数据库，P2只有2篇文献在当前数据库。此时，P1相对更有参考价值，因为它引用了大量的和你的研究相关的文献。

THANKS 谢谢！