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Knowledge structure and emerging trends of AR variants in prostate cancer: a bibliometric analysis based on CiteSpace and VOSviewer

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Abstract

Prostate cancer is a common malignancy in urology which often develops into castrationresistant prostate cancer (CRPC) after hormone therapy. Studies have shown that the mechanism of its occurrence is related to androgen receptor splice variants (AR splice variants). This work employs a bibliometric approach to explore the knowledge structure and emerging trends of AR variants in prostate cancer. The literature from 2000 to 2021 was obtained from Web of Science Core Collection (WoSCC), and the results were analyzed and visualized via CiteSpace and VOSviewer regarding publication number, citation number, country, region, institution, journal, author, keyword and reference. A total of 1503 publications were obtained. The number of publications and citations in this field is increasing. United States is the most prominent country, and University of Washington is the most influential institution in terms of this research field. European Urology is the most authoritative journal regarding this field. Gleave, Martin is the most productive author, and collaborated closely with others having centrality >0.1. The keywords abiraterone and AR-V7 have the strongest citation bursts in recent years and continuing to the end of 2021 which indicate the trend and further research directions. The most cited papers and co-cited references were related to the clinical significance of AR-V7 and its mechanism of promoting CRPC. This study used visualization software CiteSpace and VOSviewer to analyze the current status and the research trends of AR variants in prostate cancer (PCa) over the past 20 years. The findings can identify research hotspots and reveal future research directions.

Keywords

Prostate cancer; Androgen receptor splice variant; CiteSpace; VOSviewer; Bibliometric analysis

1. Introduction

Prostate cancer (PCa) is a common malignant cancer among elderly men. PCa incidence increases yearly with the aging of population, which results in heavy disease burden [1]. PCa mortality rate is decreasing with the therapies development however it still poses threat to the health of middle-aged and elderly men [2]. In western countries, PCa remains a public health concern. PCa has strong hormone sensitivity at the early stage. Most PCa progress to the castration-resistance stage with androgen deprivation therapy (ADT) and/or androgen receptor signaling inhibitors [3]. Patients of this cancer stage develop primary or secondary resistance to androgen- or androgen receptor-targeted agents which are associated with multiple underlying mechanisms [4-7]. It is also reported that incidence of PCa is associated with germline mutations and polymorphisms of AR gene [8]. One of the mechanisms contributing to castration resistance is AR splice variants which are truncated isoforms of AR, lacking a ligand-binding domain in C-terminus [9]. AR variants have thus been focused regarding prostate cancer research. Studies have demonstrated the expression of AR variants, the correlation of their existence, and castration resistance progression *in vitro* and *in vivo* [10–12]. The overexpression of AR-V7 in metastatic castration-resistant prostate cancer (mCRPC) patients is associated with shorter survival and resistance toward enzalutamide and abiraterone treatments [7]. Patients with AR aberrations, including AR-V7, demonstrate worse outcomes compared to patients without AR aberrations upon treatment by ADT with or without apalutamide for metastatic castration-sensitive PCa [13]. No study based on bibliometric analysis has summarized the knowledge structure and research progress of AR variants in PCa from recent decades.

Bibliometrics analyze the current research status and future trends in a field. The method has been used in life science fields such as medicine, biology, and public health to provide ideas for exploring current status, content, and priority research hotspots [14–16]. The results from bibliometric analyses can help investigators in understanding the research field regarding current and future perspectives.

In this study, the publications related to AR variants in PCa are collected from 2000 to 2021. Bibliometric analysis is conducted for identifying the current research concerns, global trends and future challenges of this field.

2. Materials and methods

Web of Science Core Collection (WoSCC) was used as original data source, and "androgen receptor splice variant" and "prostate cancer" as search terms. The detailed search strategy was ((Title Set) TS = (androgen receptor variant*) OR TS = (AR variant*)) AND ((TS = (prostate cancer*) OR TS = (prostate carcinoma*) OR TS = (prostate neoplasm*)). The number of relevant studies in WoS was 1682 in the era from 01 January 2000 to 31 December 2021. Only original research and reviews were included in the literature research type to ensure data accuracy and validity. After screening, 1503 original studies were obtained for downloading in plain text format as "full records with cited references".

The data from WoSCC were imported to Excel, CiteSpace and VOSviewer(Excel, Microsoft Office LTSC Professional Plus 2021, Microsoft, Washington State, United States; CiteSpace, 6.1.R6, Redsell University, PA, United States; VOSviewer, 1.6.18, Science and Technology Research Centre, Leiden University, the Netherlands). These articles were analyzed regarding author, country, research institution, journal, keywords, and citations. In CiteSpace 6.1. R6, the time selection was set to 2000 to 2021 and time slice to 1 year. The number of citations or occurrences was set to top 10% per year, and the maximum number of studies in each time slice was set to 100. Pathfinder, pruning sliced networks and pruning merged networks were simultaneously selected in the pruning algorithm. The default settings were maintained in the construction algorithm in case there were uncountable cases. Author, institution, country, keyword, reference, cited author, and cited journal were analyzed separately for the node types. In the burst-word detection, γ (0,1) was set to 0.5, and minimum duration was set to 1. If fewer than 20 burst words are retrieved, the γ value is incrementally decreased by 0.1 until 20 words are retrieved. The graph generated by CiteSpace consisted of nodes and the corresponding lines. The node size is related to its frequency in corresponding analytical module; its color, its time; its links to the co-occurrence or co-citation between the nodes; the link thickness represents closeness of connection between the two nodes; and the link color represents time of connection between the nodes. The outer circle of nodes with centrality >0.1 is purple which indicates the node having significant influence in the module.

Data source in VOSviewer was the bibliographic data. The threshold was set to 5 articles, and only the linked nodes were retained in VOSviewer-generated graphs. The layout settings were set to "2" for attraction and "-1" for repulsion, and the rest were set to their defaults.

3. Results

3.1 Publication distribution by year

Of 1503 papers, 1172 (77.98%) were the original articles and 331 (22.02%) the reviews. Based on the data analysis, publication numbers by year and trends regarding AR splice variants in PCa are shown in Fig. 1. The number of papers in this field has gradually increased since 2000 and the exponential growth from 2010 to 2016. This number was 4.8 times higher in 2017 than in 2010. The number of peaks in 2019 was 161. The trend line shows that the publication number fluctuated slightly from 2000 to 2021 but generally increased yearly. A binomial function from 2000 to 2021 was generated based on the correlation coefficient R² to further predict the publication trend of AR splice variants in PCa. The binomial function, $y = 0.3307x^2 - 0.0036x + 11.318$ (R² = 0.9067, where Y and X are the annual publications and year, respectively), was analyzed where the inflection point had not yet appeared. The results indicate rapid publication growth in recent years. The number of related citations has also increased annually. The frequency of this research has grown since 2018 resulting in the exponential increase of number of annual citations.

3.2 Countries and regions

These studies were published in 65 countries and regions. Table 1 lists the top 10 countries and regions according to the number of publications. Centrality, citation number, and total link strength are also included in the table. The United States had the most publications (n = 661, 43.98%), followed by China (n = 164, 10.91%), and Canada (n = 88, 8.09%). United States thus published the largest number of papers with the highest number of citations and the strongest total link strength which indicate the great influence of country's research within the field.

Collaborations between countries and regions are shown in Fig. 2 where there are 65 nodes and 402 links. The node size represents number of papers published in each country, and red circles outside the nodes represent number of papers published in that country in recent years. The purple circles outside nodes indicate that the node has high centrality which reveals its important position in cooperative relationship. The top five countries in terms of centrality are Australia (centrality = 0.56), England (0.47), Denmark (0.44), France (0.41), and Germany (0.40). Table 1 and Fig. 2 show that United States dominates the research on AR splice variants in PCa and exceeds other countries regarding publication number. However, in terms of collaborations between countries, Australia had the highest centrality and cooperations with multiple countries.

3.3 Organizations

According to the CiteSpace analysis, 503 organizations worldwide were involved in this field. Table 2 shows the top 10 organizations with the highest number of publications, centrality, citations, and total link strength. University of Washington had the highest publication number (n = 75, 4.99%), followed by University of British Columbia (n = 60, 3.99%), and the Fred Hutchinson Cancer Research Center (n = 49, 3.26%).

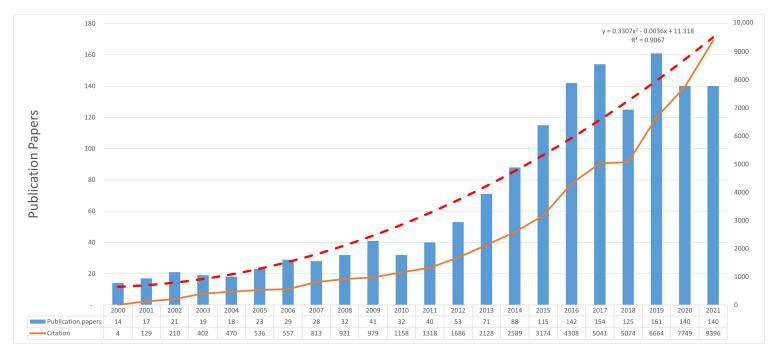


FIGURE 1. Annual publications and citation trends of androgen receptor splice variants in prostate cancer. The left vertical coordinate represents publication number, the right vertical coordinate represents citation number, and the red dashed line represents curve of binomial function.

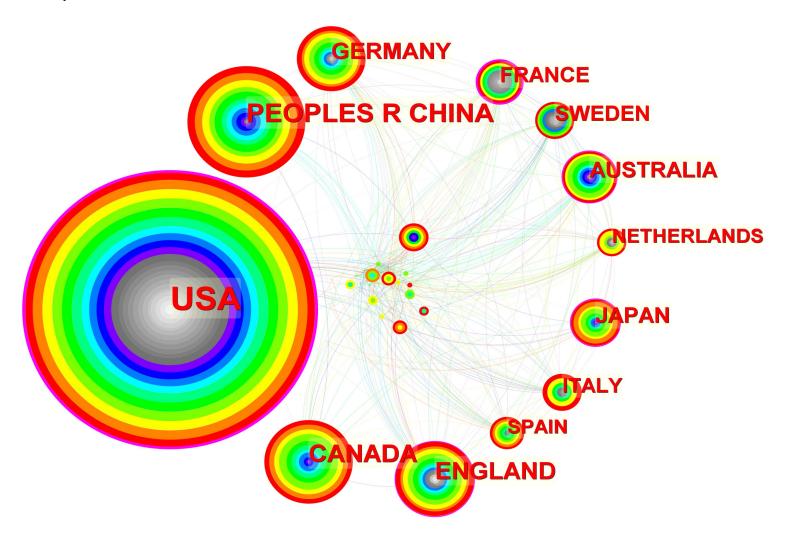


FIGURE 2. Map of countries and regions with publications on androgen receptor splice variants in prostate cancer from 2000 to 2021.

No.	Country	Centrality	Documents	Citations	Total link strength
1	United States	0.23	661	45,123	12,614
2	People's Republic of China	0.00	164	4812	4183
3	Canada	0.04	120	7182	3314
4	England	0.47	95	5802	2600
5	Germany	0.40	80	3045	1799
6	Japan	0.17	65	2298	1369
7	Australia	0.56	62	2990	1642
8	France	0.41	51	2515	831
9	Sweden	0.06	50	2094	1207
10	Italy	0.15	49	3458	1645

TABLE 1. Top 10 countries and regions involved in the research on androgen receptor splice variants in prostate cancer from 2000 to 2021.

TABLE 2. Top 10 institutions with research on androgen receptor splice variants in prostate cancer from 2000 to 2021.

No.	Institution	Centrality	Documents	Citations	Total link strength	Country
1	Univ Washington	0.00	75	7987	287	United States
2	Univ British Columbia	0.04	60	3679	174	Canada
3	Fred Hutchinson Canc Res Ctr	0.07	49	2407	173	United States
4	Johns Hopkins Univ	0.03	45	3909	167	United States
5	Univ Minnesota	0.02	33	6677	148	United States
6	Harvard Med Sch	0.17	31	1385	137	United States
7	Dana Farber Canc Inst	0.01	31	3781	132	United States
8	Baylor Coll Med	0.06	29	2338	130	United States
9	Univ Calif San Francisco	0.11	29	3025	124	United States
10	Mem Sloan Kettering Canc Ctr	0.04	26	2288	111	United States

Two of the top 10 institutions with more than 5000 citations were University of Washington (n = 7987) and University of Minnesota (n = 6677). University of Washington had the highest total number of links (n = 287). University of Minnesota published less than 40 articles but its citations exceeded 6500, suggesting that this institution is in developmental stage but with significant impact. The top 10 organizations were from United States except the University of British Columbia from Canada. This finding is consistent with the results shown in Fig. 2.

A map of institutional collaboration is shown in Fig. 3 which includes 503 nodes and 1359 node links. The node size represents number of papers published by that institution. The red circles outside nodes show number of papers published by that institution in recent years. The purple circles outside nodes indicate that the institution has high centrality, suggesting its important position in collaboration map. Brigham and Women's Hospital (n = 0.26), China Medical University (n = 0.18), Dana-Farber Cancer Institute (n = 0.17), Second Military Medical University (n = 0.17), and Harvard University (n = 0.17) were important in terms of institutional collaborations.

3.4 Journals and co-cited journals

From 2000 to 2021, 714 journals published studies on AR splice variants in PCa. The most influential journals were identified by analyzing the related journals. The 10 journals with the highest publication numbers and their impact factors (IFs) in 2022 are listed in Table 3. The top three journals were Prostate (n = 98), Clinical Cancer Research (n = 51), and Oncotarget (n = 50). European Urology (IF = 24.267), Clinical Cancer Research (IF = 13.801), and Cancer Research (IF = 13.312) were the leading journals in this field.

Co-cited journals refer to journals cited by publications in AR splice variant field that constitute the knowledge base of relevant studies. Top 10 co-cited journals in the field are listed in Table 4. Top three co-cited journals were Cancer Research (n = 940), Clinical Cancer Research (n = 696), and Proceedings of the National Academy of Sciences of the United States of America (n = 678). The journal with the highest IF was the New England Journal of Medicine (IF = 176.079). The analysis of centrality of cited journals showed that no journal had a centrality >0.1, indicating that this field has no representative journal regarding co-citations.

No.	Journals	Documents	Rate in total/%	2022 IF (JCR)
1	The Prostate	98	6.52%	4.012 (Q2)
2	Clinical Cancer Research	51	3.39%	13.801 (Q1)
3	Oncotarget	50	3.33%	4.345 (None)
4	Cancer Research	47	3.13%	13.312 (Q1)
5	PLOS One	34	2.26%	3.752 (Q2)
6	European Urology	33	2.20%	24.267 (Q1)
7	Journal of Biological Chemistry	28	1.86%	5.486 (Q2)
8	Cancers	27	1.80%	6.575 (Q1)
9	Endocrine-Related Cancer	25	1.66%	5.900 (Q1)
10	Scientific Reports	23	1.53%	4.996 (Q2)

TABLE 3. Top 10 journals related to androgen receptor splice variants in prostate cancer from 2020 to 2021.

IF: impact factors. JCR: Journal Citation Reports.

TABLE 4. Top 10 co-cited journals related to androgen receptor splice variants in prostate cancer from 2020 to 2021.

No.	Journals	Documents	Centrality	2022 IF (JCR)
1	Cancer Research	940	0.01	13.31 (Q1)
2	Clinical Cancer Research	696	0.03	13.80 (Q1)
3	Proceedings of the National Academy of Sciences of the United States of America	678	0.01	12.78 (Q1)
4	The Prostate	640	0.00	4.01 (Q2)
5	The New England Journal of Medicine	571	0.01	176.08 (Q1)
6	Journal of Biological Chemistry	538	0.00	5.49 (Q2)
7	Oncogene	483	0.01	8.76 (Q1)
8	Journal of Clinical Oncology	447	0.01	50.72 (Q1)
9	PLOS One	436	0.03	3.75 (Q2)

IF: impact factors. JCR: Journal Citation Reports.

TABLE 5. Top 10 authors regarding publication number in research on androgen receptor splice variants in prostate cancer from 2000 to 2021.

No.	Author	Affiliation	Documents	h-index	Centrality	Citation
1	Gleave, M [17]	University of British Columbia	30	78	0.11	1603
2	Dehm, S [18]	Masonic Cancer Center	27	36	0.02	2762
3	Luo, J [7]	Johns Hopkins University	26	41	0.04	5000
4	Plymate, S [11]	University of Washington	25	70	0.05	3240
5	Antonarakis, E [19]	University of Minnesota System	24	46	0.05	3911
6	Nelson, P [20]	Fred Hutchinson Cancer Center	24	89	0.01	3037
7	Corey, E [21]	University of Washington	16	87	0.03	825
8	Balk, S [22]	Beth Israel Deaconess Medical Center	14	74	0.04	603
9	de Bono, J [23]	Institute of Cancer Research—UK	14	121	0.01	748
10	Evans, C [24]	University of California Davis	13	56	0.03	692

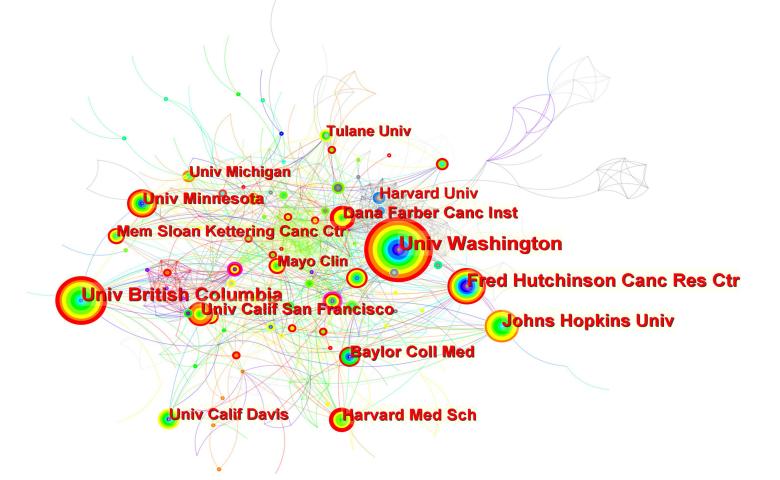


FIGURE 3. Map of the institutions having publications on androgen receptor splice variants in prostate cancer from 2000 to 2021.

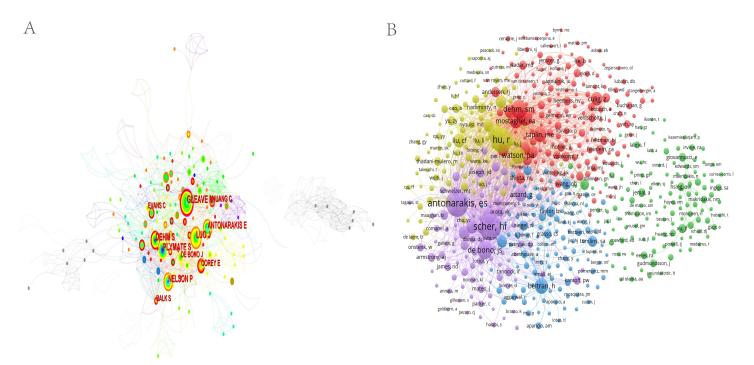


FIGURE 4. Analysis of authors and co-cited authors in the research on androgen receptor splice variants in prostate cancer from 2000 to 2021. (A) Map of the authors and collaborations. (B) Cluster analysis of the co-cited authors.

3.5 Authors and co-cited authors

In total, 691 authors contributed to 1503 publications. Table 5 shows the top 10 authors in terms of publication number (sorted by centrality if the number is same), latest affiliation, centrality, and h-index. Martin Gleave from University of British Columbia had the highest publication number (n = 30, 2.00%), followed by Scott M. Dehm from Masonic Cancer Center (n = 27, 1.80%), and Jun Luo from Johns Hopkins University (n = 26, 1.73%).

The collaborations between these authors are shown in Fig. 4A which include 691 nodes and 2247 internode links. The size of author nodes represents number of papers published by that author. Red circles outside the nodes represent number of publications by that author in recent years. The purple circles outside nodes represent author with the highest centrality. Nodes with purple circles indicate the author having important position in the collaborations. Martin Gleave had a node degree >0.1 (n = 0.11) which indicated that he collaborated closely with other authors. Two authors' articles were cited simultaneously in one study, revealing that they had a co-citation relationship. Fig. 4B shows a map of cited authors in related publications. Cluster analysis in VOSviewer identified five clusters with Dehm, S (red); Hu, R (yellow); Scher, Hi (purple); Beltran, H (blue); and Hsing, Aw (green) as the core authors. The authors in same cluster had similar research directions.

3.6 Keywords

Keywords are a high-level summary and condensation of literature. Their occurrence frequency reflects research directions and hotspots. Keywords with case differences, hyphen differences, abbreviations, or similar in meanings were merged.

After screening and merging, 646 relevant keywords were extracted. Keyword co-occurrence analysis revealed 646 nodes and 2829 internode links (Fig. 5A). Five keywords appeared more than 200 times: "androgen receptor" (n = 461), "prostate cancer" (n = 400), "expression" (n = 328), "progression" (n = 207), and "splice variant" (n = 206). Fourteen keywords appeared more than 100 times. A keyword centrality over 0.1 indicated that the keyword was more likely to be a hotspot for research, and the following 10 keywords exceeded 0.1: "apoptosis" (n = 0.22), "receptor" (n = 0.18), "cell" (n = 0.17), "activation" (n = 0.16), "cancer" (n = 0.14), "proliferation" (n = 0.12).

A cluster analysis places the related keywords in the same cluster which indicates the current research status. A keyword timeline spectrum analysis shows the time of keyword appearance and the continuation time. Identifying keywords that change in frequency in a certain period can be used to reflect the hotspot phase of research field. Combining with the research zone keyword map to summarize hotspot evolution is significant for future research directions. In Fig. 5B, the module value (modularity Q) was 0.4509 > 0.3, and the average profile value (weight mean silhouette S value) was 0.748 > 0.7, indicating that the structure of this clustered association is significant and efficient. The keyword clustering map (Fig. 5B) and timeline spectrum (Fig. 5C) revealed the fol-

lowing eight class clusters: "polymorphism", "growth", "binding", "neuroendocrine prostate cancer", "abiraterone", "differentiation", "androgen" and "gene expression". Keyword clustering and timeline spectrum analyses revealed that the research time of clusters ranged from 2000 to 2021. The clusters "polymorphism" and "growth" were concentrated before 2004, and the clusters "binding", "neuroendocrine prostate cancer", and "neuroendocrine prostate cancer" were concentrated after 2004. "Neuroendocrine prostate cancer" and "abiraterone" were mainly studied after 2010 and have continued to be studied, whereas the rest of clusters were studied more evenly over time.

Burst-word detection refers to the number of times a particular topic appears in a given period. It can be used to study past research hotspots and predict potential future directions. The top 25 keywords with the strongest citation bursts are shown in Fig. 6. "Androgen receptor gene" had the longest burst time from 2000 to 2008 with a burst strength of 8.31. The top two keywords with the strongest burst strength were "abiraterone" and "AR-V7", with strengths of 15.72 and 15.14, respectively, and the burst time was from 2018 to the present. Other recent keywords with citation bursts included degradation, liquid biopsy, and resistance.

3.7 Publications and co-cited references

An analysis of co-cited references reveals current research hotspots in the field. Table 6 lists 10 most-cited papers between 2000 and 2021. Table 7 includes the 10 most-cited references between 2000 and 2021 with the number of citations ranging from 249 to 486. The top two cited papers in both tables were the same: Antonarakis ES [7] and Hu R [25]. Both authors are from the same organization, Johns Hopkins University School of Medicine, and the same corresponding author, Jun Luo.

The top 50 references with the strongest citation bursts from 2000 to 2021 are presented in Fig. 7. The length of blue line segment represents study time frame and portion of red line segment indicates the time period of citation burst. Citation procedures usually lasted for 5 years. This length can be related to authors' writing habits as most authors cite articles published in the last 5 years. Article with the strongest citation burst was Antonarakis ES [7] in 2014 with strength of 41.9 and citation burst period from 2016 to 2019. Other references with strong citation bursts (*i.e.*, >25) were Guo Z [10], Hu R [17], Watson P [12], and Sun S [11]. Most of these references focused on identifying novel AR splice variants in PCa and explored the mechanisms by which variants promoted castration resistance in PCa in preclinical studies. The most recent references with high citation bursts were Antonarakis ES [7], Sharp A, Armstrong A, Scher H, and Abida W [36-40]. These studies focused on evaluating AR-V7 status in clinical specimens of PCa and elucidated its clinical significance.

4. Discussion

Bibliometrics is important for researchers and scholars as a discipline that studies research trends, directions and hotspots over time. PCa is the second most diagnosed malignancy and the fifth leading cause of cancer mortality in men worldwide

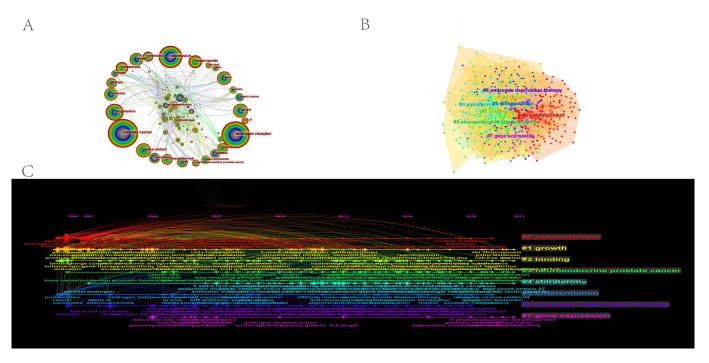


FIGURE 5. Analysis of keywords in the research on androgen receptor splice variants in prostate cancer from 2000 to 2021. (A) Map of the keyword co-occurrence. (B) Cluster analysis of the keywords. (C) Timeline spectrogram analysis of keywords.

[41]. It not only causes physical impairment but also increases society's economic burden and making it a public health issue in most countries. In recent years, AR variants have become an important hotspot in PCa research. In this study, the past and current research hotspots are elucidated in this field and future research trends and directions are predicted. It is the first study that summarizes the knowledge structure and research progress of AR variants in PCa through bibliometric analysis.

The publication distribution by years show increase in the number of papers in this research area over 20 years which peaked in 2019. The citation number of these publications has continued to increase from 2000 to 2021. The analysis of publication growth curve shows that the inflection point has yet not appeared which indicate that the number of publications in related fields will continue to increase. Eight of the top-10 countries publishing these articles are the Western countries which is consistent with the high incidence of PCa in these areas. United States published most publications with the highest number of citations as well as the strongest total link strength. The country with the second highest number of publications is China. The incidence and mortality of PCa are relatively low in China, however PCa incidence has been increasing in recent years [42]. Canada is contributing ~8% of the publications in this field. Australia has the highest centrality having more collaborations with other countries.

Among the 714 journals publishing articles related to AR variants in PCa, the top-10 journals published 416 (27.8%) documents. Prostate published the most articles (98, 6.52%). Most of the top-10 journals specialized in the fields of urology and oncology suggest that the specialized journals are more focused on this research area. European Urology had the highest IF and it was also one of the highest IF in the field of urology. This finding indicates the importance of AR variants

in PCa in urology. All top-10 co-cited journals are listed in JCR Q1 and Q2, including the New England Journal of Medicine, Cell, and the Journal of Clinical Oncology with recent IFs of 176.079, 66.85 and 50.717, respectively. This finding suggests that the references in field of AR variants in PCa are from high impact journals.

The author and co-cited author analysis helps in identifying the scholars with high influence on AR variants in PCa. Gleave Martin was the most productive author with 30 documents and collaborated most closely with other authors. Five clusters of co-cited authors were identified. Authors in the same clusters may have similar research direction or strong cooperative relationship. These authors have focused on molecular biology, immunology, genetics, cell biology mechanisms and related therapeutic aspects of prostate cancer.

The research focus related to AR variants in PCa can be explored from cooccurring keywords. The five most common cooccurring keywords are identified as: "androgen receptor", "prostate cancer", "expression", "progression" and "splice variant". "Expression" and "progression" indicate the research focus of over 20 years in this field. "Expression" related to AR variants includes two aspects. One aspect is the expression of AR splice variants in PCa tissue and cell lines. In the past few years, at least 20 isoforms of AR variant mRNAs have been expressed in PCa. Another aspect is the gene expression profiles identified in PCa-expressing AR variants. Many gene expression profiles have been identified due to different isoforms of AR variants and the heterogeneity [9]. Another research focus includes the role and mechanism of AR variants in PCa progression. AR splice variants lack Cterminal ligand-binding domain in their amino acid structure owing to the truncation by cryptic exon. They thus do not respond to conventional ADT and are correlated with PCa

TABLE 6. Top 10 cited publications in the research on androgen receptor splice variants in prostate cancer from 2000
to 2021.

No.	Title	First author	Journal	Citation	Year
1	AR-V7 and Resistance to Enzalutamide and Abiraterone in Prostate Cancer	Antonarakis, ES [7]	The New England Journal of Medicine	1786	2014
2	Ligand-Independent Androgen Receptor Variants Derived from Splicing of Cryptic Exons Signify Hormone-Refractory Prostate Cancer	Hu, R [25]	Cancer Research	769	2009
3	Emerging Mechanisms of Resistance to Androgen Receptor Inhibitors in Prostate Cancer	Watson, P [26]	Nature Reviews Cancer	714	2015
4	Hormonal Carcinogenesis	Henderson, BE [27]	Carcinogenesis	663	2000
5	A Novel Androgen Receptor Splice Variant Is Upregulated During Prostate Cancer Progression and Promotes Androgen Depletion-Resistant Growth	Guo, Z [10]	Cancer Research	642	2009
6	Castration Resistance in Human Prostate Cancer is Conferred by a Frequently Occurring Androgen Receptor Splice Variant	Sun, S [11]	Journal of Clinical Investigation	532	2010
7	<i>RB1</i> and <i>TRP53</i> Cooperate to Suppress Prostate Cancer Lineage Plasticity, Metastasis, and Antiandrogen Resistance	Ku, SY [28]	Science	508	2017
8	RNA-Seq of Single Prostate CTCs Implicates Noncanonical WNT Signaling in Antiandrogen Resistance	Miyamoto, DT [29]	Science	479	2015
9	Constitutively Active Androgen Receptor Splice Variants Expressed in Castration-Resistant Prostate Cancer Require Full-Length Androgen Receptor	Watson, P [12]	Proceedings of the National Academy	466	2010
10	Androgen Receptor Splice Variant 7 and Efficacy of Taxane Chemotherapy in Patients with Metastatic Castration-Resistant Prostate Cancer	Antonarakis ES [30]	JAMA Oncology	464	2015

AR-V7: androgen receptor splice variants 7. RB1: Transcriptional Corepressor 1. TRP53: Transformation related protein 53. CTCs: Circulating tumor cell.

progression, especially CRPC. The keyword clustering and timeline analysis reveal "binding", "neuroendocrine prostate cancer" and "abiraterone" as the three clusters identified since 2010. AR binding with DNA is a fundamental mechanism by which AR modulates PCa development and progression. The differential binding mechanism of AR variants may promote CRPC progression and abiraterone resistance [43].

The most cited publications and co-cited references suggest the current research focus on AR variants in PCa. Antonarakis ES and Hu R published the most cited articles which were also the most co-cited references. In Antonarakis' article, AR-V7 as one of AR variant isoforms was detected in circulating tumor cells (CTCs) of CRPC patients. AR-V7 positive patients had lower prostate specific antigen (PSA) response and shorter PSA progression-free survival, clinical or radiographic progression-free survival, and overall survival which suggested that AR-V7 may be associated with the resistance toward enzalutamide and abiraterone [7]. Hu R *et al.* [17] identified seven AR variant transcripts and uncovered higher AR-V7 expression, predicting the biochemical recurrence after surgical treatment. AR-V7 was constitutively active in driving the expression of canonical androgen-responsive genes, suggesting a novel mechanism for CRPC development. Both studies were from Dr. Jun Luo's team at Johns Hopkins University School of Medicine. These findings suggested the mechanism by which AR-V7 promoted CRPC progression, and the association of AR-V7 with enzalutamide and abiraterone resistance in mCRPC patients.

Predicting future research directions is vital for research. Citation burst detection can be used to explore research trends, and recent ongoing bursts can partially reveal future trends [44]. Abiraterone and AR-V7 were identified as the top two keywords with the strongest citation bursts of strengths >15. This finding suggests that the association of AR-V7 and resistance to abiraterone in PCa is an emerging trend. Abiraterone is an inhibitor of cytochrome P450, family 17, subfamily A, polypeptide 1 (*CYP17A*) which depletes adrenal and intertumoral androgen by suppressing androgen synthesis [45]. Studies have shown improved survival with abiraterone. The agents are approved by the Food and Drug Administration for treating mCRPC [35, 46]. Abiraterone had made a breakthrough in mCRPC treatment, however some patients had primary re-

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No.	Title	First author	Journal	Citation	Year
1	AR-V7 and Resistance to Enzalutamide and Abiraterone in Prostate Cancer	Antonarakis, ES [7]	New England Journal of Medicine	486	2014
2	Ligand-Independent Androgen Receptor Variants Derived from Splicing of Cryptic Exons Signify Hormone-Refractory Prostate Cancer	Hu, R [25]	Cancer Research	393	2009
3	A Novel Androgen Receptor Splice Variant Is Upregulated During Prostate Cancer Progression and Promotes Androgen Depletion-Resistant Growth	Guo, Z [10]	Cancer Research	360	2009
4	Increased Survival with Enzalutamide in Prostate Cancer after Chemotherapy	Scher, HI [31]	New England Journal of Medicine	329	2012
5	Distinct Transcriptional Programs Mediated by the Ligand-Dependent Full-Length Androgen Receptor and its Splice Variants in Castration-Resistant Prostate Cancer	Hu, R [32]	Cancer Research	292	2012
6	Splicing of a Novel Androgen Receptor Exon Generates a Constitutively Active Androgen Receptor that Mediates Prostate Cancer Therapy Resistance	Dehm, SM [33]	Cancer Research	286	2008
7	Androgen Receptor Splice Variants Mediate Enzalutamide Resistance in Castration-Resistant Prostate Cancer Cell Lines	Li, YM [34]	Cancer Research	282	2013
8	Castration Resistance in Human Prostate Cancer is Conferred by a Frequently Occurring Androgen Receptor Splice Variant	Sun, S [11]	Journal of Clinical Investigation	267	2010
9	Constitutively Active Androgen Receptor Splice Variants Expressed in Castration-Resistant Prostate Cancer Require Full-Length Androgen Receptor	Watson, PA [10]	Proceedings of the National Academy of Sciences of the United States of America	261	2010
10	Abiraterone and Increased Survival in Metastatic Prostate Cancer	de Bono, JS [35]	New England Journal of Medicine	249	2011
10 1					

TABLE 7. Top 10 co-cited references in the research on androgen receptor splice variants in prostate cancer from 2000 to 2021

AR-V7: androgen receptor splice variants 7.

sistance to the agent with no response to PSA levels, while other patients acquired secondary resistance after an initial response [7]. One plausible explanation for this abiraterone resistance may involve the presence of AR variants, a finding supported by some preclinical studies [47]. AR-V7, also known as AR3, was another identified citation burst keyword. It has truncated C-terminal ligand binding domain (LBD) of AR protein encoded by novel cryptic exon 3 in mRNA [9, 40]. It is the common isoform among androgen receptor-splicing variants and the only known variant encoding a functional protein product that is detectable in clinical samples [10, 25]. Many studies have been reported investigating the association of AR-V7 and abiraterone resistance. In the past 20 years, reference with the strongest citation burst was Antonarakis ES, which was also the most cited paper and co-cited reference. The authors reported the association between AR-V7 and resistance to abiraterone and enzalutamide in mCRPC through a prospective study [7].

The recent burst keyword, "liquid biopsy" lasted till the

end of 2021. A liquid biopsy refers to the non-invasive analysis of biomarkers in biological fluids (such as blood, plasma, urine, liquor, and saliva) to allow the detection and longitudinal follow up of cancers [48, 49]. Compared with traditional tissue biopsies for prostate cancer, liquid biopsies are non-invasive, depict PCa heterogeneity, and convenient to follow up and evaluate efficacy during PCa treatment. The biomarkers obtained from liquid biopsy include cell-free tumor DNA (ctDNA), circulating cell-free tumor RNA (ctRNA), proteins, peptides, metabolites, CTCs and extracellular vesicles (EVs). In studies investigating the clinical significance of AR-V7 in PCa patients, the majority of measured AR-V7 mRNA or protein were from liquid biopsies, especially from CTCs [38, 39, 50, 51]. In a systematic review and meta-analysis of 37 studies, the clinical significance was investigated for AR-V7 detected from liquid biopsies. The analysis found that AR-V7 positivity detected from liquid biopsy was associated with poor Overall Survival (OS), Progression-Free-Survival (PFS), and prostate specific antigen progression-free survival (PSA-

Keywords	Year	Strength	Begin	End	2000 - 2021
androgen receptor gene	2000	8.31	2000	2008	
vitamin d receptor	2000	7.14	2001	2006	
susceptibility locus	2000	3.98	2001	2002	
srd5a2 gene	2000	5.72	2002	2005	
allelic variant	2000	7.52	2003	2009	
men	2000	4.45	2003	2005	
population	2000	7.57	2004	2009	
polymorphism	2000	11.25			
risk	2000				
association	2000	5.43	2008	2010	
growth	2000	6.55			
testosterone	2000	4.31			
identification	2000	4.22	2010	2013	
prostate cancer cell	2000	3.94	2012	2013	
resistant prostate cancer	2000	4.15	2014	2016	
receptor splice variant	2000	6.20	2015	2018	
chemotherapy	2000	8.33	2016	2018	
induction	2000	4.27	2016	2018	
enzalutamide	2000	7.99	2017	2021	
abiraterone	2000	15.72	2018	2021	
ar v7	2000	15.14	2018	2021	
lineage plasticity	2000	6.23	2018	2021	
degradation	2000	5.24	2019	2021	
liquid biopsy	2000	4.96	2019	2021	
resistance	2000	4.51	2020	2021	

FIGURE 6. Top 25 keywords with the strongest citation bursts.

PFS). A subgroup analysis showed that AR-V7 positivity was associated with poor prognosis of mCRPC treated with androgen receptor signaling inhibitors (ARSi) than with taxene [52].

Recent co-cited references, especially the ones still in bursting period, indicated the current trends and future directions. Antonarakis ES represented the strongest burst from 2018 to the end of 2021 (Fig. 7) [36]. This work explored the clinical significance of AR-V7 mRNA detection in CTCs of men with mCRPC. CTC+/AR-V7+ patients were more likely to have poor clinical characteristics, including Gleason scores \geq 8 and metastatic disease at diagnosis. These results confirmed the negative prognostic impact of AR-V7 detection in CTCs of CRPC patients treated with abiraterone and enzalutamide therapy. The results further suggested that CTC/AR-V7 biomarker panel might be useful in predicting the response to first- and second-line hormonal therapy settings. The second strongest burst citation by the end of 2021 was the study by sharp an in 2019 [37]. The authors performed cross-institutional study to determine nuclear AR-V7 protein expression in tissue biopsies and autopsies from primary and metastatic PCa rather than in liquid biopsies. AR-V7 expression was rare in primary PCa but emerged in response to primary ADT and further increased in response to abiraterone or enzalutamide therapy. AR-V7-negative disease was associated with better PSA response and overall survival, indicating it as an important marker of response to endocrine therapy and related to prognosis.

Another paper with high strength burst citation in past 3 years was Armstrong, 2019 [38]. The authors reported PROPHECY study as a multicenter, and prospective-blinded study of 118 men with high risk mCRPC starting abiraterone or enzalutamide treatment. The detection of AR-V7 in CTCs

References	Year	Strength	Begin	End 2000 - 2021
Ingles S, 1997, J NATL CANCER I, V89, P166, DOI 10.1093/jnci/89.2.166, DOI		12.38	-	
Giovannucci E, 1997, P NATL ACAD SCI USA, V94, P3320, DOI 10.1073/pnas.94.7.3320, DOI	1997		2000	and the second
Makridakis N, 1999, LANCET, V354, P975, DOI 10.1016/S0140-6736(98)11282-5, DOI	1999	9.57	2000	2003
Modugno F, 2001, CLIN CANCER RES, V7, P3092	2001	9.88	2002	2006
tsing A. 2000. CANCER RES. V60. P5111	2000	9.45	2002	
atil A, 2001, CANCER-AM CANCER SOC, V92, P1130, DOI 10.1002/1097-0142(20010901)92:5<1130::AID-CNCR1430>3.0.CC	Contraction and the second second second	10.80		
lu R, 2009, CANCER RES, V69, P16, DOI 10.1158/0008-5472.CAN-08-2764, DOI	2009		2009	A CONTRACTOR OF
ehm S. 2008. CANCER RES. V68. P5469. DOI 10.1158/0008-5472.CAN-08-0594. DOI	2008	24.10	2009	2013
eles R. 2008, NAT GENET, V40, P316, DOI 10.1038/ng.90, DOI	2008	9.81	2009	2013
uo Z, 2009, CANCER RES, V69, P2305, DOI 10.1158/0008-5472.CAN-08-3795, DOI		33.90	2010	2014
atson P, 2010, P NATL ACAD SCI USA, V107, P16759, DOI 10.1073/pnas.1012443107, DOI	2010	29.57	2011	2015
un S, 2010, J CLIN INVEST, V120, P2715, DOI 10.1172/JCI41824, DOI		28.68		
an C, 2009, SCIENCE, V324, P787, DOI 10.1126/science.1168175, DOI	2009	15.48	2011	2014
ndersen R, 2010, CANCER CELL, V17, P535, DOI 10.1016/j.ccr.2010.04.027, DOI	2010	12.08	2011	2015
ang Q. 2009, CELL, V138, P245, DOI 10.1016/j.cell.2009.04.056, DOI		10.46		
ostaghel E, 2011, CLIN CANCER RES, V17, P5913, DOI 10.1158/1078-0432.CCR-11-0728, DOI		22.77		
proberg E, 2011, PLOS ONE, V6, P0, DOI 10.1371/journal.pone.0019059, DOI		22.43		
8, 2011, NEW ENGL J MED, V364, P1995, DOI 10.1056/NEJMoa1014618, DOI		19.23		
ang X, 2011, PLOS ONE, V6, P0, DOI 10.1371/journal.pone.0027970, DOI		13.95		2012
I R. 2011. PROSTATE, V71, P1656. DOI 10.1002/pros.21382. DOI		12.54		
hm S, 2011, ENDOCR-RELAT CANCER, V18, P0, DOI 10.1530/ERC-11-0141, DOI		11.28		
vlor B, 2010, CANCER CELL, V18, P11, DOI 10.1016/j.ccr.2010.05.026, DOI		10.88		the second s
(, 2011, CANCER RES, V71, P2108, DOI 10.1158/0008-5472.CAN-10-1998, DOI		10.44		
nal A, 2011, CA-CANCER J CLIN, V61, P69, DOI 10.3322/caac.20107, DOI		10.41		
R, 2012, CANCER RES, V72, P3457, DOI 10.1158/0008-5472.CAN-11-3892, DOI		28.81		100.36
an S, 2012, J BIOL CHEM, V287, P19736, DOI 10.1074/jbc.M112.352930, DOI		12.87		
her H, 2012, NEW ENGL J MED, V367, P1187, DOI 10.1056/NEJMoa1207506, DOI		24.68		
Y, 2013, CANCER RES, V73, P483, DOI 10.1158/0008-5472.CAN-12-3630, DOI		24.00		
i C, 2011, CANCER RES, V71, P6503, DOI 10.1158/0008-5472.CAN-12-5650, <u>DOI</u>		9.28		
		12.56		
an C, 2013, NEW ENGL J MED, V368, P138, DOI 10.1056/NEJMoa1209096, <u>DOI</u> Z. 2014, CLIN CANCER RES, V20, P1590, DOI 10.1158/1078-0432,CCR-13-1863, DOI		10.15		
		10.15		
asso C, 2012, NATURE, V487, P239, DOI 10.1038/nature11125, <u>DOI</u>				100 A A A A A A A A A A A A A A A A A A
tonarakis E, 2014, NEW ENGL J MED, V371, P1028, DOI 10.1056/NEJMoa1315815, DOI		41.90		
tonarakis E, 2015, JAMA ONCOL, V1, P582, DOI 10.1001/jamaoncol.2015.1341, <u>DOI</u>		13.01		
an S, 2015, NUCLEIC ACIDS RES, V43, P5880, DOI 10.1093/nar/gkv262, <u>DOI</u>		9.65		
binson D, 2015, CELL, V161, P1215, DOI 10.1016/j.cell.2015.05.001, DOI		16.72		
her H, 2016, JAMA ONCOL, V2, P1441, DOI 10.1001/jamaoncol.2016.1828, <u>DOI</u>		15.92		
tson P, 2015, NAT REV CANCER, V15, P701, DOI 10.1038/nrc4016, DOI		11.43		
er T, 2014, NEW ENGL J MED, V371, P424, DOI 10.1056/NEJMoa1405095, DOI		11.13		
beshouse A, 2015, CELL, V163, P1011, DOI 10.1016/j.cell.2015.10.025, <u>DOI</u>		9.73		
tonarakis E, 2017, J CLIN ONCOL, V35, P2149, DOI 10.1200/JCO.2016.70.1961, <u>DOI</u>		21.51		
ltran H, 2016, NAT MED, V22, P298, DOI 10.1038/nm.4045, <u>DOI</u>		14.56		1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1
u P, 2017, SCIENCE, V355, P84, DOI 10.1126/science.aah4307, <u>DOI</u>		10.19		
arp A, 2019, J CLIN INVEST, V129, P192, DOI 10.1172/JCI122819, <u>DOI</u>		17.34		
mstrong A, 2019, J CLIN ONCOL, V37, P1120, DOI 10.1200/JCO.18.01731, DOI		11.24		
her H, 2018, JAMA ONCOL, V4, P1179, DOI 10.1001/jamaoncol.2018.1621, <u>DOI</u>		10.37		
ggarwal R, 2018, J CLIN ONCOL, V36, P2492, DOI 10.1200/JCO.2017.77.6880, DOI			2019	
her H, 2017, EUR UROL, V71, P874, DOI 10.1016/j.eururo.2016.11.024, <u>DOI</u>		9.51	2019	
oida W, 2019, P NATL ACAD SCI USA, V116, P11428, DOI 10.1073/pnas.1902651116, <u>DOI</u>		12.88		
ay F, 2018, CA-CANCER J CLIN, V68, P394, DOI 10.3322/caac.21492, <u>DOI</u>	2018	10.23	2020	2021

FIGURE 7. Top 50 references with the strongest citation bursts.

by two blood-based assays was associated with shorter PFS and OS, and AR-V7 positive mCRPC had fewer PSA or soft tissue responses to abiraterone or enzalutamide. This study suggested that AR-V7 was a strong predictor of clinical outcomes in mCRPC treated with abiraterone or enzalutamide.

Scher H evaluated whether a nuclear localized AR-V7 protein in CTCs was a treatment selection marker for mCRPC in two articles having high strength burst citations in the past 3 years. First, the positivity criteria of AR-V7 were investigated. Patients who had AR-V7 proteins in cellular nuclei of CTCs were likely to survive longer on taxane-based chemotherapy; however, AR-V7 with unknown protein localization was less predictive. Thus, the nuclear-specific localization of AR-V7 was required to evaluate its predictive benefit. Scher H reported the association of AR-V7 status and prognosis of mCRPC treated with either ARSis or taxanes in multiinstitutional cohort study. Patients with AR-V7 positive CTCs had better overall survival with taxanes *vs.* ARSis, and *vice versa.* Nuclear-localized AR-V7 could be used to select an ARSi or taxane and provided individual patient benefits. Abida W described a comprehensive integrative analysis of genomic and transcriptomic profiles, histology, and clinical outcomes for 429 mCRPC patients [53]. A high frequency was identified for genomic alterations in AR, including AR splice variants. AR-V7 levels were increased in tumors exposed to taxanes and ARSi therapy, but no association was identified between AR-V7 expression and shorter time of first-line ARSi or overall survival.

This bibliometric study of prostate cancer-related androgen receptor splice variants between 2000 and 2021 analyzes research hotspots and future trends in this field, however limitations remain. Because of publication time limitations, the value of some newly published or high-impact literature may not have been disseminated, which has certain delaying effect on research analysis. Only Web of Science Core Collection database was selected as the data source. This database has considerable influence in academic journal databases, however some high quality papers may not be included. In future, more literature may be included for the detailed study in this field.

5. Conclusions

This bibliometric study of AR variants in PCa between 2000 and 2021 visualizes the literature from Web of Science Core Collection to better understand the research trends and directions. Over the 22 years investigated, the number of publications substantially increased and still its inflection point was not achieved. The most influential country is USA, and the most influential institution is The University of Washington. About 27% articles were published in 1% top-10 journals, most of which are specialized in urology and oncology. Moreover, the research foundation of this field is from high impact journals. Keyword co-occurrence analysis shows that research has been focused on "expression" and "progression" in the past two decades. The most cited papers and co-cited references are related to the clinical significance of AR-V7 and its mechanism of promoting CRPC. Citation bursts analysis indicate that the research focus has transferred from the bench to bedside, with keywords, abiraterone and AR-V7 having the strongest citation burst. The current citation burst suggests that the trend and future direction are focusing on the evaluation of AR-V7 status in PCa clinical samples and elucidating its contribution to CRPC progression.

AVAILABILITY OF DATA AND MATERIALS

Data presented in this study are available on request from corresponding author.

AUTHOR CONTRIBUTIONS

LW, TTJ and SMC—conducted data collection. KY, FB and ZJL—were involved in data analysis. KY, JL and XL—wrote the manuscript. JL and XL—were involved in project development and supervision.

ETHICS APPROVAL AND CONSENT TO PARTICIPATE

Not applicable.

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CONFLICT OF INTEREST

The authors declare no conflict of interest.

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