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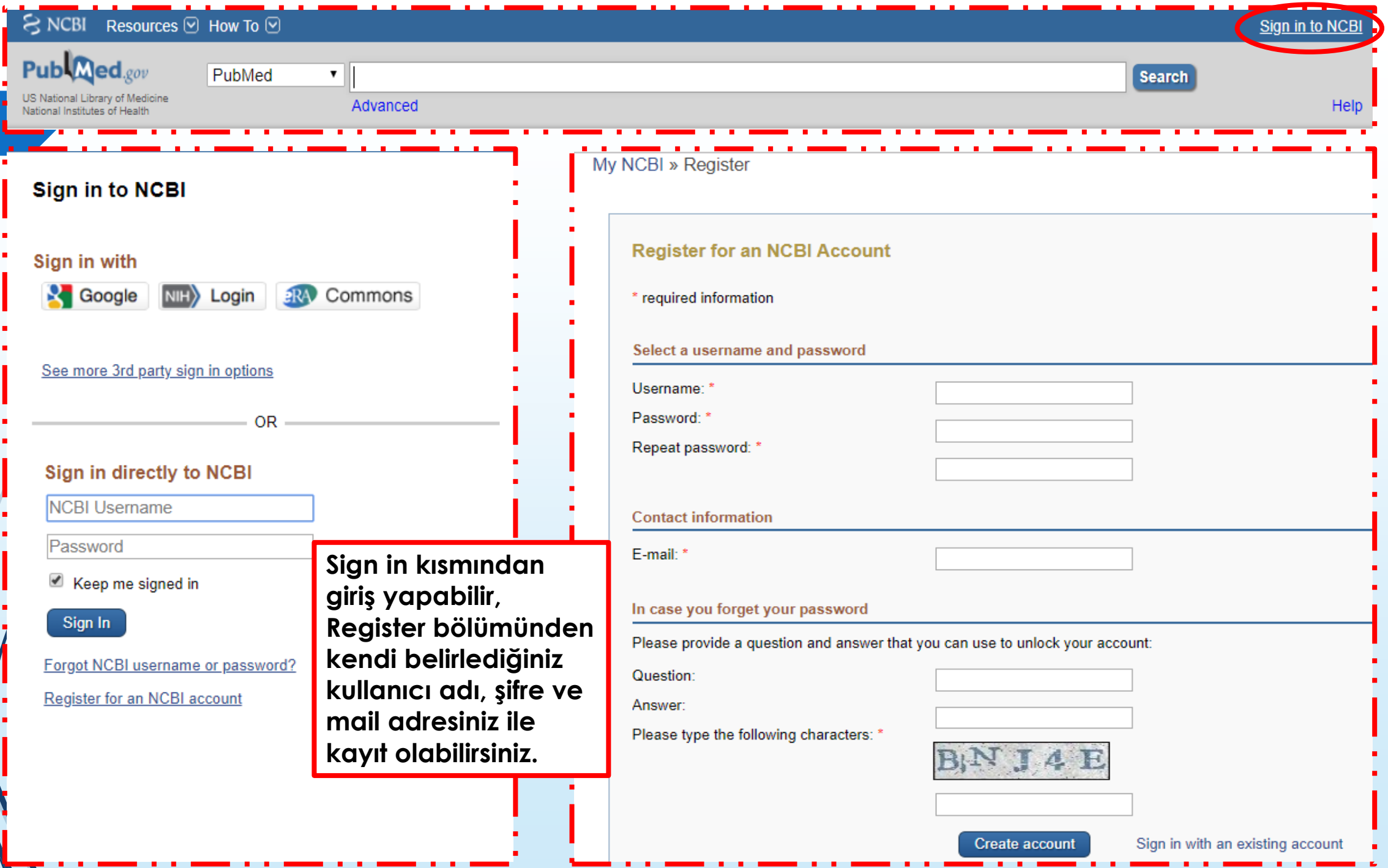

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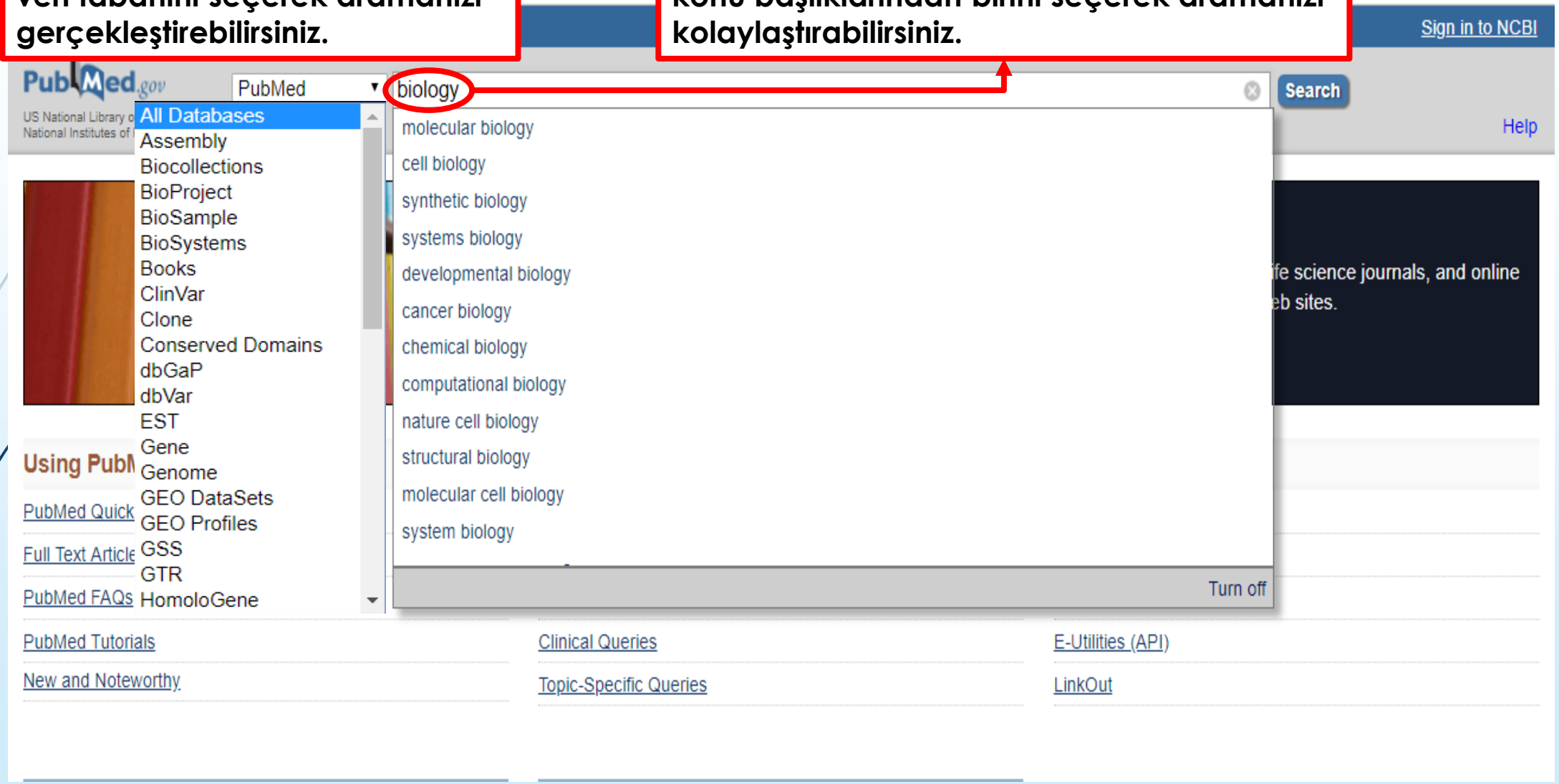

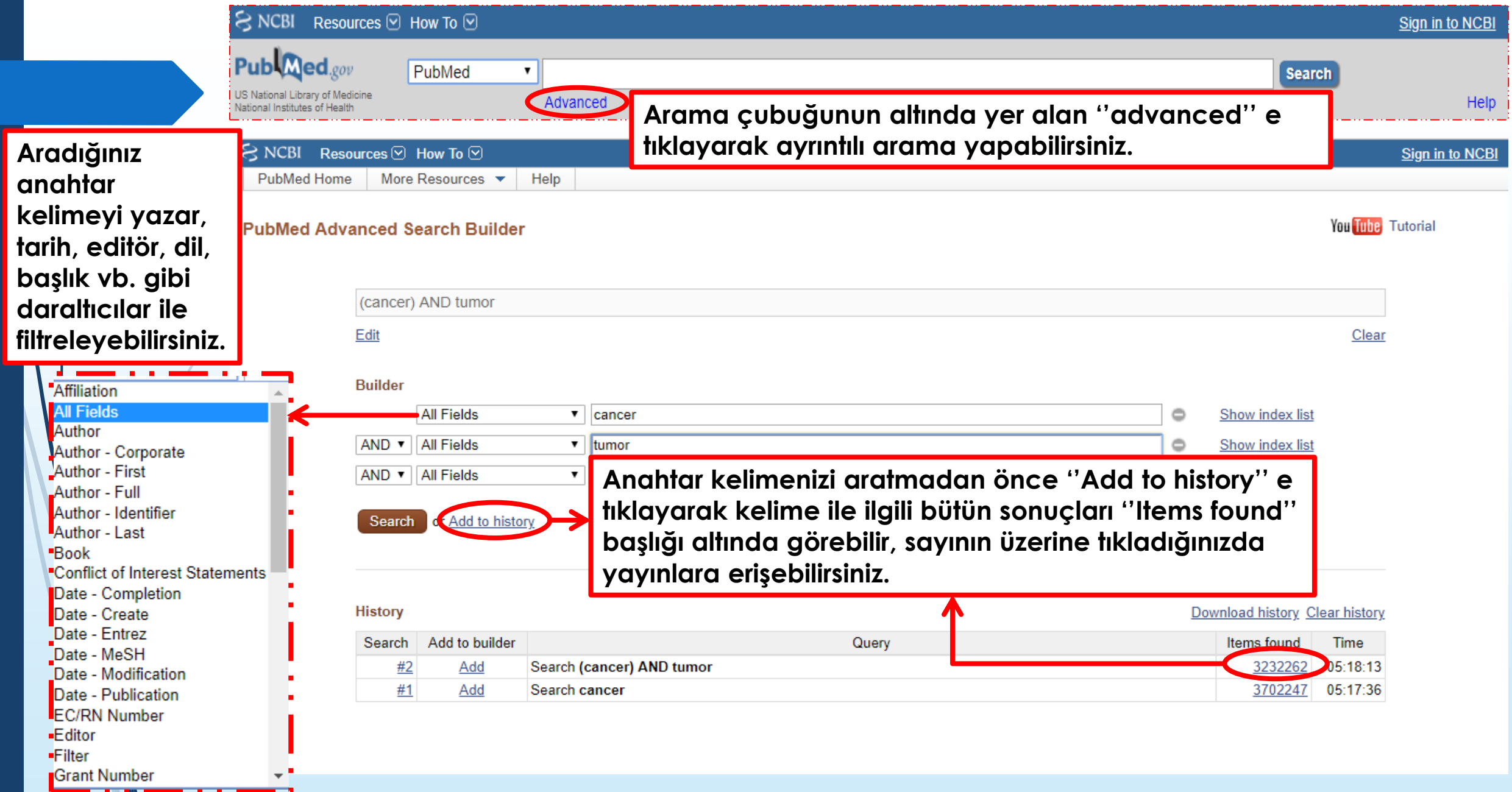
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tumor
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or Add to history
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Author
Author - Corporate
Author - First
Author - Full
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Query
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Time
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Add
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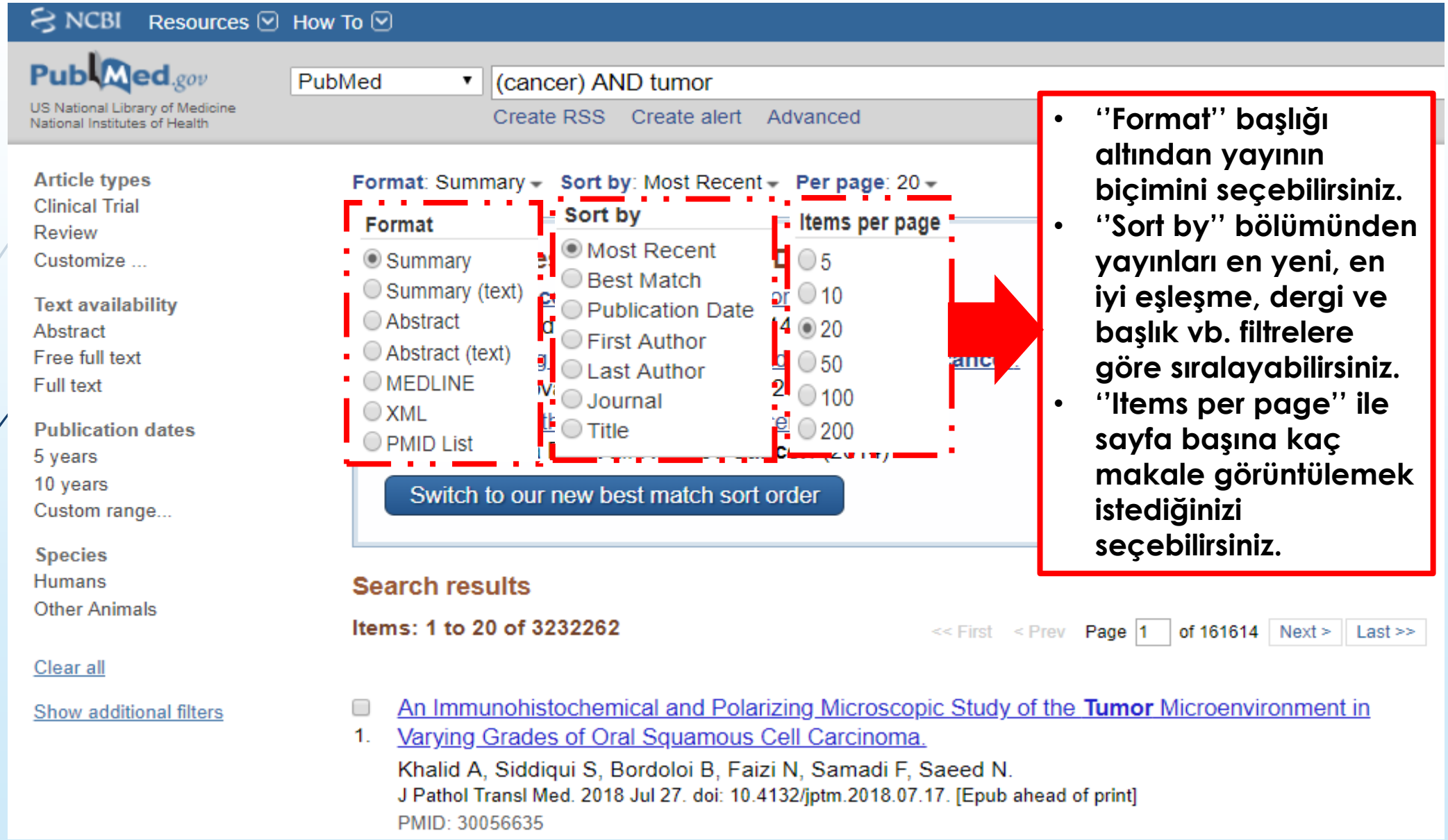
05:18:13
#1
Add
Search cancer
3702247
05:17:36



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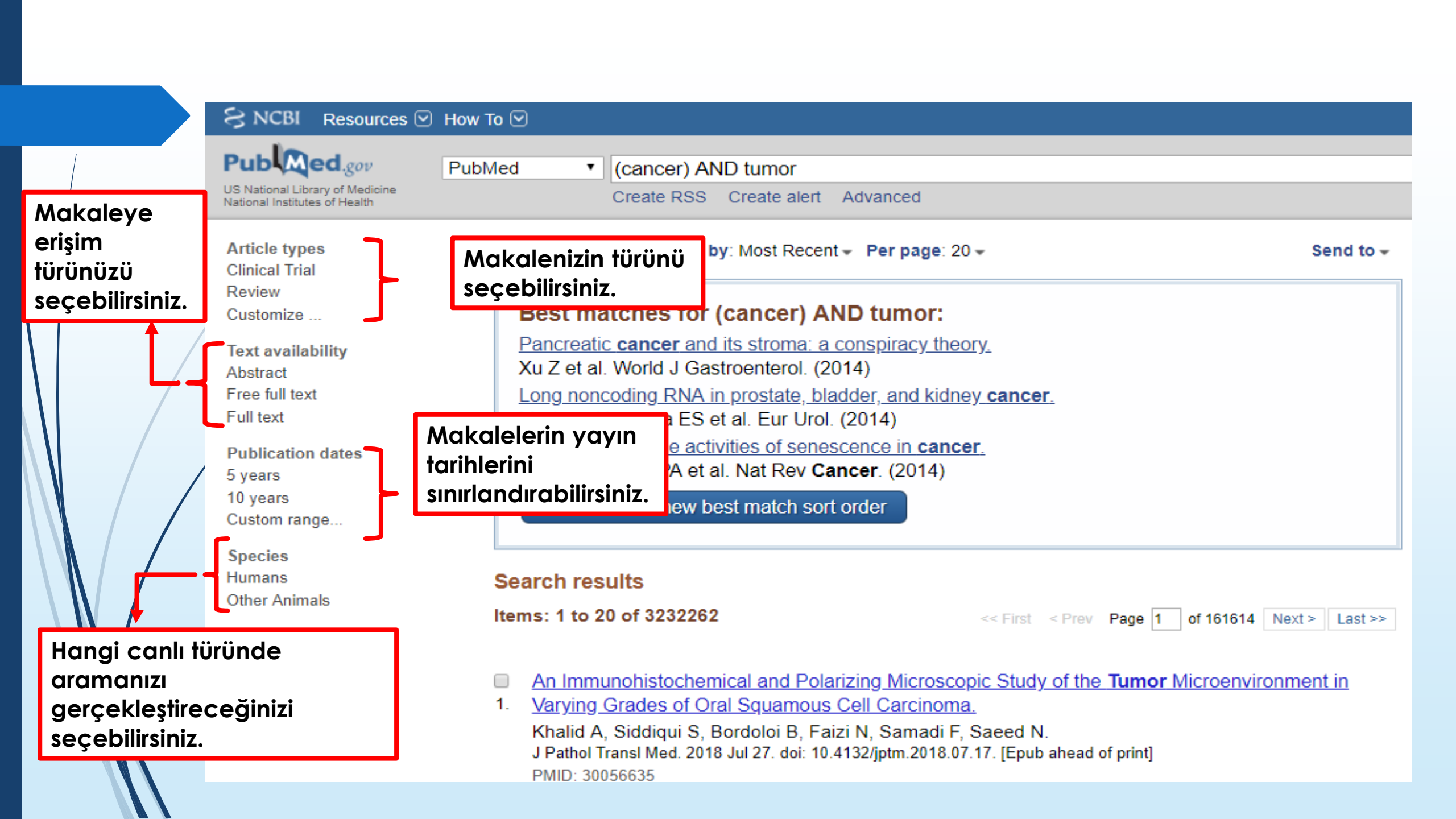
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Pancreatic cancer and its stroma: a conspiracy theory.
Xu Z et al. World J Gastroenterol. (2014)
Long noncoding RNA in prostate, bladder, and kidney cancer.
ES et al. Eur Urol. (2014)
activities of senescence in cancer.
A et al. Nat Rev Cancer. (2014)
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1. An Immunohistochemical and Polarizing Microscopic Study of the Tumor Microenvironment in Varying Grades of Oral Squamous Cell Carcinoma.
Khalid A, Siddiqui S, Bordoloi B, Faizi N, Samadi F, Saeed N.
J Pathol Transl Med. 2018 Jul 27. doi: 10.4132/jptm.2018.07.17. [Epub ahead of print]
PMID: 30056635


Format: Summary
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Best matches for cancer:
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Torre LA et al. CA Cancer J Clin. (2015)
Metabolic syndrome and risk of cancer: which link?
Mendonça FM et al. Metabolism. (2015)
LRIG and cancer prognosis.
Lindquist D et al. Acta Oncol. (2014)
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2018 Aug 9.
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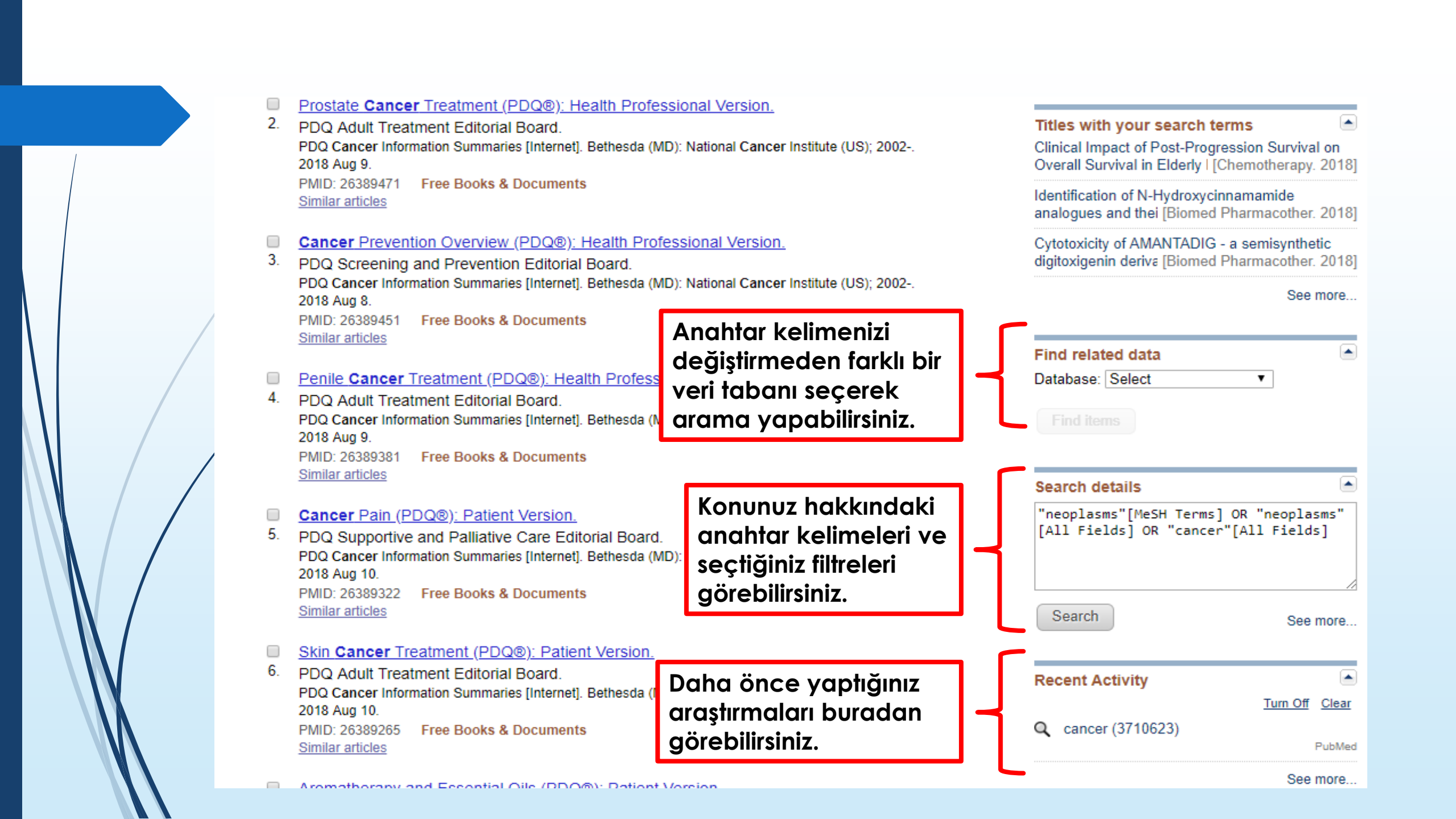
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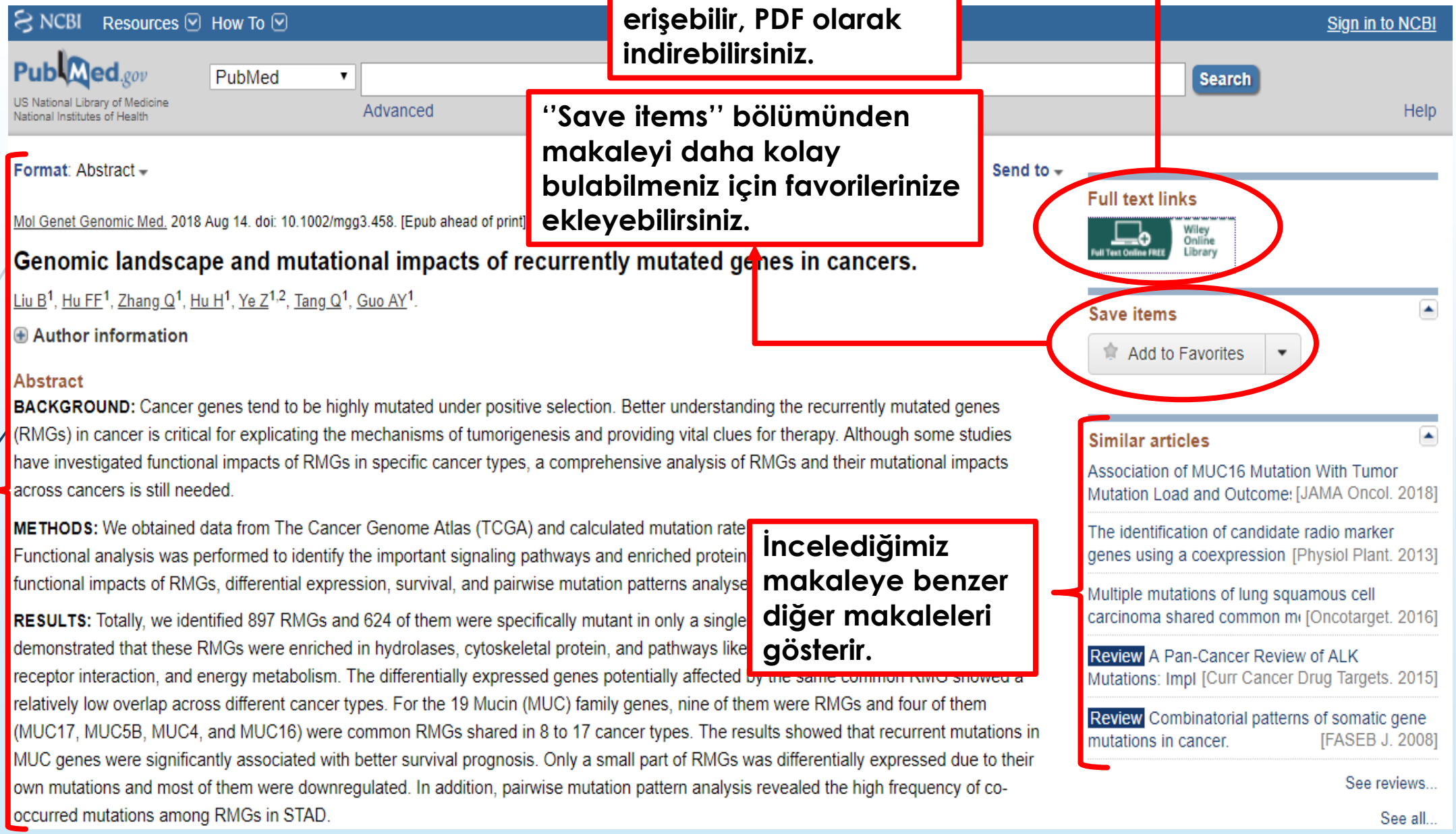

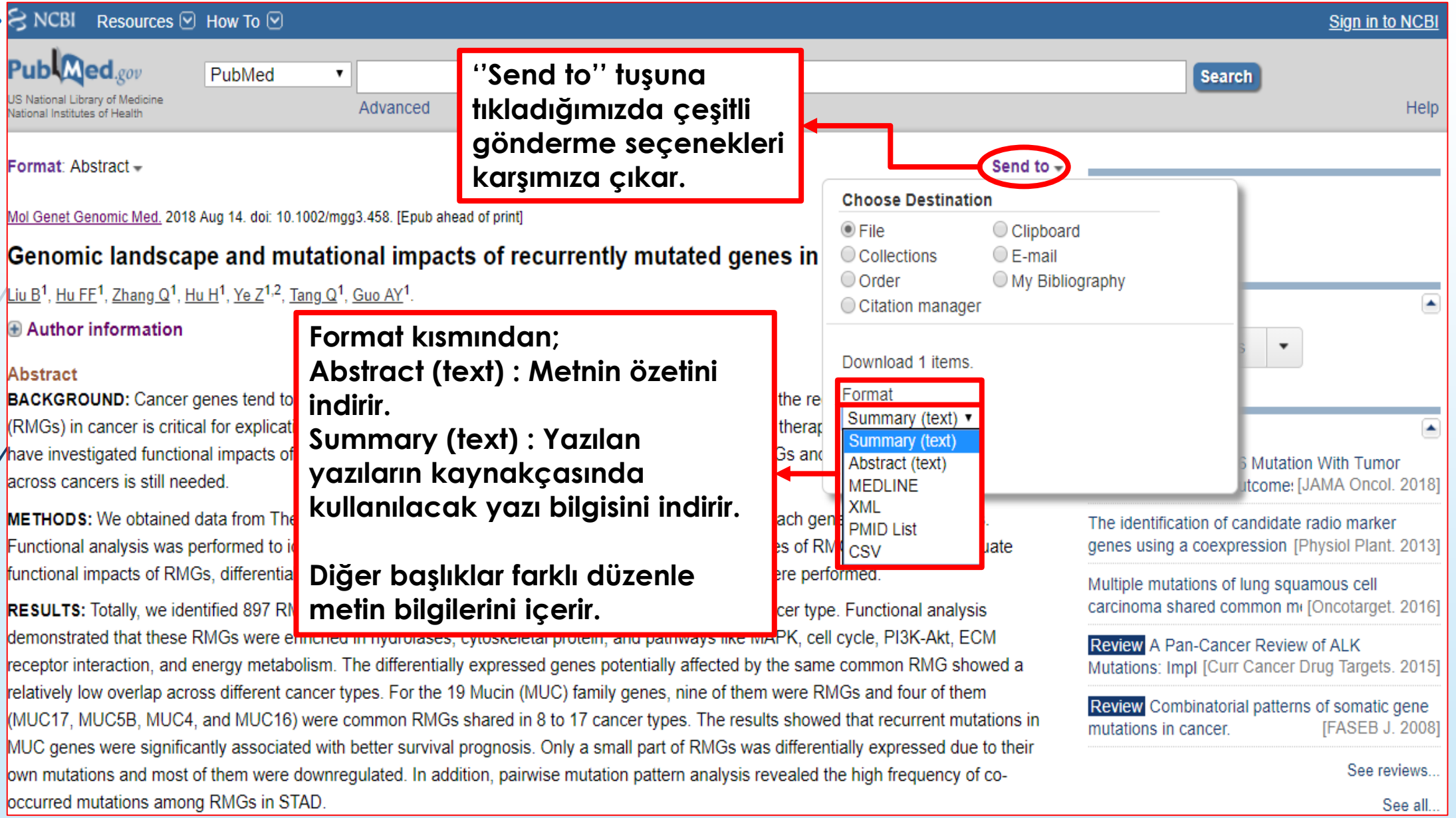
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Mol Genet Genomic Med. 2018 Aug 14. doi: 10.1002/mgg3.458. [Epub ahead of print]
Genomic landscape and mutational impacts of recurrently mutated genes in
Liu B[1], Hu FF[1], Zhang Q[1], Hu H[1], Ye Z[1,2], Tang Q[1], Guo AY[1].
Author information
Abstract
BACKGROUND: Cancer genes tend to
(RMGs) in cancer is critical for explicati
have investigated functional impacts of
across cancers is still needed.
METHODS: We obtained data from The
Functional analysis was performed to i
functional impacts of RMGs, differentia
RESULTS: Totally, we identified 897 RM
demonstrated that these RMGs were enriched in hydrolases, cytoskeletal protein, and pathways like MAPK, cell cycle, PI3K-Akt, ECM
receptor interaction, and energy metabolism. The differentially expressed genes potentially affected by the same common RMG showed a
relatively low overlap across different cancer types. For the 19 Mucin (MUC) family genes, nine of them were RMGs and four of them
(MUC17, MUC5B, MUC4, and MUC16) were common RMGs shared in 8 to 17 cancer types. The results showed that recurrent mutations in
MUC genes were significantly associated with better survival prognosis. Only a small part of RMGs was differentially expressed due to their
own mutations and most of them were downregulated. In addition, pairwise mutation pattern analysis revealed the high frequency of co-
occurred mutations among RMGs in STAD.
Mutation With Tumor
[JAMA Oncol. 2018]
The identification of candidate radio marker genes using a coexpression [Physiol Plant. 2013]
Multiple mutations of lung squamous cell carcinoma shared common m [Oncotarget. 2016]
Review A Pan-Cancer Review of ALK Mutations: Impl [Curr Cancer Drug Targets. 2015]
Review Combinatorial patterns of somatic gene mutations in cancer. [FASEB J. 2008]
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1. Mol Genet Genomic Med. 2018 Aug 14. doi: 10.1002/mgg3.458. [Epub ahead of print]

Genomic landscape and mutational impacts of recurrently mutated genes in cancers.

Liu B(1), Hu FF(1), Zhang Q(1), Hu H(1), Ye Z(1)(2), Tang Q(1), Guo AY(1).

Author information:
(1)Department of Bioinformatics and Systems Biology, Key Laboratory of Molecular Biophysics of the Ministry of Education, College of Life Science and Technology, Huazhong University of Science and Technology, Wuhan, China.
(2)Department of Biochemistry and Molecular Biology, Tianjin Key Laboratory of Medical Epigenetics, Tianjin Medical University, Tianjin, China.

BACKGROUND: Cancer genes tend to be highly mutated under positive selection. Better understanding the recurrently mutated genes (RMGs) in cancer is critical for explicating the mechanisms of tumorigenesis and providing vital clues for therapy. Although some studies have investigated functional impacts of RMGs in specific cancer types, a comprehensive analysis of RMGs and their mutational impacts across cancers is still needed.

Abstract (text) bölümünden indirdiğimiz metnin özeti.

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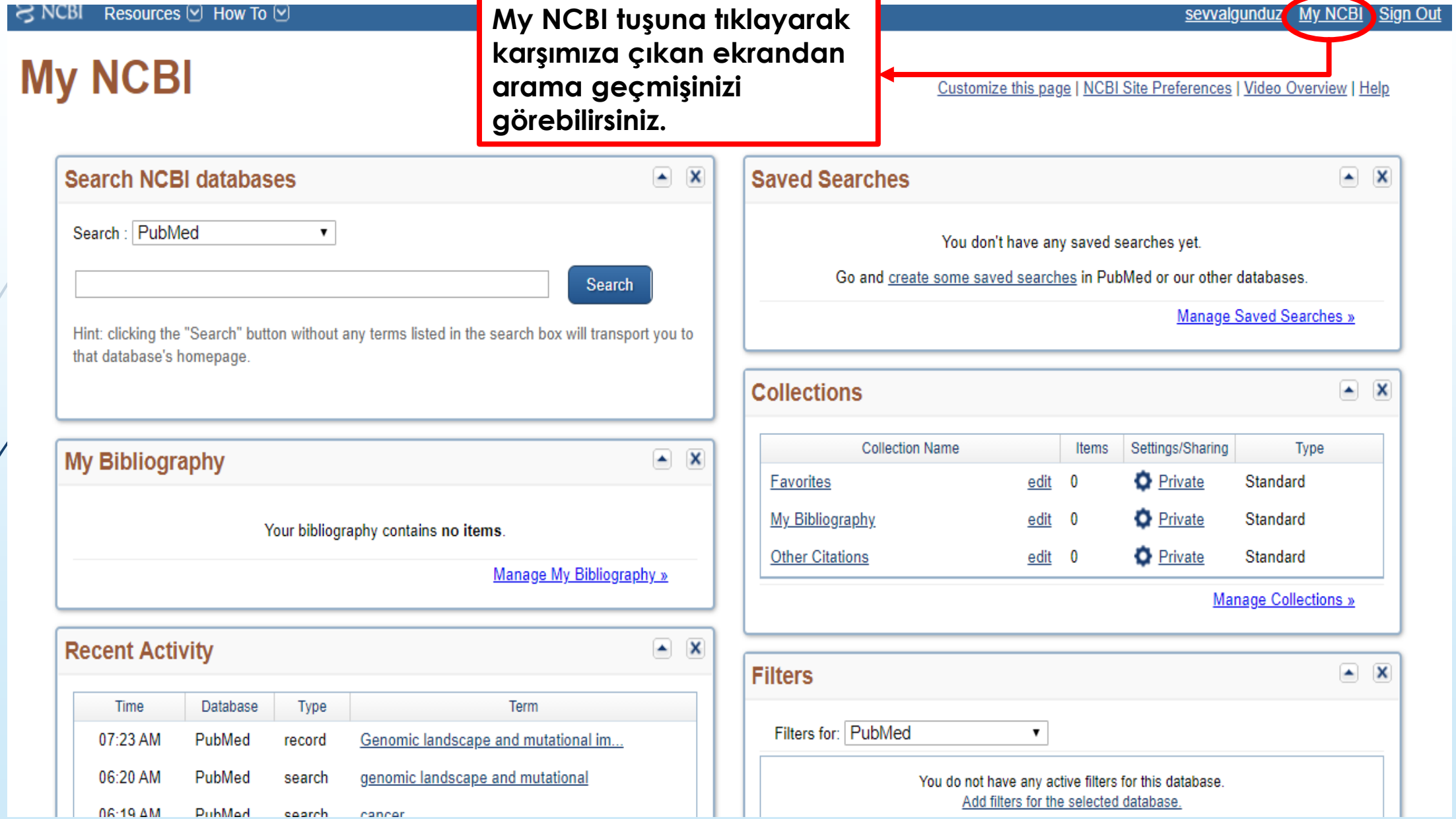
Dosya Düzen Biçim Görünüm Yardım

1: Liu B, Hu FF, Zhang Q, Hu H, Ye Z, Tang Q, Guo AY. Genomic landscape and mutational impacts of recurrently mutated genes in cancers. Mol Genet Genomic Med. 2018 Aug 14. doi: 10.1002/mgg3.458. [Epub ahead of print] PubMed PMID: 30107644.

Summary (text) bölümünden indirdiğimiz yazı bilgisi.



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