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Website: https://pubmed.ncbi.nlm.nih.gov/ https://archive.connect.h1.co/

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• 一、背景介绍

• 历史沿革

• 海量文献与文献评估

•二、利用在PubMed使用H1 Connect获取重

要的科学文献,提升科研效率

经验告诉我们,往往对我们有所启发,真正具有指 导作用的只是少量的重要文献 利用H1 Connect 的智囊,帮我们筛选经典文献



在充满挑战的科研道路上,分享"大 牛"们的宝贵经验

H1 Connect 可以提供什么帮助

帮助我找出所有重要的关键文章。

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- 节省我节选阅读重要文献的时间。
- 经过两次以上的同行评审更能确认文章的重要性及可读性。

- 1. Article Recommendations: 遴选出近期某领域研究中有着重要价值和
 - 意义的文章,并从意义、创新点、方法等方面进行评价
- 3. Collections:为用户提供了少量基于主题的文章推荐数据集,对文章 推荐进行了分类聚合。
- 4. Faculty Reviews: 学科专家不定期的对各自亚学科专业领域的题目 进行综述,提供给研究同行更加系统全面的专业内容。

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1974 2020 TEXT AVAILABILITY Abstract Free full text Full text ARTICLE ATTRIBUTE Associated data ARTICLE TYPE Books and Documents	 Pembrolizumab versus docetaxel for previously treated, non-small-cell lung cancer (KEYNOTE-010): a randomisi Cite Herbst RS, Baas P, Kim DW, Felip E, Pérez-Gracia JL, Han JY, Mo MJ, Majem M, Fidler MJ, de Castro G Jr, Garrido M, Lubiniecki C Filhart M, Garon EB. Lancet. 2016 Apr 9;387(10027):1540-1550. doi: 10.1016/S0140- Dec 19. PMID: 26712084 Clinical Trial. BACKGROUND: Despite recent advances in the treatment of ad cancer, there remains a need for effective treatments for progr the efficacy of pembrolizumab for patients with previously trea 	PD-L1-positive, advanced ed controlled trial. Jina J, Kim JH, Arvis CD, Ahn 3M, Shentu Y, Im E, Dolled- 6736(15)01281-7. Epub 2015 dvanced non-small-cell lung ressive disease. We assessed ated, PD-L1-positive, advanc
Clinical Trial Meta-Analysis	Metastatic-niche labelling reveals parenchymal cells with Ombrato L, Nolan E, Kurelac I, Mavousian A, Bridgeman VL, He Horswell S, Gonzalez-Gualda E, Matacchione G, Weston A, Kirke	h stem features. inze I, Chakravarty P, natrick I, Husain F, Speirs V.

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The genome of the obligate intracellular parasite Trachipleistophora hominis: new insights into microsporidian genome dynamics and reductive evolution.

Heinz E et al.

https://doi.org/10.1371/journal.ppat.1002979

PLoS Pathogens. 2012; 8(10):e1002979

PMID: 23133373

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Classifications

New Finding

Evaluations

Exceptional

08 Nov 2012

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William Martin

Microsporidians are central to our current understanding of eukaryote evolution, and this new microsporidian genome contains crucial information that pins the origin of pathogenicity deep within this group. It wasn't long ago that microsporidians were thought to be 'early branching eukaryotes' but that view has changed dramatically, due in no small part to the circumstance that microsporidians were surprisingly found to harbour highly reduced mitochondria, or mitosomes {1-3}. This new genome adds significantly to our understanding of pathogen evolution and points out new avenues to help battle this parasite, which was isolated from an HIV-infected patient.

References

1. A mitochondrial remnant in the microsporidian Trachipleistophora hominis.

Williams BA et al.

Evaluations	
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