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BioMed Central Open Access 期刊會員服務及投稿說明

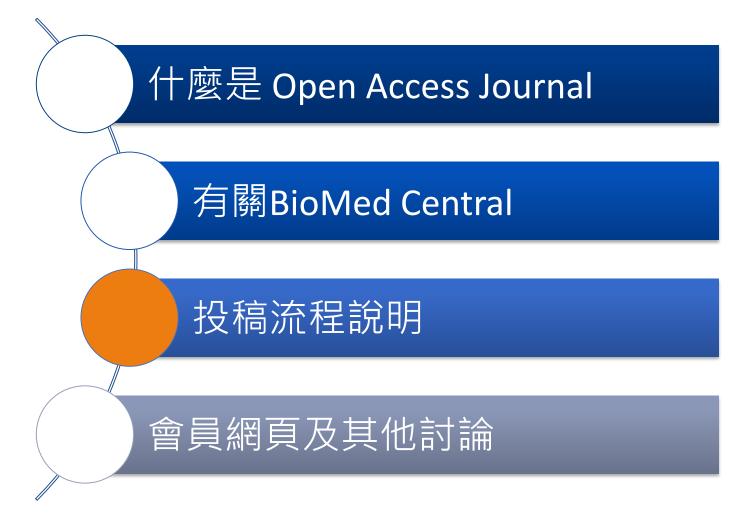
飛資得醫學 2019年10月21日







Agenda







開放取用 Open Access Journal的好處

- 增加文章的能見度、影響力和引用率
- 沒有訂閱要求,所有研究員都可充份利用研究的成果
- 預防重複性的研究
- 文章更被廣泛索引和更容易被查找
- 公開取用可以增加社會對科學和醫學研究的認知,促進科技創新
- 改善期刊的影響指數





能見度與影響力 Ahead of print

Dated: April 20, 2015



RESEARCH ARTICLE

What's in your next-generation sequence data? An exploration of unmapped DNA and RNA sequence reads from the bovine reference individual

Lynsey K. Whitacre, Polyana C. Tizioto, JaeWoo Kim, Tad S. Sonstegard, Steven G. Schroeder, Leeson J. Alexander, Juan F. Medrano, Robert D. Schnabel, Jeremy F. Taylor 🖾 and Jaced F. Decker 🖾 💿

sMC Genomics 2015 16:1114 | DOI: 10.1186/s12864-015-2313-7 | © Whitacre et al. 2015 Received: 26 August 2015 | Accepted: 15 December 2015 | Published: 29 December 2015

Abstract

Background

Next-generation sequencing projects commonly commence by aligning reads to a reference genome assembly. While improvements in alignment algorithms and computational hardware have greatly enhanced the efficiency and accuracy of alignments, a significant percentage of reads often remain unmapped.

Results

We generated *de novo* assemblies of unmapped reads from the DNA and RNA sequencing of the *Bos taurus* reference individual and identified the closest matching sequence to each contig by alignment to the NCBI non-redundant nucleotide database using BLAST. As expected, many of these contigs represent vertebrate sequence that is absent, incomplete, or misassembled in the UMD3.1 reference assembly. However, numerous additional contigs represent invertebrate species. Most prominent were several species of Spirurid nematodes and a blood-borne parasite, *Babesia bigemina*. These species are either not present in the US or are not known to infect taurine cattle and the reference animal appears to have been host to unsequenced sister species.

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Abstract

Background

Results

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Availability of data and

materials

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ETRICS

Article accesses: 496



Altmetric score from Atmetric.com

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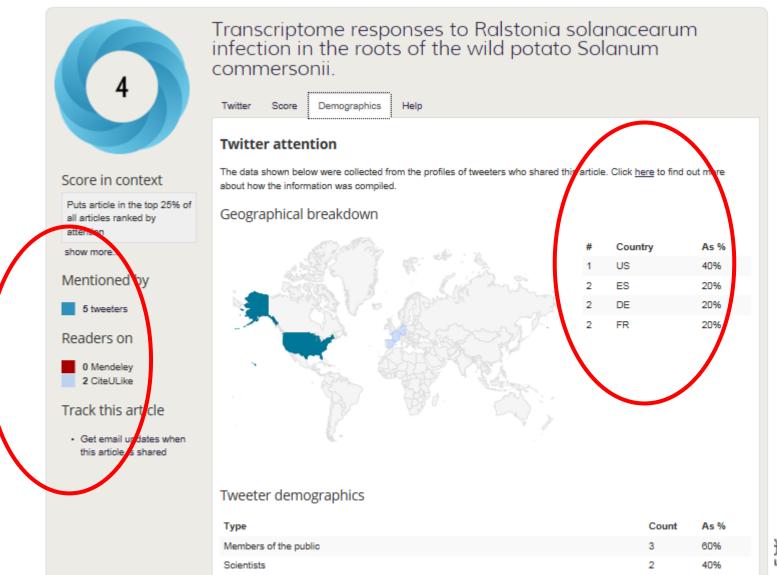






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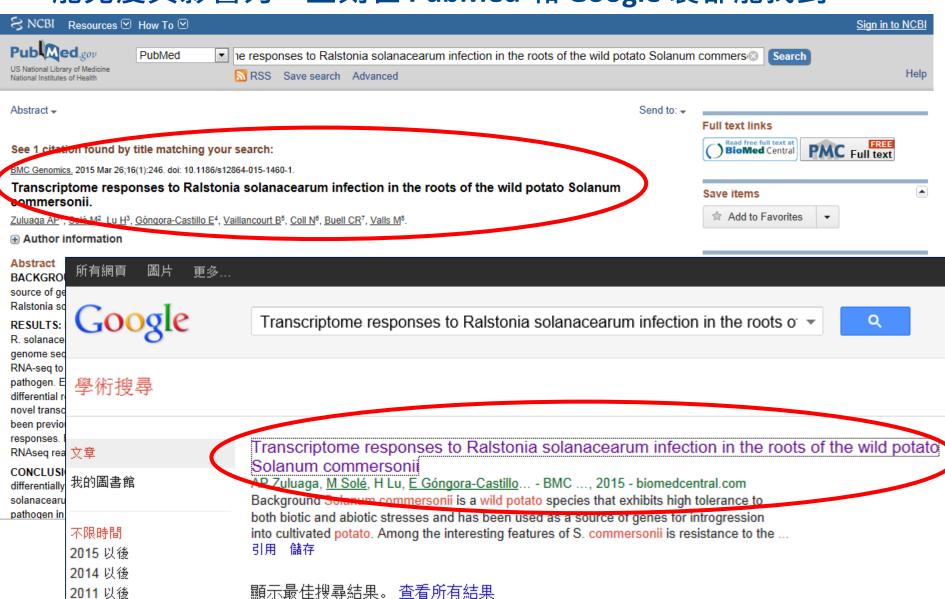


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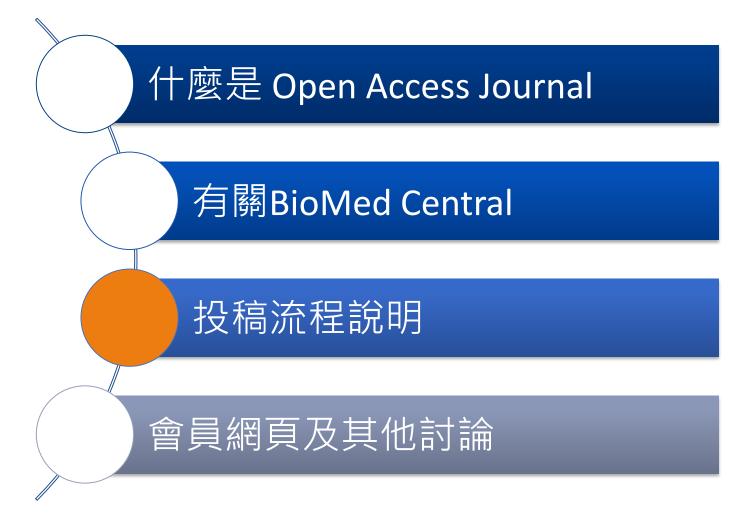








Agenda







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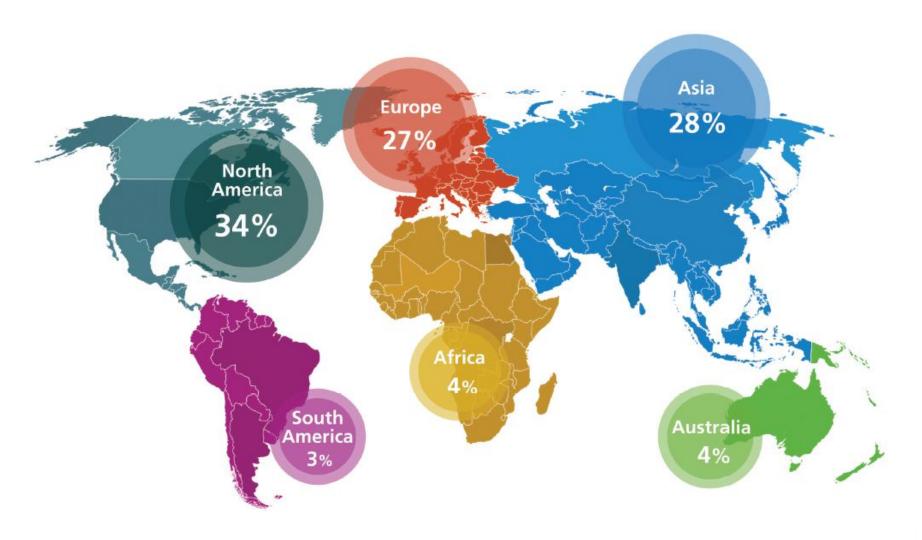








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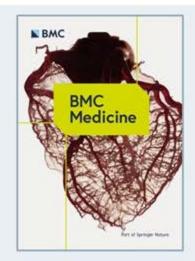
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Highlights







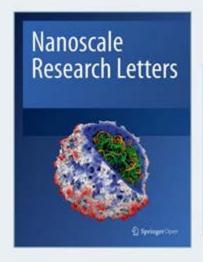
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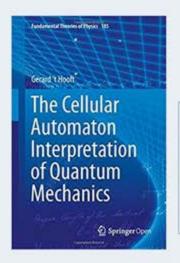
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BMC Medicine	8.005
BMC Biology	6.967
Molecular Neurodegeneration	6.51
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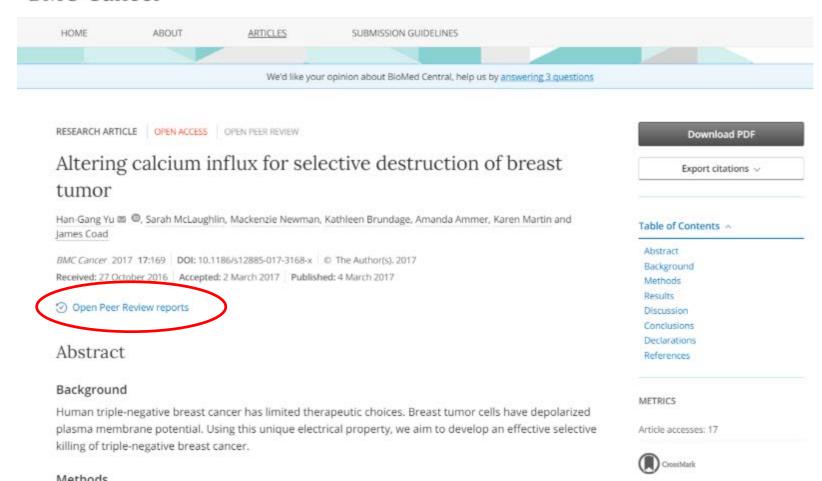
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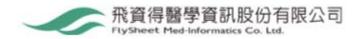




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7 Jan 2017	Author responded	Author comments - Wilfred de Vega	
Resubmission - Ve	ersion 2		
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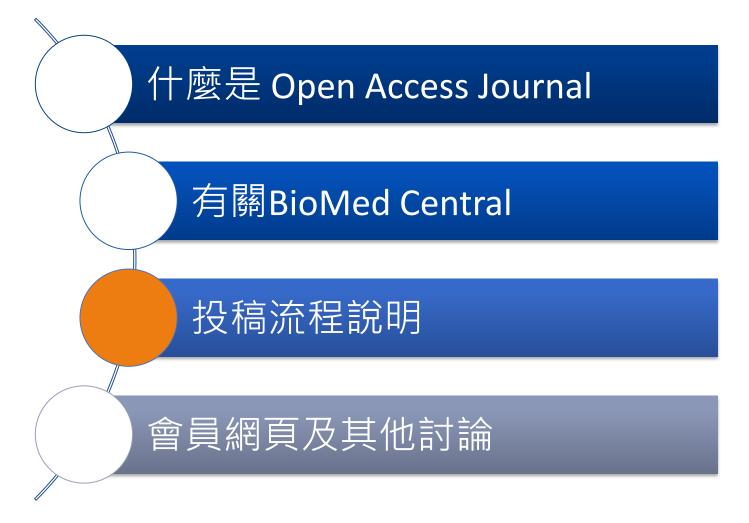
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Agenda







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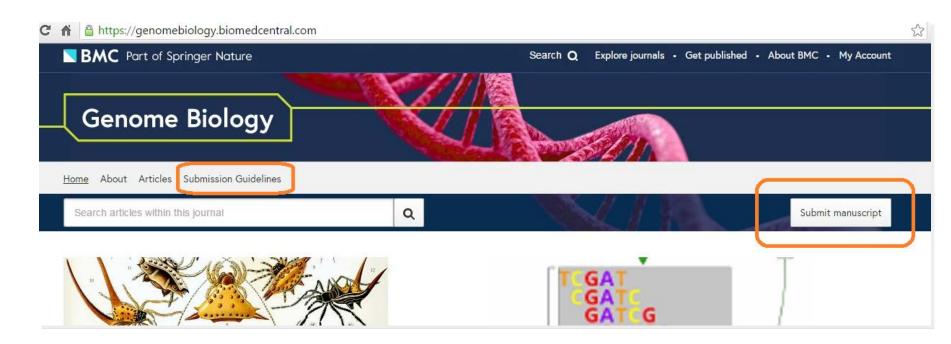
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I. 選刊: 閱讀投稿指南

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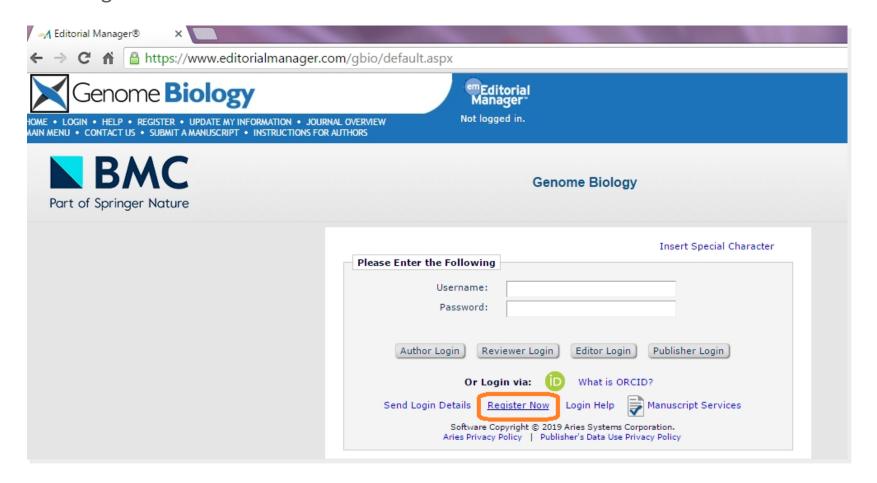






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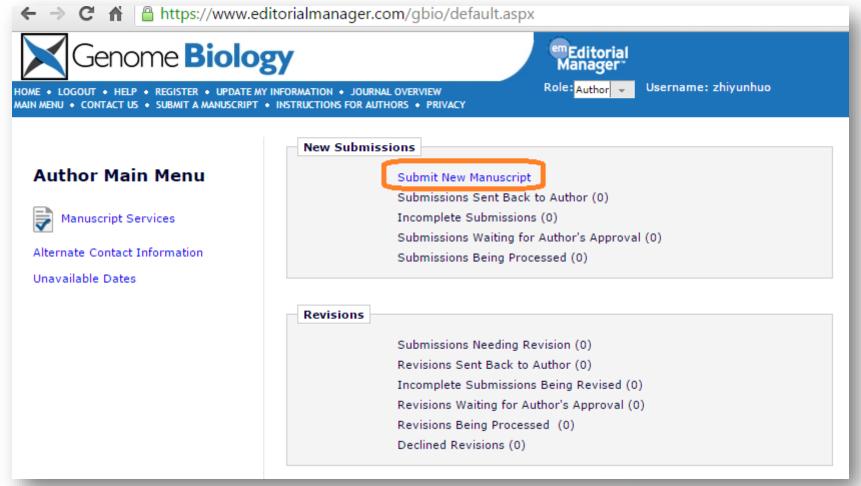






Ⅱ. 註冊

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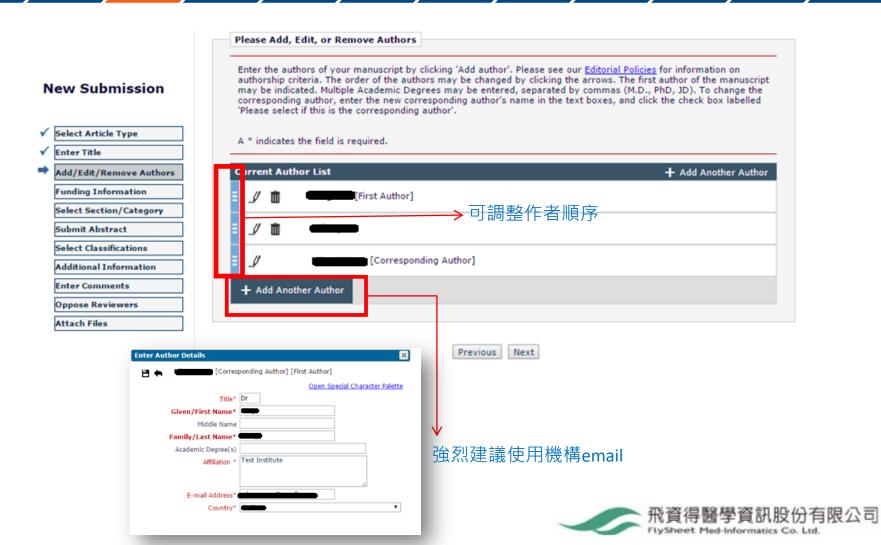
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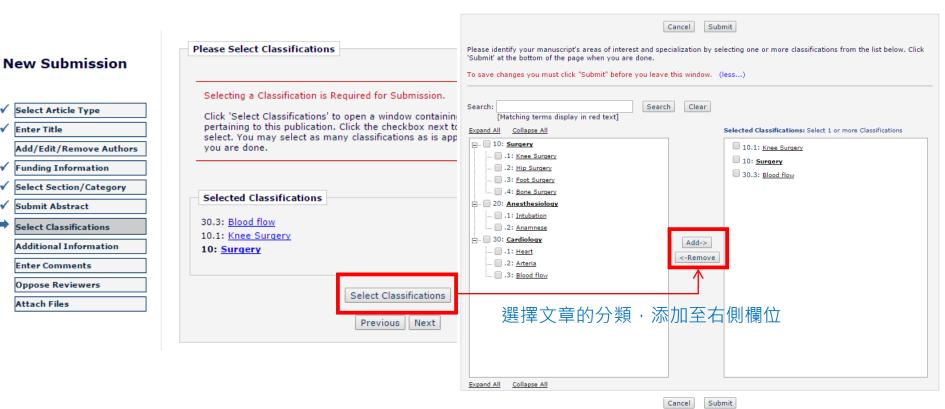
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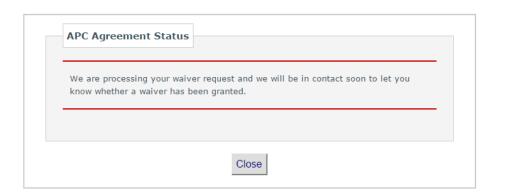
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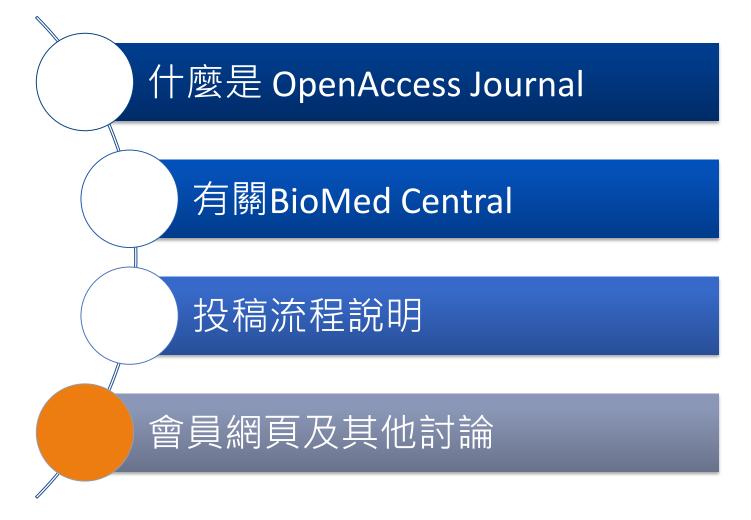
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