

احتمال علاقة معمل اوهان بوباء المنكي بوكس وما  
هو احتمالية الوباء التالي له الذي يتوقعه بيل جيتس  
قريبا؟





PHOTO FROM THE DAILY BEAST.

EXCLUSIVE

## EXC: The Infamous Wuhan Lab Recently Assembled Monkeypox Strains Using Methods Flagged For Creating 'Contagious Pathogens'.

ARE WE HERE BECAUSE OF CHINA'S EXPERIMENTS AGAIN?

خبر كارثي عن ان معمل اوهان له علاقة بالمنكي بوكس

<https://thenationalpulse.com/2022/05/22/wuhan-assembled-monkeypox-strains/>



هذا الامر ليس إشاعة بل نشره معمل اوهان نفسه في مجلته الدورية في  
فبراير 2022

Outline

Highlights

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Virologica Sinica

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In Press, Corrected Proof



Research Article

## Efficient assembly of a large fragment of monkeypox virus genome as a qPCR template using dual-selection based transformation-associated recombination

Lei Yang<sup>a, b</sup>, Lingqian Tian<sup>a, b</sup>, Leshan Li<sup>a, b</sup>, Qihong Liu<sup>a, b</sup>, Xiang Guo<sup>a, b</sup>, Yuan Zhou<sup>a</sup>, Rongjuan Pei<sup>a</sup>, Xinwen Chen<sup>a</sup> & Yun Wang<sup>a</sup>

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معمل اوهان الصيني في فبراير كان اعلن انه ركب فيروس المنكي بوكس في المعمل لكي ينتج اختبار  
للإصابة PCR

<https://www.sciencedirect.com/science/article/pii/S1995820X22000414>

فتخيلوا معى كم المصادفات هنا

معمل اوهان الذي لا يزال المتهم الأول في الكوفيد الذي قتل 6.2 مليون شخص  
ودمر صحة مئات الملايين وعلى الأقل معروف ان اول حالة إصابة ظهرت في سوق  
السّمك بجواره مباشرة وباعتراف رئيس المعمل رسميا انه كانوا يجروا تجارب على  
الكرونا بتدعيم مالي من فاوتشي واشياء كثيرة قدمتها سابقا، هو نفس المعمل  
بالصدفة البحتة قرر ان يصنع PCR للمنكي بوكس!!! وبالصدفة البحتة لأنه لا يملك  
الفيروس كامل فيقرر ان يجمعه في المعمل!!!

**“Since [Monkey pox viruses] infection has never been associated with an outbreak in China, the viral genomic material required for qPCR detection is unavailable.**

**In this report, we employed dual-selective**

“[Monkey pox viruses have strains that are] more pathogenic and [have] been reported to infect humans in various parts of the world...

وبالصدفة البحتة في تقريرهم انهم ركزوا على الأجزاء التي تصيب البشر !!!  
وبالصدفة البحتة ركزوا على اهم جينين يجعلوا المنكي بوكس يصيب البشر اكثر !!!

[transformation-associated recombination] TAR to assemble a 55-kb [Monkey pox viruses] genomic fragment that encompasses E9L and C3L,

As an efficient tool for assembling large DNA fragments up to 592 kb in length, [transformation-associated recombination] assembly has become essential for preparing infectious clones of large DNA/RNA viruses.”

وبالصدفة البحتة استخدموا طريقة TAR لنسخ وتصنيع الفيروسات المعدية كبيرة الحجم!!!  
وبالصدفة البحتة يقرّوا بأن هذه التجارب على المنكي بوكس هي خطيرة لأنها تحتوى على كل الجينات  
الخطيرة فبكل سهولة يكتمل الفيروس المركب جينيا وينشر فيروس وبائي جديد!!!

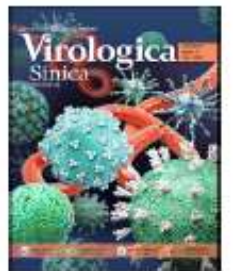
“applied in virological research could also raise potential security concerns, especially when the

assembled product contains a full set of genetic material that can be recovered into a contagious pathogen.”





ما هو الوباء التالي الذي يتوقعه اشخاص مثل بيل جيتس؟



Since mid-2016, the low pathogenic H7N9 influenza virus has evolved into a highly pathogenic (HP) phenotype in China, raising many concerns about public health and poultry industry. The insertion of a "KRTA" motif at hemagglutinin cleavage site (HACS) occurred in the early stage of HP H7N9 variants. During the co-circulation, the HACS of HP-H7N9 variants were more polymorphic in birds and humans. Although HP-H7N9 variants, unlike the H5 subtype virus, exhibited the insertions of basic and non-basic amino acids, the underlying function of those insertions and substitutions remains unclear. The results of bioinformatics analysis indicated that the PEVPKRKRTAR/G motif of HACS had become the dominant motif in China. Then, we generated six H7N9 viruses bearing the PEIPKGR/G, PEVPKGR/G, PEVPKRKRTAR/G, PEVPKGKRTAR/G, PEVPKGKRIAR/G, and PEVPKRKRRR/G motifs. Interestingly, after the deletion of threonine and alanine (TA) at HACS, the H7N9 viruses manifested decreased thermostability and virulence in mice, and the PEVPKRKRTAR/G-motif virus is prevalent in birds and humans probably due to its increased transmissibility and moderate virulence. By contrast, the insertion of non-basic amino acid isoleucine and alanine (IA) decreased the transmissibility in chickens and virulence in mice. Remarkably, the I335V substitution of H7N9 virus enhanced infectivity and transmission in chickens, suggesting that the combination of mutations and insertions of amino acids at the HACS promoted replication and pathogenicity in chickens and mice. The ongoing evolution of H7N9 increasingly threatens public health and poultry industry, so, its comprehensive surveillance and prevention of H7N9 viruses should be pursued.

Research Article

Combined insertion of basic and non-basic amino acids at hemagglutinin cleavage site of highly pathogenic H7N9 virus promotes replication and pathogenicity in chickens and mice

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ABSTRACT

Since mid-2016, the low pathogenic H7N9 influenza virus has evolved into a highly pathogenic (HP) phenotype in China, raising many concerns about public health and poultry industry. The insertion of a "KRTA" motif at hemagglutinin cleavage site (HACS) occurred in the early stage of HP H7N9 variants. During the co-circulation,

الكوارث للأسف لا تتوقف بل الكارثة الأخرى في نفس المجلة ان معمل اوهان يجري  
تجارب على انفلونزا الطيور H7N9 الخطير جدا الذي يخشى العلماء انه ينتشر  
للشعر ويصبح مميت جدا ولكن معمل اوهان يجري تجارب بإضافة جينات أكثر عدائية  
gain of function

The I335V substitution of H7N9 virus enhanced infectivity and transmission in chickens, suggesting that the combination of mutations and insertions of amino acids at the HAAS promoted replication and pathogenicity in chickens and mice.”

أي بمعنى آخر تجعله ينتشر أكثر ويميت أكثر وتجربته على الدجاج  
كطيور والفئران كتحديات أيضا

“highly transmissible in ferrets,  
posing the threats of pandemic  
potential in humans.”

بل الكارثة انهم يعترفوا ان بهذا سيكون انتقاله للبشر اسهل وممكن

يسبب Pandemic

“Surprisingly, the monobasic mutants showed clear symptoms of infection, including shaggy hair and abdominal breathing, after the second day of infection,

followed by 15% of weight loss compared to the initial weight and death, with a mortality rate of 35%. In addition, each H7N9 viruses can replicate efficiency in lungs of the infected mice...

significantly increased viral titers in the lungs of mice compared to that of the [the control] group, indicating that the insertion of KRTA increased the pathogenicity of mice.”

بل يعترفوا بانهم نجحوا ان يجعلوه مميت للفئران بنسبة 35% عن طريق  
الإضافات الجديدة

“Notably, we observed that the insertion of non-basic amino acids TA and IA in H7N9 viruses caused death in mice despite having the same viral titer,

which suggested that HPAIV H7N9 inserted with a non-basic amino acid posed a potential threat to humans.”

وفي احد التجارب جعلوه يقتل الفئران رغم انه نفس نسبة الفيروسات  
وبهذا يكون خطير جدا على البشر

ولهذا سأجعل نفسي مكان النبي بيل جيتس واتبأ بان الوباء القادم بعد  
كوفيد ومنكي بوكس هو غالباً نوع H7N9 من انفلونزا الطيور وقد يكون  
قاتل بنسبة مرتفعة اعلى بكثير من الكوفيد

وربنا يرحمنا

# والمجد لله دائماً

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