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Abstract The 1918 pandemic is still unique in the history of flu pandemics. The pathogenicity of the virus was extreme, and young adults more than infants and old people were its main victims. Many a death was caused by complications.

### [1918 lurks in everybody's mind. The "Spanish" flu and us]

For the first time, this deadly 1918 Spanish flue virus has been reconstructed and characterized. This research, reported in this week's issue of Science, is part of a larger research initiative being led by Adolfo Garcia-Sastre, PhD, Professor of Microbiology and Peter Palese, PhD, Professor and Chairman of Microbiology at Mount Sinai School of Medicine.

### Deadly 1918 Spanish flue virus has been reconstructed and ...

Efficacy of Whole-Cell Killed Bacterial Vaccines in Preventing Pneumonia and Death during the 1918 Influenza Pandemic Yu-Wen Chien , Emory University Keith P. Klugman , Emory University

### Efficacy of Whole-Cell Killed Bacterial Vaccines in ...

The basic reproductive rate (R<sub>0</sub>) for SARS-CoV-2 is estimated to be 2.5 (range 1.8-3.6) compared with 2.0-3.0 for SARS-CoV and the 1918 influenza pandemic, 0.9 for MERS-CoV, and 1.5 for the 2009 influenza pandemic. SARS-CoV-2 causes mild or asymptomatic disease in most cases; however, severe to critical illness occurs in a small ...

### Comparing SARS-CoV-2 with SARS-CoV and influenza pandemics

The devastating influenza pandemic known as 'Spanish flu', which killed at least 20 million people all over the world in 1918, was responsible for the first bitter blow inflicted on triumphant bacteriology, fortified by the series of resounding successes achieved in identifying the pathogenic agents of terrible diseases such as anthrax, cholera, tuberculosis, plague, and syphilis.

### Scientific Triumphalism and Learning from Facts ...

In 1943, EHK completed his Ph.D. in Medical Bacteriology at the University of Wisconsin, and received his MD from the University of California at San Francisco in 1947. EHK was an intern at the Mallory Institute of Pathology at Boston City Hospital (BCH) from 1947 to 1948, beginning a 43-year association with Harvard Medical School (HMS).

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Based on ClustalW v2.1 sequence alignment, the rRNA gene amplified from a pink bacterial colony had a 99% pairwise identity with that of *Roseimicrobium gellanilyticum*, a species of *Verrucomicrobia* isolated from a soil consortium in Japan (10). Therefore, we designated our isolate *Roseimicrobium* sp. strain ORNL1.

### Complete Genome Sequence of the Novel *Roseimicrobium* sp ...

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### (PDF) Traces of SARS-CoV-2 RNA in the Blood of COVID-19 ...

Bacterial culture results in autopsy series involving 96 postmortem cultures of lung tissue from victims of the 1918-1919 influenza pandemic. In the 14 of 96 autopsy series that did not report the predominance of lung pneumopathogens [29, 36, 41-53], pneumopathogens accounted collectively for 37.4% of pneumonia deaths.

### Predominant Role of Bacterial Pneumonia as a Cause of ...

Genetic relationships between the 1918 pandemic strain and strains that caused the clinically mild first wave of epidemics in 1918 and pandemics before 1918 remain undefined (9-11). It is commonly believed that the 1918 pandemic resulted from the sudden emergence and worldwide spread of an inherently hypervirulent influenza strain.

### Pathogenic Responses among Young Adults during the 1918 ...

SUMMARY Identification of gram-negative bacilli, both enteric and nonenteric, by conventional methods is not realistic for clinical microbiology laboratories performing routine cultures in today's world. The use of commercial kits, either manual or automated, to identify these organisms is a common practice. The advent of rapid or "spot" testing has eliminated the need for some commonly ...

### Manual and Automated Instrumentation for Identification of ...

Thirty strains of fermentative coryneform-like bacteria designated CDC fermentative coryneform group 3 and coryneform group 5 were compared biochemically by cellular fatty acid analysis and by DNA relatedness with the type strain of *Dermabacter hominis*, ATCC 49369. DNA from 22 strains of both CDC groups showed 69 to 96% relatedness (hydroxyapatite method) to labeled DNA from ATCC 49369 and to ...

### Recognition of *Dermabacter hominis*, formerly CDC ...

Posttranscriptional modification in tRNA is known to play a multiplicity of functional roles, including maintenance of tertiary structure and cellular adaptation to environmental factors such as temperature. Nucleoside modification has been studied in unfractionated tRNA from three psychrophilic bacteria (ANT-300 and *Vibrio* sp. strains 5710 and 29-6) and one psychrotrophic bacterium ...

### Posttranscriptional modification of tRNA in psychrophilic ...

*Nocardia* species are aerobic gram-positive bacteria of the Actinomycetales order, soil saprophytes involved in the decomposition of plant material (2, 7, 10). Isolation from respiratory samples may be indicative of colonization or invasive infection. The criteria used to determine the clinical significance of a culture positive for a *Nocardia* sp. include signs and symptoms in the patient and ...

### First Spanish Case of Nocardiosis Caused by *Nocardia* ...

----- EPA-600/2-77-092 May 1977 LIVESTOCK AND THE ENVIRONMENT A Bibliography with Abstracts Volume IV by M. L. Rowe Linda Merryman Animal Waste Technical Information Center School of Environmental Science East Central Oklahoma State University Ada, Oklahoma 74820 Grant No. R801454-04 Project Officer R. Douglas Kreis Robert S. Kerr Environmental Research Laboratory Ada, Oklahoma 74820 ROBERT S ...

**Livestock and the Environment: a Bibliography With ...**

An infectious coronavirus disease 2019 (COVID-19) emerged in the city of Wuhan (China) in December 2019, causing a pandemic that has dramatically impacted public health and socioeconomic activities worldwide. A previously unknown coronavirus, Severe Acute Respiratory Syndrome CoV-2 (SARS-CoV-2), has been identified as the causative agent of COVID-19.

**Rescue of SARS-CoV-2 from a single bacterial artificial ...**

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**LibGuides: Center for the History of Microbiology/ASM ...**

In this announcement, we present the complete genome sequence of *Bifidobacterium adolescentis* strain ZJ2, which was isolated from a female centenarian in Anhui, China. The final genome consists of a 2,401,766-bp chromosome with a G+C content of 59.90%.

**Complete Genome Sequence of *Bifidobacterium adolescentis* ...**

Peptidoglycan (PG) is essential in most bacteria. Thus, it is often targeted by various assaults, including interbacterial attacks via the type VI secretion system (T6SS). Here, we report that the Gram-negative bacterium *Acinetobacter baumannii* strain ATCC 17978 produces, secretes, and incorporates the noncanonical d-amino acid d-lysine into its PG during stationary phase.

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