

Grippe Aviaire

Février 2007

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10 questions sur la grippe aviaire

Titre : 10 questions sur la grippe aviaire

Auteur(s) : COISNE Sophie

Source : Recherche Paris 1970. 2006; (393) : 42-45

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CODEN : RCCHBV

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Type de document : Serial

Nombre de références : 6 ref.

Code(s) de classement : 002B05C03

Descripteur(s) anglais

Descripteur(s) : Avian influenza; Influenza; Human; Chicken; Risk factor; Mutation; Vaccine; Antiviral; Treatment; Avian influenza; Influenzavirus AH5N1

Desc. génériques : Virology; Infectious diseases; Medical sciences; Influenzavirus A; Orthomyxoviridae; Virus; Viral disease; Infection; Aves; Vertebrata

Descripteur(s) français

Descripteur(s) : Influenzavirus aviaire; Grippe; Homme; Poulet; Facteur risque; Mutation; Vaccin; Antiviral; Traitement; Pandemie; Grippe aviaire; Influenzavirus AH5N1

Desc. génériques : Virologie; Maladies infectieuses; Sciences médicales; Influenzavirus A; Orthomyxoviridae; Virus; Virose; Infection; Aves; Vertebrata

Localisation : INIST, Shelf number 6273, INIST No. 354000134348330010

Origine de la notice : INIST

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Transcutaneous immunization with inactivated influenza virus induces protective immune responses

Titre : Transcutaneous immunization with inactivated influenza virus induces protective immune responses

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Source : Vaccine . 2006; 24 (35-36) : 6110-6119

ISSN : 0264-410X

CODEN : VACCDE

Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Nombre de références : 48 ref.

Résumé : The recent outbreaks of highly pathogenic avian influenza in Asia and spread of the disease worldwide highlight the need to redefine conventional immunization approaches and establish effective mass vaccination strategies to face global pandemics. Transcutaneous immunization (TCI) is a novel route for vaccination, which uses the topical application of vaccine antigens on the skin. In this study, we investigated the potential of TCI using inactivated whole influenza virus. We found that TCI with whole inactivated influenza virus induced influenza virus-specific antibodies with hemagglutination inhibition and neutralizing activities as well as cellular immune responses, even without an adjuvant, and conferred protective immunity to virus challenge. Co-administration with cholera toxin (CT), a potent adjuvant for TCI, significantly enhanced immune responses against the influenza virus antigen. To enhance penetration of the skin barrier to the particulate influenza viral antigens, we tested the effects of the potential penetration enhancers/immunomodulators oleic acid (OA) and retinoic acid (RA). Pretreatment of mouse skin with OA elicited increased levels of influenza virus-specific binding and neutralizing antibodies to levels equivalent to those induced by intranasal immunization with inactivated influenza virus. OA and RA treatments differentially affected the pattern of cytokine production upon stimulation with influenza viral antigen and provided enhanced protection. These results reveal a promising perspective for the application of transcutaneous immunization to prevent influenza epidemics as well as a range of other infectious diseases.

Code(s) de classement : 002A05F04; 002A05C10

Descripteur(s) anglais

Descripteur(s) : Influenzavirus; Percutaneous route; Immunization; Vaccine; Immunoprotection

Desc. génériques : Immunology; Pharmacology; Applied microbiology; Microbiology; Biological sciences; Virology; Microbiology; Biological sciences; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Influenzavirus; Voie percutanee; Immunisation; Vaccin; Immunoprotection

Desc. génériques : Immunologie; Pharmacologie; Microbiologie appliquée; Microbiologie; Sciences biologiques; Virologie; Microbiologie; Sciences biologiques; Orthomyxoviridae; Virus

Localisation : INIST, Shelf number 20289, INIST No. 354000133528780090

Origine de la notice : INIST

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Chemical-biological terrorism and its impact on children

Titre : Chemical-biological terrorism and its impact on children

Auteur(s) : Committee on Environmental Health and Committee on Infectious Diseases, Unknown

Source : Pediatrics Evanston. 2006; 118 (3) : 1267-1278

ISSN : 0031-4005

CODEN : PEDIAU

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 53 ref.

Résumé : Children remain potential victims of chemical or biological terrorism. In recent years, children have even been specific targets of terrorist acts. Consequently, it is necessary to address the needs that children would face after a terrorist incident. A broad range of public health initiatives have occurred since September 11, 2001. Although the needs of children have been addressed in many of them, in many cases, these initiatives have been inadequate in ensuring the protection of children. In addition, public health and health care system preparedness for terrorism has been broadened to the so-called all-hazards approach, in which response plans for terrorism are blended with plans for a public health or health care system response to unintentional disasters (eg, natural events such as earthquakes or pandemic flu or manmade catastrophes such as a hazardous-materials spill). In response to new principles and programs that have appeared over the last 5 years, this policy statement provides an update of the 2000 policy statement. The roles of both the pediatrician and public health agencies continue to be emphasized; only a coordinated effort by pediatricians and public health can ensure that the needs of children, including emergency protocols in schools or child care centers, decontamination protocols, and mental health interventions, will be successful.

Code(s) de classement : 002B01

Descripteur(s) anglais

Descripteur(s) : Biology; Terrorism; Child; Emergency; Preparation; Dragging; Teaching; Pediatrics

Desc. génériques : Medical sciences; Human

Descripteur(s) français

Descripteur(s) : Biologie; Terrorisme; Enfant; Urgence; Preparation; Entrainement; Enseignement; Pédiatrie; Etat de preparation

Desc. génériques : Sciences médicales; Homme

Localisation : INIST, Shelf number 6967, INIST No. 354000133590140500

Origine de la notice : INIST

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A microsphere immunoassay for detection of antibodies to avian influenza virus

Titre : A microsphere immunoassay for detection of antibodies to avian influenza virus

Auteur(s) : DEREGT Dirk; FURUKAWA STOFFER Tara L; TOKARYK Kara L; PASICK John; BURTON HUGHES Kimberley M; HOOPER MCGREYVY Kathleen; BAXI Shailja; BAXI Mohit K

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Source : Journal of virological methods. 2006; 137 (1) : 88-94

ISSN : 0166-0934

CODEN : JMVMDH

Date de publication : 2006

Pays de publication : Netherlands

Langue(s) : English

Type de document : Serial

Nombre de références : 19 ref.

Résumé : A microsphere immunoassay (MIA) was developed for the detection of serum antibodies to avian influenza virus. A recombinant influenza A nucleoprotein expressed in baculovirus was conjugated to microspheres and incubated with antibodies. High median fluorescent intensities (MFIs) were obtained with a monoclonal antibody and positive chicken sera. Chickens were inoculated with 10 strains of avian influenza virus representing different subtypes, including high and low pathogenic H5 and H7 subtypes. Three hundred and fifty-four samples from experimentally infected chickens and controls were tested with a competitive ELISA (cELISA) and the MIA. MFIs were converted to positive/negative (PN) ratios. The results of both tests, as percentage inhibition and PN ratio, showed a high correlation ($R^{sup>2} = 0.77$). From the comparison data, a ratio of ≥ 4.5 was selected as the cut-off value for positivity in the MIA. Using this cut-off value, the sensitivity and specificity of the MIA relative to the cELISA when all discordant experimental samples were retested was 99.3 and 93.1%, respectively. The relative specificity increased to 94.7% when additional negative sera ($n = 68$) were tested. The MIA may be useful for surveillance testing and as a screening test for flocks infected with low pathogenic avian influenza virus and could be expanded for simultaneous detection of antibodies against other avian infectious disease agents.

Code(s) de classement : 002A05C09

Descripteur(s) anglais

Descripteur(s) : Avian influenza virus; Detection; Antibody; Microbiology; Method; Virology

Desc. génériques : Virology; Microbiology; Biological sciences; Influenza virus A; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Influenza virus aviaire; Detection; Anticorps; Microbiologie; Methode; Virologie

Desc. génériques : Virologie; Microbiologie; Sciences biologiques; Influenza virus A; Orthomyxoviridae; Virus

Localisation : INIST, Shelf number 18295, INIST No. 354000133543610130

Origine de la notice : INIST

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Is the price of cheap chicken bird flu?

Titre : Is the price of cheap chicken bird flu?

Auteur(s) : PERKINS Eva C

Source : Searcher Medford NJ. 2006; 14 (6) : 8-16 [6 p.]

ISSN : 1070-4795

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Résumé : Un certain nombre d'informations et de statistiques utiles concernant la grippe aviaire peuvent être trouvées sur les sites Web des organismes suivants : Centers for Diseases Control and Prevention, le Ministère de l'Agriculture des États-Unis, l'Organisation Mondiale de la Santé, la base de données Pubmed de la National Library of Medicine, Grain et BirdLife International.

Code(s) de classement : 001A01D02B; 205

Descripteur(s) anglais

Descripteur(s) : Information source; Biomedical information; Communicable disease; Poultry

Desc. génériques : Information sciences; Documentation

Descripteur(s) français

Descripteur(s) : Source information; Information biomédicale; Maladie contagieuse; Volaille; Grippe aviaire

Desc. génériques : Sciences de l'information; Documentation

Localisation : INIST, Shelf number 27485, INIST No. 354000153150220010

Origine de la notice : INIST

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A household survey to assess the burden of influenza in rural Thailand

Titre : A household survey to assess the burden of influenza in rural Thailand

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Source : Southeast Asian journal of tropical medicine and public health. 2006; 37 (3) : 488-493

ISSN : 0125-1562

CODEN : SJTMAK

Date de publication : 2006

Pays de publication : Thailand

Langue(s) : English

Type de document : Serial

Nombre de références : 13 ref.

Résumé : Little is known about the disease burden of influenza in middle-income tropical countries like Thailand. The recent outbreak of avian influenza (H5N1) and studies on influenza from neighboring countries highlight the need for data on incidence, access to care, and health care cost. In May/ June 2003, we conducted a province-wide household survey using two-stage cluster sampling to determine the burden of influenza-like illness in Sa Kaeo Province. We used the total number of reported influenza that occurred in May 2003 and a prospective study of outpatient influenza in clinic patients to develop an estimate of the annualized incidence of influenza. Of 718 subjects, 16 (2.2%) suffered an episode of influenza-like illness in the preceding month; 14 sought care, of whom 7 went to a hospital facility. Fifty percent reported missing on average 3 days of work or school. The total individual cost per illness episode was 663 baht (US\$15.78). The proportion of outpatients with influenza-like illness caused by an influenza virus in May was 16% and the annualized influenza incidence was estimated to be 5,941/100,000 in Sa Kaeo Province. This survey adds to information indicating that in rural Thailand, the burden of influenza is substantial and costs associated with an illness episode are up to 20% of an average monthly income.

Code(s) de classement : 002B01; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Influenza; Surveillance; Rural environment; Thailand; Tropical medicine

Desc. génériques : Medical sciences; Virology; Infectious diseases; Medical sciences; Viral disease; Infection; Asia

Descripteur(s) français

Descripteur(s) : Grippe; Surveillance; Milieu rural; Thaïlande; Médecine tropicale

Desc. génériques : Sciences médicales; Virologie; Maladies infectieuses; Sciences médicales; Virose; Infection; Asie

Localisation : INIST, Shelf number 19778, INIST No. 354000133569110090

Origine de la notice : INIST

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Management de la securite des aliments. De l' HACCP a l' ISO 22000

Titre : Management de la securite des aliments. De l' HACCP a l' ISO 22000

Auteur(s) : BOUTOU Olivier

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Éditeur : AFNOR, Paris

ISBN : 2124401106

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Type de document : Book

Nombre de références : dissem.

Résumé : La qualite irreprochable des produits exigee par le consommateur final, la maitrise des risques lies a la securite alimentaire et l' internationalisation des echanges sont autant d' enjeux auxquels doivent faire face les acteurs du marche agroalimentaire. De plus, les crises de ces dernieres annees (maladie de la vache folle, grippe aviaire, listeria...) ont entraine une defiance de la part des consommateurs. Afin de rendre plus clairs et plus accessibles les travaux du Codex Alimentarius et les nombreux textes reglementaires, et afin d' aider ces professionnels de l' alimentation a mettre en place des dispositifs de management oriente securite alimentaire, un ensemble de partenaires ont choisi de faire partager leurs experiences au travers des referentiels dont le plus recent est l' ISO 22000, "Systemes de management de la securite des denrees alimentaires - Exigences pour tout organisme appartenant a la chaine alimentaire". Ce referentiel, regroupant les essentiels en la matiere, est un ensemble de regles de bonnes pratiques d' hygiene, qui complete les principes de l' HACCP (Hazard Analysis Critical Contrat Point). Olivier Boutou decrypte dans cet ouvrage l' ISO 22000, ses fondements, ses principes et sa mecanique. La norme ainsi explicitee devient facilement applicable, et permet de mettre en place un management de securite des denrees alimentaires... efficient !

Code(s) de classement : 002B30A11

Descripteur(s) anglais

Descripteur(s) : Feeding; Standards; Quality; Regulation; Hygiene; Food industry; Product; Safety

Desc. génériques : Public health; Medical sciences

Descripteur(s) français

Descripteur(s) : Alimentation; Norme; Qualite; Reglementation; Hygiene; Industrie alimentaire; Produit; Securite; International standards organisation

Desc. génériques : Sante publique; Sciences medicales

Localisation : BDSP/ENSP, Shelf number 152120, CK50/0188

Origine de la notice : BDSP

Global patterns of influenza A virus in wild birds

Titre : Global patterns of influenza A virus in wild birds

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Source : Science Washington DC. 2006; 312 (5772) : 384-388

ISSN : 0036-8075

CODEN : SCIEAS

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Notes : 1/4 p. ref. et notes

Résumé : The outbreak of highly pathogenic avian influenza of the H5N1 subtype in Asia, which has subsequently spread to Russia, the Middle East, Europe, and Africa, has put increased focus on the role of wild birds in the persistence of influenza viruses. The ecology, epidemiology, genetics, and evolution of pathogens cannot be fully understood without taking into account the ecology of their hosts. Here, we review our current knowledge on global patterns of influenza virus infections in wild birds, discuss these patterns in the context of host ecology and in particular birds' behavior, and identify some important gaps in our current knowledge.

Code(s) de classement : 002A05C04

Descripteur(s) anglais

Descripteur(s) : Avian influenza virus; Public health; Fauna; Wild animal; Aves; Ecology; Epidemiology; Genetics; Evolution; Pathogenicity; Migratory; Transmission from animal to animal; Europe; Asia; Review

Desc. génériques : Virology; Microbiology; Biological sciences; Influenzavirus A; Orthomyxoviridae; Virus; Vertebrata

Descripteur(s) français

Descripteur(s) : Influenzavirus aviaire; Sante publique; Faune; Animal sauvage; Aves; Ecologie; Epidemiologie; Genetique; Evolution; Pouvoir pathogene; Migrateur; Transmission animal animal; Europe; Asie; Article synthese

Desc. génériques : Virologie; Microbiologie; Sciences biologiques; Influenzavirus A; Orthomyxoviridae; Virus; Vertebrata

Localisation : INIST, Shelf number 6040, INIST No. 354000142715070130

Origine de la notice : INIST

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The structure of H5N1 avian influenza neuraminidase suggests new opportunities for drug design

Titre : The structure of H5N1 avian influenza neuraminidase suggests new opportunities for drug design

Auteur(s) : RUSSELL Rupert J; HAIRE Lesley F; STEVENS David J; COLLINS Patrick J; YI PU LIN; BLACKBURN G Michael; HAY Alan J; GAMBLIN Steven J; SKEHEL John J

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Source : Nature London. 2006; 443 (7107) : 45-49

ISSN : 0028-0836

CODEN : NATUAS

Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Nombre de références : 34 ref.

Résumé : The worldwide spread of H5N1 avian influenza has raised concerns that this virus might acquire the ability to pass readily among humans and cause a pandemic. Two anti-influenza drugs currently being used to treat infected patients are oseltamivir (Tamiflu) and zanamivir (Relenza), both of which target the neuraminidase enzyme of the virus. Reports of the emergence of drug resistance make the development of new anti-influenza molecules a priority. Neuraminidases from influenza type A viruses form two genetically distinct groups: group-1 contains the N1 neuraminidase of the H5N1 avian virus and group-2 contains the N2 and N9 enzymes used for the structure-based design of current drugs. Here we show by X-ray crystallography that these two groups are structurally distinct. Group-1 neuraminidases contain a cavity adjacent to their active sites that closes on ligand binding. Our analysis suggests that it may be possible to exploit the size and location of the group-1 cavity to develop new anti-influenza drugs.

Code(s) de classement : 002B02S05; 002A05C04

Descripteur(s) anglais

Descripteur(s) : Avian influenza virus; Subtype; Exo <alpha> sialidase; Development; New product; Drug; Antiviral; Mechanism of action; Design

Desc. génériques : Virology; Infectious diseases; Pharmacology; Medical sciences; Virology; Microbiology; Biological sciences; Influenzavirus A; Orthomyxoviridae; Virus; O Glycosidases; Glycosidases; Hydrolases; Enzyme

Descripteur(s) français

Descripteur(s) : Influenzavirus aviaire; Soustype; Exo <alpha> sialidase; Developpement; Produit nouveau; Médicament; Antiviral; Mécanisme action; Conception; Virus H5N1

Desc. génériques : Virologie; Maladies infectieuses; Pharmacologie; Sciences médicales; Virologie; Microbiologie; Sciences biologiques; Influenzavirus A; Orthomyxoviridae; Virus; O Glycosidases; Glycosidases; Hydrolases; Enzyme

Localisation : INIST, Shelf number 142, INIST No. 354000142259820120

Origine de la notice : INIST

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Neuraminidase pharmacophore model derived from diverse classes of inhibitors

Titre : Neuraminidase pharmacophore model derived from diverse classes of inhibitors

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Source : Bioorganic and medicinal chemistry letters Print. 2006; 16 (11) : 3009-3014

ISSN : 0960-894X

Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Nombre de références : 25 ref.

Résumé : A three-dimensional pharmacophore model was developed based on 22 currently available inhibitors, which were carefully selected with great diversity in both molecular structure and bioactivity, for discovering new potent neuraminidase (NA) inhibitors to fight against avian influenza virus. The best hypothesis (Hypol), consisting of five features, namely, one positive ionizable group, one negative ionizable group, one hydrophobic point, and two hydrogen-bond donors, has a correlation coefficient of 0.902, a root mean square deviation of 1.392, and a cost difference of 72.88, suggesting that a highly predictive pharmacophore model was successfully obtained. The application of the model shows great success in predicting the activities of 88 known NA inhibitors in our test set with a correlation coefficient of 0.818 with a cross-validation of 98% confidence level. Accordingly, our model should be reliable in identifying structurally diverse compounds with desired biological activity.

Code(s) de classement : 002B02S05

Descripteur(s) anglais

Descripteur(s) : Exo <alpha> sialidase; Modeling; Three dimensional model; Enzyme inhibitor; Molecular structure; Antiviral; Avian influenza virus; Hydrogen bond; Correlation coefficient; Prediction; Cross validation; Structure activity relation

Desc. génériques : Virology; Infectious diseases; Pharmacology; Medical sciences; O Glycosidases; Glycosidases; Hydrolases; Enzyme; Influenzavirus A; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Exo <alpha> sialidase; Modelisation; Modele 3 dimensions; Inhibiteur enzyme; Structure moleculaire; Antiviral; Influenzavirus aviaire; Liaison hydrogene; Coefficient correlation; Prediction; Validation croisee; Relation structure activite; ; QSAR

Desc. génériques : Virologie; Maladies infectieuses; Pharmacologie; Sciences medicales; O Glycosidases; Glycosidases; Hydrolases; Enzyme; Influenzavirus A; Orthomyxoviridae; Virus

Localisation : INIST, Shelf number 22446, INIST No. 354000142732140390

Origine de la notice : INIST

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The Spanish influenza of 1918 in St. Louis, Missouri

Titre : The Spanish influenza of 1918 in St. Louis, Missouri

Auteur(s) : KALNINS Irene

Affiliation(s) : Saint Louis University School of Nursing, St. Louis, Missouri, United States

Source : Public health nursing Boston Mass. 2006; 23 (5) : 479-483

ISSN : 0737-1209

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 21 ref.

Résumé : In view of current concern about a possible pandemic of virulent avian influenza, it is timely to revisit the public health response to the "Spanish" influenza of 1918. St. Louis, Missouri, was the most successful of nine largest cities in limiting the death toll from influenza and pneumonia through the use of public health measures during the first 8 weeks of the epidemic. A second wave of cases increased the final death rate, but it remained below that of other major cities. Public health officials attributed the lower death rate to the early and rigorous ban on public gatherings.

Code(s) de classement : 002B30A11; 002B05C02C; 002B30A05

Descripteur(s) anglais

Descripteur(s) : Influenza; Spanish; Spain; Missouri; Nurse; Nursing; Professional practice; History; Antecedent; Case history; Public health

Desc. génériques : Public health; Medical sciences; Virology; Infectious diseases; Medical sciences; Public health; Medical sciences; Viral disease; Infection; Europe; United States; North America; America

Descripteur(s) français

Descripteur(s) : Grippe; Espagnol; Espagne; Missouri; Infirmier; Nursing; Pratique professionnelle; Histoire; Antecedent; Historique; Sante publique

Desc. génériques : Sante publique; Sciences medicales; Virologie; Maladies infectieuses; Sciences medicales; Sante publique; Sciences medicales; Virose; Infection; Europe; Etats Unis; Amerique du Nord; Amerique

Localisation : INIST, Shelf number 22029, INIST No. 354000157135800100

Origine de la notice : INIST

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Safety and immunogenicity of an inactivated adjuvanted whole-virion influenza A (H5N1) vaccine : a phase I randomised controlled trial. Commentary

Titre : Safety and immunogenicity of an inactivated adjuvanted whole-virion influenza A (H5N1) vaccine : a phase I randomised controlled trial. Commentary

Auteur(s) : STEPHENSON Join, comment; JIANGTAO LIN; JIANSAN ZHANG; XIAOPING DONG; HANHUA FANG; JIANGTING CHEN; NAN SU; QIANG GAO; ZHENSHAN ZHANG; YUXUAN LIU; ZHIHONG WANG; MENG YANG; RUIHUA SUN; CHANGGUI LI; SU LIN; MEI JI; YAN LIU; XU WANG; WOOD John; ZIJIAN FENG; YU WANG; WEIDONG YIN

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Source : Lancet British edition. 2006; 368 (9540) : 965-966,991-997 [9 p.]

ISSN : 0140-6736

CODEN : LANCAO

Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Type de document : article; comments

Nombre de références : 40 ref.

Résumé : Background Avian influenza A virus H5N1 has caused widespread infections that have resulted in severe disease or death in poultry and wild birds as well as human beings. This virus has the potential to emerge as a pandemic threat and H5N1 vaccines are being developed in many countries. Our aim was to assess the safety and immunogenicity of an inactivated adjuvanted whole-virion H5N1 vaccine. Methods A stratified randomised, placebo-controlled, double-blind phase I clinical trial was done in 120 volunteers aged 18-60 years. Volunteers were assigned to receive two doses of placebo (n=24) or an inactivated whole-virion influenza A (H5N1) vaccine with 1.25 pg (24), 2.5 pg (24), 5 pg (24), or 10 pg (24) haemagglutinin per dose with aluminium hydroxide adjuvant on day 0 and 28. Serum samples were obtained on day 0, 14, 28, 42, and 56 for haemagglutination inhibition and virus neutralisation assays. This trial is registered with the ClinicalTrials.gov registry with the number NCT00356798. Findings All four formulations of vaccines were well tolerated. No serious adverse event was reported and most local and systemic reactions were mild and transient. All formulations induced antibody responses after the first dose; the highest immune response of 78% seropositivity was seen in the 10 pg group after two vaccine doses. Two individuals dropped out: one in the 1.25 pg group (withdrew consent) and one in the 10 pg group (discontinued); one individual was also excluded from the final analysis. Interpretation A two-dose regimen of an adjuvanted 10 pg inactivated whole-virion H5N1 vaccine met all European regulatory requirements for annual licensing of seasonal influenza vaccine. Lower doses of this vaccine could achieve immune responses equivalent to those elicited by adjuvanted or non-adjuvanted split-virion vaccines. The use of a whole virion vaccine could be more adaptable to the antigen-sparing strategy recommended by WHO for protection against an influenza pandemic.

Code(s) de classement : 002B01

Descripteur(s) anglais

Descripteur(s) : Immunoprophylaxis; Toxicity; Safety; Immunogenicity; Immune response; Adjuvant; Virion; Prevention; Vaccine; Vaccination; Phase I trial; Clinical trial; Critical study; Medicine; Influenzavirus AH5N1;

Randomised controlled trial

Desc. génériques : Medical sciences

Descripteur(s) français

Descripteur(s) : Immunoprofylaxie; Toxicite; Securite; Immunogenicite; Reponse immune; Adjuvant; Virion; Prevention; Vaccin; Vaccination; Essai clinique phase I; Essai clinique; Etude critique; Medecine; Influenzavirus AH5N1; Essai randomise controle

Desc. génériques : Sciences medicales

Localisation : INIST, Shelf number 5004, INIST No. 354000143591120140

Origine de la notice : INIST

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Surveillance de la grippe aviaire chez l' Homme en France. Numero thematique. Les zoonoses en France

Titre : Surveillance de la grippe aviaire chez l' Homme en France. Numero thematique. Les zoonoses en France

Auteur(s) : BONMARIN I; LEVY BRUHL D

Auteur(s) : Institut de Veille Sanitaire INVS Saint Maurice, France; Institut de Veille Sanitaire INVS Departement des maladies infectieuses Saint Maurice, France

Source : BULLETIN EPIDEMIOLOGIQUE HEBDOMADAIRE. 2006-07-04; (27-28) : 208-

ISSN : 0245-7466

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Type de document : Serial

Code(s) de classement : 002B30A11

Descripteur(s) anglais

Descripteur(s) : Epidemiology; Sanitary surveillance; Travel; Population; High risk; France

Desc. génériques : Public health; Medical sciences; Europe

Descripteur(s) français

Descripteur(s) : Epidemiologie; Surveillance sanitaire; Voyage; Population; Risque eleve; France; Institut veille sanitaire

Desc. génériques : Sante publique; Sciences medicales; Europe

Localisation : BDSP/InVS

Origine de la notice : BDSP

Surveillance des infections a influenzavirus chez les oiseaux en France. Numero thematique. Les zoonoses en France

Titre : Surveillance des infections a influenzavirus chez les oiseaux en France. Numero thematique. Les zoonoses en France

Auteur(s) : JESTIN V; SCHMITZ A; HARS J; CHERBONNEL M; LE GALL RECLE G; PICAULT JP; FRANCAULT J

Auteur(s) : Agence Francaise de Securite Sanitaire des Aliments AFSSA Ploufragan, France; Office national de la chasse et de la faune sauvage Gieres, France; Direction generale de l' alimentation Paris, France

Source : BULLETIN EPIDEMIOLOGIQUE HEBDOMADAIRE. 2006-07-04; (27-28) : 208-209

ISSN : 0245-7466

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Type de document : Serial

Nombre de références : 4 ref.

Résumé : L' influenza aviaire est une infection des oiseaux (qu' ils soient sauvages, domestiques ou d' ornement) par des influenzavirus A. L' infection, quand elle s' exprime, se manifeste par un large eventail de signes cliniques variables selon l' hote (espece, age, statut sanitaire), la souche virale, les conditions environnementales. Neanmoins, aucun signe n' etant pathognomonique, un diagnostic de laboratoire est indispensable. Au plan de la virulence, les souches virales sont classees en deux categories : faiblement pathogenes (FP) et hautement pathogenes (HP) sur la base de criteres standardises internationaux. Au sein d' un vaste repertoire de sous-types connus, les virus influenza des oiseaux (AIV) HP ont ete associes a ce jour, a 2 exceptions pres, avec les seuls sous-types H5 et H7. Aussi, bien que l' immense majorite de ces derniers soit FP, tous les virus H5 et H7, en raison de leur propension a devenir HP apres mutation, font obligatoirement depuis 2002 l' objet d' une surveillance reguliere dans les Etats membres et sont maintenant soumis a declaration au plan international. En vue de detecter/confirmer a la fois les formes inapparentes et les formes plus ou moins cliniquement exprimees, quatre modalites de surveillance de l' influenza aviaire sont donc en place en France : - chez les volailles avec : un reseau d' epidemiovigilance clinique; des enquetes annuelles, au minimum serologiques; - dans l' avifaune sauvage avec une surveillance virologique : sur oiseaux captures ou tues a la chasse; lors de mortalites groupees (depuis septembre 2005). (Introduction)

Code(s) de classement : 002B30A11

Descripteur(s) anglais

Descripteur(s) : Surveillance; Zoonosis; Mortality; Virology; France

Desc. génériques : Public health; Medical sciences; Europe

Descripteur(s) français

Descripteur(s) : Surveillance; Zoonose; Mortalite; Virologie; France

Desc. génériques : Sante publique; Sciences medicales; Europe

Localisation : BDSP/InVS

Origine de la notice : BDSP

Les sentinelles de la vie. Le monde des vaccins

Titre : Les sentinelles de la vie. Le monde des vaccins

Auteur(s) : BERTRAND Jean Jacques; SALIOU Pierre; SEYTRE Bernard

Source : 2006; 219 p.; tabl.

Éditeur : Albin Michel, Paris

ISBN : 2226172637

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Type de document : Book

Nombre de références : dissem.

Résumé : Les menaces sont nombreuses : pandémie de grippe aviaire, chikungunya, sida, SRAS, sans parler du bioterrorisme. Les fleaux font la une de l'actualité. Un espoir : la prévention par les vaccins. Mais l'industrie pharmaceutique est-elle prête à faire face aux pandémies annoncées ? Les décideurs ont-ils une réelle volonté de vacciner la planète ? Depuis les premières inoculations des médecins chinois au Xe siècle, en passant par Jenner et Pasteur, jusqu'aux recherches actuelles qui explorent les ressources prodigieuses de la biologie moléculaire et préparent les vaccins de demain, les auteurs racontent la lutte contre les grandes épidémies et retracent l'histoire de la vaccination. Qu'est-ce qu'une maladie infectieuse ? Comment agissent les vaccins ? Pourquoi vacciner nos enfants contre la polio, le tétanos, la coqueluche et bien d'autres maladies ? Si ce livre apporte des réponses précises et nuancées à ces questions, il évoque également la monstrueuse injustice que représente la non-distribution des vaccins dans le monde : des millions d'enfants des pays en développement meurent chaque année de maladies que la vaccination systématique a fait disparaître ailleurs. Face à ce bilan dramatique, de nouvelles actions sont entreprises entre institutions publiques et initiatives privées. Les espoirs sont donc réels pour que des sentinelles puissent enfin veiller sur la vie à travers le monde

Code(s) de classement : 002B30A01

Descripteur(s) anglais

Descripteur(s) : Vaccine; Vaccination; Policy; Case history; Immunology; Microorganism; Infection; Industrialized country; Developing countries; Pharmaceutical industry; Epidemic; Influenza

Desc. génériques : Public health; Medical sciences; Viral disease

Descripteur(s) français

Descripteur(s) : Vaccin; Vaccination; Politique; Historique; Immunologie; Microorganisme; Infection; Pays industrialisé; Pays en développement; Industrie pharmaceutique; Epidémie; Grippe

Desc. génériques : Santé publique; Sciences médicales; Virose

Localisation : BDSP/ENSP, Shelf number 153639, DM00/0065

Origine de la notice : BDSP

Agir face aux crises. Katrina, grippe aviaire, tsunami

Titre : Agir face aux crises. Katrina, grippe aviaire, tsunami

Auteur(s) : Groupe Solben, Unknown

Source : 2006; 111 p.

Éditeur : Plon, Paris

ISBN : 225920399X

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Type de document : Book

Résumé : La réponse aux crises sanitaires, environnementales, financières et internationales constitue aujourd'hui un enjeu majeur de gouvernance. Qu'il s'agisse de catastrophes naturelles - le Tsunami ou l'ouragan Katrina - de crises financières - comme celle de l'Argentine - de conflits politico-sécuritaires à répercussion internationale - le conflit afghan - ou de menaces pour la santé publique - la grippe aviaire - responsables politiques et institutionnels doivent réussir à les gérer en confiance avec la société civile et en coordination avec les autres gouvernements et les institutions internationales. L'internationalisation des crises et de leurs impacts expose de plus en plus les dirigeants nationaux et les organisations internationales au jugement de l'opinion publique. Cette dernière, grâce à l'accélération des moyens de communication et d'information, cherche de plus en plus les responsables des défaillances. C'est pourquoi cette progression soudaine de la notion de responsabilité de l'action publique a fait évoluer le concept de gouvernance : elle met au premier plan la capacité, individuelle et collective, à anticiper et à faire face aux crises de toutes natures. Cet ouvrage, élaboré par le groupe "SOLBEN" composé de personnes ayant ou ayant eu des responsabilités opérationnelles, s'appuie sur l'expérience de situations réelles. Il a pour vocation de formuler des pistes de réflexion et d'action pour améliorer la prévention et la gestion des crises en remettant en question certaines idées reçues

Code(s) de classement : 002B30A11

Descripteur(s) anglais

Descripteur(s) : Crisis; Management; Environment; Natural disaster; Economic crisis; Europe; UNO; Responsibility; International cooperation; Afghanistan; France; Argentina; United States

Desc. génériques : Public health; Medical sciences; Asia; South America; America; North America

Descripteur(s) français

Descripteur(s) : Crise; Gestion; Environnement; Cataclysme; Crise économique; Europe; ONU; Responsabilité; Coopération internationale; Afghanistan; France; Argentine; États Unis

Desc. génériques : Santé publique; Sciences médicales; Asie; Amérique du Sud; Amérique; Amérique du Nord

Localisation : BDSP/ENSP, Shelf number 152122, CA00/0373

Origine de la notice : BDSP

Emergency preparedness for pandemic influenza

Titre : Emergency preparedness for pandemic influenza

Auteur(s) : SCARFONE Richard J; ALEXANDER Sharon; COFFIN Susan E; JOHN Keith H St; SULLIVAN Frank; WAGNER Jacqueline; ZAOUTIS Theoklis

Affiliation(s) : University of Pennsylvania School of Medicine, Philadelphia PA, United States; Infection Prevention and Control Professional, Infection Prevention and Control Department, The Children's Hospital of Philadelphia, Philadelphia, PA, United States; Infection Prevention/Control and Occupational Health, The Children's Hospital of Philadelphia, Philadelphia, PA, United States; Emergency Preparedness Program, The Children's Hospital of Philadelphia, Philadelphia, PA, United States; Environmental Health and Safety, The Children's Hospital of Philadelphia, Philadelphia, PA, United States

Source : Pediatric emergency care. 2006; 22 (9) : 661-668

ISSN : 0749-5161

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 32 ref.

Code(s) de classement : 002B27B; 002B27B14C; 002B27B15

Descripteur(s) anglais

Descripteur(s) : Resuscitation; Intensive care; Child; Emergency; Influenza

Desc. génériques : Resuscitation; Intensive care medicine; Medical sciences; Resuscitation; Intensive care medicine; Medical sciences; Resuscitation; Intensive care medicine; Medical sciences; Human; Viral disease; Infection

Descripteur(s) français

Descripteur(s) : Réanimation; Soins intensifs; Enfant; Urgence; Grippe

Desc. génériques : Réanimation; Soins intensifs; Sciences médicales; Réanimation; Soins intensifs; Sciences médicales; Réanimation; Soins intensifs; Sciences médicales; Homme; Virose; Infection

Localisation : INIST, Shelf number 20958, INIST No. 354000157191240130

Origine de la notice : INIST

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Development of an internal positive control for rapid diagnosis of avian influenza virus infections by real-time reverse transcription-PCR with lyophilized reagents

Titre : Development of an internal positive control for rapid diagnosis of avian influenza virus infections by real-time reverse transcription-PCR with lyophilized reagents

Auteur(s) : DAS Amaresh; SPACKMAN Erica; SENNE Dennis; PEDERSEN Jan; SUAREZ David L

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Source : Journal of clinical microbiology Print. 2006; 44 (9) : 3065-3073

ISSN : 0095-1137

CODEN : JCMIDW

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 41 ref.

Résumé : We developed an internal positive control (IPC) RNA to help ensure the accuracy of the detection of avian influenza virus (AIV) RNA by reverse transcription (RT)-PCR and real-time RT-PCR (RRT-PCR). The IPC was designed to have the same binding sites for the forward and reverse primers of the AIV matrix gene as the target amplicon, but it had a unique internal sequence used for the probe site. The amplification of the viral RNA and the IPC by RRT-PCR were monitored with two different fluorescent probes in a multiplex format, one specific for the AIV matrix gene and the other for the IPC. The RRT-PCR test was further simplified with the use of lyophilized bead reagents for the detection of AIV RNA. The RRT-PCR with the bead reagents was more sensitive than the conventional wet reagents for the detection of AIV RNA. The IPC-based RRT-PCR detected inhibitors in blood, kidney, lungs, spleen, intestine, and cloacal swabs, but not allantoic fluid, serum, or tracheal swabs. The accuracy of RRT-PCR test results with the lyophilized beads was tested on cloacal and tracheal swabs from experimental birds inoculated with AIV and compared with virus isolation (VI) on embryonating chicken eggs. There was 97 to 100% agreement of the RRT-PCR test results with VI for tracheal swabs and 81% agreement with VI for cloacal swabs, indicating a high level of accuracy of the RRT-PCR assay. The same IPC in the form of armored RNA was also used to monitor the extraction of viral RNA and subsequent detection by RRT-PCR.

Code(s) de classement : 002A05C10; 002B05

Descripteur(s) anglais

Descripteur(s) : Avian influenza virus; Diagnosis; Real time; Reverse transcription polymerase chain reaction; Microbiology

Desc. génériques : Virology; Microbiology; Biological sciences; Infectious diseases; Medical sciences; Influenzavirus A; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Influenzavirus aviaire; Diagnostic; Temps reel; Reaction chaine polymerase RT; Microbiologie

Desc. génériques : Virologie; Microbiologie; Sciences biologiques; Maladies infectieuses; Sciences médicales; Influenzavirus A; Orthomyxoviridae; Virus

Localisation : INIST, Shelf number 17088, INIST No. 354000157105450020

Origine de la notice : INIST

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Noninvasive positive-pressure ventilation : An experimental model to assess air and particle dispersion

Titre : Noninvasive positive-pressure ventilation : An experimental model to assess air and particle dispersion

Auteur(s) : HUI David S; HALL Stephen D; CHAN Matthew T V; CHOW Benny K; TSOU Jin Y; JOYNT Gavin M; SULLIVAN Colin E; SUNG Joseph J Y

Affiliation(s) : Departments of Medicine and Therapeutics, The Chinese University of Hong Kong, Hong Kong; School of Mechanical Engineering, The University of New South Wales, Sydney, NSW, Australia; Department of Anesthesia and Intensive Care, The Chinese University of Hong Kong, Hong Kong; Department of Architecture, The Chinese University of Hong Kong, Hong Kong; Department of Medicine, The University of Sydney, NSW, Australia

Source : Chest . 2006; 130 (3) : 730-740

ISSN : 0012-3692

CODEN : CHETBF

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 45 ref.

Résumé : Background: Health-care workers are concerned about the risk of acquiring contagious diseases such as severe acute respiratory syndrome and avian influenza after recent outbreaks. We studied exhaled air and particle dispersion through an oronasal mask attached to a human-patient simulator (HPS) during noninvasive positive-pressure ventilation (NPPV). Methods: Airflow was marked with intrapulmonary smoke for visualization. Therapy with inspiratory positive airway pressure (IPAP) was started at 10 cm H₂O and gradually increased to 18 cm H₂O, whereas expiratory positive airway pressure was maintained at 4 cm H₂O. A leakage jet plume was revealed by a laser light sheet and images captured by video. Smoke concentration in the plume was estimated from the light scattered by smoke particles. Findings: A jet plume of air leaked through the mask exhaust holes to a radial distance of 0.25 m from the mask during the application of IPAP at 10 cm H₂O with some leakage from the nasal bridge. The leakage plume exposure probability was highest about 60 to 80 mm lateral to the median sagittal plane of the HPS. Without nasal bridge leakage, the jet plume from the exhaust holes increased to a 0.40-m radius from the mask, whereas exposure probability was highest about 0.28 m above the patient. When IPAP was increased to 18 cm H₂O, the vertical plume extended to 0.45 m above the patient with some horizontal spreading along the ward ceiling. Conclusion: Substantial exposure to exhaled air occurs within a 0.5-m radius of patients receiving NPPV. Medical wards should be designed with an architectural aerodynamics approach and knowledge of air/particle dispersion from common mechanical ventilatory techniques.

Code(s) de classement : 002B11; 002B12; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Influenza; Cardiovascular disease; Respiratory disease; Positive pressure; Artificial ventilation; Mechanical ventilation; Experimental study; Models; Air; Particle; Dispersion

Desc. génériques : Pneumology; Respiratory system; Medical sciences; Cardiovascular system; Medical sciences; Virology; Infectious diseases; Medical sciences; Viral disease; Infection

Descripteur(s) français

Descripteur(s) : Grippe; Appareil circulatoire pathologie; Appareil respiratoire pathologie; Pression positive; Ventilation artificielle; Ventilation mécanique; Etude expérimentale; Modèle; Air; Particule; Dispersion

Desc. génériques : Pneumologie; Appareil respiratoire; Sciences médicales; Système cardiovasculaire; Sciences médicales; Virologie; Maladies infectieuses; Sciences médicales; Virose; Infection

Localisation : INIST, Shelf number 7627, INIST No. 354000142885350140

Origine de la notice : INIST

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Genetic immunity and influenza pandemics

Titre : Genetic immunity and influenza pandemics

Auteur(s) : RUMYANTSEV Sergey N

Affiliation(s) : Andent Inc, Waukegan, IL, United States

Source : FEMS immunology and medical microbiology. 2006; 48 (1) : 1-10

ISSN : 0928-8244

Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Nombre de références : 31 ref.

Résumé : In addition to the great number of publications focused on the leading role of virus mutations and reassortment in the origin of pandemic influenza, general opinion emphasizes the victim side of the epidemic process. Based on the analysis and integration of relevant ecological, epidemiological, clinical, genetic and experimental data, the present article is focused on the evolution of 'virus - victim' ecological systems resulting in the formation of innate (i.e. genetic, constitutional) immunity in the involved species and populations. This kind of immunity functions today as the greatest natural barrier to the pandemic spread of influenza among humans and ecologically related kinds of animals. Global influenza pandemics can arise when the worldwide population contains at least a minimum number of people susceptible to a known or mutant influenza virus. Special attention is paid in this article to individual tests for the presence of this barrier, including the implications of specific findings for public health policy. Such tests could be based on in vitro observation of the action of relevant virus strains on primary cell cultures or on their cellular or molecular components extracted from individuals. The resources of the Human Genome Project should also be utilized.

Code(s) de classement : 002A05

Descripteur(s) anglais

Descripteur(s) : Genetics; Public health; Microbiology; Immunology; Influenza

Desc. génériques : Microbiology; Biological sciences; Viral disease; Infection

Descripteur(s) français

Descripteur(s) : Génétique; Santé publique; Microbiologie; Immunologie; Grippe

Desc. génériques : Microbiologie; Sciences biologiques; Virose; Infection

Localisation : INIST, Shelf number 17567B, INIST No. 354000142227570010

Origine de la notice : INIST

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Structure and receptor specificity of the hemagglutinin from an H5N1 influenza virus

Titre : Structure and receptor specificity of the hemagglutinin from an H5N1 influenza virus

Auteur(s) : STEVENS James; BLIXT Ola; TUMPEY Terrence M; TAUBENBERGER Jeffery K; PAULSON James C; WILSON Ian A

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Source : Science Washington DC. 2006; 312 (5772) : 404-410

ISSN : 0036-8075

CODEN : SCIEAS

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Notes : 3/4 p. ref. et notes

Résumé : The hemagglutinin (HA) structure at 2.9 angstrom resolution, from a highly pathogenic Vietnamese H5N1 influenza virus, is more related to the 1918 and other human H1 HAs than to a 1997 duck H5 HA. Glycan microarray analysis of this Viet04 HA reveals an avian α 2-3 sialic acid receptor binding preference. Introduction of mutations that can convert H1 serotype HAs to human α 2-6 receptor specificity only enhanced or reduced affinity for avian-type receptors. However, mutations that can convert avian H2 and H3 HAs to human receptor specificity, when inserted onto the Viet04 H5 HA framework, permitted binding to a natural human α 2-6 glycan, which suggests a path for this H5N1 virus to gain a foothold in the human population.

Code(s) de classement : 002B05C02C; 002A05C04

Descripteur(s) anglais

Descripteur(s) : Influenzavirus A; Subtype; Avian influenza virus; Molecular structure; Hemagglutinin; Mutation; Biological receptor; Human; Species specificity

Desc. génériques : Virology; Infectious diseases; Medical sciences; Virology; Microbiology; Biological sciences; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Influenzavirus A; Soustype; Influenzavirus aviaire; Structure moléculaire; Hemagglutinine; Mutation; Recepteur biologique; Homme; Specificité espèce; Virus H5N1

Desc. génériques : Virologie; Maladies infectieuses; Sciences médicales; Virologie; Microbiologie; Sciences biologiques; Orthomyxoviridae; Virus

Localisation : INIST, Shelf number 6040, INIST No. 354000142715070190

Origine de la notice : INIST

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H5N1 virus attachment to lower respiratory tract

Titre : H5N1 virus attachment to lower respiratory tract

Auteur(s) : VAN RIEL Debby; MUNSTER Vincent J; DE WIT Emmie; RIMMELZWAAN Guus F; FOUCHIER Ron A M; OSTERHAUS Ab D M E; KUIKEN Thijs

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ISSN : 0036-8075

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Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Notes : 1/4 p. ref. et notes

Code(s) de classement : 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Influenzavirus A; Subtype; Pathogenesis; Human; Animal; Comparative study; Respiratory tract; Trachea; Pulmonary alveolus; Respiratory disease; Avian influenza virus

Desc. génériques : Virology; Infectious diseases; Medical sciences; Orthomyxoviridae; Virus; Respiratory system

Descripteur(s) français

Descripteur(s) : Influenzavirus A; Soustype; Pathogenie; Homme; Animal; Etude comparative; Voie respiratoire; Trachee; Alveole pulmonaire; Appareil respiratoire pathologie; Influenzavirus aviaire; Virus H5N1

Desc. génériques : Virologie; Maladies infectieuses; Sciences medicales; Orthomyxoviridae; Virus; Appareil respiratoire

Localisation : INIST, Shelf number 6040, INIST No. 354000142715070170

Origine de la notice : INIST

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Host species barriers to influenza virus infections

Titre : Host species barriers to influenza virus infections

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ISSN : 0036-8075

CODEN : SCIEAS

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Notes : 1/4 p. ref. et notes

Résumé : Most emerging infectious diseases in humans originate from animal reservoirs; to contain and eradicate these diseases we need to understand how and why some pathogens become capable of crossing host species barriers. Influenza virus illustrates the interaction of factors that limit the transmission and subsequent establishment of an infection in a novel host species. Influenza species barriers can be categorized into virus-host interactions occurring within individuals and host-host interactions, either within or between species, that affect transmission between individuals. Viral evolution can help surmount species barriers, principally by affecting virus-host interactions; however, evolving the capability for sustained transmission in a new host species represents a major adaptive challenge because the number of mutations required is often large.

Code(s) de classement : 002A05C04; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Avian influenza virus; Transmission from animal to man; Host virus relation; Host specificity; Epidemic; Evolution; Adaptation; Mutation; Human; Review

Desc. génériques : Virology; Microbiology; Biological sciences; Virology; Infectious diseases; Medical sciences; Influenza virus A; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Influenzavirus aviaire; Transmission animal homme; Relation hôte virus; Spécificité hôte; Epidémie; Evolution; Adaptation; Mutation; Homme; Article synthèse; Barrière des espèces

Desc. génériques : Virologie; Microbiologie; Sciences biologiques; Virologie; Maladies infectieuses; Sciences médicales; Influenzavirus A; Orthomyxoviridae; Virus

Localisation : INIST, Shelf number 6040, INIST No. 354000142715070160

Origine de la notice : INIST

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Predictability and preparedness in influenza control

Titre : Predictability and preparedness in influenza control

Auteur(s) : SMITH Derek J

Source : Science Washington DC. 2006; 312 (5772) : 392-394

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CODEN : SCIEAS

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Notes : 1/4 p. ref. et notes

Résumé : The threat of pandemic human influenza looms as we survey the ongoing avian influenza pandemic and wonder if and when it will jump species. What are the risks and how can we plan? The nub of the problem lies in the inherent variability of the virus, which makes prediction difficult. However, it is not impossible; mathematical models can help determine and quantify critical parameters and thresholds in the relationships of those parameters, even if the relationships are nonlinear and obscure to simple reasoning. Mathematical models can derive estimates for the levels of drug stockpiles needed to buy time, how and when to modify vaccines, whom to target with vaccines and drugs, and when to enforce quarantine measures. Regardless, the models used for pandemic planning must be tested, and for this we must continue to gather data, not just for exceptional scenarios but also for seasonal influenza.

Code(s) de classement : 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Influenza; Public health; Transmission from man to man; Epidemic; Evaluation; Risk; Prediction; Mathematical model; Evolution; Thailand; Influenzavirus; Influenzavirus A; Human

Desc. génériques : Virology; Infectious diseases; Medical sciences; Viral disease; Infection; Asia; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Grippe; Sante publique; Transmission homme homme; Epidemie; Evaluation; Risque; Prediction; Modele mathematique; Evolution; Thaïlande; Influenzavirus; Influenzavirus A; Homme; Pandemie

Desc. génériques : Virologie; Maladies infectieuses; Sciences médicales; Virose; Infection; Asie; Orthomyxoviridae; Virus

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Origine de la notice : INIST

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Les pandémies dans l'histoire depuis le Vème siècle av. J. - C

Titre : Les pandémies dans l'histoire depuis le Vème siècle av. J. - C

Auteur(s) : ZAJDENWEBER Daniel; MARTIN Pierre; MOUTON Francois; SEGOVIA KUENY Sandrine; GIN Stephane; LAVAUX Magalie; BEAUCHENE Vinciane; CHOW KOO Roseline; DIAZ Jean Pierre

Source : RISQUES . 2006-06; (66) : 21-57; tabl.

ISSN : 1152-9253

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Type de document : Serial

Nombre de références : dissem.

Résumé : Sommaire du dossier : Les pandémies dans l'Histoire - De panzootie a pandémie - Les pouvoirs publics face a la pandémie - Couverture du risque mortalité du bétail : le rôle des pouvoirs publics, l'intervention de l'assureur - Pandémie de grippe aviaire : les entreprises sont-elles prêtes ? - Impact d'une pandémie de grippe d'origine aviaire sur les assureurs de personnes

Code(s) de classement : 002B30A01

Descripteur(s) anglais

Descripteur(s) : Contagion; Communicable disease; Epidemic; History; Health; Epidemiology; Sanitary surveillance; Civil service; Responsibility; Insurance; Private sector; France; Epizootics; Spongiform encephalopathy; Risk management

Desc. génériques : Public health; Medical sciences; Europe; Prion disease; Infection

Descripteur(s) français

Descripteur(s) : Contagion; Maladie contagieuse; Epidémie; Histoire; Santé; Épidémiologie; Surveillance sanitaire; Administration publique; Responsabilité; Assurance; Secteur privé; France; Épizootie; Encephalopathie spongiforme; Gestion risque

Desc. génériques : Santé publique; Sciences médicales; Europe; Prion maladie; Infection

Localisation : BDSP/ENSP, Shelf number 155105

Origine de la notice : BDSP

Rapport fait au nom de la mission d'information sur la grippe aviaire : mesures preventives. Tome 3. Plan pandémie : une strategie de gestion de crise

Titre : Rapport fait au nom de la mission d'information sur la grippe aviaire : mesures preventives. Tome 3. Plan pandémie : une strategie de gestion de crise

Auteur(s) : LE GUEN Jean Marie; DOOR Jean Pierre, rapp

Auteur(s) : Assemblée Nationale Paris, France

Source : 2006 01 26; 581 p.; pdf

Éditeur : Assemblée Nationale, Paris

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Type de document : Book

Résumé : Apres deux rapports, l'un consacre aux moyens medicaux disponibles en cas de pandémie, l'autre relatif à l'étendue de l'épizootie de grippe aviaire dans le monde et en France, la mission d'information sur la grippe aviaire constituée en octobre 2005 publie un troisième rapport dédié plus particulièrement à l'évaluation du plan gouvernemental de préparation à la pandémie. Ce plan gouvernemental de prévention et de lutte "pandémie grippale" dont la France s'est dotée s'inscrit dans la ligne de plans antérieurs, eux-mêmes inspirés d'un plan mondial OMS de préparation à une pandémie de grippe publiée en 1999 et mise à jour en 2005. Il propose une stratégie générale qui consiste à faire fonctionner l'Etat et la société en "mode dégradé" jusqu'à la mise au point d'un vaccin, afin de freiner la propagation du virus. La première partie du rapport, après avoir rappelé l'histoire du plan français, expose les orientations générales du plan gouvernemental et présente les mesures qui devront être mises en œuvre : organisation d'un dispositif de gestion de crise à l'échelon national ou local, déploiement de "mesures barrières" susceptibles de réduire le risque de contagion tels que masques respiratoires et médicaments antiviraux, et enfin, préservation de la continuité des services indispensables à la population comme les transports collectifs ou l'enseignement. La seconde partie du rapport aborde l'organisation du système de santé en période pré-pandémique, puis en phase pandémique, avec un éclairage particulier sur le rôle de l'hôpital tout au long de la crise et une contribution de Mme Berengere Poletti, membre de la mission, sur la télémédecine. L'auteur y met en évidence les avantages d'un recours plus grand aux nouvelles technologies dans la perspective d'une pandémie. Le rapport se termine par une liste de recommandations concernant six points : les structures administratives de gestion de crise, les collectivités territoriales, l'InVS, l'information de la population et l'aide aux malades soignés à domicile, la médecine de ville et les structures hospitalières

Code(s) de classement : 002B30A01

Descripteur(s) anglais

Descripteur(s) : Influenza; Vector; Animal; Plane; Case history; Antiviral; Protection safety equipment; Individual safety equipment; Vaccine; Service; Transport; Teaching; Firm; Economic sector; Epidemiology; Sanitary surveillance; Hospitalization; Hospital; Organization; Emergency; Hospital environment; Health staff; General practitioner; Resuscitation; Balance; Proposition; Prevention; Health; France; Epizootics; Risk management; Telemedicine

Desc. génériques : Public health; Medical sciences; Viral disease; Infection; Europe

Descripteur(s) français

Descripteur(s) : Grippe; Vecteur; Animal; Plan; Historique; Antiviral; Equipement protection securite; Protection individuelle; Vaccin; Service; Transport; Enseignement; Entreprise; Secteur economique; Epidemiologie; Surveillance sanitaire; Hospitalisation; Hopital; Organisation; Urgence; Milieu hospitalier; Personnel sanitaire; Medecin generaliste; Reanimation; Bilan; Proposition; Prevention; Sante; France; Epizootie; Gestion risque; Telemedecine

Desc. génériques : Sante publique; Sciences medicales; Virose; Infection; Europe

Localisation : BDSP/ENSP, Shelf number 154729

Origine de la notice : BDSP

La gestion de la grippe aviaire

Titre : La gestion de la grippe aviaire

Auteur(s) : BRICQ Nicole, rapp

Auteur(s) : Senat Paris, France

Source : 2006 07 04; 130 p.; pdf, ann.

Éditeur : Senat, Paris

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Type de document : Book

Résumé : Au nom de la commission des finances du Senat et en application de l' article 57 de la LOLF, Nicole BRICQ, rapporteure speciale de la mission "securite sanitaire" a conduit, au cours du premier semestre de l' annee 2006, une mission de controle sur la mise en oeuvre des mesures de lutte contre la grippe aviaire. Apres avoir procede a de nombreuses auditions et deplacements dans des departements francais, elle fait part de ses observations dans ce rapport. Dans une premiere partie, elle estime que le dispositif national de lutte qui comprend deux volets, un volet relatif a l' influenza aviaire et un volet relatif a la pandémie humaine, peut etre considere comme globalement efficace malgre certains points faibles qui devront etre corriges : difficultes liees a la surveillance et au recensement des basses-cours familiales, nombre de laboratoires de criblage sur le territoire ou probleme de formation des veterinaires en ce qui concerne le premier volet, faible association des medecins liberaux, limite des capacites hospitalieres nationales ou inegale mobilisation des services deconcentres et des collectivites territoriales s' agissant du deuxieme volet. Elle met egalement en avant une architecture administrative generale inadaptée et peu structuree ainsi qu' un financement initial sous evalue et manquant de lisibilite a plusieurs niveaux. Puis, s' interessant aux enjeux majeurs, la rapporteure speciale estime que la dimension internationale de la crise de grippe aviaire est un element essentiel de sa gestion et doit etre pleinement prise en compte au niveau national. Elle appelle donc, dans une deuxieme partie, a un renforcement de la cooperation europeenne dans le domaine de la sante humaine et a une prise en compte coordonnee, a l' echelle mondiale, de la situation des pays les plus exposes. Enfin, en conclusion, la rapporteure estime qu' il faut se preparer a vivre avec un risque durable, au niveau national et international. La durabilite de ce risque impose de reflechir a l' evolution du dispositif mis en place et a l' emergence d' une gouvernance mondiale de la securite sanitaire

Code(s) de classement : 002B30A01

Descripteur(s) anglais

Descripteur(s) : Influenza; Vector; Animal; Plane; Rearing; Veterinary center; Epidemiology; Sanitary surveillance; Risk analysis; Vaccination; Ministry; Financing; Cooperation; Europe; International cooperation; Developing countries; Balance; Proposition; France; Epizootics; Risk management

Desc. génériques : Public health; Medical sciences; Viral disease; Infection

Descripteur(s) français

Descripteur(s) : Grippe; Vecteur; Animal; Plan; Elevage; Centre veterinaire; Epidemiologie; Surveillance sanitaire; Analyse risque; Vaccination; Ministere; Financement; Cooperation; Europe; Cooperation internationale; Pays en developpement; Bilan; Proposition; France; Epizootie; Gestion risque; Agence francaise de securite sanitaire des aliments

Desc. génériques : Sante publique; Sciences medicales; Virose; Infection

Localisation : BDSP/ENSP, Shelf number 154533

Origine de la notice : BDSP

Infections virales émergentes. Discussion : Les zoonoses, passe, present et avenir; Emerging viral diseases. Discussion

Titre : Infections virales émergentes. Discussion : Les zoonoses, passe, present et avenir; Emerging viral diseases. Discussion

Auteur(s) : BRICAIRE Francois; BOSSI Philippe; PENE M Pierre; TUBIANA M Maurice

Affiliation(s) : Service des Maladies infectieuses et tropicales Groupe Hospitalier de la Pitie-Salpetriere, 47 bld de l'Hopital, 75015 Paris, France

Source : Bulletin de l'Academie nationale de medecine. 2006; 190 (3) : 597-609

ISSN : 0001-4079

CODEN : BANMAC

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Langue(s) du résumé : English

Type de document : Serial

Type de document : article; comments

Nombre de références : 24 ref.

Résumé : Les infections émergentes ou reémergentes sont depuis quelques années revenues dans l'actualité. Plusieurs raisons y concourent: outre des faits tenant à nos habitudes de vie, à des modifications écologiques, et à des faits socio-politiques, des possibilités techniques actuelles permettent de mieux diagnostiquer ou mettre en évidence des agents infectieux. Ces faits s'expriment à travers de nombreux pathogènes, essentiellement des virus de divers types: le virus West Nile, le virus Chikungunya et en expansion le virus de l'encéphalite japonaise. On peut insister sur les virus responsables des fièvres hémorragiques virales représentés surtout par les virus Ebola et Marburg. Plus récemment, a pu être isolé le Coronavirus responsable du SRAS, infection ayant provoqué une épidémie qui a suffisamment inquiété pour modifier les mentalités et le concept de prise en charge des épidémies. C'est dans cet esprit que la grippe aviaire, phénomène animal, fait craindre et annoncer une future pandémie avec un virus adapté à l'homme. Doivent être mis à part la variole et le Monkeypox qui sont des virus virtuellement émergents dans le cadre d'un éventuel acte bioterroriste. La prise en charge de ces infections émergentes justifie réflexions et organisation d'un certain nombre de structures pour répondre au mieux aux exigences actuelles de notre société.

Code(s) de classement : 002B30A11; 002B05C02C; 002B05C02B

Descripteur(s) anglais

Descripteur(s) : Viral disease; Discussion; Arbovirus; Emerging disease; Severe acute respiratory syndrome; Bioterrorism; Influenza; Medicine; Human; Hemorrhagic fever; Review; Smallpox; Monkey pox; Chikungunya virus; Japanese encephalitis virus; Reemerging disease

Desc. génériques : Public health; Medical sciences; Virology; Infectious diseases; Medical sciences; Virology; Infectious diseases; Medical sciences; Infection; Virus; Arbovirus disease; Alphavirus; Togaviridae; Japanese encephalitis group virus; Flavivirus; Flaviviridae; Respiratory disease; Lung disease; Skin disease

Descripteur(s) français

Descripteur(s) : Virose; Discussion; Arbovirus; Maladie émergente; Syndrome respiratoire aigu sévère; Bioterrorisme; Grippe; Médecine; Homme; Fièvre hémorragique; Article synthèse; Variole; Variole singe; Virus Chikungunya; Virus encéphalite japonaise; Maladie réémergente

Desc. génériques : Santé publique; Sciences médicales; Virologie; Maladies infectieuses; Sciences médicales; Virologie; Maladies infectieuses; Sciences médicales; Infection; Virus; Arbovirose; Alphavirus; Togaviridae; Virus groupe encéphalite japonaise; Flavivirus; Flaviviridae; Appareil respiratoire pathologie; Poumon pathologie; Peau pathologie

Localisation : INIST, Shelf number 740, INIST No. 354000157159810030

Origine de la notice : INIST

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A global pandemic influenza vaccine action plan

Titre : A global pandemic influenza vaccine action plan

Auteur(s) : KIENY Marie Paule; COSTA Alejandro; HOMBACH Joachim; CARRASCO Peter; PERVIKOV Yuri; SALISBURY David; GRECO Michel; GUST Ian; LAFORCE Marc; FRANCO PAREDES Carlos; SANTOS Jose Ignacio; D' HONDT Eric; RIMMELZWAAN Guus; KARRON Ruth; FUKUDA Keiji

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ISSN : 0264-410X

CODEN : VACCDE

Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Nombre de références : 3 ref.

Résumé : In case of an influenza pandemic, the world will be in a situation where potential vaccine supply will fall short by several billion doses from global needs. The World Health Organization (WHO) convened in Geneva on May 2-3, 2006 a consultation of all stakeholders in influenza vaccines and immunization to identify practical solutions to fill this gap. The consultation resulted in a global action plan outlining promising specific strategies to increase influenza vaccine production and surge-capacity before and during an influenza pandemic. Although the timing and severity of the next influenza pandemic cannot be predicted, vaccines are considered the one of the most important medical interventions for reducing morbidity and mortality if and when such an event occurs. Despite this acknowledged role, current limitations on influenza vaccine manufacturing capacity mean that, should a pandemic virus emerge in the near future, vaccine supplies would fall short of the anticipated global demand by several billion doses. Concern about this situation was formally acknowledged in May 2005, when the World Health Assembly approved a resolution [1] on strengthening pandemic influenza preparedness and response. That resolution called on the World Health Organization (WHO) to seek solutions with international and national partners, including the private sector, to reduce the present global shortage of influenza vaccines. More specifically, the resolution asked WHO to look at strategies for economizing on the use of antigen and transferring production technologies from industrialized to developing countries. In response to this request, WHO convened a consultation from 2-3 May 2006 attended by representatives of the major stakeholders in the area of influenza vaccines and immunization. The consultation had two main objectives: (1) To prepare a global action plan with specific short-, medium-, and long-term activities designed to increase influenza vaccine production and surge-capacity, to identify key obstacles and driving forces, and to estimate funding needs. (2) To strengthen the engagement and collaboration of key partners and stakeholders.

Code(s) de classement : 002A05F04

Descripteur(s) anglais

Descripteur(s) : Vaccine; Immunization; Influenza A

Desc. génériques : Immunology; Pharmacology; Applied microbiology; Microbiology; Biological sciences; Viral disease; Infection

Descripteur(s) français

Descripteur(s) : Vaccin; Immunisation; Grippe A

Desc. génériques : Immunologie; Pharmacologie; Microbiologie appliquée; Microbiologie; Sciences biologiques; Virose; Infection

Localisation : INIST, Shelf number 20289, INIST No. 354000157170860010

Origine de la notice : INIST

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Genotype turnover by reassortment of replication complex genes from avian Influenza A virus

Titre : Genotype turnover by reassortment of replication complex genes from avian Influenza A virus

Auteur(s) : MACKEN Catherine A; WEBBY Richard J; BRUNO William J

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Source : Journal of general virology. 2006; 87 (p.10) : 2803-2815

ISSN : 0022-1317

CODEN : JGVIAY

Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Nombre de références : 29 ref.

Résumé : Reassortment among the RNA segments of Influenza A virus caused the two most recent human influenza pandemics; recently, reassortment has generated viral genotypes associated with outbreaks of avian H5N1 influenza in Asia and Europe. A statistical analysis has been developed for the systematic identification and characterization of reassortant viruses. The analysis was applied to the genes of the replication complex of 152 avian influenza A viruses isolated between 1966 and 2004 from predominantly terrestrial and domestic aquatic avian species. The results indicated that reassortment among these genes was pervasive throughout this period and throughout both the Eurasian and North American lineages of the virus. Evidence is presented that the circulating genotypes of the replication complex are being replaced continually by novel genotypes created by reassortment. No constraints for coordinated reassortment among genes of the replication complex were evident; rather, reassortment almost always proceeded one segment at a time. A maximum-likelihood estimate of the rate of reassortment was derived. For significantly diverged Asian avian influenza A viruses from the period 1991-2004, it was estimated that the median duration between creation of a new genotype and its next segment reassortment was 3 years. Reassortments that introduced previously unobserved influenza genetic material were detected. These findings point to substantial potential for rapid generation of novel avian influenza A viruses, emphasizing the importance of intensive surveillance of these host species in preparation for a possible pandemic.

Code(s) de classement : 002A05C10

Descripteur(s) anglais

Descripteur(s) : Avian influenza virus; Influenza A virus; Genotype; Turnover; Replication; Gene; Microbiology; Virology

Desc. génériques : Virology; Microbiology; Biological sciences; Influenzavirus A; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Influenzavirus aviaire; Virus grippal A; Genotype; Turnover; Replication; Gene; Microbiologie; Virologie

Desc. génériques : Virologie; Microbiologie; Sciences biologiques; Influenzavirus A; Orthomyxoviridae; Virus

Localisation : INIST, Shelf number 13533, INIST No. 354000142921150050

Origine de la notice : INIST

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Genetic characterization of the H9N2 influenza viruses circulated in the poultry population in Israel

Titre : Genetic characterization of the H9N2 influenza viruses circulated in the poultry population in Israel

Auteur(s) : PERK Shimon; BANET NOACH Caroline; SHIHMANter Ester; POKAMUNSKI Shimon; PIRAK Michael; LIPKIND Michael; PANSHIN Alexander

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Source : Comparative immunology microbiology and infectious diseases. 2006; 29 (4) : 207-223

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Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Nombre de références : 34 ref.

Résumé : The partial nucleotide sequences of the hemagglutinin (HA) genes of 72 H9N2 influenza viruses isolated from chickens and turkeys in Israel during the period 2000-2005 were genetically analyzed. The isolates possessed the three types of amino acid motif -R-S-S-R/ G-L-, -R-S-N-R/G-L-, and -R-S-K-R/G-L- at the cleavage site of HA. Phylogenetic analyses showed that all Israeli isolates belonged to the same group which further divided into three closely related sub-groups. The HA genes of these isolates were related to the HA gene of A/chicken/Germany/R45/98 isolated from chicken in Germany in 1998.

Code(s) de classement : 002A05C10

Descripteur(s) anglais

Descripteur(s) : Avian influenza virus; Genetics; Poultry; Israel; Subtype; Hemagglutinin; Gene; Cleavage site; Phylogeny; Influenza; Microbiology; Immunology

Desc. génériques : Virology; Microbiology; Biological sciences; Influenzavirus A; Orthomyxoviridae; Virus; Asia; Farming animal; Viral disease; Infection; Veterinary

Descripteur(s) français

Descripteur(s) : Influenzavirus aviaire; Génétique; Volaille; Israël; Soustype; Hemagglutinine; Gène; Site clivage; Phylogénèse; Grippe; Microbiologie; Immunologie

Desc. génériques : Virologie; Microbiologie; Sciences biologiques; Influenzavirus A; Orthomyxoviridae; Virus; Asie; Animal élevage; Virose; Infection; Vétérinaire

Localisation : INIST, Shelf number 16817, INIST No. 354000152256350030

Origine de la notice : INIST

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Grippe aviaire : les medecins du travail impliqués : De la grippe saisonniere au risque de pandémie

Titre : Grippe aviaire : les medecins du travail impliqués : De la grippe saisonniere au risque de pandémie

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Source : Concours medical Paris. 2006; 128 (31-32) : 1322-1324

ISSN : 0010-5309

CODEN : COMEAO

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Type de document : Serial

Nombre de références : 9 ref.

Code(s) de classement : 002B01; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Physician; Public health; Work; Avian influenza

Desc. génériques : Medical sciences; Virology; Infectious diseases; Medical sciences

Descripteur(s) français

Descripteur(s) : Medecin; Sante publique; Travail; Grippe aviaire

Desc. génériques : Sciences medicales; Virologie; Maladies infectieuses; Sciences medicales

Localisation : INIST, Shelf number 10949, INIST No. 354000139308220090

Origine de la notice : INIST

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Virus respiratoires émergents : virus du sras et virus influenza A/H5N1 hautement pathogène; Emergent viruses : SARS-associated coronavirus and H5N1 influenza virus

Titre : Virus respiratoires émergents : virus du sras et virus influenza A/H5N1 hautement pathogène; Emergent viruses : SARS-associated coronavirus and H5N1 influenza virus

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Source : Annales de biologie clinique Paris. 2006; 64 (3) : 195-208

ISSN : 0003-3898

CODEN : ABCLAI

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Langue(s) du résumé : English

Type de document : Serial

Nombre de références : 30 ref.

Résumé : Cette synthèse a pour but de faire le point des connaissances concernant le virus du syndrome respiratoire aigu sévère (Sras) et le virus A/H5N1 hautement pathogène. En cas de suspicion d'infection par SARS-CoV, les recherches virologiques sont réalisées à partir d'une aspiration rhinopharyngée, d'une expectoration, d'un écouvillonnage de gorge ou d'un liquide de lavage broncho-alvéolaire (LBA) mais aussi des selles, des urines voire du sérum d'un patient venant d'une région où le virus du Sras aurait réémergé (ou travaillant dans un laboratoire étudiant le virus du Sras). L'isolement du virus sur culture cellulaire ou l'extraction de l'ARN viral en vue d'une RT-PCR doivent être réalisés dans un laboratoire de niveau de confinement BSL3 (Biosafety level 3). Une RT-PCR spécifique du virus du Sras permet le diagnostic virologique. En raison de la moindre sensibilité de la RT-PCR spécifique du Sras, l'analyse des échantillons cliniques doit être répétée pendant plusieurs jours. La prise en charge d'échantillons biologiques pour une suspicion d'infection par le virus A/H5N1 hautement pathogène (tableau clinique de grippe chez une personne revenant d'une région où circule le virus A/H5N1), respecte la même procédure. En effet, le diagnostic virologique repose sur la mise en évidence du virus ou de son ARN génomique. Les échantillons biologiques permettant le diagnostic sont des prélèvements respiratoires, des selles, du sérum ou du liquide céphalo-rachidien (LCR). L'isolement du virus sur culture cellulaire doit être réalisé dans un laboratoire de niveau de confinement BSL3. Pour la recherche de l'ARN génomique, l'extraction de l'ARN doit, elle aussi, être réalisée dans un laboratoire de niveau de confinement BSL3. Une RT-PCR en temps réel ciblant un fragment du gène de l'hémagglutinine H5 permet un diagnostic spécifique de la grippe A/H5N1. Ces procédures sont actuellement en place dans les laboratoires de virologie afin de diagnostiquer les éventuels premiers cas de grippe à virus A/H5N1 ou la réémergence du virus du Sras.

Code(s) de classement : 002B24; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Emerging disease; Human; Respiratory system; Severe acute respiratory syndrome; Pathogenic; Coronavirus; Influenzavirus; Clinical biology; Virological exploration; Laboratory investigations; Influenzavirus AH5N1; Avian influenza

Desc. génériques : Medical sciences; Virology; Infectious diseases; Medical sciences; Viral disease; Infection; Coronaviridae; Nidovirales; Virus; Orthomyxoviridae; Respiratory disease; Lung disease

Descripteur(s) français

Descripteur(s) : Maladie émergente; Homme; Appareil respiratoire; Syndrome respiratoire aigu sévère; Pathogène; Coronavirus; Influenzavirus; Biologie clinique; Exploration virologique; Examen laboratoire; Influenzavirus AH5N1; Grippe aviaire

Desc. génériques : Sciences médicales; Virologie; Maladies infectieuses; Sciences médicales; Virose; Infection; Coronaviridae; Nidovirales; Virus; Orthomyxoviridae; Appareil respiratoire pathologie; Poumon pathologie
Localisation : INIST, Shelf number 1014, INIST No. 354000153186200010

Origine de la notice : INIST

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An in vitro fluorescence screen to identify antivirals that disrupt hepatitis B virus capsid assembly

Titre : An in vitro fluorescence screen to identify antivirals that disrupt hepatitis B virus capsid assembly

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Source : Nature biotechnology. 2006; 24 (3) : 358-362

ISSN : 1087-0156

CODEN : NABIF9

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Type de document : correspondence,-letters

Nombre de références : 29 ref.

Résumé : Virus assembly has not been routinely targeted in the development of antiviral drugs, in part because of the lack of tractable methods for screening in vitro. We have developed an in vitro assay of hepatitis B virus (HBV) capsid assembly, based on fluorescence quenching of dye-labeled capsid protein, for testing potential inhibitors. This assay is adaptable to high-throughput screening and can identify small-molecule inhibitors of virus assembly that prevent, inappropriately accelerate and/or misdirect capsid formation to yield aberrant particles. An in vitro primary screen has the advantage of identifying promising lead compounds affecting assembly without the requirement that they be taken up by cells in culture and be nontoxic. Our approach may facilitate the identification of antivirals targeting viruses other than HBV, such as avian influenza and HIV.

Code(s) de classement : 002A31; 215

Descripteur(s) anglais

Descripteur(s) : In vitro; Fluorescence; Antiviral; Capsid; Targeting; Hepatitis B virus; Method; Joining; Screening; Inhibitor

Desc. génériques : Biotechnology; Biological sciences; Orthohepadnavirus; Hepadnaviridae; Virus

Descripteur(s) français

Descripteur(s) : In vitro; Fluorescence; Antiviral; Capside; Ciblage; Virus hépatite B; Methode; Assemblage; Criblage; Inhibiteur

Desc. génériques : Biotechnologie; Sciences biologiques; Orthohepadnavirus; Hepadnaviridae; Virus

Localisation : INIST, Shelf number 19676, INIST No. 354000115188760240

Origine de la notice : INIST

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Respiratory hygiene in the emergency department

Titre : Respiratory hygiene in the emergency department

Auteur(s) : ROTHMAN Richard E; IRVIN Charlene B; MORAN Gregory J; SAUER Lauren; BRADSHAW Ylisabth S; FRY Robert B JR; JOSEPHINE Elaine B; LEDYARD Holly K; HIRSHON Jon Mark

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Source : Annals of emergency medicine. 2006; 48 (5) : 570-582

ISSN : 0196-0644

CODEN : AEMED3

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 63 ref.

Résumé : The emergency department (ED) is an essential component of the public health response plan for control of acute respiratory infectious threats. Effective respiratory hygiene in the ED is imperative to limit the spread of dangerous respiratory pathogens, including influenza, severe acute respiratory syndrome, avian influenza, and bioterrorism agents, particularly given that these agents may not be immediately identifiable. Sustaining effective respiratory control measures is especially challenging in the ED because of patient crowding, inadequate staffing and resources, and ever-increasing numbers of immunocompromised patients. Threat of contagion exists not only for ED patients but also for visitors, health care workers, and inpatient populations. Potential physical sites for respiratory disease transmission extend from out-of-hospital care, to triage, waiting room, ED treatment area, and the hospital at large. This article presents a summary of the most current information available in the literature about respiratory hygiene in the ED, including administrative, patient, and legal issues. Wherever possible, specific recommendations and references to practical information from the Centers for Disease Control and Prevention are provided. The "Administrative Issues" section describes coordination with public health departments, procedures for effective facility planning, and measures for health care worker protection (education, staffing optimization, and vaccination). The patient care section addresses the potentially infected ED patient, including emergency medical services concerns, triage planning, and patient transport. "Legal Issues" discusses the interplay between public safety and patient privacy. Emergency physicians play a critical role in early identification, treatment, and containment of potentially lethal respiratory pathogens. This brief synopsis should help clinicians and administrators understand, develop, and implement appropriate policies and procedures to address respiratory hygiene in the ED.

Code(s) de classement : 002B27B; 002B27B14C; 002B27B02

Descripteur(s) anglais

Descripteur(s) : Resuscitation; Intensive care; Hygiene; Emergency department

Desc. génériques : Resuscitation; Intensive care medicine; Medical sciences; Resuscitation; Intensive care medicine; Medical sciences; Pneumology; Respiratory system; Resuscitation; Intensive care medicine; Medical sciences

Descripteur(s) français

Descripteur(s) : Reanimation; Soins intensif; Hygiene; Service urgence

Desc. génériques : Reanimation; Soins intensifs; Sciences medicales; Reanimation; Soins intensifs; Sciences medicales; Pneumologie; Appareil respiratoire; Reanimation; Soins intensifs; Sciences medicales

Localisation : INIST, Shelf number 19670, INIST No. 354000158840760120

Origine de la notice : INIST

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Acute respiratory distress syndrome induced by avian influenza a (H5N1) virus in mice

Titre : Acute respiratory distress syndrome induced by avian influenza a (H5N1) virus in mice

Auteur(s) : TONG XU; JIAN QIAO; LIHONG ZHAO; GUIRONG WANG; GUIMEI HE; KAI LI; YONG TIAN; MINGYU GAO; JIANLIN WANG; HUIYU WANG; CHANGGUI DONG

Affiliation(s) : Department of Pathophysiology, College of Veterinary Medicine, China Agricultural University, Beijing, China

Source : American journal of respiratory and critical care medicine. 2006; 174 (9) : 1011-1017

ISSN : 1073-449X

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 31 ref.

Résumé : Rationale and Objective: The acute respiratory distress syndrome (ARDS) caused by avian influenza H5N1 viral infection has been reported in many humans since this virus was found to infect humans in Hong Kong in 1997, but no studies regarding an animal model of ARDS with H5N1 viral infection have been found in the literature. Here we present a mouse model of ARDS induced by H5N1 virus. Methods: Six- to 8-wk-old BALB/c mice were inoculated intranasally ($50 \mu\text{l}$) with $1 \times 10^{5.5}$ mouse infectious doses of A/Chicken/Hebei/108/2002 (H5N1) virus. Lung injury was assessed by observation of lung water content and histopathology. Arterial blood gas, white blood cell count in bronchial alveolar lavage fluid, and tumor necrosis factor- α and interleukin-6 in bronchoalveolar lavage fluid and serum were measured at the indicated time points. Results: Our data showed that H5N1 viral infection in mice resulted in typical ARDS, which was characterized by the following features: (1) about 80% of mice (13 of 16) dead on Days 6 to 8 postinoculation; (2) highly edematous lungs and dramatically increased lung wet:dry weight ratios and lung wet weight:body weight ratios; (3) inflammatory cellular infiltration, alveolar and interstitial edema, and hemorrhage in lungs; (4) progressive and severe hypoxemia; and (5) significant increase in neutrophils, tumor necrosis factor- α , and interleukin-6 in BALF. Conclusion: These results suggested that we successfully established a mouse model of ARDS with H5N1 viral infection, which may benefit further investigation into the pathogenesis of human ARDS induced by H5N1 virus.

Code(s) de classement : 002B27B; 002B27B02; 002B27D01

Descripteur(s) anglais

Descripteur(s) : Respiratory distress; Resuscitation; Intensive care; Influenza A virus; Animal; Mouse; Cytokine; Avian influenza

Desc. génériques : Resuscitation; Intensive care medicine; Medical sciences; Pneumology; Respiratory system; Resuscitation; Intensive care medicine; Medical sciences; Transfusion; Medical sciences; Influenzavirus A; Orthomyxoviridae; Virus; Rodentia; Mammalia; Vertebrata; Respiratory disease

Descripteur(s) français

Descripteur(s) : Detresse respiratoire; Reanimation; Soins intensif; Virus grippal A; Animal; Souris; Cytokine; Grippe aviaire

Desc. génériques : Reanimation; Soins intensifs; Sciences medicales; Pneumologie; Appareil respiratoire; Reanimation; Soins intensifs; Sciences medicales; Transfusion; Sciences medicales; Influenzavirus A; Orthomyxoviridae; Virus; Rodentia; Mammalia; Vertebrata; Appareil respiratoire pathologie

Localisation : INIST, Shelf number 2013, INIST No. 354000158834330100

Origine de la notice : INIST

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Emerging respiratory viruses : Challenges and vaccine strategies

Titre : Emerging respiratory viruses : Challenges and vaccine strategies

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Affiliation(s) : Laboratory of Infectious Diseases, National Institute of Allergy and Infectious Diseases, National Institutes of Health, Bethesda, Maryland 20892, United States

Source : Clinical microbiology reviews Print. 2006; 19 (4) : 592, 614-636 [24 p.]

ISSN : 0893-8512

CODEN : CMIREX

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 245 ref.

Résumé : The current threat of avian influenza to the human population, the potential for the reemergence of severe acute respiratory syndrome (SARS)-associated coronavirus, and the identification of multiple novel respiratory viruses underline the necessity for the development of therapeutic and preventive strategies to combat viral infection. Vaccine development is a key component in the prevention of widespread viral infection and in the reduction of morbidity and mortality associated with many viral infections. In this review we describe the different approaches currently being evaluated in the development of vaccines against SARS-associated coronavirus and avian influenza viruses and also highlight the many obstacles encountered in the development of these vaccines. Lessons learned from current vaccine studies, coupled with our increasing knowledge of the host and viral factors involved in viral pathogenesis, will help to increase the speed with which efficacious vaccines targeting newly emerging viral pathogens can be developed.

Code(s) de classement : 002A05F04; 002B05

Descripteur(s) anglais

Descripteur(s) : Vaccine; Review

Desc. génériques : Immunology; Pharmacology; Applied microbiology; Microbiology; Biological sciences; Infectious diseases; Medical sciences

Descripteur(s) français

Descripteur(s) : Vaccin; Article synthèse

Desc. génériques : Immunologie; Pharmacologie; Microbiologie appliquée; Microbiologie; Sciences biologiques; Maladies infectieuses; Sciences médicales

Localisation : INIST, Shelf number 21815, INIST No. 354000158744400020

Origine de la notice : INIST

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Meta-analysis: Convalescent blood products for spanish influenza pneumonia : A future H5N1 treatment? Commentary

Titre : Meta-analysis: Convalescent blood products for spanish influenza pneumonia : A future H5N1 treatment? Commentary

Auteur(s) : LUKE Thomas C; KILBANE Edward M; JACKSON Jeffrey L; HOFFMAN Stephen L; TREATOR John J, comment

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Source : Annals of internal medicine. 2006; 145 (8) : 599-609,631-632 [13 p.]

ISSN : 0003-4819

CODEN : AIMEAS

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Type de document : article; comments

Nombre de références : 74 ref.

Résumé : Background: Studies from the Spanish influenza era reported that transfusion of influenza-convalescent human blood products reduced mortality in patients with influenza complicated by pneumonia. Treatments for H5N1 influenza are unsatisfactory, and convalescent human plasma containing H5N1 antibodies could be an effective therapy during outbreaks and pandemics. Purpose: To determine whether transfusion with influenza-convalescent human blood products reduced the risk for death in patients with Spanish influenza pneumonia. Data Sources: Manual search of English-language journals from 1918 to 1925. Citations from retrieved studies were also searched. Study Selection: Published English-language studies that had at least 10 patients in the treatment group, used convalescent blood products to treat Spanish influenza pneumonia in a hospital setting, and reported on a control or comparison group. Data Extraction: Two investigators independently extracted data on study characteristics, outcomes, adverse events, and quality. Data Synthesis: Eight relevant studies involving 1703 patients were found. Treated patients, who were often selected because of more severe illness, were compared with untreated controls with influenza pneumonia in the same hospital or ward. The overall crude case-fatality rate was 16% (54 of 336) among treated patients and 37% (452 of 1219) among controls. The range of absolute risk differences in mortality between the treatment and control groups was 8% to 26% (pooled risk difference, 21% [95% CI, 15% to 27%]). The overall crude case-fatality rate was 19% (28 of 148) among patients who received early treatment (after <4 days of pneumonia complications) and 59% (49 of 83) among patients who received late treatment (after ≥4 days of pneumonia complications). The range of absolute risk differences in mortality between the early treatment group and the late treatment group was 26% to 50% (pooled risk difference, 41% [CI, 29% to 54%]). Adverse effects included chill reactions and possible exacerbations of symptoms in a few patients. Limitations: Studies were few and had many methodologic limitations. No study was a blinded, randomized, or placebo-controlled trial. Some pertinent studies may have been missed. Conclusions: Patients with Spanish influenza pneumonia who received influenza-convalescent human blood products may have experienced a clinically important reduction in the risk for death. Convalescent human H5N1 plasma could be an effective, timely, and widely available treatment that should be studied in clinical trials.

Code(s) de classement : 002B30A11; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Pneumonia; Metaanalysis; Evidence based medicine; Immunotherapy; Blood product; Influenza; Bibliographic review; Treatment; Convalescence; Serum; Immunomodulation; Prospective; Human; Prevention; Epidemiology; Antibody; Feasibility; History; Influenzavirus AH5N1; Spanish flu

Desc. génériques : Public health; Medical sciences; Virology; Infectious diseases; Medical sciences; Viral disease; Infection; Respiratory disease; Lung disease

Descripteur(s) français

Descripteur(s) : Pneumonie; Metaanalyse; Medecine factuelle; Immunotherapie; Constituant sang; Grippe; Revue bibliographique; Traitement; Convalescence; Serum; Immunomodulation; Prospective; Homme; Prevention; Epidemiologie; Anticorps; Faisabilite; Histoire; 1918 1925; Influenzavirus AH5N1; Grippe espagnole

Desc. génériques : Sante publique; Sciences medicales; Virologie; Maladies infectieuses; Sciences medicales; Virose; Infection; Appareil respiratoire pathologie; Poumon pathologie

Localisation : INIST, Shelf number 2014, INIST No. 354000158730460060

Origine de la notice : INIST

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Cross-protectiveness and immunogenicity of influenza A/duck/singapore/3/97(H5) vaccines against infection with A/vietnam/1203/04(H5N1) virus in ferrets

Titre : Cross-protectiveness and immunogenicity of influenza A/duck/singapore/3/97(H5) vaccines against infection with A/vietnam/1203/04(H5N1) virus in ferrets

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Source : The Journal of infectious diseases. 2006; 194 (8) : 1040-1043

ISSN : 0022-1899

CODEN : JIDIAQ

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Type de document : short-communication

Nombre de références : 15 ref.

Résumé : Ferrets were immunized with two 7- μ g doses of hemagglutinin from inactivated whole-virus vaccines containing the hemagglutinin gene of A/Duck/Singapore/3/97(H5N3) then inoculated with a lethal dose of A/Vietnam/1203/04(H5N1) (Viet/1203/04). Serum samples did not react with Viet/1203/04 in hemagglutination-inhibition (HI) or virus-neutralization (VN) tests. All vaccinated ferrets survived the challenge, whereas all mock-immunized ferrets died. Immunized ferrets had significantly lower virus titers in the upper respiratory tract and less-severe disease. Vaccine generated from antigenically different H5 virus protects against infection by a highly pathogenic H5 strain. Neither HI nor VN testing provides correlates of cross-protection in ferrets.

Code(s) de classement : 002A05F04; 002B05; 002A05C10

Descripteur(s) anglais

Descripteur(s) : Virus; Immunogenicity; Singapore; Vaccine; Vietnam; Microbiology; Infection; Influenza A; Avian influenza

Desc. génériques : Immunology; Pharmacology; Applied microbiology; Microbiology; Biological sciences; Infectious diseases; Medical sciences; Virology; Microbiology; Biological sciences; Asia; Viral disease

Descripteur(s) français

Descripteur(s) : Virus; Immunogenicite; Singapour; Vaccin; Vietnam; Microbiologie; Infection; Grippe A; Grippe aviaire

Desc. génériques : Immunologie; Pharmacologie; Microbiologie appliquee; Microbiologie; Sciences biologiques; Maladies infectieuses; Sciences medicales; Virologie; Microbiologie; Sciences biologiques; Asie; Virose

Localisation : INIST, Shelf number 2052, INIST No. 354000157260790020

Origine de la notice : INIST

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Avian influenza a (H5N1) infection in a child

Titre : Avian influenza a (H5N1) infection in a child

Auteur(s) : WITAYATHAWORNWONG Prasonk

Affiliation(s) : Pediatric Department, Petchabun Hospital, Petchabun, Thailand

Source : Southeast Asian journal of tropical medicine and public health. 2006; 37 (4) : 684-689

ISSN : 0125-1562

CODEN : SJTMAK

Date de publication : 2006

Pays de publication : Thailand

Langue(s) : English

Type de document : Serial

Type de document : case-report,-clinical-case

Nombre de références : 1 p.1/4

Résumé : A previously healthy, 9-year-old girl was admitted to the hospital with respiratory insufficiency. She had mild and severe respiratory symptoms for 3 weeks and 4 days before admission, respectively. She had a history of close contact with her domestic poultry, which was infected with avian influenza A (H5N1). She was isolated with the air-borne transmission prevention mode of treatment. Acute respiratory distress syndrome (ARDS) was documented from the time of admission and mechanical ventilation was introduced without improvement. She had multiple episodes of diarrhea for 2 days. Her condition deteriorated and she expired in 4 days. Throat swab RT-PCR and viral culture for avian influenza A (H5N1) were positive.

Code(s) de classement : 002B01; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Influenza A; Child; Tropical medicine; Avian influenza

Desc. génériques : Medical sciences; Virology; Infectious diseases; Medical sciences; Viral disease; Infection; Human

Descripteur(s) français

Descripteur(s) : Grippe A; Enfant; Médecine tropicale; Grippe aviaire

Desc. génériques : Sciences médicales; Virologie; Maladies infectieuses; Sciences médicales; Virose; Infection; Homme

Localisation : INIST, Shelf number 19778, INIST No. 354000157264000130

Origine de la notice : INIST

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Regional collaboration in the Middle East to deal with H5N1 avian flu

Titre : Regional collaboration in the Middle East to deal with H5N1 avian flu

Auteur(s) : LEVENTHAL Alex; RAMLAWI Assad; BELBIESI Adel; BALICER Ran D

Affiliation(s) : Israel Ministry of Health, PO Box 1176, Jerusalem 91010, Israel; Palestinian Authority Ministry of Health, Ramallah, West Bank, Jordan; Jordan Ministry of Health, Amman, Jordan; Steering Committee for Pandemic Influenza Preparedness, Ramat-Gan 52392, Israel

Source : BMJ British medical journal International ed. 2006; 333 (7573) : 856-858

ISSN : 0959-8146

Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Nombre de références : 10 ref.

Résumé : In 2005-6 Arab and Israeli collaboration contained outbreaks of avian flu in the Middle East This initiative shows how building relationships through joint efforts creates an infrastructure for cross border collaboration during emergencies.

Code(s) de classement : 002B01; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Regional; Middle east; Medicine; Avian influenza; Influenzavirus AH5N1

Desc. génériques : Medical sciences; Virology; Infectious diseases; Medical sciences; Asia

Descripteur(s) français

Descripteur(s) : Regional; Moyen Orient; Medecine; Grippe aviaire; Influenzavirus AH5N1

Desc. génériques : Sciences médicales; Virologie; Maladies infectieuses; Sciences médicales; Asie

Localisation : INIST, Shelf number 5002A, INIST No. 354000139227040180

Origine de la notice : INIST

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Zoonotic viral diseases and the frontier of early diagnosis, control and prevention

Titre : Zoonotic viral diseases and the frontier of early diagnosis, control and prevention

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Source : Journal of internal medicine. 2006; 260 (5) : 399-408

ISSN : 0954-6820

Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Nombre de références : 83 ref.

Résumé : Public awareness of the human health risks of zoonotic infections has grown in recent years. Currently, concern of H5N1 flu transmission from migratory bird populations has increased with foci of fatal human cases. This comes on the heels of other major zoonotic viral epidemics in the last decade. These include other acute emerging or re-emerging viral diseases such as severe acute respiratory syndrome (SARS), West-Nile virus, Ebola virus, monkeypox, as well as the more inapparent insidious slow viral and prion diseases. Virus infections with zoonotic potential can become serious killers once they are able to establish the necessary adaptations for efficient human-to-human transmission under circumstances sufficient to reach epidemic proportions. The monitoring and early diagnosis of these potential risks are overlapping frontiers of human and veterinary medicine. Here, current viral zoonotics and evolving threats are reviewed.

Code(s) de classement : 002B01; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Viral disease; Early stage; Diagnosis; Emerging disease; Surveillance; Check; West Nile encephalitis; Prevention; Ebola virus; Severe acute respiratory syndrome; Vaccine; Immunoprophylaxis; Zoonosis; Review; Bibliographic review; Medicine; Avian influenza; Influenzavirus AH5N1

Desc. génériques : Medical sciences; Virology; Infectious diseases; Medical sciences; Infection; Arbovirus disease; Filovirus; Filoviridae; Mononegavirales; Virus; Respiratory disease; Lung disease; Cerebral disorder; Central nervous system disease; Nervous system diseases

Descripteur(s) français

Descripteur(s) : Virose; Stade precoce; Diagnostic; Maladie emergente; Surveillance; Controle; Encephalite West Nile; Prevention; Virus Ebola; Syndrome respiratoire aigu severe; Vaccin; Immunoprophylaxie; Zoonose; Article synthese; Revue bibliographique; Medecine; Grippe aviaire; Influenzavirus AH5N1

Desc. génériques : Sciences medicales; Virologie; Maladies infectieuses; Sciences medicales; Infection; Arbovirose; Filovirus; Filoviridae; Mononegavirales; Virus; Appareil respiratoire pathologie; Poumon pathologie; Encephale pathologie; Systeme nerveux central pathologie; Systeme nerveux pathologie

Localisation : INIST, Shelf number 893A, INIST No. 354000158759340010

Origine de la notice : INIST

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A computer evaluation of ventilation performance in a negative-pressure operating theater

Titre : A computer evaluation of ventilation performance in a negative-pressure operating theater

Auteur(s) : CHOW Tin Tai; KWAN Anne; ZHANG LIN; WEI BAI

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Source : Anesthesia and analgesia. 2006; 103 (4) : 913-918

ISSN : 0003-2999

CODEN : AACRAT

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 11 ref.

Résumé : BACKGROUND: A negative-pressure operating theater is required to limit the spread of respiratory diseases in patients with severe acute respiratory syndrome, tuberculosis, avian influenza, or similar infectious diseases. In Hong Kong, we converted a conventional operating theater into a negative-pressure operating theater that has been in service for more than a year. In this article, we introduce its ventilation design and evaluate the airflow performance in relation to different combinations of medical lamp configurations and modes of launching infectious particles into the room air. METHODS: We used a computational fluid dynamics technique for the numerical analysis. RESULTS: Our analyses showed that the airflow performance in the negative-pressure operating theater was satisfactory and comparable to the original positive-pressure design. The airflow pattern effectively controlled the dispersion of infectious particles. Our calculations demonstrated that the airflow contained the dispersion of infectious particles released from the patient sufficiently to protect the surgical team, and vice versa. CONCLUSIONS: Computational fluid dynamics can be used to assess airflow in a negative-pressure operating room and model the dispersion of infectious particles from the patient.

Code(s) de classement : 002B27A

Descripteur(s) anglais

Descripteur(s) : Negative pressure; Anesthesia

Desc. génériques : Anesthesia; Medical sciences

Descripteur(s) français

Descripteur(s) : Pression négative; Anesthésie

Desc. génériques : Anesthésie; Sciences médicales

Localisation : INIST, Shelf number 3213, INIST No. 354000152276640180

Origine de la notice : INIST

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Factors involved in the aerosol transmission of infection and control of ventilation in healthcare premises

Titre : Factors involved in the aerosol transmission of infection and control of ventilation in healthcare premises

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Source : The Journal of hospital infection. 2006; 64 (2) : 100-114

ISSN : 0195-6701

Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Nombre de références : 155 ref.

Résumé : The epidemics of severe acute respiratory syndrome (SARS) in 2003 highlighted both short- and long-range transmission routes, i.e. between infected patients and healthcare workers, and between distant locations. With other infections such as tuberculosis, measles and chicken-pox, the concept of aerosol transmission is so well accepted that isolation of such patients is the norm. With current concerns about a possible approaching influenza pandemic, the control of transmission via infectious air has become more important. Therefore, the aim of this review is to describe the factors involved in: (1) the generation of an infectious aerosol, (2) the transmission of infectious droplets or droplet nuclei from this aerosol, and (3) the potential for inhalation of such droplets or droplet nuclei by a susceptible host. On this basis, recommendations are made to improve the control of aerosol-transmitted infections in hospitals as well as in the design and construction of future isolation facilities.

Code(s) de classement : 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Severe acute respiratory syndrome; Influenza; Aerosols; Transmission; Check; Ventilation

Desc. génériques : Virology; Infectious diseases; Medical sciences; Viral disease; Infection; Prevention; Respiratory disease; Lung disease

Descripteur(s) français

Descripteur(s) : Syndrome respiratoire aigu severe; Grippe; Aerosol; Transmission; Controle; Ventilation

Desc. génériques : Virologie; Maladies infectieuses; Sciences medicales; Virose; Infection; Prevention; Appareil respiratoire pathologie; Poumon pathologie

Localisation : INIST, Shelf number 18802, INIST No. 354000142926930010

Origine de la notice : INIST

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Les solutions vaccinales chez l' homme et l' animal face aux virus influenza aviaire H₅N₁; H₅N₁ : vaccine solutions for humans and other animals

Titre : Les solutions vaccinales chez l' homme et l' animal face aux virus influenza aviaire H₅N₁; H₅N₁ : vaccine solutions for humans and other animals

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Affiliation(s) : Academie nationale de medecine, France

Source : Bulletin de l'Academie nationale de medecine. 2006; 190 (4-5) : 963-972

ISSN : 0001-4079

CODEN : BANMAC

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Langue(s) du résumé : English

Type de document : Serial

Nombre de références : 21 ref.

Résumé : L' hypervirulence du virus H₅N₁ rend impossible la production classique du virus grippal sur oeuf embryonne; il existe des solutions alternatives : en pathologie aviaire, un virus recombinant vivant variole du canari (fowl-pox)-H₅ est utilise. Beaucoup plus largement encore un vaccin inactif adjuve H₅N₂, immunogene par son hemagglutinine H₅ est commercialise (technique DIVA). La prophylaxie est differente entre pays europeens et asiatiques. En pathologie humaine, la solution retenue par l' OMS consiste a preparer un vaccin a partir d' un virus apathogene PR8 auquel H₅ modifiee et Nj d' un virus pandemique ont ete greffees. Le mediocre pouvoir immunogene du vaccin impliquera une augmentation de la quantite d' antigene et l' utilisation d' immunostimulants. Pourra-t-on satisfaire une demande forte en cas de pandémie ? Recemment un vaccin vivant a base d' adenovirus couple au gene H₅ a ete propose. Enfin des auteurs chinois soulignent l' interet potentiel d' un serum hyperimmun prepare sur animal en utilisation preventive et curative.

Code(s) de classement : 002B30A11; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Immunoprophylaxis; Human; Animal; Face; Prevention; Vaccine; Medicine; Scientific research; Avian influenza; Influenzavirus AH5N1

Desc. génériques : Public health; Medical sciences; Virology; Infectious diseases; Medical sciences

Descripteur(s) français

Descripteur(s) : Immunoprophylaxie; Homme; Animal; Face; Prevention; Vaccin; Medecine; Recherche scientifique; Virus H5N1; Grippe aviaire; Influenzavirus AH5N1

Desc. génériques : Sante publique; Sciences medicales; Virologie; Maladies infectieuses; Sciences medicales

Localisation : INIST, Shelf number 740, INIST No. 354000143004210150

Origine de la notice : INIST

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Safety and immunogenicity of nonadjuvanted and MF59-adjuvanted influenza A/H9N2 vaccine preparations

Titre : Safety and immunogenicity of nonadjuvanted and MF59-adjuvanted influenza A/H9N2 vaccine preparations

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Source : Clinical infectious diseases. 2006; 43 (9) : 1135-1142

ISSN : 1058-4838

CODEN : CIDIEL

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 23 ref.

Résumé : Background. Influenza A/H9N2 viruses can infect humans and are considered to be a pandemic threat. Effective vaccines are needed for these and other avian influenza viruses. Methods. We performed a phase I, randomized, double-blind trial to evaluate the safety and immunogenicity of a 2-dose schedule (administered on days 0 and 28) of 4 dose levels (3.75, 7.5, 15, and 30 μ g of hemagglutinin) of inactivated influenza A/chicken/Hong Kong/G9/97 (H9N2) vaccine with and without MF59 adjuvant. Vaccine safety was assessed with a diary and selected blood tests. Immunogenicity was measured using serum hemagglutination inhibition (HAI) and microneutralization (MNt) antibody assays. Results. Ninety-six healthy adults (age, 18-34 years) were enrolled in the study. Arm discomfort was more common in groups that received adjuvant, but adverse effects of the vaccination were generally mild. Geometric mean serum HAI and MNt antibody titers to the influenza A/chicken/Hong Kong/G9/97 (H9N2) virus strain for all vaccine groups were similar on day 0 but were significantly higher ($P < .001$) on both days 28 and 56 for the MF59-adjuvanted vaccine groups than for groups given nonadjuvanted vaccine. Other measures of immunogenicity were also higher in the adjuvanted vaccine groups. HAI and MNt geometric mean titers measured after the administration of a single dose of MF59-adjuvanted vaccine were similar to those measured after 2 doses of nonadjuvanted vaccine. Conclusions. The combination of MF59 adjuvant with a subunit vaccine was associated with improved immune responses to an influenza A/H9N2 virus. The adjuvanted vaccine was immunogenic even after a single dose, raising the possibility that a 1-dose vaccination strategy may be attainable with the use of adjuvanted vaccine.

Code(s) de classement : 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Influenza A; Immunoprophylaxis; Immunogenicity; Prevention; Vaccine

Desc. génériques : Virology; Infectious diseases; Medical sciences; Viral disease; Infection

Descripteur(s) français

Descripteur(s) : Grippe A; Immunoprophylaxie; Immunogenicité; Prévention; Vaccin

Desc. génériques : Virologie; Maladies infectieuses; Sciences médicales; Virose; Infection

Localisation : INIST, Shelf number 18407, INIST No. 354000158813460020

Origine de la notice : INIST

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Integrated molecular signature of disease : Analysis of influenza virus-infected macaques through functional genomics and proteomics

Titre : Integrated molecular signature of disease : Analysis of influenza virus- infected macaques through functional genomics and proteomics

Auteur(s) : BAAS T; BASKIN C R; DIAMOND D L; GARCIA SASTRE A; BIELEFELDT OHMANN H; TUMPEY T M; THOMAS M J; CARTER V S; TEAL T H; VAN HOEVEN N; PROLL S; JACOBS J M; CALDWELL Z R; GRITSENKO M A; HUKKANEN R R; CAMP D G II; SMITH R D; KATZE M G

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Source : Journal of virology. 2006; 80 (21) : 10813-10828

ISSN : 0022-538X

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 85 ref.

Résumé : Recent outbreaks of avian influenza in humans have stressed the need for an improved nonhuman primate model of influenza pathogenesis. In order to further develop a macaque model, we expanded our previous in vivo genomics experiments with influenza virus-infected macaques by focusing on the innate immune response at day 2 postinoculation and on gene expression in affected lung tissue with viral genetic material present. Finally, we sought to identify signature genes for early infection in whole blood. For these purposes, we infected six pigtailed macaques (*Macaca nemestrina*) with reconstructed influenza A/Texas/36/91 virus and three control animals with a sham inoculate. We sacrificed one control and two experimental animals at days 2, 4, and 7 postinfection. Lung tissue was harvested for pathology, gene expression profiling, and proteomics. Blood was collected for genomics every other day from each animal until the experimental endpoint. Gross and microscopic pathology, immunohistochemistry, viral gene expression by arrays, and/or quantitative real-time reverse transcription-PCR confirmed successful yet mild infections in all experimental animals. Genomic experiments were performed using macaque-specific oligonucleotide arrays, and high-throughput proteomics revealed the host response to infection at the mRNA and protein levels. Our data showed dramatic differences in gene expression within regions in influenza virus-induced lesions based on the presence or absence of viral mRNA. We also identified genes tightly coregulated in peripheral white blood cells and in lung tissue at day 2 postinoculation. This latter finding opens the possibility of using gene expression arrays on whole blood to detect infection after exposure but prior to onset of symptoms or shedding.

Code(s) de classement : 002A05C10

Descripteur(s) anglais

Descripteur(s) : Influenzavirus; Genomics; Proteomics; Microbiology; Virology; Infection

Desc. génériques : Virology; Microbiology; Biological sciences; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Influenzavirus; Genomique; Proteomique; Microbiologie; Virologie; Infection

Desc. génériques : Virologie; Microbiologie; Sciences biologiques; Orthomyxoviridae; Virus

Localisation : INIST, Shelf number 13592, INIST No. 354000158805950510

Origine de la notice : INIST

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Sense and the science of childhood immunization : Can we achieve more with less?

Titre : Sense and the science of childhood immunization : Can we achieve more with less?

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Affiliation(s) : Children's Hospital of Pittsburgh, 3705 Fifth Avenue, Pittsburgh, PA 15213-2583, United States; Department of Molecular Microbiology & Immunology, Johns Hopkins University Bloomberg School of Public Health, 615 N. Wolfe Street, Baltimore, MD 21205, United States

Source : Vaccine . 2006; 24 (42-43) : 6460-6467

ISSN : 0264-410X

CODEN : VACCDE

Date de publication : 2006

Pays de publication : United Kingdom

Langue(s) : English

Type de document : Serial

Nombre de références : 73 ref.

Résumé : The threat of biological terrorism with small pox virus and a global influenza pandemic in the face of limited vaccine supply recently stimulated research into the evaluation of fractional dose vaccine regimens, with promising immunogenicity results. While this approach is not new, it has been less applied to vaccines for less sensational but nevertheless, significant killer diseases. This manuscript provides an overview of the basics of immunization as it applies to the current practice of immunization in children, comments on the untapped avenues for cost reduction of vaccine delivery, and the potential for saving more lives with currently available resources.

Code(s) de classement : 002A05F04

Descripteur(s) anglais

Descripteur(s) : Immunization; Vaccine; Antigen

Desc. génériques : Immunology; Pharmacology; Applied microbiology; Microbiology; Biological sciences

Descripteur(s) français

Descripteur(s) : Immunisation; Vaccin; Antigène

Desc. génériques : Immunologie; Pharmacologie; Microbiologie appliquée; Microbiologie; Sciences biologiques

Localisation : INIST, Shelf number 20289, INIST No. 354000158786230020

Origine de la notice : INIST

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Place des antiviraux dans la lutte contre une pandémie grippale. Apport des travaux récents de modélisation Synthèse préparée par le groupe "Epidémiologie" 1 InVs-Inserm novembre 2005; Role of antiviral drugs in containing pandemic influenza. Contribution of recent modelling exercises Synthesis prepared by the InVs/Inserm "Epidemiology" group - November 2005

Titre : Place des antiviraux dans la lutte contre une pandémie grippale. Apport des travaux récents de modélisation Synthèse préparée par le groupe "Epidémiologie" 1 InVs-Inserm novembre 2005; Role of antiviral drugs in containing pandemic influenza. Contribution of recent modelling exercises Synthesis prepared by the InVs/Inserm "Epidemiology" group - November 2005

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Source : Médecine et maladies infectieuses. 2006; 36 (9) : 449-453

ISSN : 0399-077X

CODEN : MMAIB5

Date de publication : 2006

Pays de publication : France

Langue(s) : French

Langue(s) du résumé : English

Type de document : Serial

Nombre de références : 6 ref.

Résumé : La menace d'une pandémie grippale a conduit de nombreux pays à constituer des stocks d'antiviraux et tout particulièrement d'oseltamivir, qui peut être utilisé en prophylaxie comme en curatif. Plusieurs travaux récents, fondés sur des modélisations mathématiques, ont étudié l'impact de l'utilisation à large échelle de ce médicament pour contrôler une pandémie naissante, ralentir sa diffusion dans un pays ou en limiter l'impact épidémiologique. Ils concluent que la combinaison de l'utilisation des antiviraux en prophylaxie en anneau autour des premiers cas, combinée à des mesures de distanciation sociale, pourraient permettre de contrôler les premiers foyers de transmission d'un virus à potentiel pandémique, à la double condition d'une transmissibilité modérée du virus et d'une très grande rapidité et couverture de mise en œuvre des mesures de contrôle. En cas d'échec, ces mêmes stratégies seraient susceptibles de retarder la diffusion du virus dans un pays. L'utilisation des antiviraux en curatif devrait diminuer substantiellement l'impact épidémiologique de la pandémie, voire la proportion de la population atteinte, dans l'hypothèse où le traitement curatif réduirait la durée de la phase infectieuse. Ces travaux sont en faveur de la constitution de stocks d'antiviraux destinés à couvrir, au minimum les besoins dans le traitement des cas et au mieux permettant de contribuer à l'effort international de contrôle des premiers foyers épidémiques et d'intervenir autour des premières chaînes de transmission autochtones, pour ralentir la diffusion nationale du virus. Ces résultats sont conditionnés à l'efficacité de l'oseltamivir sur le virus pandémique et confirment l'importance d'une surveillance épidémiologique sensible et réactive.

Code(s) de classement : 002B05C02C; 002B02S05

Descripteur(s) anglais

Descripteur(s) : Influenza; Antiviral; Modeling; Synthesis; Epidemiology; Surveillance

Desc. génériques : Virology; Infectious diseases; Medical sciences; Virology; Infectious diseases; Pharmacology; Medical sciences; Viral disease; Infection

Descripteur(s) français

Descripteur(s) : Grippe; Antiviral; Modelisation; Synthese; Epidemiologie; Surveillance; Pandemie

Desc. génériques : Virologie; Maladies infectieuses; Sciences medicales; Virologie; Maladies infectieuses; Pharmacologie; Sciences medicales; Virose; Infection

Localisation : INIST, Shelf number 15434, INIST No. 354000139220350030

Origine de la notice : INIST

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Influenza, grippe aviaire et pandémie

Titre : Influenza, grippe aviaire et pandémie

Auteur(s) : RIOUX Yolaine

Source : PERSPECTIVE INFIRMIERE. 2006-09/2006-10; 4 (1) : 29-33

ISSN : 1708-1890

Date de publication : 2006

Pays de publication : Canada

Langue(s) : French

Type de document : Serial

Résumé : Le ministre de la Santé et des Services sociaux du Québec a lancé en 2006 le dernier Plan québécois de lutte face à une pandémie d'influenza. Cet article permet de faire le point sur la grippe aviaire, la pandémie d'influenza, le plan québécois et le rôle des infirmières dans ce cadre

Code(s) de classement : 002B30A11

Descripteur(s) anglais

Descripteur(s) : Occupational role; Nurse; Quebec; Health policy

Desc. génériques : Public health; Medical sciences; Canada; North America; America

Descripteur(s) français

Descripteur(s) : Rôle professionnel; Infirmier; Québec; Politique sanitaire

Desc. génériques : Santé publique; Sciences médicales; Canada; Amérique du Nord; Amérique

Localisation : BDSP/APHPDOC

Origine de la notice : BDSP

Detection and isolation of highly pathogenic H5N1 avian influenza A viruses from blow flies collected in the vicinity of an infected poultry farm in Kyoto, Japan, 2004

Titre : Detection and isolation of highly pathogenic H5N1 avian influenza A viruses from blow flies collected in the vicinity of an infected poultry farm in Kyoto, Japan, 2004

Auteur(s) : SAWABE Kyoko; HOSHINO Keita; ISAWA Haruhiko; SASAKI Toshinori; HAYASHI Toshihiko; TSUDA Yoshio; KURAHASHI Hiromu; TANABAYASHI Kiyoshi; HOTTA Akitoyo; SAITO Takehiko; YAMADA Akio; KOBAYASHI Mutsuo

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Source : The American journal of tropical medicine and hygiene. 2006; 75 (2) : 327-332

ISSN : 0002-9637

CODEN : AJTHAB

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Type de document : case-report,-clinical-case

Nombre de références : 23 ref.

Résumé : During the outbreak of highly pathogenic avian influenza that occurred in Tamba Town, Kyoto Prefecture in 2004, a total of 926 flies were collected from six sites within a radius of 2.3 km from the poultry farm. The H5 influenza A virus genes were detected from the intestinal organs, crop, and gut of the two blow fly species, *Calliphora nigribarbis* and *Aldrichina grahami*, by reverse transcription-polymerase chain reaction for the matrix protein (M) and hemagglutinin (HA) genes. The HA gene encoding multiple basic amino acids at the HA cleavage site indicated that this virus is a highly pathogenic strain. Based on the full-length sequences of the M, HA, and neuraminidase (NA) segments of virus isolates through embryonated chicken eggs, the virus from *C. nigribarbis* (A/blow fly/Kyoto/93/2004) was characterized as H5N1 subtype influenza A virus and shown to have > 99.9% identities in all three RNA segments to a strain from chickens (A/chicken/Kyoto/3/2004) and crows (A/crows/Kyoto/53/2004) derived during this outbreak period in Kyoto in 2004. Our results suggest it is possible that blow flies could become a mechanical transmitter of H5N1 influenza virus.

Code(s) de classement : 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Infection; Detection; Isolation; Japan; Tropical medicine; Diptera; Poultry; Avian influenza

Desc. génériques : Virology; Infectious diseases; Medical sciences; Asia; Insecta; Arthropoda; Invertebrata

Descripteur(s) français

Descripteur(s) : Infection; Detection; Isolement; Japon; Médecine tropicale; Diptera; Volaille; Virus H5N1; Grippe aviaire

Desc. génériques : Virologie; Maladies infectieuses; Sciences médicales; Asie; Insecta; Arthropoda; Invertebrata

Localisation : INIST, Shelf number 6817, INIST No. 354000133473080270

Origine de la notice : INIST

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Rapid diagnosis of avian influenza (AIH and assessment of pathogenicity of avian H5 and H7 subtypes by molecular methods. New diagnostic technology: Applications in animal health and biologicals controls, Saint-Malo, France, October 3-5, 2005, proceedings of an international conference

Titre : Rapid diagnosis of avian influenza (AIH and assessment of pathogenicity of avian H5 and H7 subtypes by molecular methods. New diagnostic technology: Applications in animal health and biologicals controls, Saint-Malo, France, October 3-5, 2005, proceedings of an international conference

Auteur(s) : SENNE D A; PEDERSEN J C; SUAREZ D L; PANIGRAHY B; VANNIER Philippe, ed; ESPESETH David, ed

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Affiliation(s) : U.S. Department of Agriculture, Animal and Plant Health Inspection Service, Veterinary Services, National Veterinary Services Laboratories, Ames, Iowa, United States; U.S. Department of Agriculture, Agriculture Research Service, Southeast Poultry Research Laboratory, Athens, Georgia, United States; AFSSA, Ploufragan, France; IABs, Perkasie, PA, United States

Source : Developments in biologicals. 2006; 126 : 171-177

Informations congrès : *New diagnostic technology: Applications in animal health and biologicals controls. International conference, *Saint-Malo France, *2005-10-03

ISSN : 1424-6074

Date de publication : 2006

Pays de publication : Switzerland

Langue(s) : English

Type de document : Serial; *Conference-Meeting

Nombre de références : 14 ref.

Résumé : Avian influenza (AI) is a highly contagious viral disease of poultry that is found worldwide. There are two forms of AI: a mild form called low pathogenicity avian influenza (LPAI), and a severe form called highly pathogenic avian influenza (HPAI). HPAI is associated with the H5 and H7 subtypes of AI virus (AIV) and is subject to Federal control and International reporting. A real-time reverse transcription-polymerase chain reaction (rRT-PCR) assay has been developed and validated that can help in the early detection of AI outbreaks. The rRT-PCR assay can also be used to identify infections caused by H5 and H7 subtypes of AIV. New isolates of AIV must be characterized as LPAI or HPAI for reporting and control purposes. The criteria for classification of an AI virus as HPAI are defined by the World Organization for Animal Health (OIE); the definition includes a virulence and a molecular criterion. The virulence requirement for HPAI is defined as an AIV killing 75% or more of eight inoculated chickens within 10 days. The molecular criterion is the presence of multiple dibasic amino acids at the proteolytic cleavage site of the haemagglutinin (H) protein. All HPAI viruses isolated before 2002 fulfilled both the virulence and molecular criteria. Consequently, nucleotide sequencing of the H gene and deduction of the amino acid motif at the H cleavage site has been successfully used to assess the virulence of H5 and H7 AIVs rapidly. Since 2002, however, there have been three outbreaks of HPAI where the viruses responsible for the outbreaks have either fulfilled the virulence criterion or the molecular criterion, but not both.

Code(s) de classement : 002A01D

Descripteur(s) anglais

Descripteur(s) : Aves; Virus; Chicken; Diagnosis; Pathogenicity; Subtype; Method; Poultry; Severe; Real time;

Reverse transcription polymerase chain reaction; Influenza A; Polymerase chain reaction; Detection; Identification; Isolate; Avian influenza

Desc. génériques : Biological sciences; Vertebrata; Veterinary; Viral disease; Infection

Descripteur(s) français

Descripteur(s) : Aves; Virus; Poulet; Diagnostic; Pouvoir pathogene; Soustype; Methode; Volaille; Grave; Temps reel; Reaction chaine polymerase RT; Grippe A; Reaction chaine polymerase; Detection; Identification; Isolat; Grippe aviaire

Desc. génériques : Sciences biologiques; Vertebrata; Veterinaire; Virose; Infection

Localisation : INIST, Shelf number 13557, INIST No. 354000157238940260

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Detection of animal viruses using nucleic acid sequence-based amplification (NASBA). New diagnostic technology: Applications in animal health and biologics controls, Saint-Malo, France, October 3-5, 2005, proceedings of an international conference

Titre : Detection of animal viruses using nucleic acid sequence-based amplification (NASBA). New diagnostic technology: Applications in animal health and biologics controls, Saint-Malo, France, October 3-5, 2005, proceedings of an international conference

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Source : Developments in biologicals. 2006; 126 : 7-15

Informations congrès : *New diagnostic technology: Applications in animal health and biologics controls. International conference, *Saint-Malo France, *2005-10-03

ISSN : 1424-6074

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Pays de publication : Switzerland

Langue(s) : English

Type de document : Serial; *Conference-Meeting

Nombre de références : 19 ref.

Résumé : As seen in recent avian influenza outbreaks in Asia, prevention is the key to fighting infectious disease successfully. Efficient disease surveillance systems on the basis of molecular diagnostics will help monitor the emergence of viruses in the early stage and thus prompt containment measures can be in place to minimize disease spread. Here we describe and review molecular diagnostics focusing on nucleic acid sequence-based amplification (NASBA) technology in detecting viruses causing animal diseases, such as avian influenza, foot-and-mouth disease, and Newcastle disease. NASBA offers high sensitivity, specificity, accuracy, and speed of availability of results, and NASBA would be the most applicable molecular diagnostics for disease surveillance and control.

Code(s) de classement : 002A01D

Descripteur(s) anglais

Descripteur(s) : Detection; Nucleic acid; Amplification; Asia; Prevention; Early stage; Review; Newcastle disease; Foot and mouth disease; Sensitivity; Specificity; Avian influenza

Desc. génériques : Biological sciences; Viral disease; Infection; Veterinary

Descripteur(s) français

Descripteur(s) : Detection; Acide nucleique; Amplification; Asie; Prevention; Stade precoce; Article synthese; Newcastle maladie; Fievre aphteuse; Sensibilite; Specificite; Grippe aviaire

Desc. génériques : Sciences biologiques; Virose; Infection; Veterinaire

Localisation : INIST, Shelf number 13557, INIST No. 354000157238940020

Origine de la notice : INIST

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Three Indonesian clusters of H5N1 virus infection in 2005

Titre : Three Indonesian clusters of H5N1 virus infection in 2005

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CODEN : NEJMAG

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 23 ref.

Résumé : BACKGROUND Since 2003, the widespread ongoing epizootic of avian influenza A (H5N1) among poultry and birds has resulted in human H5N1 cases in 10 countries. The first case of H5N1 virus infection in Indonesia was identified in July 2005. METHODS We investigated three clusters of Indonesian cases with at least two ill persons hospitalized with laboratory evidence of H5N1 virus infection from June through October 2005. Epidemiologic, clinical, and virologic data on these patients were collected and analyzed. RESULTS Severe disease occurred among all three clusters, including deaths in two clusters. Mild illness in children was documented in two clusters. The median age of the eight patients was 8.5 years (range, 1 to 38). Four patients required mechanical ventilation, and four of the eight patients (50%) died. In each cluster, patients with H5N1 virus infection were members of the same family, and most lived in the same home. In two clusters, the source of H5N1 virus infection in the index patient was not determined. Virus isolates were available for one patient in each of two clusters, and molecular sequence analyses determined that the isolates were clade 2 H5N1 viruses of avian origin. CONCLUSIONS In 2005 in Indonesia, clusters of human infection with clade 2 H5N1 viruses included mild, severe, and fatal cases among family members.

Code(s) de classement : 002B01; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Indonesia; Viral disease; 2005; Public health; Medicine; Avian influenza; Influenzavirus AH5N1

Desc. génériques : Medical sciences; Virology; Infectious diseases; Medical sciences; Asia; Infection

Descripteur(s) français

Descripteur(s) : Indonesia; Virose; 2005; Sante publique; Medecine; Grippe aviaire; Influenzavirus AH5N1

Desc. génériques : Sciences medicales; Virologie; Maladies infectieuses; Sciences medicales; Asie; Infection

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Avian influenza a (H5N1) infection in eastern Turkey in 2006

Titre : Avian influenza a (H5N1) infection in eastern Turkey in 2006

Auteur(s) : ONER Ahmet Faik; BAY Ali; ARSLAN Sukru; AKDENIZ Hayrettin; SAHIN Huseyin Avni; CESUR Yasar; EPCACAN Serdar; YILMAZ Neziha; DEGER Ibrahim; KIZILYILDIZ Baran; KARSEN Hasan; CEYHAN Mehmet

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Source : The New England journal of medicine. 2006; 355 (21) : 2179-2185

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CODEN : NEJMAG

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 14 ref.

Résumé : BACKGROUND An outbreak of highly pathogenic avian influenza A (H5N1) that had previously been detected throughout Asia, with major economic and health repercussions, extended to eastern Turkey in late December 2005 and early January 2006. METHODS We documented the epidemiologic, clinical, and radiologic features of all cases of confirmed H5N1 virus infection in patients who were admitted to Yuzuncu Yil University Hospital in Van, Turkey, between December 31, 2005, and January 10, 2006. RESULTS H5N1 virus infection was diagnosed in eight patients. The patients were 5 to 15 years of age, and all eight had a history of close contact with diseased or dead chickens. The mean (<plus or minus sign>SD) time between exposure and the onset of illness was 5.0<plus or minus sign>1.3 days. All the patients had fever, and seven had clinical and radiologic evidence of pneumonia at presentation; four patients died. Results of enzyme-linked immunosorbent assay and rapid influenza tests were negative in all patients, and the diagnosis was made by means of a polymerase-chain-reaction assay. CONCLUSIONS H5N1, which causes a spectrum of illnesses in humans, including severe and fatal respiratory disease, can be difficult to diagnose.

Code(s) de classement : 002B01; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Influenza A; Turkey; 2006; Public health; Medicine; Avian influenza; Influenzavirus AH5N1

Desc. génériques : Medical sciences; Virology; Infectious diseases; Medical sciences; Viral disease; Infection; Asia

Descripteur(s) français

Descripteur(s) : Grippe A; Turquie; 2006; Sante publique; Medecine; Grippe aviaire; Influenzavirus AH5N1

Desc. génériques : Sciences médicales; Virologie; Maladies infectieuses; Sciences médicales; Virose; Infection; Asie

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Origine de la notice : INIST

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Natural variation can significantly alter the sensitivity of influenza A (H5N1) viruses to oseltamivir

Titre : Natural variation can significantly alter the sensitivity of influenza A (H5N1) viruses to oseltamivir

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Source : Antimicrobial agents and chemotherapy. 2006; 50 (11) : 3809-3815

ISSN : 0066-4804

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Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 34 ref.

Résumé : Geographic spread of highly pathogenic avian H5N1 influenza viruses may give rise to an influenza pandemic. During the first months of a pandemic, control measures would rely mainly on antiviral drugs, such as the neuraminidase (NA) inhibitors oseltamivir and zanamivir. In this study, we compare the sensitivities to oseltamivir of the NAs of several highly pathogenic H5N1 viruses isolated in Asia from 1997 to 2005. The corresponding 50% inhibitory concentrations were determined using a standard in vitro NA inhibition assay. The K_{m} for the substrate and the affinity for the inhibitor (K_{i}) of NA were determined for a 1997 and a 2005 virus, using an NA inhibition assay on cells transiently expressing the viral enzyme. Our data show that the sensitivities of the NAs of H5N1 viruses isolated in 2004 and 2005 to oseltamivir are about 10-fold higher than those of earlier H5N1 viruses or currently circulating H1N1 viruses. Three-dimensional modeling of the N1 protein predicted that Glu248Gly and Tyr252His changes could account for increased sensitivity. Our data indicate that genetic variation in the absence of any drug-selective pressure may result in significant variations in sensitivity to anti-NA drugs. Although the clinical relevance of a 10-fold increase in the sensitivity of NA to oseltamivir needs to be investigated further, the possibility that sensitivity to anti-NA drugs could increase (or possibly decrease) significantly, even in the absence of treatment, underscores the need for continuous evaluation of the impact of genetic drift on this parameter, especially for influenza viruses with pandemic potential.

Code(s) de classement : 002B02S

Descripteur(s) anglais

Descripteur(s) : Sensitivity; Oseltamivir; Antiviral; Influenzavirus AH5N1

Desc. génériques : Infectious diseases; Pharmacology; Medical sciences; Exo α sialidase; O Glycosidases; Glycosidases; Hydrolases; Enzyme; Enzyme inhibitor; Neuraminidase inhibitor

Descripteur(s) français

Descripteur(s) : Sensibilité; Oseltamivir; Antiviral; Influenzavirus AH5N1

Desc. génériques : Maladies infectieuses; Pharmacologie; Sciences médicales; Exo α sialidase; O Glycosidases; Glycosidases; Hydrolases; Enzyme; Inhibiteur enzyme; Inhibiteur neuraminidase

Localisation : INIST, Shelf number 13334, INIST No. 354000158897780380

Origine de la notice : INIST

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Inhibition of multiple subtypes of influenza A virus in cell cultures with morpholino oligomers

Titre : Inhibition of multiple subtypes of influenza A virus in cell cultures with morpholino oligomers

Auteur(s) : QING GE; PASTEY Manoj; KOBASA Darwyn; PUTHAVATHANA Piliapan; LUPFER Christopher; BESTWICK Richard K; IVERSEN Patrick L; JIANZHU CHEN; STEIN David A

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Source : Antimicrobial agents and chemotherapy. 2006; 50 (11) : 3724-3733

ISSN : 0066-4804

CODEN : AACHAX

Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 49 ref.

Résumé : Peptide-conjugated phosphorodiamidate morpholino oligomers (P-PMO) are single-stranded nucleic acid-like antisense agents that can reduce gene expression by sterically blocking complementary RNA sequence. P-PMO are water soluble and nuclease resistant, and they readily achieve uptake into cells in culture under standard conditions. Eight P-PMO, each 20 to 22 bases in length, were evaluated for their ability to inhibit influenza A virus (FLUAV) A/PR/8/34 (H1N1) replication in cell culture. The P-PMO were designed to base pair with FLUAV RNA sequences that are highly conserved across viral subtypes and considered critical to the FLUAV biological-cycle, such as gene segment termini and mRNA translation start site regions. Several P-PMO were highly efficacious, each reducing viral titer in a dose-responsive and sequence-specific manner in A/PR/8/34-infected cells. Two P-PMO, one designed to target the AUG translation start site region of PB1 mRNA and the other the 3'-terminal region of nucleoprotein viral genome RNA, also proved to be potent against several other FLUAV strains, including A/WSN/33 (H1N1), A/Memphis/8/88 (H3N2), A/Eq/Miami/63 (H3N8), A/Eq/Prague/56 (H7N7), and the highly pathogenic A/Thailand/1(KAN-1)/04 (H5N1). The P-PMO exhibited minimal cytotoxicity in cell viability assays. High efficacy by two of the P-PMO against multiple FLUAV subtypes suggests that these oligomers represent a broad-spectrum therapeutic approach against a high percentage of known FLUAV strains.

Code(s) de classement : 002B02S

Descripteur(s) anglais

Descripteur(s) : Inhibitor; Multiple; Subtype; Typing; Influenza A virus; In vitro; Cell culture; Oligomer

Desc. génériques : Infectious diseases; Pharmacology; Medical sciences; Influenzavirus A; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Inhibiteur; Multiple; Soustype; Typage; Virus grippal A; In vitro; Culture cellulaire; Oligomere

Desc. génériques : Maladies infectieuses; Pharmacologie; Sciences médicales; Influenzavirus A; Orthomyxoviridae; Virus

Localisation : INIST, Shelf number 13334, INIST No. 354000158897780280

Origine de la notice : INIST

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The NS1 gene contributes to the virulence of H5N1 avian influenza viruses

Titre : The NS1 gene contributes to the virulence of H5N1 avian influenza viruses

Auteur(s) : ZEJUN LI; YONGPING JIANG; PEIRONG JIAO; AIQIN WANG; FENGJU ZHAO; GUOBIN TIAN; XIJUN WANG; KANGZHEN YU; ZHIGAO BU; HUALAN CHEN

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Date de publication : 2006

Pays de publication : United States

Langue(s) : English

Type de document : Serial

Nombre de références : 34 ref.

Résumé : In the present study, we explored the genetic basis underlying the virulence and host range of two H5N1 influenza viruses in chickens. A/goose/Guangdong/1/96 (GS/GD/1/96) is a highly pathogenic virus for chickens, whereas A/goose/Guangdong/2/96 (GS/GD/2/96) is unable to replicate in chickens. These two H5N1 viruses differ in sequence by only five amino acids mapping to the PA, NP, M1, and NS1 genes. We used reverse genetics to create four single-gene recombinants that contained one of the sequence-differing genes from nonpathogenic GS/GD/2/96 and the remaining seven gene segments from highly pathogenic GS/GD/1/96. We determined that the NS1 gene of GS/GD/2/96 inhibited the replication of GS/GD/1/96 in chickens, while the substitution of the PA, NP, or M gene did not change the highly pathogenic properties of GS/GD/1/96. Conversely, of the recombinant viruses generated in the GS/GD/2/96 background, only the virus containing the NS1 gene of GS/GD/1/96 was able to replicate and cause disease and death in chickens. The single-amino-acid difference in the sequence of these two NS1 genes resides at position 149. We demonstrate that a recombinant virus expressing the GS/GD/1/96 NS1 protein with Ala149 is able to antagonize the induction of interferon protein levels in chicken embryo fibroblasts (CEFs), but a recombinant virus carrying a Val149 substitution is not capable of the same effect. These results indicate that the NS1 gene is critical for the pathogenicity of avian influenza virus in chickens and that the amino acid residue Ala149 correlates with the ability of these viruses to antagonize interferon induction in CEFs.

Code(s) de classement : 002A05C05

Descripteur(s) anglais

Descripteur(s) : Avian influenza virus; Gene; Virulence; Host range; Infectivity; Property structure relationship

Desc. génériques : Genetics; Virology; Microbiology; Biological sciences; Influenza virus A; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Influenza virus aviaire; Gene; Virulence; Spectre hôte; Pouvoir infectant; Relation structure propriété

Desc. génériques : Génétique; Virologie; Microbiologie; Sciences biologiques; Influenza virus A; Orthomyxoviridae; Virus

Localisation : INIST, Shelf number 13592, INIST No. 354000158896380210

Origine de la notice : INIST

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Genomic analysis of influenza A viruses, including avian Flu (H5N1) strains

Titre : Genomic analysis of influenza A viruses, including avian Flu (H5N1) strains

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Date de publication : 2006

Pays de publication : Netherlands

Langue(s) : English

Type de document : Serial

Nombre de références : 27 ref.

Résumé : This study was designed to conduct genomic analysis in two steps, such as the overall relative synonymous codon usage (RSCU) analysis of the five virus species in the orthomyxoviridae family, and more intensive pattern analysis of the four subtypes of influenza A virus (H1N1, H2N2, H3N2, and H5N1) which were isolated from human population. All the subtypes were categorized by their isolated regions, including Asia, Europe, and Africa, and most of the synonymous codon usage patterns were analyzed by correspondence analysis (CA). As a result, influenza A virus showed the lowest synonymous codon usage bias among the virus species of the orthomyxoviridae family, and influenza B and influenza C virus were followed, while suggesting that influenza A virus might have an advantage in transmitting across the species barrier due to their low codon usage bias. The ENC values of the host-specific HA and NA genes represented their different HA and NA types very well, and this reveals that each influenza A virus subtype uses different codon usage patterns as well as the amino acid compositions. In NP, PA and PB2 genes, most of the virus subtypes showed similar RSCU patterns except for H5N1 and H3N2 (A/HK/1774/1999) subtypes which were suspected to be transmitted across the species barrier, from avian and porcine species to human beings, respectively. This distinguishable synonymous codon usage patterns in non-human origin viruses might be useful in determining the origin of influenza A viruses in genomic levels as well as the serological tests. In this study, all the process, including extracting sequences from GenBank flat file and calculating codon usage values, was conducted by Java codes, and these bioinformatics-related methods may be useful in predicting the evolutionary patterns of pandemic viruses.

Code(s) de classement : 002B30A11; 002B05C02C

Descripteur(s) anglais

Descripteur(s) : Genomics; Influenza A; Correspondence analysis; Influenza A virus; Codon; Public health; Epidemiology; Avian influenza; Influenzavirus AH5N1

Desc. génériques : Public health; Medical sciences; Virology; Infectious diseases; Medical sciences; Viral disease; Infection; Influenzavirus A; Orthomyxoviridae; Virus

Descripteur(s) français

Descripteur(s) : Genomique; Grippe A; Analyse correspondance; Virus grippal A; Codon; Sante publique; Epidemiologie; Souche H5N1; Grippe aviaire; Influenzavirus AH5N1

Desc. génériques : Sante publique; Sciences medicales; Virologie; Maladies infectieuses; Sciences medicales; Virose; Infection; Influenzavirus A; Orthomyxoviridae; Virus

Localisation : INIST, Shelf number 20856, INIST No. 354000158837220040

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