



Bioaerosols and Transmission, a Diverse and Growing Community of Practice

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Mubareka S, Groulx N, Savory E, Cutts T, Theriault S, Scott JA, Roy CJ, Turgeon N, Bryce E, Astrakianakis G, Kirychuk S, Girard M, Kobinger G, Zhang C and Duchaine C (2019) Bioaerosols and Transmission, a Diverse and Growing Community of Practice. Front. Public Health 7:23. doi: 10.3389/fpubh.2019.00023 The transmission of infectious microbes via bioaerosols is of significant concern for both human and animal health. However, gaps in our understanding of respiratory pathogen transmission and methodological heterogeneity persist. New developments have enabled progress in this domain, and one of the major turning points has been the recognition that cross-disciplinary collaborations across spheres of human and animal health, microbiology, biophysics, engineering, aerobiology, infection control, public health, occupational health, and industrial hygiene are essential. Collaborative initiatives support advances in topics such as bioaerosol behavior, dispersion models, risk assessment, risk/exposure effects, and mitigation strategies in clinical, experimental, agricultural, and other field settings. There is a need to enhance the knowledge translation for researchers, stakeholders, and private partners to support a growing network of individuals and agencies to achieve common goals to mitigate inter- and intra-species pathogen transmission via bioaerosols.

Keywords: bioaerosols, microbes, virus, infections, viral dissemination, network, CANIBAN, collaborations

MODERN MIASMAS

The importance of infectious bioaerosols in disease transmission has been long-acknowledged, yet poorly understood. Paltry data and methodological heterogeneity limit many related studies. Effective ventilation and infection prevention and control (IP & C) measures in the form of droplet and airborne isolation in healthcare institutions underscore the contribution of these modes of pathogen dispersion. Moreover, recent outbreaks such as the Severe Acute Respiratory Syndrome coronavirus (SARS-CoV) and Middle East Respiratory Syndrome coronavirus (MERS-CoV) outbreaks highlight major gaps in our ability to assess and determine risk and to mitigate patient and healthcare worker (HCW) exposure alike (1, 2).

The SARS outbreak was eventually controlled in the absence of an effective vaccine or antiviral, strictly through public health interventions and IP & C measures aimed at controlling, among other things, bioaerosols emitted by infected patients (3). A decade on from this experience, MERS-CoV has caused community and healthcare-associated severe acute respiratory infections in the Middle East and South Korea, spreading by similar mechanism(s) (4–6).

From an animal health perspective, pathogens such as porcine reproductive and respiratory syndrome virus may be transmitted through the air for significant distances and have significant economic impact on the agricultural sector (7). Efforts to characterize this and other relevant organisms have been undertaken previously, specifically looking at the burden of inoculum related to transmission (8) and the prevention of spread of aerosols through various adequate ventilation strategies (9–11). Emission of African swine fever virus into the air by infected pigs raises the possibility for droplet and/or airborne transmission of this virus, which has recently produced significant alarm after documented spread in China (12–14). This hemorrhagic virus is associated with high mortality and significant loss for producers through depopulation and trade restrictions (15, 16).

In recent years, new developments have enabled progress in bioaerosol research, thus establishing a community of practice in the field. Although many developments have been technical, one of the major catalysts has been the recognition that crossdisciplinary collaborations across the various spheres of human and animal health, microbiology, engineering, aerobiology, infection control, public health, occupational health, and industrial hygiene are essential. A network approach has proven successful in other cross-disciplinary fields, including One Health and eco-health whereby wildlife, computational and evolutionary microbiologists, virologists, biologists, epidemiologists, ecologists, environmental scientists, climatologists, and human, animal, and public health practitioners are collaborating to address challenges in zoonotic diseases research and control (17, 18). This has enhanced surveillance efforts and launched ambitious, large scale projects such as the Global Virome Project, though gaps remain in stakeholder engagement and monitoring activities (19, 20). Translation of bioaerosol research stands to benefit in a similar fashion, provided coordinated and committed efforts.

Potentially infectious bioaerosols are pertinent to a wide range of pathogens, some of which may be endemic and cause sporadic infections and outbreaks (e.g., *Mycobacterium tuberculosis* and *Legionella pneumophila*, the causative agents of tuberculosis and legionnaires' disease, respectively) and/or have potential to cause epidemics or pandemics (e.g., influenza virus A) (21–24). In addition, the importance of potentially infectious bioaerosols across different settings is underscored. These include, but are not limited to, agriculture (both crop and livestock), wastewater treatment plants, environmental reservoirs (soil and water), acute and long-term healthcare institutions, and shared public spaces (transportation hubs, recreational areas, etc.). Here, we discuss the state of the science for this nascent field and identify gaps requiring urgent attention.

VALUE PROPOSITION FOR THE STUDY OF POTENTIALLY INFECTIOUS BIOAEROSOLS

Progress in the field has been stimulated by advances in other areas which have been applied to the study of infectious bioaerosols. These include: (a) enhanced detection by molecular methods, principally real-time and quantitative PCR, next-generation sequencing, metagenomics, and biosensors (25–30), and (b) establishment of both conventional and novel infrastructure, such as small and large-scale wind tunnels, biocontained rotating drums for aging aerosols, and field-ready aerosol samplers (31–33). Ongoing research has also facilitated the development and dissemination of procedures and protocols for experimental work, including artificial aerosols, as well as animal models of transmission including the ferret model for influenza virus transmission and macaque model for Ebola virus transmission (34, 35).

Whilst aerosols can be used as a means of delivery of therapeutic agents directly to the respiratory tract, they may also be used for the nefarious dispersion of human or animal pathogens. Scientists focused on bioaerosol generation, pathogen survival in air and aerobiological fitness must be acutely cognizant of any potential for dual use and routinely re-assess projects and proposed experiments with this perspective in mind. Oversight by institutional biosafety officers and committees may also consider this aspect of bioaerosol research during risk assessments and other internal approval processes.

Given the role of potentially infectious bioaerosols to transmit infectious agents to human and animal populations, it stands to reason that their detection and characterization will ultimately contribute to mitigating the spread of disease. This is possible through efforts focused on the following themes:

1. Fundamental biological and physical properties of bioaerosols and their generation

Unresolved fundamental questions may be answered by studying infectious bioaerosols. Areas meriting attention are numerous and a few are briefly discussed here.

In the case of agents for which airborne transmission is well-recognized (i.e., M. tuberculosis), associations between particle size and generation, pathogen content and virulence, as well as the deposition within the respiratory tract may be determined. The physical and chemical aspects of aerosols including the effects of relative humidity can affect pathogen viability and understanding these aspects of bioaerosol behavior may be beneficial to understanding conditions for controlling bioaerosol dispersion (36-38). Moreover, pathogens such as viruses may vary with respect to isoelectric points, possibly imparting different charges to infectious bioaerosols which may affect their behavior (39, 40). Finally, although climate change and pollution have been major issues within our biosphere, little is known about the impact of pollution particulates and gases, such as ozone, on the biology and physical properties of bioaerosols (41).

Our understanding of bioaerosol production from expulsion events such as breathing, talking, sneezing,

and coughing have generally been extrapolated from models, though more recent empiric evidence has become available, significantly enhancing our understanding of the transmission potential of bioaerosol emissions from naturallyinfected hosts (42, 43). Awareness of factors contributing to particle velocity and penetration into space enables modeling strategies that inform engineering controls in a multitude of settings. For example, in healthcare, understanding the dispersion of potential pathogens in the environment can inform infection prevention and control practices (44). There is also potential benefit to determining pathogen characteristics. If enhanced infectious bioaerosol survival in air is associated with certain strains, genotypes or mutations, this provides possibilities for follow-on work to (a) determine the mechanism and (b) enhance surveillance. The latter would have implications for public and animal health. Studying these factors individually or in combination can optimize air handling and other mechanical, environmental or chemical means of neutralizing bioaerosols prior to host exposure, thus alleviating dependence on personal protective equipment, which is the last means of protection prior to exposure.

2. Detection, surveillance, and early warning

New understanding of the role of bioaerosols in the propagation of human and animal pathogens could lead to new practices. As it stands, it remains uncertain which bioaerosol particles are predominantly responsible for personto-person transmission. There may also be differences between pathogen populations in the infected host vs. what is emitted into the air. This poses an excellent opportunity to leverage advances in metagenomics with developments in aerobiology to extract more sophisticated data from aerosol samples and, potentially, determine genetic bottlenecks for transmission. This would feed back into more fundamental work on bacterial or viral determinants of transmission, as well as lead to novel means for surveillance and mitigation.

Similarly, translational studies enhance our understanding of risk and determinants of exposure. For many settings and situations, several components of risk assessments are based on limited evidence. This is of particular concern from an occupational health perspective, in healthcare (e.g., purported aerosol-generating procedures), agricultural, or specific settings such as wastewater treatment plants (45–47). Developing the capacity to generate empiric data on pathogen content and biology from bioaerosols can lead to tools for outbreak investigation, surveillance, and development of risk assessment strategies and policies. As the technology for biosensors accelerates, the possibilities for rapid point-of-care testing and even remote sampling are highlighted. Integration of biosensors with bioaerosol sampling (26) has significant potential for early warning and public safety.

3. Research and development of mitigation strategies

As advances are made, capacity is built to optimize and evaluate known and novel means to control the dispersion of infectious bioaerosols. These include the use of germicidal and pulsed ultraviolet light, mechanical air filtration and respiratory protection which have both mechanical and electrostatic filtering systems (48, 49). In addition, the potential to use ozone to control indoor bioaerosols in urban and rural settings is being examined. Ozone is a strong oxidizing agent having high redox potential and a short half-life, dependent on temperature. It has been used extensively in the food industry (specifically water treatment, washing of produce, and food preparation), as well as in domestic restoration to remediate odors and smoke damage. Its historical use and proven efficacy demonstrate its potential to remediate contaminated indoor air (50, 51). This would present a novel application of a known treatment modality.

GAPS AND CHALLENGES

A number of important gaps have been identified (52) and these can be prioritized based on potential benefit:

Infrastructure

Biocontainment of infectious particles is the first consideration when aerosolizing human and/or animal pathogens in an experimental setting. These systems are generally customdesigned and constructed, with few facilities capable of conducting aerosol work with risk group 2 pathogens, and fewer still with the capacity to work with risk group 3 or 4 agents.

Technical

As challenges in infrastructure are overcome, opportunities to optimize and improve methods and techniques in bioaerosol research must be taken. A limitation of many studies focused on viral bioaerosols is the pervasive use of nucleic acid detection, rather than infectious virus isolation. The latter is a much more accurate indicator of the infectious potential of a bioaerosol but is infrequently performed due to poor sensitivity and other technical issues (53). The development of sampling devices and techniques optimized to preserve pathogen viability would considerably advance the utility of studying bioaerosols for risk assessment and management. For other applications, a more rapid, field-ready point of care test would be useful and would offer remote sampling possibilities. Biosensors have the potential to fill this gap and integration into aerosol sampling devices is under development (26). Finally, the collection of nucleic acid may be leveraged to obtain more sophisticated information than is available by PCR and Sanger sequencing. Metagenomics on environmental samples has been well-described in other spheres and is currently being explored for air samples (54-57).

Standardization

To advance bioaerosol-related data interpretation, an effort must be made to standardize and share protocols and reagents to reduce the degree of data heterogeneity to enable meaningful analyses across studies. This may apply to animal models, artificial aerosolizations, collection strategies in the field and clinical settings, processing of samples, and detection methods. Developing or adopting a standardized approach to aerosol sampling is challenging and requires substantial efforts to select the best sampling strategy to minimize sampling biases, optimize sample concentration and organism retrieval whilst preserving integrity (58).

Establishing standard approaches also enables training and implementation and eases knowledge translation amongst



different groups (**Figure 1**). Multiple investigators have published small studies on the recovery of viral RNA emitted by naturally infected humans using different approaches for recruitment, sampling, processing, and detection. Unfortunately, the results are difficult to compare and, standing alone, are of limited statistical significance (45, 59–61). By standardizing approaches, a more feasible method can be used to compare separate studies allowing for larger multi-center studies that are substantially more conclusive and impactful.

Training

Additional educational and operational needs, such as training of research personnel, proper use of personal protective equipment and developing decontamination protocols must be addressed. As more highly qualified personnel are properly trained, the greater the capacity for designing and utilizing experimental systems and for completing meaningful clinical and field studies. The need for cross-disciplinary experience is underscored, since robust knowledge of physics, mechanical engineering, and microbiology are required. Fostering this expertise requires a collaborative effort as each discipline offers unique insight. Currently, there are a limited number of workshops or accredited courses available to cultivate both interest and baseline knowledge. Although the profile of bioaerosol research is rising, few trainees are exposed to the field early in their careers. Courses such as *Bioaérosols et aérobiologie* (Bioaerosols and Aerobiology) developed at the Université Laval in Québec may help close this gap further given the opportunity to expand reach.

Knowledge Translation and Response

The relevance of bioaerosol data to occupational health and infection prevention and control requires attention. Validation

of bioaerosol data for risk assessments and risk management is needed. Air sampling was conducted during both SARS-CoV and MERS-CoV outbreaks, accruing important insight into the environmental distribution and persistence of these coronaviruses (4, 62, 63). The absence of data for more common, lower consequence pathogens represents missed opportunities to build a contextual and impactful body of knowledge during inter-outbreak and inter-pandemic periods.

To develop evidence-based policy and ensure the relevance of this work, early stakeholder engagement is needed. Consultation with human and animal public health, infection prevention and control, industrial hygiene, and occupational health stakeholders will ensure that the work is germane to the challenges presented by bioaerosol exposures. This pull also increases the likelihood that there will be data generated for risk assessment, policy development and implementation.

THE ROAD AHEAD; SOLUTIONS AND RECOMMENDATIONS

Part of the chaos that characterized the SARS epidemic can be attributed to a lack of knowledge regarding the route(s) of transmission. This remains true for high consequence coronaviruses, as well as for a range of other respiratory pathogens, both novel and established. We propose the following to address the challenges outlined above and to further develop the field of applied bioaerosol research:

1. An open network approach

Working in isolation, microbiologists, aerobiologists, engineers, and epidemiologists have made only incremental progress. A synergistic and pluralistic approach is required for action-driven research which effects change. As capacity grows, so will opportunities for training and response. A networked approach will lead to context-specific follow-on research to ultimately inform policy for the mitigation of respiratory pathogens spread in healthcare institutions, agricultural settings and public spaces. Toward this end, the present authors have established the Canadian Infectious Bioaerosols Network (CANIBAN) that brings together members of each of these disciplines with the primary objectives of understanding transmission of pathogens.

2. Shared infrastructure, technical protocols, and training programs

A network would form a hub around which key resources such as infrastructure, equipment, and operational procedures could be shared, thus also enabling training and underscoring best practices in biosafety and biosecurity. Shared resources may encompass wind tunnels, cough chambers and mannequins, animal exposure and other biosafety enclosures, and rotating drums to examine pathogen survival in air,

REFERENCES

1. Chowell G, Abdirizak F, Lee S, Lee J, Jung E, Nishiura H, et al. Transmission characteristics of MERS and SARS in the healthcare setting: together with instrumentation and analysis tools for fluid flow measurement and biochemical assays.

3. Identifying and engaging knowledge-users

Limited awareness and understanding of infectious bioaerosols among potential knowledge users, coupled with equally limited outreach by bioaerosol researchers has restricted knowledge transfer and applications. Knowledgeuser engagement in research planning from the outset and ongoing involvement through to dissemination is key for effective projects. Potential knowledge users include infection prevention and control practitioners, infectious disease specialists, veterinarians and animal health epidemiologists, industrial hygienists, and public health agencies.

4. Enhanced capacity-building for response measures Increasingly, bioaerosol sampling has been proposed and implemented in outbreak investigations and pathogen surveillance using *ad hoc* approaches. The development of best practices in these areas is essential to generating valid and actionable data.

In summary, a collective path for researchers, stakeholders, and private partners is needed to support a network of individuals and agencies to achieve common goals. As the field of bioaerosol studies grows, applications will diversify, along with novel technologies for remote sampling and point of care testing yielding results in real time. This stands to mitigate the spread of nefarious pathogens and contribute to early warning and response measures, thus ultimately benefiting human and animal health.

AUTHOR CONTRIBUTIONS

SM framed the manuscript and contributed content. NG contributed content and generated the figure. ES contributed content and supported manuscript organization. TC, ST, JS, CR, NT, EB, GA, SK, MG, GK, and CZ contributed content and expertise and CD helped to frame and contributed content.

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a comparative study. BMC Med. (2015) 13:210. doi: 10.1186/s12916-015-0450-0

2. Kim SH, Chang SY, Sung M, Park JH, Bin Kim H, Lee H, et al. Extensive Viable Middle East Respiratory Syndrome (MERS) *Coronavirus* contamination in air and surrounding environment in MERS isolation wards. Clin Infect Dis. (2016) 63:363–9. doi: 10.1093/cid/ciw239

- Yassi A, Moore D, Fitzgerald JM, Bigelow P, Hon CY, Bryce E, et al. Research gaps in protecting healthcare workers from SARS and other respiratory pathogens: an interdisciplinary, multi-stakeholder, evidence-based approach. J Occup Environ Med. (2005) 47:41–50. doi: 10.1097/01.jom.0000150207.18085.41
- Bin SY, Heo JY, Song MS, Lee J, Kim EH, Park SJ, et al. Environmental contamination and viral shedding in MERS patients during MERS-CoV outbreak in South Korea. *Clin Infect Dis.* (2016) 62:755–60. doi: 10.1093/cid/civ1020
- Cho SY, Kang JM, Ha YE, Park GE, Lee JY, Ko JH, et al. MERS-CoV outbreak following a single patient exposure in an emergency room in South Korea: an epidemiological outbreak study. *Lancet* (2016) 388:994–1001. doi: 10.1016/S0140-6736(16)30623-7
- Al-Omari A, Rabaan AA, Salih S, Al-Tawfiq JA, Memish ZA. MERS coronavirus outbreak: implications for emerging viral infections. *Diagn Microbiol Infect Dis.* (2019) 93:265–85. doi: 10.1016/j.diagmicrobio.2018.10.011
- Brito B, Dee S, Wayne S, Alvarez J, Perez A. Genetic diversity of PRRS virus collected from air samples in four different regions of concentrated swine production during a high incidence season. *Viruses* (2014) 6:4424–36. doi: 10.3390/v6114424
- Anderson BD, Lednicky JA, Torremorell M, Gray GC. The use of bioaerosol sampling for airborne virus surveillance in swine production facilities: a mini review. *Front Vet Sci.* (2017) 4:121. doi: 10.3389/fvets.2017.00121
- 9. Dee S, Otake S, Deen J. Use of a production region model to assess the efficacy of various air filtration systems for preventing airborne transmission of porcine reproductive and respiratory syndrome virus and *Mycoplasma hyopneumoniae*: results from a 2-year study. *Virus Res.* (2010) 154:177–84. doi: 10.1016/j.virusres.2010.07.022
- Alonso C, Otake S, Davies P, Dee S. An evaluation of interventions for reducing the risk of PRRSV introduction to filtered farms via retrograde air movement through idle fans. *Vet Microbiol.* (2012) 157:304–10. doi: 10.1016/j.vetmic.2012.01.010
- Alonso C, Murtaugh MP, Dee SA, Davies PR. Epidemiological study of air filtration systems for preventing PRRSV infection in large sow herds. *Prev Vet Med.* (2013) 112:109–17. doi: 10.1016/j.prevetmed.2013.06.001
- de Carvalho Ferreira HC, Weesendorp E, Quak S, Stegeman JA, Loeffen WL. Quantification of airborne African swine fever virus after experimental infection. *Vet Microbiol.* (2013) 165:243–51. doi: 10.1016/j.vetmic.2013.03.007
- Ge S, Li J, Fan X, Liu F, Li L, Wang Q, et al. Molecular characterization of African swine fever virus, China, 2018. *Emerg Infect Dis.* (2018) 24:2131–3. doi: 10.3201/eid2411.181274
- Zhou X, Li N, Luo Y, Liu Y, Miao F, Chen T, et al. Emergence of African swine fever in China, 2018. *Transbound Emerg Dis.* (2018) 65:1482–4. doi: 10.1111/tbed.12989
- Guinat C, Porphyre T, Gogin A, Dixon L, Pfeiffer DU, Gubbins S. Inferring within-herd transmission parameters for African swine fever virus using mortality data from outbreaks in the Russian Federation. *Transbound Emerg Dis.* (2018) 65:e264–71. doi: 10.1111/tbed.12748
- Wang T, Sun Y, Qiu HJ. African swine fever: an unprecedented disaster and challenge to China. *Infect Dis Poverty* (2018) 7:111. doi: 10.1186/s40249-018-0495-3
- Machalaba C, Romanelli C, Stoett P, Baum SE, Bouley TA, Daszak P, et al. Climate change and health: transcending silos to find solutions. *Ann Glob Health* (2015) 81:445–58. doi: 10.1016/j.aogh.2015.08.002
- Plowright RK, Parrish CR, Mccallum H, Hudson PJ, Ko AI, Graham AL, et al. Pathways to zoonotic spillover. *Nat Rev Microbiol.* (2017) 15:502–10. doi: 10.1038/nrmicro.2017.45
- Carroll D, Daszak P, Wolfe ND, Gao GF, Morel CM, Morzaria S, et al. The global virome project. *Science* (2018) 359:872–4. doi: 10.1126/science.aap7463
- Khan MS, Rothman-Ostrow P, Spencer J, Hasan N, Sabirovic M, Rahman-Shepherd A, et al. The growth and strategic functioning of One Health networks: a systematic analysis. *Lancet Planet Health* (2018) 2:e264–73. doi: 10.1016/S2542-5196(18)30084-6
- 21. Wurie FB, Lawn SD, Booth H, Sonnenberg P, Hayward AC. Bioaerosol production by patients with tuberculosis during normal tidal

breathing: implications for transmission risk. *Thorax* (2016) 71:549–54. doi: 10.1136/thoraxjnl-2015-207295

- 22. Montagna MT, De Giglio O, Cristina ML, Napoli C, Pacifico C, Agodi A, et al. Evaluation of *Legionella* air contamination in healthcare facilities by different sampling methods: an italian multicenter study. *Int J Environ Res Public Health* (2017) 14:E670. doi: 10.3390/ijerph14070670
- Hamilton KA, Hamilton MT, Johnson W, Jjemba P, Bukhari Z, Lechevallier M, et al. Health risks from exposure to Legionella in reclaimed water aerosols: toilet flushing, spray irrigation, and cooling towers. *Water Res.* (2018) 134:261–79. doi: 10.1016/j.watres.2017.12.022
- 24. Rule AM, Apau O, Ahrenholz SH, Brueck SE, Lindsley WG, De Perio MA, et al. Healthcare personnel exposure in an emergency department during influenza season. *PLoS ONE* (2018) 13:e0203223. doi: 10.1371/journal.pone.0203223
- Prussin AJ II, Marr LC, Bibby KJ. Challenges of studying viral aerosol metagenomics and communities in comparison with bacterial and fungal aerosols. FEMS Microbiol Lett. (2014) 357:1–9. doi: 10.1111/1574-6968.12487
- Fronczek CF, Yoon JY. Biosensors for monitoring airborne pathogens. J Lab Autom. (2015) 20:390–410. doi: 10.1177/2211068215580935
- Degois J, Clerc F, Simon X, Bontemps C, Leblond P, Duquenne P. First metagenomic survey of the microbial diversity in bioaerosols emitted in waste sorting plants. *Ann Work Expo Health* (2017) 61:1076–86. doi: 10.1093/annweh/wxx075
- Ladhani L, Pardon G, Meeuws H, Van Wesenbeeck L, Schmidt K, Stuyver L, et al. Sampling and detection of airborne influenza virus towards point-of-care applications. *PLoS ONE* (2017) 12:e0174314. doi: 10.1371/journal.pone.0174314
- Coleman KK, Nguyen TT, Yadana S, Hansen-Estruch C, Lindsley WG, Gray GC. Bioaerosol Sampling for respiratory viruses in singapore's mass rapid transit network. *Sci Rep.* (2018) 8:17476. doi: 10.1038/s41598-018-35896-1
- Serrano-Silva N, Calderon-Ezquerro MC. Metagenomic survey of bacterial diversity in the atmosphere of Mexico City using different sampling methods. *Environ Pollut.* (2018) 235:20–9. doi: 10.1016/j.envpol.2017.12.035
- Fabian P, Mcdevitt JJ, Houseman EA, Milton DK. Airborne influenza virus detection with four aerosol samplers using molecular and infectivity assays: considerations for a new infectious virus aerosol sampler. *Indoor Air* (2009) 19:433–41. doi: 10.1111/j.1600-0668.2009.00609.x
- Su WC, Tolchinsky AD, Chen BT, Sigaev VI, Cheng YS. Evaluation of physical sampling efficiency for cyclone-based personal bioaerosol samplers in moving air environments. *J Environ Monit.* (2012) 14:2430–7. doi: 10.1039/c2em30299c
- Verreault D, Killeen SZ, Redmann RK, Roy CJ. Susceptibility of monkeypox virus aerosol suspensions in a rotating chamber. J Virol Methods (2013) 187:333-7. doi: 10.1016/j.jviromet.2012.10.009
- Richard M, Fouchier RA. Influenza A virus transmission via respiratory aerosols or droplets as it relates to pandemic potential. *FEMS Microbiol Rev.* (2016) 40:68–85. doi: 10.1093/femsre/fuv039
- Harbourt DE, Johnston SC, Pettitt J, Warren TK, Dorman WR. Detection of ebola virus RNA through aerosol sampling of animal biosafety Level 4 rooms housing challenged nonhuman primates. *J Infect Dis.* (2017) 215:554–8. doi: 10.1093/infdis/jiw610
- Yang W, Elankumaran S, Marr LC. Relationship between humidity and influenza A viability in droplets and implications for influenza's seasonality. *PLoS ONE* (2012) 7:e46789. doi: 10.1371/journal.pone.0046789
- Yang W, Marr LC. Mechanisms by which ambient humidity may affect viruses in aerosols. *Appl Environ Microbiol.* (2012) 78:6781–8. doi: 10.1128/AEM.01658-12
- Vejerano EP, Marr LC. Physico-chemical characteristics of evaporating respiratory fluid droplets. J R Soc Interface (2018) 15. doi: 10.1098/rsif.2017.0939. [Epub ahead of print].
- Michen B, Graule T. Isoelectric points of viruses. J Appl Microbiol. (2010) 109:388–97. doi: 10.1111/j.1365-2672.2010.04663.x
- Groulx N, Lecours C, Turgeon N, Volckens J, Tremblay ME, Duchaine C. Nanoscale aerovirology: an efficient yet simple method to analyze the viral distribution of single bioaerosols. *Aerosol Sci Technol.* (2016) 50:732–9. doi: 10.1080/02786826.2016.1184223
- 41. Groulx N, Urch B, Duchaine C, Mubareka S, Scott JA. The Pollution Particulate Concentrator (PoPCon): a platform to investigate the effects of

particulate air pollutants on viral infectivity. *Sci Total Environ.* (2018) 628-9:1101-07. doi: 10.1016/j.scitotenv.2018.02.118

- Fabian P, Mcdevitt JJ, Dehaan WH, Fung RO, Cowling BJ, Chan KH, et al. Influenza virus in human exhaled breath: an observational study. *PLoS ONE* (2008) 3:e2691. doi: 10.1371/journal.pone.00 02691
- 43. Yan J, Grantham M, Pantelic J, Bueno De Mesquita PJ, Albert B, Liu F, et al. Infectious virus in exhaled breath of symptomatic seasonal influenza cases from a college community. *Proc Natl Acad Sci USA*. (2018) 115:1081–6. doi: 10.1073/pnas.1716561115
- Tang JW. The effect of environmental parameters on the survival of airborne infectious agents. J R Soc Interface (2009) 6 (Suppl. 6):S737–46. doi: 10.1098/rsif.2009.0227.focus
- Bischoff WE, Swett K, Leng I, Peters TR. Exposure to influenza virus aerosols during routine patient care. J Infect Dis. (2013) 207:1037–46. doi: 10.1093/infdis/jis773
- 46. Thompson KA, Pappachan JV, Bennett AM, Mittal H, Macken S, Dove BK, et al. Influenza aerosols in UK hospitals during the H1N1 (2009) pandemic-the risk of aerosol generation during medical procedures. *PLoS ONE* (2013) 8:e56278. doi: 10.1371/journal.pone.0056278
- Brisebois E, Veillette M, Dion-Dupont V, Lavoie J, Corbeil J, Culley A, et al. Human viral pathogens are pervasive in wastewater treatment center aerosols. *J Environ Sci.* (2018) 67:45–53. doi: 10.1016/j.jes.2017. 07.015
- Lee SA, Grinshpun SA, Reponen T. Respiratory performance offered by N95 respirators and surgical masks: human subject evaluation with NaCl aerosol representing bacterial and viral particle size range. *Ann Occup Hyg.* (2008) 52:177–85. doi: 10.1093/annhyg/men005
- Miaskiewicz-Peska E, Lebkowska M. Comparison of aerosol and bioaerosol collection on air filters. *Aerobiologia* (2012) 28:185–93. doi: 10.1007/s10453-011-9223-1
- Martinelli M, Giovannangeli F, Rotunno S, Trombetta CM, Montomoli E. Water and air ozone treatment as an alternative sanitizing technology. J Prev Med Hyg. (2017) 58:E48–52.
- Jiang L, Li M, Tang J, Zhao X, Zhang J, Zhu H, et al. Effect of different disinfectants on bacterial aerosol diversity in poultry houses. *Front Microbiol.* (2018) 9:2113. doi: 10.3389/fmicb.2018.02113
- Herfst S, Bohringer M, Karo B, Lawrence P, Lewis NS, Mina MJ, et al. Drivers of airborne human-to-human pathogen transmission. *Curr Opin Virol.* (2017) 22:22–9. doi: 10.1016/j.coviro.2016.11.006
- Nikiforuk AM, Cutts TA, Theriault SS, Cook BWM. Challenge of liquid stressed protective materials and environmental persistence of ebola virus. *Sci Rep.* (2017) 7:4388. doi: 10.1038/s41598-017-04137-2
- Hall RJ, Leblanc-Maridor M, Wang J, Ren X, Moore NE, Brooks CR, et al. Metagenomic detection of viruses in aerosol samples from workers in animal slaughterhouses. *PLoS ONE* (2013) 8:e72226. doi: 10.1371/journal.pone.0072226

- Bashir M, Ahmed M, Weinmaier T, Ciobanu D, Ivanova N, Pieber TR, et al. Functional metagenomics of spacecraft assembly cleanrooms: presence of virulence factors associated with human pathogens. *Front Microbiol.* (2016) 7:1321. doi: 10.3389/fmicb.2016.01321
- Meta S.U.B.I.C. The Metagenomics and Metadesign of the Subways and Urban Biomes (MetaSUB) International Consortium inaugural meeting report. *Microbiome* (2016) 4:24. doi: 10.1186/s40168-016-0168-z
- Be NA, Avila-Herrera A, Allen JE, Singh N, Checinska Sielaff A, Jaing C, et al. Whole metagenome profiles of particulates collected from the International Space Station. *Microbiome* (2017) 5:81. doi: 10.1186/s40168-017-0292-4
- Mbareche H, Veillette M, Bonifait L, Dubuis ME, Benard Y, Marchand G, et al. A next generation sequencing approach with a suitable bioinformatics workflow to study fungal diversity in bioaerosols released from two different types of composting plants. *Sci Total Environ.* (2017) 601–2:1306–14. doi: 10.1016/j.scitotenv.2017.05.235
- 59. Lindsley WG, Blachere FM, Davis KA, Pearce TA, Fisher MA, Khakoo R, et al. Distribution of airborne influenza virus and respiratory syncytial virus in an urgent care medical clinic. *Clin Infect Dis.* (2010) 50:693–8. doi: 10.1086/650457
- Mubareka S, Granados A, Naik U, Darwish I, Cutts TA, Astrakianakis G, et al. Influenza virus emitted by naturally-infected hosts in a healthcare setting. J Clin Virol. (2015) 73:105–7. doi: 10.1016/j.jcv.2015.11.002
- Leung NH, Zhou J, Chu DK, Yu H, Lindsley WG, Beezhold DH, et al. Quantification of influenza virus RNA in aerosols in patient rooms. *PLoS ONE* (2016) 11:e0148669. doi: 10.1371/journal.pone.0148669
- Booth TF, Kournikakis B, Bastien N, Ho J, Kobasa D, Stadnyk L, et al. Detection of airborne severe acute respiratory syndrome (SARS) coronavirus and environmental contamination in SARS outbreak units. *J Infect Dis.* (2005) 191:1472–7. doi: 10.1086/429634
- Tsai YH, Wan GH, Wu YK, Tsao KC. Airborne severe acute respiratory syndrome coronavirus concentrations in a negative-pressure isolation room. *Infect Cont Hosp Epidemiol.* (2006) 27:523–5. doi: 10.1086/504357

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