Modeling of hantavirus cardiopulmonary syndrome

Juan Pablo Gutiérrez Jara^ª*© María Teresa Muñoz Quezada^b©

^a Centro de Investigación de Estudios Avanzados del Maule (CIEAM), Vicerrectoría de Investigación y Postgrado, Universidad Católica del Maule, Talca, Chile

^b Centro de Investigación en Neuropsicología y Neurociencias Cognitivas (CINPSI Neurocog), Facultad de Ciencias de la Salud, Universidad Católica del Maule, Talca, Chile

* Corresponding author jpgutierrezjara@gmail.com

Citation

Gutiérrez Jara JP, Muñoz Quezada MT. Modeling of hantavirus cardiopulmonary syndrome. *Medwave* 2022;22(3):002526

DOI

10.5867/ medwave.2022.03.002526

> Submission date Sep 7, 2021 Acceptance date Mar 1, 2022 Publication date Apr 26, 2022

Keywords Hantavirus, Epidemiology, Mathematics

Postal address Avenida San Miguel 3605 Talca, Chile

Abstract

Introduction

Hantavirus cardiopulmonary syndrome is a disease caused by a virus belonging to the *Bunyanvirales* order and transmitted to humans through rodents. This disease is considered endemic in Chile and has a high lethality rate. There is evidence of person-to-person contagion of the Andes virus in Argentina and Chile.

Objectives

To analyze the possibility of hantavirus transmission between humans through a SIER mathematical model.

Methods

A SEIR mathematical model (susceptible, exposed, infectious, and recovered) is proposed to express the dynamics of hantavirus disease, including the possibility of person-to-person transmission and risk perception.

Results

The maximum person-to-person transmission decreases by about 25% after increasing people's risk perception by reducing the resistance to change and increasing the speed of reaction.

Conclusions

It is urgent to review risk communication strategies and prevention measures, considering the possibility of massive person-to-person transmission. In addition, we should strengthen research and plan the development of a vaccine to protect populations exposed to this highly lethal disease.

MAIN MESSAGES

- The hantavirus disease is endemic in some countries, including Argentina and Chile.
- The spread of the Andes virus between people is becoming more recurrent in Argentina and Chile.
- The proposed model shows that the delay of prevention measures may cause a significant increase in hantavirus infection. Its high lethality rate can become a severe public health problem.
- The scarce information on person-to-person hantavirus transmission and the generalist nature of the model are the main limitations of this work.

INTRODUCTION

Hantavirus cardiopulmonary syndrome is an infection caused by rodents of the *Bunyavirales* order [1]. The *Sin Nombre orthohantavirus* and Andes virus are the main agents infecting humans within this order. *Peromyscus maniculatus* (deer mouse) and *Oligoryzomys* (long-tailed mouse) rodents are the natural reservoirs of hantavirus [1–3]. The transmission from rodents to humans is through inhalation of viral particles of urine, feces, or saliva [4,5]. Although isolated, there are reported cases of transmission via bites.

In Chile, Hantavirus is an endemic disease [4] that produced 91, 33, 70, and 30 cases from 2017 to 2020, respectively [6]. Although its incidence is not high compared to other diseases, it has a fatality rate between 30 and 60% [7].

Person-to-person transmission of the Andes virus has been found in Argentina and Chile. In Argentina, the first confirmed case occurred in 1996, and in 2014 three other cases were reported [8]. In Chile in 2019, a possible person-to-person transmission was recorded from one person (index case) to her children, including a newborn, and to her daughter's caregiver [4]. In addition, the virus was detected in breast milk, so transmission by this route cannot be ruled out. In Argentina, a study conducted between November 2018 and February 2019 recorded 34 confirmed person-to-person transmission cases in the province of Chubut [9]. The investigators found that patients with a high viral load and liver injury were more likely to spread the infection than other patients.

Mathematical models have contributed to understanding the dynamics of hantavirus transmission among rodents [10–16]. Some studies have modeled the demographic effect and associated environmental variables [13–16], and others have analyzed the rodent-to-rodent [10–12] and rodent-to-human transmission [17,18]. However, there is a paucity of models incorporating person-to-person hantavirus transmission – which is the main novelty of this work.

For rodent-to-human transmission dynamics of hantavirus, the studies mentioned above work with the premise of four infection states: susceptible (S), exposed (E), infected (I), and recovered (R), commonly denoted by SEIR [19–21]. Infection in humans presents an incubation period of one to six weeks [6],

then the host becomes symptomatic and may later acquire an immunity or die. Therefore, its disease evolution can be extrapolated through the SEIR model.

A factor determining disease control is people's risk perception [22–25]. This aspect is supported by the decrease in hantavirus disease during years of a strong prevention publicity campaign in Chile [4,26,27].

This paper presents the dynamics of hantavirus disease using a mathematical SEIR model [19–21], which incorporates person-to-person transmission of the Andes virus.

METHODS

MATHEMATICAL MODEL

We based the dynamics of human-associated disease (h) on a SEIR model, which includes susceptible (S), exposed (E), infected (I), and recovered (R) states. An infected individual begins an incubation period (E) of one and six weeks and subsequently presents a symptomatic period (I) where the disease can be transmitted to other humans. Finally, the individual acquires immunity (R) or dies.

We also analyzed the rodent-to-rodent transmission (m) through the SEIR model. The transmission process is similar to humans but with different infection periods. Thus, to distinguish the SIER states associated with the disease between humans and rodents, the subscript (b) is used for humans and (m) for rodents (Table 1).

A predetermining factor of rodent-to-human transmission is the risk perception (P) [28], which refers to people's subjective judgments about adverse outcomes of disease, accidents, or death. The risk perception has a cognitive (i.e., what they know about hantavirus) and an emotional component (i.e., what they

Table 1. Notation of the disease states.

	Susceptible	Exposed	Infected	Recovered
Human	S_b	E_b	I_b	R _b
Rodent	S _m	E_m	I_m	R _m

Source: Prepared by the authors of this study.

feel about hantavirus). Its assessment is essential to create practical public health interventions and improve risk communication since it informs about their perceived dangers and coping strategies. By taking the appropriate measures and generating an effective risk communication campaign, the probability of contagion can be significantly reduced. The risk perception (P) depends on a wide variety of factors, including resistance to change (λ_1) and speed of reacting (λ_2) to positive cases.

The model involving hantavirus dynamics between humans and rodents is given by the system of differential equations in (1).

$$\begin{cases} S'_{b} = b_{b}N_{b} - S_{b} \left(\beta I_{m} + \beta_{b}I_{b}\right) / N_{b} - d_{b}S_{b} \\ E'_{b} = S_{b} \left(\beta I_{m} + \beta_{b}I_{b}\right) / N_{b} - \left(\delta_{b} + d_{b}\right) E_{b} \\ I'_{b} = \delta_{b}E_{b} - \left(\gamma_{b} + d_{b} + \epsilon\right) I_{b} \\ R'_{b} = \gamma_{b}I_{b} - d_{b}R_{b} \end{cases}$$
(1)
$$\begin{aligned} S'_{m} = b_{m}N_{m} - \beta_{m}S_{m}I_{m} / N_{m} - d_{m}S_{m} \\ E'_{m} = \beta_{m}S_{m}I_{m} / N_{m} - \left(\delta_{m} + d_{m}\right) E_{m} \\ I'_{m} = \delta_{m}E_{m} - \left(\gamma_{m} + d_{m}\right) I_{m} \\ R'_{m} = \gamma_{m}R_{m} - d_{m}R_{m} \end{aligned}$$

In this model, β_h , β_m y β corresponds to person-to-person, rodent-to rodent, and rodent-to-human contagion rates, respectively. Regarding person-to-person transmission rates (β_b and β), these are dependent on the risk perception, with $\beta_b = \beta_b^* (P_*/P)$ and $\beta = \beta^* (P_*/P)$, where β_b^* , β^* y P_* correspond to the average person-to-person and rodent-to-human transmission rates and the average quantified risk perception, respectively. Then, N_b and N_m denote the human and rodent populations, respectively. It is important to note that the rodent-to-human transmission $\beta S_b I_m / N_m$ is affected by the number of rodents per human $K = N_m / N_b$ [29–31]. Thus, it finally remains $\beta S_b (I_m / N_m) K = \beta S_b I_m / N_b$.

The transition rates from exposure (E) to infectious (I) states and from infectious to recovered (R) are given by δ and γ , respectively. Thus δ^{-1} and γ^{-1} determine the average time they remain in their incubated and infectious condition, respectively. The subscripts *b* and *m* differentiate the rates associated with humans and rodents, respectively. Birth and non-disease mortality rates are expressed by b_x and d_x with $x \in \{b, m\}$, respectively. The disease lethality rate is given by ϵ .

In mathematical epidemiology, an essential value to determine threshold conditions is the basic reproductive number R_0 [32,33], which is defined as "the expected number of secondary cases produced by a first infectious agent in a susceptible population" [32], which establishes that if $R_0 > 1$ the disease spreads, and if $R_0 < 1$ it will not spread. In a local outbreak of the Andes virus in Argentina [9], the estimated reproductive number of transmissions between humans was 2.12 before announcing isolation measures for asymptomatic patients and self-quarantine for high-risk patients. Once control measures were applied, this estimate decreased to 0.96 [9]. From system (1), using the following generation method, the basic reproductive number in the person-to-person transmission is given by (2).

$$R_0 = \beta_b / \left(\gamma_b + d_b + \epsilon \right) \tag{2}$$

BACKGROUND

According to the Ministry of Health in Chile [6], 13 cases of hantavirus were reported between January and March 2021, compared to 30 cases in the entire 2020 year (Table 2). It should be noted that – due to the COVID-19 pandemic and health measures proposed by the authorities – the mobility of people in these two years was reduced. In contrast, previous to the restriction measures in 2019, 70 cases were recorded.

NUMERICAL SIMULATIONS

Table 3 shows the values used in the numerical simulations. These values are extracted from other research, and some have been estimated from the available data.

We considered $N_h = 100\ 000$ as the initial human population without infected cases, and we considered $N_m = 100$ as the initial population of rodents, with only one infected case.

The numerical simulations expressed in graphs were elaborated via MATLAB software.

RESULTS

NO PERSON-TO-PERSON TRANSMISSION

Before looking at the impact of person-to-person transmission, it is essential to visualize the rodent-to-human transmission. Through the model simulation and canceling out the rates associated with the person-to-person transmission, Figure 1 shows that in one year goes to 5.8 cases per 100 000 inhabitants (Figure 1b), with a variation during this period of +0.06 (Figure 1a). Thus, the simulations align with the real dynamics, where agriculture and forestry are the primary sources of labor.

The number of infected rodents is challenging to estimate since there is no systematized registry of rodents carrying the Andes virus in Chile. However, Figure 2 shows the dynamics of the number of infected rodents that the proposed model yields. Figure 2a shows a slight variation of +2 cases, and Figure 2b shows 12 accumulated cases for one year.

	2019		2020		20211	
Region	Number of cases	Rate of incidence	Number of cases	Rate of incidence	Number of cases	Rate of incidence
Valparaíso	2	0.1	0	0.2	0	0.2
Metropolitan	2	0.2	0	0.0	0	0.0
O'Higgins	1	0.1	1	0.1	0	0.1
Maule	15	1.3	3	0.3	0	0.0
Ñuble	8	1.6	3	0.6	2	0.4
Biobío	7	0.4	1	0.1	1	0.1
Araucanía	9	0.9	4	0.4	1	0.1
Los Ríos	12	3.0	5	1.2	4	1.0
Los Lagos	11	1.2	8	0.9	4	0.4
Aysén	1	0.9	3	2.8	0	0.0
Magallanes	0	0.0	0	0.0	0	0.0
In study	2		2		1	
Total	70	0.4	30	0.2	13	0.07

Table 2. Number of hantavirus cases in Chile.

¹Data as of March 21, 2021.

There are no cases from the region of Arica and Parinacota to Coquimbo.

Incidence rate per hundred thousand inhabitants.

Source: table extracted from the Department of Epidemiology, Ministry of Health, Minuta 23/03/2021 (http://epi.minsal.cl/wp-content/uploads/2021/03/ Minuta_hanta_2021_3_23.pdf) [6].

PERSON-TO-PERSON TRANSMISSION

Figure 3 shows humans' disease dynamics and risk perception, considering values shown in Table 3. Figure 3a illustrates the dynamics of hantavirus infection in humans during a year in which initially there were no infections. The maximum number of infections occurs almost at the end of one year due to the long incubation periods. It is evident from Figure 3b that risk

perception has a behavior similar to the infectious curve, with daily variations.

Compared to Figures 3 and 4 shows a decimal positional decrease in the rate of resistance to change $(\lambda_1/10)$, and an increase in the same magnitude to the reaction rate $(\lambda_2 * 10)$. It is observed from Figure 4a that the maximum contagion decreased by almost 25%, which is highly significant after

Table 3. Summary and	parameter values	for numerical	simulations.
----------------------	------------------	---------------	--------------

	Parameter	Definition	Value	Ref.
HUMAN	b_b	Birth rate	0.0001243	[34]
	d_b	Mortality rate	0.0001243	EA
	ϵ	Lethality rate due to hantavirus	[0.2 to 0.6]	[4]
	β^*	Rodent-to-human transmission rate	0.01	[17]
	β_b^*	Person-to-person transmission rate	$2.12(\gamma_b + d_b + \epsilon)$	[9]
	$1/\delta_b$	Average incubation time in humans	[7 to 42]	[6]
	$1/\gamma_b$	Average infectious time in humans	[1 to 5]	[6]
	P_*	Average quantified risk perception	[0 to 1]	[25]
	λ_1	Rate of resistance to change	0.65	[35]
	λ_2	Reaction rate	0.05	[24]
RODENT	b_m	Rodent birth rate	0.00139	[36]
	d_b	Rodent mortality rate	0.00139	EA
	β_m	Rodent-to-rodent transmission rate	1.38 $(\gamma_m + d_m)$	[10]
	$1/\delta_m$	Average incubation in rodents	[7 to 30]	[10]
	$1/\gamma_b$	Average infectious time in rodents	[7 to 60]	[10]

CA: chosen by authors.;

Source: Prepared by the authors of this study.



Figure 1. People with hantavirus without considering person-to-person transmission.

Source: Prepared by the authors of this study.

increasing the risk perception considering λ_1 and λ_2 values. Another factor to note is that in Figure 4b (unlike Figure 3b), the daily variability in a "sawtooth" does not appear, increasing risk perception. Thus, the higher the risk perception, the lower the number of human infections. Therefore, prevention campaigns play a fundamental role in this problem.

The sawtooth presented in Figure 3b, as opposed to Figure 4b, occurs because risk perception depends mainly on two factors of our model: the rate of resistance to change and people's reaction speed. The latter also depends on the number of human hantavirus infections. Therefore, with a higher rate of resistance to change and lower reaction speed, there is greater variability in risk perception after an increase in human infections. In other words, the higher the risk perception, the "smoother" the curve.

Figure 2. Rodent dynamics.



After decreasing the rate of resistance to change and increasing reaction speed, leaving $\lambda_1 = 0,0065$ and $\lambda_2 = 1, 5$, a highly significant decrease in the maximum number of contagions is evident (Figure 5a), following the increase in risk perception (Figure 5b). Unlike the other findings associated with risk perception (Figures 3 and 4b), Figure 5b shows that the behavior of the curve is not bell-shaped during the described period.

DISCUSSION

In this paper, we presented a model that expresses the dynamics of hantavirus transmission, including the possibility of personto-person transmission and the general population's risk perception, showing how the latter variable, associated with



Source: Prepared by the authors of this study.



Figure 3. Hantavirus cases and risk perception, with $\lambda_1 = 0,65$ and $\lambda_2 = 0,05$.

Resistance to change (λ_1) equal to 0.65. Reaction speed (λ_2) equal to 0.05. Source: Prepared by the authors of this study.

people's behavior, is a determining factor in the spread of the disease. The scarce available information on person-to-person transmission and the model's generality are the main limitations of this study. However, the model and its respective graphs associated with cases of human infection (Figures 1 and 3–5) show that surveillance and prevention strategies should be increased considering the possibility of a virus mutation. Suppose person-to-person transmission occurs and the number of infected people increases significantly. In that case, the risk of crossing infectious disease routes (such as the reported cases of COVID-19 and hantavirus) may increase [37]. Considering the high lethality rate of hantavirus, the population would be at serious risk in areas with rodents hosting the virus [38].

Previous studies have confirmed person-to-person transmission of hantavirus, particularly of the Andes virus [4,8,9]. A study in Argentina [9] estimated that the basic reproductive number was 2.12 before the quarantine was announced in the outbreak locality. Therefore, prevention and constant monitoring measures should be taken in the face of a possible virus mutation [39]. Otherwise, a new pandemic may occur, such as those already experienced by SARS-CoV-2 and H1N1 [38].

Climate change, the accelerated expansion of agribusiness in natural reserves, the increase in temperatures, and anthropic disasters such as forest fires, produce the migration of wild rodents following the scarcity of food that occurs in their



Figure 4. Hantavirus cases and risk perception, with $\lambda_1 = 0.065$ and $\lambda_2 = 0.5$.

Resistance to change (λ_1) is equal to 0.065. Reaction speed (λ_2) is equal to 0.5. Source: Prepared by the authors of this study.



Figure 5. Hantavirus cases and risk perception, with $\lambda_1 = 0.0065$ and $\lambda_2 = 1.5$.

Resistance to change (λ_1) is equal to 0.0065. Reaction speed (λ_2) is equal to 1.5. Source: Prepared by the authors of this study.

natural habitat [40,41]. Therefore, their migration may increase the risk of rodent-to-human transmission [42]. .

On the other hand, a study conducted in rural Chilean communities [43] evaluated the risk perception of hantavirus and showed that a high percentage of the participants knew about the disease and had heard of rodents as a natural reservoir for the virus. However, less than 50% knew that rodents and their droppings transmit the disease, and less than half of the participants mentioned fever as the main symptom. In this same study [43], it was noted that television had the highest communicational impact, which may be due to low connectivity and internet access in rural areas. Participants knew that hantavirus disease could be severe and cause death. They perceived themselves, their families, and communities as at risk of contracting this disease because of their location and occupations. Most fear the disease due to living in small communities where cases and deaths have occurred. In summary, the perception of risk towards the disease was mainly due to the impact caused by hantavirus deaths, with some - but scarce- knowledge of the disease, learned through prevention and risk communication campaigns.

At the beginning of March 2020, the first case of COVID-19 was reported in Chile. Given the significant increase in cases, control measures were taken, mainly by reducing mobility, mandatory use of masks, and physical distancing. The reduction of people's mobility impacted the number of hantavirus cases, as 30 cases were reported that year, representing less than half of cases reported in 2019. In 2021 restriction measures were relaxed, giving greater freedom in people's mobility. This freedom translated into 13 cases only for the first eleven epidemiological weeks, between January and March 2021 [6].

Although a series of vaccine and antiviral treatments have been developed worldwide and tested in animal models and human

clinical trials for hantavirus, there are no vaccines or treatments officially approved by international organizations [44].

Ongoing prevention campaigns and effective risk communication strategies are essential for controlling or reducing hantavirus infections in humans. These should involve community participation and be attentive to the local and cultural realities of the potentially affected populations [43,45]. Basic research from universities and infectious disease research centers should be strengthened, focusing on implementing new strategies for viral genomic epidemiological surveillance of natural reservoirs. Multidisciplinary work, including epidemiologists, ecologists, physicians, immunologists, behavioral science researchers, should be enhanced to identify infectious sources of hantavirus [39]. The decrease in cases after increasing the perception of risk from the proposed mathematical model highlights the importance of these measures. Studies show a significant decrease in the number of cases in Chile during the periods of greater prevention campaigns through various media, posters in clinics, and educational establishments [4,26,27].

CONCLUSION

In this paper, we have presented a mathematical SEIR model to express the dynamics of hantavirus infection. We have included the possibility of person-to-person transmission and the risk perception of the population toward the disease.

The model results, via numerical simulations, were elaborated from the scarce data on person-to-person transmission and, mainly, from the information provided by the Chilean Ministry of Health, among other studies.

Through the simulations, we observed that the risk perception determines a sensitivity in the peak of the infectious curve after the variation of this latter curve. Therefore, it plays a fundamental role in the spread of the disease.

Increasing risk perception in the general population through mitigation measures by health authorities is relevant to reducing hantavirus cases and alerting a possible new epidemic outbreak through person-to-person transmission.

Notes

Contributor roles

JPGJ: conceptualization, methodology, formal analysis, research, writing (revisions and edits), data management, data presentation, project management. MTMQ: conceptualization, methodology, formal analysis, research, writing (revisions and edits), supervision.

Acknowledgments

We thank the Vicerrectoría de Investigación y Postgrado (VRIP) and the Centro de Investigación de Estudios Avanzados del Maule (CIEAM), both belonging to the Universidad Católica del Maule.

Competing interests

The authors declare no competing interests for this article.

Funding

The authors declare no external sources of funding.

Provenance and peer review

Not commissioned. Externally peer-reviewed by three reviewers, double-blind.

Language of submission

Spanish.

References

- Kuhn JH, Adkins S, Alioto D, Alkhovsky SV, Amarasinghe GK, Anthony SJ, et al. 2020 taxonomic update for phylum Negarnaviricota (Riboviria: Orthornavirae), including the large orders Bunyavirales and Mononegavirales. Arch Virol. 2020;165: 3023–3072. http://link.springer.com/10.1007/ s00705-020-04731-2 https://doi.org/10.1007/s00705-020-04731-2
- 2. Pan american health organization. Hantavirus [Internet]. 2017. https://www3.paho.org/hq/index.php?option=com_content& view=article&id=14911:hantavirus&Itemid=40721&lang=en
- Filippone C, Castel G, Murri S, Ermonval M, Korva M, Avšič-Županc T, et al. Revisiting the genetic diversity of emerging hantaviruses circulating in Europe using a pan-viral resequencing microarray. Sci Rep. 2019;9. http://www.nature.com/articles/ s41598-019-47508-7 https://doi.org/10.1038/s41598-019-47508-7
- Ministerio de Salud del Gobierno de Chile. Boletín Epidemiológico Trimestral, Hantavirus. 2020. http://epi.minsal. cl/wp-content/uploads/2020/02/BET_HANTAVIRUS_2019. pdf

- Sabino-Santos G Jr, Maia FGM, Martins RB, Gagliardi TB, Souza WM de, Muylaert RL, et al. Natural infection of Neotropical bats with hantavirus in Brazil. Sci Rep. 2018;8. http://www. nature.com/articles/s41598-018-27442-w https://doi.org/10. 1038/s41598-018-27442-w
- 6. Departamento de Epidemiología. Síndrome Cardiopulmonar por Hantavirus. Ministerio de Salud de Chile; 2021. http://epi. minsal.cl/hantavirus-materiales-relacionados/
- Reyes Zaldívar FT, Ferrés M. Hantavirus: Descripción de dos décadas de endemia y su letalidad. ARS med. 2019;44: 30–39. https://173.236.243.65/index.php/MED/issue/view/44-1-2019 https://doi.org/10.11565/arsmed.v44i1.1522
- Alonso DO, Pérez-Sautu U, Bellomo CM, Prieto K, Iglesias A, Coelho R, et al. Person-to-Person Transmission of Andes Virus in Hantavirus Pulmonary Syndrome, Argentina, 2014. Emerg Infect Dis. 2020;26: 756–759. http://wwwnc.cdc.gov/eid/ article/26/4/19-0799_article.htm https://doi.org/10.3201/ eid2604.190799
- Martínez VP, Di Paola N, Alonso DO, Pérez-Sautu U, Bellomo CM, Iglesias AA, et al. "Super-Spreaders" and Personto-Person Transmission of Andes Virus in Argentina. N Engl J Med. 2020;383: 2230–2241. http://www.nejm.org/doi/ 10.1056/NEJMoa2009040 https://doi.org/10.1056/ NEJMoa2009040
- Allen LJS, McCormack RK, Jonsson CB. Mathematical models for hantavirus infection in rodents. Bull Math Biol. 2006;68: 511–24. http://link.springer.com/10.1007/s11538-005-9034-4 https:// doi.org/10.1007/s11538-005-9034-4
- Abramson G. Mathematical modelling of hantavirus: From the mean field to the individual level. Progress in Mathematical Biology Research; 2008. https://www.researchgate.net/publication/ 228958845_Mathematical_modelling_of_Hantavirus_From_ the_mean_field_to_the_individual_level
- Bürger R, Chowell G, Gavilán E, Mulet P, Villada LM. Numerical solution of a spatio-temporal gender-structured model for hantavirus infection in rodents. Math Biosci Eng. 2018;15: 95–123. http://www.aimsciences.org/journals/displayArticlesnew.jsp? paperID=14138 https://doi.org/10.3934/mbe.2018004
- Gedeon T, Bodelón C, Kuenzi A. Hantavirus transmission in sylvan and peridomestic environments. Bull Math Biol. 2010;72: 541–64. http://link.springer.com/10.1007/s11538-009-9460-4 https://doi.org/10.1007/s11538-009-9460-4
- Liu JL. Threshold dynamics of a time-delayed hantavirus infection model in periodic environments. Math Biosci Eng. 2019;16: 4758–4776. http://www.aimspress.com/article/10. 3934/mbe.2019239 https://doi.org/10.3934/mbe.2019239
- Xiao Y, Zhang Y, Gao M. Modeling hantavirus infections in mainland China. Applied Mathematics and Computation. 2019;360: 28–41. https://linkinghub.elsevier.com/retrieve/pii/ S0096300319303960 https://doi.org/10.1016/j.amc.2019.05. 009
- Wesley CL, Allen LJS, Langlais M. Models for the spread and persistence of hantavirus infection in rodents with direct and indirect transmission. Math Biosci Eng. 2010;7: 195–211. http:// www.aimspress.com/article/10.3934/mbe.2010.7.195 https:// doi.org/10.3934/mbe.2010.7.195
- Sauvage F, Langlais M, Pontier D. Predicting the emergence of human hantavirus disease using a combination of viral dynamics and rodent demographic patterns. Epidemiol Infect. 2007;135: 46–56. https://www.cambridge.org/core/product/ identifier/S0950268806006595/type/journal_article https:// doi.org/10.1017/S0950268806006595
- Nsoesie EO, Mekaru SR, Ramakrishnan N, Marathe MV, Brownstein JS. Modeling to predict cases of hantavirus pulmonary syndrome in Chile. Bausch DG, editor. PLoS Negl Trop Dis.

2014;8. https://dx.plos.org/10.1371/journal.pntd.0002779 https://doi.org/10.1371/journal.pntd.0002779

- Kermack WO, McKendrick AG. A Contribution to the Mathematical Theory of Epidemics. Proc R Soc A Math Phys Eng Sci. 1927;1: 700–21. http://rspa.royalsocietypublishing.org/ cgi/doi/10.1098/rspa.1927.0118
- Kermack WO, McKendrick AG. II. The Problem of Endemicity. Proc R Soc A Math Phys Eng Sci [Internet]. 1932;1: 55–83. http://rspa.royalsocietypublishing.org/cgi/doi/10.1098/rspa. 1932.0171 https://doi.org/10.1098/rspa.1932.0171
- Kermack WO, McKendrick AG. III. Further Studies of the Problem of Endemicity. Proc R Soc A Math Phys Eng Sci. 1933; 94–122. http://rspa.royalsocietypublishing.org/cgi/doi/ 10.1098/rspa.1933.0106 https://doi.org/10.1098/rspa.1933. 0106
- 22. Córdova-Lepe FD, Gutiérrez Jara JP, Muñoz Quezada MT. An index to quantify individual social responsibility in the decision to be vaccinated. Medwave. 2018;18: e7237. http://www.medwave. cl/link.cgi/Medwave/Revisiones/MetodInvestReport/7237 https://doi.org/10.5867/medwave.2018.04.7237
- Córdova-Lepe F, Cabrera Hernández M, Gutiérrez-Jara JP. Modeling the epidemiological impact of a preventive behavioral group. Medwave. 2018;18: e7396–e7396. http://www.medwave. cl/link.cgi/Medwave/Revisiones/MetodInvestReport/7396 https://doi.org/10.5867/medwave.2018.08.7396
- Cabrera M, Córdova-Lepe F, Gutiérrez-Jara JP, Vogt-Geisse K. An SIR-type epidemiological model that integrates social distancing as a dynamic law based on point prevalence and socio-behavioral factors. Sci Rep. 2021;11. http://www.nature.com/articles/ s41598-021-89492-x https://doi.org/10.1038/s41598-021-89492-x
- Muñoz-Quezada MT, Lucero BA, Gutiérrez-Jara JP, Buralli RJ, Zúñiga-Venegas L, Muñoz MP, et al. Longitudinal exposure to pyrethroids (3-PBA and trans-DCCA) and 2,4-D herbicide in rural schoolchildren of Maule region, Chile. Sci Total Environ. 2020;749: 141512. https://linkinghub.elsevier.com/retrieve/pii/ S0048969720350415 https://doi.org/10.1016/j.scitotenv.2020. 141512
- Tercera L. Ministerio de Salud lanza campaña contra el hantavirus. 2008. https://www.latercera.com/noticia/ministerio-de-saludlanza-campana-contra-el-hantavirus/
- Ministerio de Salud del Gobierno de Chile. Guía clínica de prevención, diagnóstico y tratamiento del Síndrome Cardiopilmonar por Hantavirus. 2013. http://epi.minsal.cl/wpcontent/uploads/2016/03/guia_hantavirus.pdf
- Paek H-J, Hove T. Risk Perceptions and Risk Characteristics. Oxford University Press: In: Oxford Research Encyclopedia of Communication [Internet]; 2017. https://oxfordre.com/ communication/view/10.1093/acrefore/9780190228613.001. 0001/acrefore-9780190228613-e-283 https://doi.org/10.1093/ acrefore/9780190228613.013.283
- Ross R. Some Quantitative Studies in Epidemiology. Nature. 1911;87: 466–467. http://www.nature.com/articles/087466a0 https://doi.org/10.1038/087466a0
- MacDonald G. The Measurement of Malaria Transmission. Proceedings of the Royal Society of Medicine. 2016;48: 295–302. http://journals.sagepub.com/doi/10. 1177/003591575504800409 https://doi.org/10.1177/ 003591575504800409
- Simoy MI, Aparicio JP. Ross-Macdonald models: Which one should we use? Acta Trop. 2020;207: 105452. https://linkinghub. elsevier.com/retrieve/pii/S0001706X19313610 https://doi.org/ 10.1016/j.actatropica.2020.105452
- 32. van den Driessche P, Watmough J. Reproduction numbers and sub-threshold endemic equilibria for compartmental models

of disease transmission. Math Biosci. 2002;180: 29–48. http://linkinghub.elsevier.com/retrieve/pii/S0025556402001086 https://doi.org/10.1016/s0025-5564(02)00108-6

- 33. Gutiérrez-Jara JP, Córdova-Lepe FD, Muñoz-Quezada MT. Dynamics between infectious diseases with two susceptibility conditions: A mathematical model. Math Biosci. 2019;309: 66–77. https://linkinghub.elsevier.com/ retrieve/pii/S0025556418305455 https://doi.org/10.1016/j. mbs.2019.01.005
- Instituto Nacional de Estadísticas, Gobierno de Chile. Estadísticas vitales. 2018. https://www.ine.cl/estadisticas/ sociales/demografia-y-vitales/nacimientos-matrimonios-ydefunciones
- 35. Boada-Cuerva M, Boada-Grau J, Prizmic-Kuzmica AJ, De Diego NG, Vigil-Colet A. RTC-11: Adaptación de la escala de resistencia al cambio en dos países (España y Argentina). AN PSICOL-SPAIN. 2018;34: 360. http://revistas.um.es/ analesps/article/view/analesps.34.2.286721 https://doi.org/10. 6018/analesps.34.2.286721
- Organización de las Naciones Unidas para la Alimentación y la Agricultura. Biología general de los roedores. http://www.fao. org/3/x5052s/x5052S03.htm#3
- 37. Wilson TM, Paddock CD, Reagan-Steiner S, Bhatnagar J, Martines RB, Wiens AL, et al. Intersecting Paths of Emerging and Reemerging Infectious Diseases. Emerg Infect Dis. 2021;27: 1517–1519. https://wwwnc.cdc.gov/eid/article/27/5/ 20-4779_article.htm https://doi.org/10.3201/eid2705.204779
- Khan A, Khan M, Ullah S, Wei D-Q. Hantavirus: The Next Pandemic We Are Waiting For? Interdiscip Sci. 2021;13, 1:147–152. http://link.springer.com/10.1007/s12539-020-00413-4 https:// doi.org/10.1007/s12539-020-00413-4
- 39. Kim W-K, Cho S, Lee S-H, No JS, Lee G-Y, Park K, et al. Genomic Epidemiology and Active Surveillance to Investigate Outbreaks of Hantaviruses. Front Cell Infect Microbiol. 2020;10. https:// www.frontiersin.org/articles/10.3389/fcimb.2020.532388/full https://doi.org/10.3389/fcimb.2020.532388
- Prist PR, Uriarte M, Fernandes K, Metzger JP. Climate change and sugarcane expansion increase Hantavirus infection risk. Munoz-Zanzi C, editor. PLoS Negl Trop Dis. 2017;11. https:// dx.plos.org/10.1371/journal.pntd.0005705 https://doi.org/10. 1371/journal.pntd.0005705
- Carbajo AE, Vera C, González PL. Hantavirus reservoir Oligoryzomys longicaudatus spatial distribution sensitivity to climate change scenarios in Argentine Patagonia. Int J Health Geogr. 2009;8. http://ij-healthgeographics.biomedcentral.com/ articles/10.1186/1476-072X-8-44 https://doi.org/10.1186/ 1476-072X-8-44
- 42. Watson DC, Sargianou M, Papa A, Chra P, Starakis I, Panos G. Epidemiology of Hantavirus infections in humans: A comprehensive, global overview. Crit Rev Microbiol. 2014;40: 261–72. http://www.tandfonline.com/doi/full/ 10.3109/1040841X.2013.783555 https://doi.org/10.3109/ 1040841X.2013.783555
- 43. Valdivieso F, Gonzalez C, Najera M, Olea A, Cuiza A, Aguilera X, et al. Knowledge, attitudes, and practices regarding hantavirus disease and acceptance of a vaccine trial in rural communities of southern Chile. Hum Vaccin Immunother. 2017;13: 808–815. https://www.tandfonline.com/doi/full/10.1080/21645515. 2016.1250989 https://doi.org/10.1080/21645515.2016.1250989
- 44. Brocato RL, Hooper JW. Progress on the Prevention and Treatment of Hantavirus Disease. Viruses. 2019;11. https:// www.mdpi.com/1999-4915/11/7/610 https://doi.org/10. 3390/v11070610
- 45. Harris C, Armién B. Sociocultural determinants of adoption of preventive practices for hantavirus: A knowledge, attitudes, and

REVIEWS

practices survey in Tonosí, Panama. Jonsson CB, editor. PLoS

Negl Trop Dis. 2020;14. https://dx.plos.org/10.1371/journal. pntd.0008111 https://doi.org/10.1371/journal.pntd.0008111

Modelamiento del síndrome cardiopulmonar por hantavirus

Resumen

Introducción

El síndrome cardiopulmonar por hantavirus es una enfermedad causada por un virus perteneciente al orden bunyanvirales, y transmitida hacia los humanos a través de roedores. Esta enfermedad en Chile es considerada endémica, la cual tiene una alta tasa de letalidad. En la actualidad existen estudios que evidencian el contagio entre personas del virus Andes, cuya localidad se concentra en los países de Argentina y Chile.

Objetivos

Analizar la posibilidad de transmisión de hantavirus entre humanos, mediante un modelo matemático tipo SEIR.

Métodos

Se plantea un modelo matemático tipo SEIR (susceptible, expuesto, infeccioso y recuperado) para expresar la dinámica de la enfermedad por hantavirus, incluyendo la posibilidad de transmisión entre humanos y la percepción del riesgo.

Resultados

El máximo de contagio entre humanos disminuye cerca de 25% tras aumentar la percepción de riesgo de las personas, mediante la reducción de la tasa de resistencia al cambio y aumento la velocidad de reaccionar de las personas.

Conclusiones

Es urgente revisar las estrategias de comunicación de riesgo y medidas de prevención ante esta posibilidad de contagios masivos entre humanos, además de fortalecer la investigación y proyectar el desarrollo de una vacuna para proteger las poblaciones expuestas a esta enfermedad con alta tasa de letalidad.



This work is licensed under a Creative Commons Attribution 4.0 International License.