JAPANESE ENCEPHALITIS VIRUS



The mission of the Swine Health Information Center is to protect and enhance the health of the United States swine herd through coordinated global disease monitoring, targeted research investments that minimize the impact of future disease threats, and analysis of swine health data.

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SUMMARY

IMPORTANCE

- Japanese encephalitis virus (JEV) is an arbovirus that causes disease in humans throughout Asia and the western Pacific region. The virus also produces neurological disease in pigs and horses and reproductive failure in pigs.
- Japanese encephalitis (JE) is currently a foreign animal disease, but its emergence in the United States is a concern due to the presence of competent mosquito vectors and susceptible hosts.

PUBLIC HEALTH

- JEV is a zoonotic pathogen. It causes an estimated 70,000 cases of viral encephalitis annually, with a 20– 30% mortality rate. However, humans are incidental (dead-end) hosts.
- Amplification of JEV in swine often precedes human epidemics.

INFECTION IN SWINE

- Most JEV infections are subclinical in pigs.
- In sows, infection with JEV before 60–70 days of gestation can cause abortion, fetal mummification or stillbirth, or the birth of weak piglets. Infected boars can have reduced sperm motility and develop testicular edema and temporary infertility.
- Piglets born alive may develop encephalitis, displaying tremors and convulsions.

TREATMENT

• There is no available treatment for JEV infection in swine.

CLEANING AND DISINFECTION

- JEV does not survive well in the environment. The virus is inactivated at 40°C (104°F).
- JEV is inactivated by 1% hypochlorite, 3–8% paraformaldehyde, 2% glutaraldehyde, mM binary ethylenimine, ethanol, and isopropanol. JEV is also sensitive to ultraviolet light.⁴

PREVENTION AND CONTROL

- JE is reportable to the World Organization for Animal Health (OIE).
- Mosquito control is essential for the prevention of JE. Recommended practices include removing stagnant water around pig enclosures, using insecticides, installing insect screens, and using fans inside buildings where pigs are housed to disrupt mosquito activity.
- Swine vaccination is practiced in some areas (see *Immunology*).

TRANSMISSION

- JEV is transmitted by mosquitoes. One well-known vector is *Culex tritaeniorhynchus*. Wading ardeid birds (herons and egrets) are the primary natural reservoirs of the virus. Pigs act as amplifying hosts.
- Direct pig-to-pig transmission has been documented experimentally. JEV persists in the tonsils.
- Artificial insemination practices could lead to JEV spread since infected boars shed virus in the semen.

PATHOGENESIS

 JEV replicates in the skin (keratinocytes) and is transferred to the lymph nodes within dendritic cells. Then, virus disseminates to tissues including the brain via the blood and nerves. Replication occurs in the central nervous system and lymphoid tissues, especially the tonsils.

DIAGNOSIS

- Virus isolation is the reference standard. Molecular methods described for JEV include conventional and nested reverse transcriptase polymerase chain reaction (RT-PCR), quantitative RT-PCR (qRT-PCR, TaqMan and SYBR green), multiplex ligase detection—polymerase chain reaction and microarray (MLPM), multiplex ligation-dependent probe amplification (MLPA), reverse transcription loop-mediated isothermal amplification (RT-LAMP), and antigen-capture enzyme-linked immunosorbent assay (ELISA). RNA has been recovered from oronasal swabs and oral fluids experimentally.
- Serological tests include virus neutralization, hemagglutination inhibition, complement fixation, IgMcapture ELISA, and indirect IgG and IgM ELISA. Tests with potential for on-site use include latex agglutination, lateral flow assay, and protein-based dipstick IgG ELISA.

EPIDEMIOLOGY

- Pigs are the main amplifying host of JEV. In addition to humans, horses can become ill. Most other susceptible species are dead-end hosts and do not transmit JEV.
- JEV is endemic in much of Asia and the western Pacific region.

ETIOLOGY

- JEV belongs to the family *Flaviviridae*. Five genotypes have been described (GI–GV); GI-b is now the dominant JEV genotype throughout Asia, replacing GII and GIII strains.
- The E, prM, and NS1 proteins are immunogenic.

HISTORY IN SWINE

Human JE epidemics have been described in Japan since the 19th century. The pig-mosquito transmission cycle was confirmed in the late 1960s.

IMMUNITY

- Natural infection results in long lasting immunity in adult pigs and surviving piglets.
- Vaccines are based on GIII, but this genotype has been replaced by GI, leading to concern about efficacy.
- Some cross-protection is likely between strains and other flaviviruses.

GAPS IN PREPAREDNESS

- Introduction of JEV by an infected adult mosquito (likely via aircraft) is thought to be the highest risk for the United States. JEV could spread quickly and become endemic similar to West Nile virus.
- Although pig-to-pig transmission of JEV has been described, its role in natural transmission is unclear.
- Continued research on the cross-protective characteristics of flaviviruses and effective vaccines is needed.

LITERATURE REVIEW: JAPANESE ENCEPHALITIS VIRUS

IMPORTANCE

Japanese encephalitis virus (JEV) is an arbovirus that causes encephalitis in humans throughout Asia and the western Pacific region. The virus also produces neurological disease in pigs and horses and reproductive failure in pigs. Japanese encephalitis (JE) is currently a foreign animal disease, but its emergence in the United States is a concern due to the presence of competent mosquito vectors and susceptible hosts.

PUBLIC HEALTH

JEV is a zoonotic pathogen that cycles between mosquitoes, domestic and wild birds, domestic and feral pigs, and humans.¹ JEV is the greatest known cause of epidemic viral encephalitis worldwide,² and more than one-third of the world population is at risk.³ JEV predominantly infects children and travelers from non-endemic areas who have not developed immunity to the virus.¹ An estimated 70,000 human cases of JEV occur annually, with a 20–30% mortality rate.⁴ Up to 50% of survivors develop long-term neurological sequelae.⁴ Humans are considered dead-end hosts.¹ General awareness of JEV may be low in developing areas; in Cambodia, for instance, very few households knew that JEV is transmitted to pigs and people through mosquito bites.⁵

Amplification of JEV in swine often precedes human epidemics.^{6, 7} Human risk is influenced by the density, size, and spatial organization of rice paddies, swine farms, and human communities.^{2, 8} JEV is not only a rural disease. Confirmed circulation in peri-urban environments exists in Cambodia,^{9, 10} where identical JEV strains have been found in humans, pigs, and mosquitoes (*Culex tritaeniorhynchus*).¹¹ Dogs have been proposed as JEV sentinels in peri-urban and rural environments since they live with or near humans; they develop antibodies but do not transmit the virus to humans.¹²

Some studies suggest that proximity to pigs is related to increased JEV infection. Mosquitoes living near farm households with pigsties are more likely to be infected with JEV compared to those living near farm households without pigsties or intensive pig farms located 2 km away.¹³ Similarly, JEV prevalence in mosquitoes is highest in traps located close to pigs in urban Vietnam.¹⁴ In Hong Kong, absence of human cases from 2006–2010 was linked to a sudden reduction in the number of pigs, while a 2011 resurgence was related to a new strain.¹⁵ Despite this evidence, several recent studies have investigated JEV transmission in areas with little to no pig farming. In these regions, the existence of a secondary reservoir could explain JEV transmission.¹⁶ Examples include:

- Singapore: pig farming was abolished in 1992 and afterward, transmission of JEV nearly ceased. However, serosurveillance has since detected anti-JEV antibodies in migratory birds (2010–2017), wild boar (2014), and sentinel chickens (2013–2018).¹⁷
- Goa, India: despite few human JE cases in this region, about 7% of pigs were seropositive.¹⁸ The highest prevalence in swine occurred in the post-monsoon season,¹⁹ which correlated with a previous study showing that mosquito populations were high at that time.²⁰

Laboratory-acquired JEV infection has been reported in humans, and work with the virus is restricted to BSL-3 facilities and practices. Transmission occurs through needle sticks and potentially at mucosal surfaces if exposed to high concentrations of aerosolized virus.²¹ Transmission of JEV via blood transfusion was reported in two patients for the first time in 2018.²²

INFECTION IN SWINE

CLINICAL SIGNS

JEV mainly causes subclinical infection in pigs. Reproductive failure is the most common clinical manifestation. Infection with JEV before 60–70 days of gestation can cause abortion, fetal mummification or stillbirth, or the birth of weak piglets.⁴ In endemic areas, the breeding impact may be greatest in boars. Infected boars can

experience reduced numbers and motility of sperm.⁴ Testicular edema and temporary infertility in boars have also been observed.⁴ Piglets born alive may develop encephalitis, displaying tremors and convulsions.⁴

An experimental study inoculated 3-week-old domestic pigs (white-line crossbred) intravenously with GI-b, an emerging genotype (see *Etiology*). Non-specific clinical illness occurred (fever, weight loss, lethargy, depression) but resolved in one week. Mild ataxia was also seen in a few pigs.²³

No characteristic lesions are seen in sows. Lesions in boars may include thickening of the tunica vaginalis and epididymis and edema and inflammation of the epididymis, tunica vaginalis, and testis.⁴ Affected piglets develop hydrocephalus, cerebellar hypoplasia, subcutaneous edema, and spinal hypomyelinogenesis.⁴ Multifocal hepatic and splenic necrosis may also be seen grossly, while histological lesions are generally restricted to the central nervous system.^{4, 24} Brain and spinal cord changes can include serosal petechiae, bicavitary effusions, neuronophagia, glial nodules, and perivascular cuffing.⁴

TREATMENT

There is no available treatment for JEV infection in swine. Anti-flavivirus drugs have been reviewed by Pierson and Diamond.²⁵

CLEANING AND DISINFECTION

SURVIVAL

JEV does not survive well in the environment. The virus is inactivated at 40°C (104°F) and can be destroyed by heating for 30 min at 56°C (132.8°F). JEV is stable in alkaline environments (pH 7–9) and inactive in acidic environments (pH 1–3).⁴

DISINFECTION

JEV is inactivated by 1% hypochlorite, 3–8% paraformaldehyde, 2% glutaraldehyde, mM binary ethylenimine, ethanol, and isopropanol. JEV is sensitive to ultraviolet light.⁴

PREVENTION AND CONTROL

DISEASE REPORTING

JE is an OIE-listed disease and must be reported internationally according to the *Terrestrial Animal Health Code*. Any suspicious clinical or necropsy findings should always be reported to the USDA and your State Animal Health Official.

DISEASE PREVENTION AND CONTROL

Mosquito control is essential for the prevention of JE. Recommended practices include removing stagnant water around pig enclosures, using insecticides, installing insect screens, and using fans inside buildings where pigs are housed to disrupt mosquito activity.²⁶ However, the discovery of direct pig-to-pig transmission shows that JEV may spread and circulate even in areas without competent mosquito vectors.^{27, 28}

The decline of JEV in endemic areas has been attributed to large-scale human immunization, immunization of pigs (see *Immunology*), separation of pigs from human communities, alterations in agricultural practices to reduce the presence of mosquitoes, and better overall standards of living.²⁹

TRANSMISSION

JEV is usually transmitted by mosquitoes. Wading ardeid birds (such as herons and egrets) are the primary natural reservoirs of the virus. Pigs act as amplifying hosts, developing high viremia that lasts for 2–4 days.^{4, 29} Additional proposed amplifying hosts include orangutans (in Borneo) and bats.³⁰ Seropositive raccoons, raccoon dogs, and

flying foxes have also been identified.² Many species are most likely incidental (dead-end) hosts, including horses, cattle, buffalo, goats, sheep, dogs, and humans.⁴

Culex tritaeniorhynchus, found in rice fields, wells, ponds, ditches, and urban environments, is the primary JEV vector in Asia.³¹ Others include *Cx. fuscocephala, Cx. annulirostris*,⁶ *Cx. annulus, Cx. sitiens*, and potentially *Cx. quinquefasciatus*.² Important regional vectors include *Cx. vishnui* in India and *Cx. gelidus* in Indonesia.³⁰ *Anopheles sinensis* and *Aedes albopictus* are also vectors,³² along with several *Armigeres* and *Mansonia* species.³³

American mosquitoes exposed to JEV are susceptible to infection.^{34, 35} Some natural JEV vectors (*Cx. pipiens, Cx. quinquefasciatus,* and *Ae. japonicus*)³⁶ are also vectors of West Nile virus in the United States.³⁷ Environmental temperatures in the range of 22–34°C (71.6–93.2°F) have been shown to increase mosquito density,³² decrease larval development time, and reduce the extrinsic incubation period of the virus in mosquito vectors, affecting potential JEV transmission.³⁸

Vector-free JEV transmission has been documented between pigs by Ricklin et al.²⁷

- Two sentinel pigs were placed in contact with three pigs that had been infected intravenously with JEV.
 Viremia was detected by quantitative reverse transcription polymerase chain reaction (qRT-PCR) in both contact pigs for 2–4 days, but at a lower level than in needle-infected animals.²⁷
- Six sentinel pigs were housed with two needle-infected pigs. Viremia was detected by qRT-PCR in only one contact pig and lasted for three days; values were similar to those found in needle-infected pigs.²⁷
- Nine pigs were oronasally inoculated with three different doses of JEV (10³, 10⁵, or 10⁷ TCID₅₀). By qRT-PCR, viremia was detected in all groups for about four days. JEV RNA was also found in the majority of oronasal swabs between 3–6 days post-infection.²⁷
- Nine pigs were oronasally inoculated with three different doses of JEV (10¹, 10², or 10³ TCID₅₀). All pigs became infected and were viremic for 5–6 days. Viral RNA was detected in oronasal swabs as early as three days post-infection; by seven days post-infection, all swabs were positive.²⁷

Through all experiments, the highest levels of viral RNA were consistently found in the tonsil, and persistence was documented for at least 25 days.²⁷ A further study confirmed that high viral RNA loads are found in the tonsils and that JEV can persist for up to 46 days.²⁸ Isolation of JEV RNA from oronasal swabs was also replicated by further work. Three 5-week-old control pigs challenged with JEV GIII shed virus oronasally from 4–10 days post-infection. Low amounts of JEV RNA were also detected in oronasal swabs from pigs infected with a GI strain or vaccinated with TRIP/JEV.prME (see *Immunology*).²⁸

A more recent experimental study found that viremia developed at three dpi (following intravenous inoculation with GI-b), neutralizing antibodies developed by 28 dpi, and virus was shed in nasal secretions. Similar to other JEV strains, tissue tropism for nervous and lymphoid tissues was seen. ²³Another experimental study found that nasal epithelium could be a route of entry for JEV in pigs.³⁹ Aerosol transmission has been demonstrated experimentally in mice.⁴⁰

Artificial insemination practices in modern swine production could potentially lead to JEV transmission. JEV is known to cause inflammation of the genital tract in boars, which can lead to subsequent shedding of the virus in semen and possible transmission to breeding sows.⁴¹

PATHOGENESIS

Following a bite from an infected mosquito, JEV replicates in the skin (keratinocytes) and is then transferred to the lymph nodes within dendritic cells.⁴ From there, virus disseminates to many different tissues, including the brain via the blood and nerves.⁴² Replication occurs in the CNS and lymphoid tissues, especially tonsil, where

viral persistence has been documented for up to 46 days.^{27, 28, 43} In immunocompromised hosts, cell and tissue tropism and viral replication are enhanced.⁴

DIAGNOSIS

TESTS TO DETECT NUCLEIC ACIDS, VIRUS, OR ANTIGENS

Virus isolation is the reference standard for definitive diagnosis. JEV can be grown in 2–4-day-old mice, chicken embryos, or cell lines, including African green monkey kidney (Vero) cells, baby hamster kidney (BHK) cells, porcine kidney (PSEK) cells, and the *Aedes albopictus* mosquito cell line C6/36.⁴ A pig tonsil epithelial (PT) cell line has recently been established for JEV research.⁴⁴ Indirect immunofluorescence can be used to identify viral antigen using JEV-specific monoclonal antibodies or monoclonal antibodies for flaviviruses in general. Isolated viruses can also be identified by RT-PCR or serological methods.

RT-PCR is routinely used in diagnostic laboratories,³¹ but methods have been hampered by the short period of viremia and low sensitivity.⁴⁵ Numerous variations have been described for conventional and nested RT-PCR and qRT-PCR (TaqMan, SYBR green) as reviewed by Bharucha and colleagues.⁴⁵ For swine, multiplex RT-PCR assays have been developed for JEV and other swine pathogens, including pseudorabies virus (PRV), porcine parvovirus (PPV), porcine circovirus type 2 (PCV2), porcine reproductive and respiratory syndrome virus (PRRSV), classical swine fever virus (CSFV), *Actinobacillus pleuropneumoniae*, and *Mycoplasma hyopneumoniae*.⁴⁶⁻⁵²

Additional molecular techniques have been described for JEV, including:

- Multiplex ligase detection-polymerase chain reaction and microarray (MLPM) for detection of PCV, PRRSV, CSFV, PPV, PRV, and JEV⁵³
- Multiplex ligation-dependent probe amplification (MLPA) for detection of PCV, PRRSV, CSFV, PPV, PRV, and JEV⁵⁴
- Droplet digital PCR (ddPCR, JEV only)⁵⁵
- Reverse transcription loop-mediated isothermal amplification (RT-LAMP) detecting the NS1 gene,⁵⁶ NS3 gene,⁵⁷ E gene,⁵⁸ C gene^{59, 60}
- RT-LAMP plus lateral flow dipstick (LFD) to detect multiple JEV strains⁶¹

RT-LAMP assays are highly specific and do not cross-react with other clinically and/or serologically related swine diseases.^{56-58, 61} Given the efficacy, simplicity, and speed of these assays, they are an option for on-site testing in an outbreak situation.

Flavivirus-specific monoclonal antibodies (MAb) can detect JEV antigen in serum, and immunohistochemistry can be used to detect JEV antigen in fetal tissues.⁴ Antigen capture enzyme-linked immunosorbent assay (Ag-ELISA) has been developed for simple and rapid clinical detection of JEV infection as an alternative to virus isolation and RT-PCR. MAb against the E protein is used for its high specificity to JEV, followed with a polyclonal antibody (PcAb) conjugate to enhance sensitivity and detection of multiple strains. Ag-ELISA may be unable to detect very low viral loads.⁶²

TESTS TO DETECT ANTIBODY

Serological tests can be used to determine JEV prevalence in a population or to make a diagnosis in a diseased individual. Paired samples must be used to demonstrate a rise in titer from the acute to convalescent phases of disease.⁶³ The OIE recommends that diagnosis by serology be confirmed by virus isolation.

Serological methods for JEV include:

- Virus neutralization (VN), also known as the plaque reduction neutralization test (PRNT), is difficult and time-consuming, but it is the most specific serological test for JEV. VN is the gold standard for JEV diagnosis.⁶³
- Hemagglutination inhibition (HI) is widely used despite its cross-reactivity with other flaviviruses. A four-fold rise in titer is sufficient for diagnosis of flavivirus infection, though not necessarily JEV.⁶³
- An immunochromatographic test has been described that is faster and more efficient than HI. JEV E
 protein is expressed in soluble form in a bacterial expression system, eliminating the need for erythrocytes
 and mouse brain-derived JEV antigen.⁶⁴
- IgM capture-ELISA (MAC-ELISA) detecting anti-NS1 antibody is used for detection of recent JEV infection (within 2–3 days and for up to three weeks post-infection).⁴ Indirect IgG ELISA⁶⁵ and IgM ELISA⁶⁶ have been described using recombinant NS1 protein as antigen, as well as a mAb-based blocking ELISA.⁶⁷
- Complement fixation (CF) is occasionally used for diagnosis by combining test serum, antigen extracted from inoculated mouse brains with acetone/ether, and pooled fresh guinea pig serum (complement). A fourfold change in titer is required for diagnosis.⁶³
- Indirect immunofluorescence has been used to detect either anti-JEV IgG or IgM in humans, with similar sensitivity and specificity to existing serological methods.⁶⁸

The interpretation of serological results, especially in older pigs, must consider vaccination history and age since maternal antibody can persist for up to eight weeks.⁴ There is some level of cross-reactivity with other flaviviruses for all JEV serological tests. Epitope blocking ELISAs can help to determine if cross-reactions have occurred (i.e., presence of other viruses from the JEV serogroup).³

Experimentally, protein biochip technology has been described that simultaneously detects antibodies to CSFV (E2 protein), PPV (VP2 protein), JEV (domain III of the E protein), and PRRSV (N protein).⁶⁹

Tests with potential on-site application must be quick, low cost, and simple (requiring no special equipment or skills). Tests designed with these factors in mind include:

- Latex agglutination using recombinant NS1 protein as antigen^{70, 71}
- Lateral flow assay (LFA) using recombinant NS1 protein as antigen⁷²
- Protein-based dipstick IgG ELISA using recombinant NS1 protein as antigen⁷³

SAMPLES

Infected tissues (brain, spleen, liver, placenta) are preferred for virus isolation.⁴ Four to six fetuses should be sampled to avoid missing a diagnosis.^{4, 24} Samples appropriate for RT-PCR include cerebrospinal fluid, sera, tissue culture supernatants, and tissue homogenates.⁴ Ag-ELISA detects JEV in cerebrospinal fluid, brain tissue, and mosquito homogenate.⁶² Detection of anti-JEV antibodies in serum, thoracic, or abdominal fluid of stillborn piglets is considered diagnostic.²⁴

RNA has been recovered from oronasal swabs (via qRT-PCR) in pigs experimentally infected with JEV as well as contact pigs.²⁷ Further, RNA has been found in oronasal swabs from control pigs challenged with JEV, pigs previously infected with JEV, and pigs inoculated with an experimental vaccine.²⁸ JEV is also found in oral fluids in experimentally infected pigs.⁷⁴

EPIDEMIOLOGY

SPECIES AFFECTED

Wading ardeid birds (such as herons and egrets) are the primary natural reservoirs. Other reservoirs may exist (see *Transmission*). Pigs (domestic and wild boar) are the main amplifying host of JEV, exhibiting high and prolonged viremia and a natural infection rate of 98–100%.³ Sporadic infection occurs in horses, resulting in severe encephalitis, or more commonly, subclinical infection. In horses, the mortality rate is about 5%. Rarely, JEV-associated encephalitis has been described in cattle.^{75, 76} Subclinical infection has also been reported in sheep, goats, dogs, cats, chickens, ducks, wild mammals, reptiles, and amphibians, but these species are not known to contribute to the spread of the disease.

GEOGRAPHIC DISTRIBUTION

JEV is endemic in Asia and the western Pacific region. JEV occurs as far west as Pakistan, as far south as northern Australia and Papua New Guinea,²⁹ as far north as Siberia, and as far east as the island of Saipan.⁷⁷ Locally-acquired JEV and yellow fever virus co-infection was identified in a man from Angola in 2016.⁷⁸

Seasonal precipitation, humidity, and temperature changes are thought to influence JEV transmission by affecting human agricultural practices and the life cycle of mosquito vectors. Higher absolute humidity is a key factor known to influence longevity, mating, dispersal, and feeding behavior of mosquitoes.⁷⁹

In endemic areas, cases generally occur sporadically throughout the year, with occasional increases during the rainy season (e.g., Cambodia). JEV exhibits more epidemic viral activity in northern temperate Asian climates, with disease outbreaks in summer months (e.g., north India, Sri Lanka, Japan). Subtropical regions, such as Thailand, Vietnam, Laos, and Malaysia may see a combination of epidemic and endemic disease characteristics, where JE circulates year-round but peaks during the hot rainy seasons.^{1,7}

Year-round transmission occurs in tropical regions.⁸⁰ Maintenance of JEV in temperate climates might occur through overwintering mosquitoes, poikilothermic vertebrates, hibernating bats, and/or by vertical transmission from female mosquitoes to offspring.^{77, 80} Annual reintroduction into temperate climates by migrating birds, bats, or wind-borne mosquitoes is also a possibility.⁶ JEV infection has recently been documented in wild boar during winter in Japan.⁸¹ Furthermore, JEV persists in swine tonsils for weeks to months, a possible new mechanism for virus overwintering in pigs.^{27, 28}

Epizootics involving swine appear to be cyclical, consisting of two separate amplification cycles. During the first cycle, roughly 20% of pigs will become infected and develop antibodies within ten days. This is followed by a second cycle, 1–2 weeks later, in which mosquitoes transmit the virus to remaining naïve pigs, effectively raising the rate of seroconversion to almost 100%. Clinical cases in humans typically occur following this cycle of amplification in swine.⁶ As each extrinsic incubation period in mosquitoes ranges from 5–15 days, it can take up to 30 days for the virus to complete its infection process in humans.³² Direct pig-to-pig transmission of JEV has been documented experimentally, but whether it plays a role in swine epizootics is unknown (see *Transmission*).

MORBIDITY AND MORTALITY

In swine, production losses arise primarily from reproductive failure and high piglet mortality.⁵⁸ The mortality rate in infected piglets with no immunity to JEV is near 100%, while mortality in infected adult swine is close to zero. Reproductive losses can reach 50–70%.

Seroprevalence of JEV in pigs varies regionally. In monitored swine, seroconversion can reach 100%.^{9, 10} Ladreyt and colleagues⁷ recently summarized JEV seroprevalence studies in swine. Data on antibody detection via ELISA or HI (without PRNT confirmation) should be interpreted with caution since it could be confounded by cross-reactivity with other flaviviruses.⁷

ETIOLOGY

CHARACTERISTICS OF FLAVIVIRUSES

JEV is a small (40–60 nm), enveloped, single-stranded RNA virus belonging to the family *Flaviviridae*. The family contains four genera, *Flavivirus, Pestivirus, Hepacivirus*, and *Pasivirus*.⁸² JEV belongs to the genus *Flavivirus*, along with dengue virus, West Nile virus, yellow fever virus, Zika virus, St. Louis encephalitis virus, Murray Valley encephalitis virus, and others. Speciation of flaviviruses is based on nucleotide sequencing and antigens but also considers geographic distribution, vector types, host types, disease types, and ecology.⁸²

Flaviviruses contain a single long open reading frame (ORF) encoding for three structural proteins, capsid (C), envelope (E), and membrane (prM).^{4, 82} prM and E are involved in viral entry and virion assembly.⁴ During egress, prM is enzymatically cleaved to produce the M protein.⁴ Flaviviruses also encode for seven nonstructural proteins that play a role in viral protein processing, RNA synthesis, and evasion of innate immunity (NS1, NS2A, NS2B, NS3, NS4A, NS4B, and NS5).^{4, 82} The E, prM, and NS1 proteins are immunogenic.⁴

CHARACTERISTICS OF JAPANESE ENCEPHALITIS VIRUS

There is a single JEV serotype. Five genotypes have been described (GI, GII, GII, GIV, and GV); however, genotype relationship to phenotype or virus fitness is still unclear.² The basis for genotyping is phylogenetic analysis of the *E* gene, which is vitally important in viral attachment and entry into host cells.³ Global distribution of JEV genotypes includes (but is not limited to):

- GI-a: Cambodia, Thailand, Vietnam, India, Australia
- GI-b: Vietnam, China, Japan, Korea, Taiwan
- GII: Indonesia, Malaysia, southern Thailand, Papua New Guinea, Australia
- GIII: throughout Asia (not Australasia)
- GIV: Indonesia
- GV: Malaysia, China (Tibet), South Korea⁴

GIII was the most frequently isolated genogroup until the 1990s. GI-b is now believed to be the dominant JEV genotype throughout Asia, replacing GII and GIII strains.^{77, 80, 83, 84} Local transmission leads to GI genetic diversity,⁸⁵ and new variations are being described.⁸⁶ However, GIII strains are still detectable^{87, 88} and regional co-circulation of GI and GIII strains has been documented in China.⁸⁹ Despite its current isolation, GIV is believed to be ancestral to other circulating genotypes and shows the largest antigenic and phylogenetic difference of all the genotypes.³⁴ GV recently reemerged in Tibet and South Korea after remaining undetected for nearly 60 years.⁸⁰ Since its reemergence, GV has shown a high capacity for dispersal in a short time and also warrants continued monitoring given its potential for high virulence.⁹⁰

HISTORY IN SWINE

Human JEV epidemics have been described in Japan since the 19th century.²⁵ The pig-mosquito transmission cycle was first investigated in 1964 and confirmed by 1969.⁷ As previously described, GIII strains were prevalent until the 1990s. At that time, GI became dominant in pigs in thought Asia. However, GIII and GIV strains continue to circulate.⁸⁷

IMMUNITY

POST-EXPOSURE

Natural infection contributes to long-lasting immunity in adult pigs and surviving piglets. Protective antibodies recognize proteins E and NS1.^{25, 91} Both CD4+ and CD8+ T cells also have a role in flavivirus immunity.²⁵ Some sows and boars develop protective immunity prior to reaching sexual maturity, which prevents clinical disease (reproductive failure in sows, infertility in boars).⁹² As described by Ladreyt and colleagues,⁷ anti-JEV antibodies appear 3–7 dpi after experimental infection and are detectable for 7–35 dpi (study endpoints).⁷

VACCINES

Most vaccine development is focused on humans, not pigs.⁴ Vannice and colleagues have described the status of JE vaccines and vaccination programs.⁹³ Swine vaccination is not widely practiced due to cost and the limited time window for administration. However, vaccination of pigs may reduce the risk of human disease in JEV-endemic areas.⁴

Swine vaccines for JEV are mostly GIII-based. Since GI has largely replaced GIII, there is concern over vaccine efficacy. A recent experimental study found that in mice, there is partial cross-protection between GI and GIII strains. Administration of an inactivated GI vaccine led to 50–90% protection against a GII challenge, while administration of an inactivated GIII strain led to 60–80% protection against a GI challenge.⁹⁴

In swine, vaccine programs should consider seasonal variations in JEV infection rate. Based on temperature variations and month of birth, for example, it was predicted that only 80% of pigs born from July to September in Vietnam would reach immunity by sexual maturity.⁹⁵

CROSS-PROTECTION

Cross-protective antibody responses have been induced experimentally in pigs exposed to closely related flaviviruses.⁴ Flavivirus cross-reactive epitopes are found predominantly in domain II of the E protein.⁹¹ However, neutralizing epitopes have been mapped in domain III, showing a high level of conservation between multiple strains of JEV and West Nile virus, suggesting some level of cross-protection.⁹⁶ Experimentally, no cross-neutralizing activity between anti-GI and anti-G3 antibodies was observed in guinea pigs infected with two different Japanese strains of JEV.⁹⁷

GAPS IN PREPAREDNESS

As summarized by Oliveira and colleagues,⁹⁸ introduction of JEV by an infected adult mosquito (likely via aircraft) is the highest risk for the United States. Regions most at risk have climates similar to JEV-endemic areas, such as California and most of the Eastern United States (except Southern Florida and the far Northeast). Like West Nile virus, JEV could spread quickly and become endemic after only a few years.⁹⁸ Evidence suggests that JEV is already circulating in areas outside of Asia, including Europe⁹⁹ and Africa.⁷⁸ Reasons why JEV has not yet emerged in the United States could include:

- High susceptibility of JEV to heat, UV light, and common detergents
- Low numbers of mosquitoes and hosts near airports and seaports
- Short period of viremia (3–4 days) in pigs and ardeid birds
- Lack of contact between hosts (ardeid birds, pigs) and vectors (mosquitoes)
- Cross-protection between JEV and endemic flaviviruses like West Nile virus
- Limited infection capability of mosquitoes during establishment

Although pig-to-pig transmission of JEV has been described, its importance is unclear in cases of natural infection. Modeling indicates that pig-to-pig transmission plus vector-borne transmission explains swine seroprevalence data in Cambodia better than vector-borne transmission alone.¹⁰⁰ Further studies on JEV transmission are needed. Additionally, continued research on the cross-protective characteristics of flaviviruses and effective vaccines will help prepare for JEV.⁶

REFERENCES

- 1. Mackenzie J, Williams D, Smith D. Japanese encephalitis virus: the geographic distribution, incidence, and spread of a virus with a propensity to emerge in new areas. *Perspect Med Virol*. 2006:201-268.
- 2. Le Flohic G, Porphyre V, Barbazan P, Gonzalez J. Review of climate, landscape, and viral genetics as drivers of the Japanese encephalitis virus ecology. *PLoS Negl Trop Dis.* 2013;7(9)

- 3. Thapaliya D, Hanson B, Kates A, Klosterman C, Nair R, Wardyn S. Zoonotic Diseases of Swine: Foodborne and Occupational Aspects of Infection. *Zoonoses-Infections Affecting Humans and Animals*. Springer; 2015.
- 4. Williams D, Mackenzie J, Bingham J. Flaviviruses. In: Zimmerman J, Karriker L, Ramirez A, Schwartz K, Stevenson G, eds. *Diseases of Swine*. 11th ed. John Wiley & Sons; 2019:Chap 530-543.
- 5. Henriksson E, Söderberg R, Ström Hallenberg G, et al. Japanese encephalitis in small-scale pig farming in rural Cambodia: pig seroprevalence and farmer awareness. *Pathogens*. 2021;10(5). doi:10.3390/pathogens10050578
- 6. van den Hurk A, Ritchie S, Mackenzie J. Ecology and geographical expansion of Japanese encephalitis virus. *Ann Rev Entomol.* 2009;54:17-35.
- 7. Ladreyt H, Durand B, Dussart P, Chevalier V. How central is the domestic pig in the epidemiological cycle of Japanese encephalitis virus? A review of scientific evidence and implications for disease control. *Viruses.* Oct 15 2019;11(10). doi:10.3390/v11100949
- 8. Lord JS. Changes in rice and livestock production and the potential emergence of Japanese encephalitis in Africa. *Pathogens*. Mar 4 2021;10(3). doi:10.3390/pathogens10030294
- 9. Cappelle J, Duong V, Pring L, et al. Intensive circulation of Japanese encephalitis virus in peri-urban sentinel pigs near Phnom Penh, Cambodia. *PLoS Negl Trop Dis*. Dec 2016;10(12):e0005149. doi:10.1371/journal.pntd.0005149
- 10. Di Francesco J, Choeung R, Peng B, et al. Comparison of the dynamics of Japanese encephalitis virus circulation in sentinel pigs between a rural and a peri-urban setting in Cambodia. *PLoS Negl Trop Dis*. Aug 2018;12(8):e0006644. doi:10.1371/journal.pntd.0006644
- 11. Duong V, Choeung R, Gorman C, et al. Isolation and full-genome sequences of Japanese encephalitis virus genotype I strains from Cambodian human patients, mosquitoes and pigs. *J Gen Virol*. Sep 2017;98(9):2287-2296. doi:10.1099/jgv.0.000892
- 12. Pham-Thanh L, Nguyen-Tien T, Magnusson U, et al. Dogs as sentinels for flavivirus exposure in urban, peri-urban and rural Hanoi, Vietnam. *Viruses*. Mar 19 2021;13(3). doi:10.3390/v13030507
- 13. Ren X, Fu S, Dai P, et al. Pigsties near dwellings as a potential risk factor for the prevalence of Japanese encephalitis virus in adult in Shanxi, China. *Infect Dis Poverty*. Jun 8 2017;6(1):100. doi:10.1186/s40249-017-0312-4
- 14. Lindahl J, Chirico J, Boqvist S, Thu HT, Magnusson U. Occurrence of Japanese encephalitis virus mosquito vectors in relation to urban pig holdings. *Am J Trop Med Hyg*. Dec 2012;87(6):1076-82. doi:10.4269/ajtmh.2012.12-0315
- 15. Zhao S, Lou Y, Chiu APY, He D. Modelling the skip-and-resurgence of Japanese encephalitis epidemics in Hong Kong. *J Theor Biol*. Oct 7 2018;454:1-10. doi:10.1016/j.jtbi.2018.05.017
- 16. Hameed M, Wahaab A, Nawaz M, et al. Potential role of birds in Japanese encephalitis virus zoonotic transmission and genotype shift. *Viruses*. Feb 24 2021;13(3). doi:10.3390/v13030357
- 17. Yap G, Lim XF, Chan S, et al. Serological evidence of continued Japanese encephalitis virus transmission in Singapore nearly three decades after end of pig farming. *Parasit Vectors*. May 17 2019;12(1):244. doi:10.1186/s13071-019-3501-0
- Kumar HBC, Dhanze H, Bhilegaonkar KN, Chakurkar EB, Kumar A, Yathish HM. Serological evidence of Japanese encephalitis virus infection in pigs in a low human incidence state, Goa, India. *Prev Vet Med*. Feb 2020;175:104882. doi:10.1016/j.prevetmed.2020.104882
- 19. Chethan Kumar HB, Dhanze H, Bhilegaonkar KN, Chakurkar EB, Kumar A, Yathish HM. Reply to commentary on serological evidence of Japanese encephalitis virus infection in pigs in a low human incidence state, Goa, India. *Prev Vet Med.* 04 2020;177:104945. doi:10.1016/j.prevetmed.2020.104945
- 20. Korgaonkar NS, Kumar A, Yadav RS, Kabadi D, Dash AP. Mosquito biting activity on humans & detection of Plasmodium falciparum infection in Anopheles stephensi in Goa, India. *Indian J Med Res.* 2012;135:120-6. doi:10.4103/0971-5916.93434
- 21. Fischer M, Lindsey N, Staples JE, Hills S. Japanese encephalitis vaccines: recommendations of the Advisory Committee on Immunization Practices (ACIP). *MMWR Recomm Rep.* 2010;59(Rr-1):1-27.
- 22. Cheng VCC, Sridhar S, Wong SC, et al. Japanese encephalitis virus transmitted via blood transfusion, Hong Kong, China. *Emerg Infect Dis.* Jan 2018;24(1). doi:10.3201/eid2401.171297

- 23. Park SL, Huang YS, Lyons AC, et al. North American domestic pigs are susceptible to experimental infection with Japanese encephalitis virus. *Sci Rep.* May 21 2018;8(1):7951. doi:10.1038/s41598-018-26208-8
- 24. Yaeger M. Disorders of pigs. In: Njaa B, ed. *Kirkbride's Diagnosis of Abortion and Neonatal Loss in Animals*. John Wiley & Sons; 2011.
- 25. Pierson TC, Diamond MS. The continued threat of emerging flaviviruses. *Nat Microbiol*. 06 2020;5(6):796-812. doi:10.1038/s41564-020-0714-0
- 26. McOrist S. Fertility problems in pigs. *Pig Disease Identification and Diagnosis Guide*. CAB International; 2014.
- 27. Ricklin ME, García-Nicolás O, Brechbühl D, et al. Vector-free transmission and persistence of Japanese encephalitis virus in pigs. *Nat Commun.* 2016;7. doi:10.1038/ncomms10832
- 28. Garcia-Nicolas O, Ricklin ME, Liniger M, et al. A Japanese encephalitis virus vaccine inducing antibodies strongly enhancing in vitro infection is protective in pigs. *Viruses*. 2017;9(5). doi:10.3390/v9050124
- 29. Erlanger T, Weiss S, Keiser J, Utzinger J, Wiedenmayer K. Past, present, and future of Japanese encephalitis. *Emerg Infect Dis.* 2009;15(1):1-7.
- 30. Hubalek Z, Rudolf I, Nowotny N. Arboviruses pathogenic for domestic and wild animals. *Adv Virus Res.* 2014;89:201-275.
- 31. Mansfield KL, Hernandez-Triana LM, Banyard AC, Fooks AR, Johnson N. Japanese encephalitis virus infection, diagnosis and control in domestic animals. *Vet Microbiol*. Mar 2017;201:85-92. doi:10.1016/j.vetmic.2017.01.014
- 32. Tian H, Bi P, Cazelles B, et al. How environmental conditions impact mosquito ecology and Japanese encephalitis: an eco-epidemiological approach. *Environ Int*. 2015;79
- 33. Weaver S, Barrett A. Transmission cycles, host range, evolution and emergence of arboviral disease. *Nat Rev Micro*. 2004;2(10):789-801.
- 34. Huang Y, Higgs S, Horne K, Vanlandingham D. Flavivirus-mosquito interactions. *Viruses-Basel*. 2014:Chap 11.
- 35. Huang YJ, Harbin JN, Hettenbach SM, et al. Susceptibility of a North American *Culex quinquefasciatus* to Japanese encephalitis virus. *Vector Borne Zoonotic Dis*. Nov 2015;15(11):709-11. doi:10.1089/vbz.2015.1821
- 36. Government of Hong Kong Center for Health Protection (CHP). Scientific Committee on Vector-borne Diseases: Vectors of Japanese Encephalitis in Hong Kong. Accessed June 9, 2015. http://www.chp.gov.hk/files/pdf/vectors_of_japanese_encephalitis_in_hk_r.pdf
- 37. Centers for Disease Control and Prevention (CDC). Mosquito Species in Which West Nile Virus Has Been Detected, United States, 1999-2012. Accessed June 9, 2015. https://www.cdc.gov/westnile/resources/pdfs/mosquito%20species%201999-2012.pdf
- 38. Wang L, Hu W, Magalhaes R, et al. The role of environmental factors in the spatial distribution of Japanese encephalitis in mainland China. *Environ Int*. 2014;73:1-9.
- 39. García-Nicolás O, Braun RO, Milona P, et al. Targeting of the nasal mucosa by Japanese encephalitis virus for non-vector-borne transmission. *J Virol*. Dec 15 2018;92(24). doi:10.1128/jvi.01091-18
- 40. Chai C, Palinski R, Xu Y, et al. Aerosol and contact transmission following intranasal infection of mice with Japanese encephalitis virus. *Viruses*. Jan 21 2019;11(1). doi:10.3390/v11010087
- 41. Althouse G, Rossow K. The potential risk of infectious disease dissemination via artificial insemination in swine. *Reprod Dom Anim.* 2011;46:64-67.
- 42. Redant V, Favoreel HW, Dallmeier K, Van Campe W, De Regge N. Efficient control of Japanese encephalitis virus in the central nervous system of infected pigs occurs in the absence of a pronounced inflammatory immune response. *J Neuroinflammation*. Oct 23 2020;17(1):315. doi:10.1186/s12974-020-01974-3
- 43. Ricklin ME, Garcìa-Nicolàs O, Brechbühl D, et al. Japanese encephalitis virus tropism in experimentally infected pigs. *Vet Res.* Feb 24 2016;47:34. doi:10.1186/s13567-016-0319-z

- 44. Xi S, Liu K, Xiao C, et al. Establishment and characterization of the pig tonsil epithelial (PT) cell line as a new model for persist infection of Japanese encephalitis virus. *Vet Microbiol*. Mar 2020;242:108587. doi:10.1016/j.vetmic.2020.108587
- 45. Bharucha T, Sengvilaipaseuth O, Vongsouvath M, et al. Development of an improved RT-qPCR Assay for detection of Japanese encephalitis virus (JEV) RNA including a systematic review and comprehensive comparison with published methods. *PLoS One*. 2018;13(3):e0194412. doi:10.1371/journal.pone.0194412
- 46. Wu H, Rao P, Jiang Y, Opriessnig T, Yang Z. A sensitive multiplex real-time PCR panel for rapid diagnosis of viruses associated with porcine respiratory and reproductive disorders. *Mol Cell Probes*. 2014;28(5-6):264-270.
- 47. Xu X, Chen G, Huang Y, et al. Development of multiplex PCR for simultaneous detection of six swine DNA and RNA viruses. *J Virol Methods*. 2012;183(1):69-74.
- 48. Zeng Z, Liu Z, Wang W, Tang D, Liang H. Establishment and application of a multiplex PCR for rapid and simultaneous detection of six viruses in swine. *J Virol Methods*. 2014;208:102-106.
- 49. Hu L, Lin X, Nie F, et al. Simultaneous typing of seven porcine pathogens by multiplex PCR with a GeXP analyser. *J Virol Methods*. Jun 2016;232:21-8. doi:10.1016/j.jviromet.2015.12.004 10.1016/j.jviromet.2015.12.004. Epub 2015 Dec 17.
- 50. Zhang M, Xie Z, Xie L, et al. Simultaneous detection of eight swine reproductive and respiratory pathogens using a novel GeXP analyser-based multiplex PCR assay. *J Virol Methods*. Nov 2015;224:9-15. doi:10.1016/j.jviromet.2015.08.001
- 51. Jiang Y, Nie F, Jiang S, et al. Development of multiplex oligonucleotide microarray for simultaneous detection of six swine pathogens. *J Virol Methods*. Nov 2020;285:113921. doi:10.1016/j.jviromet.2020.113921
- 52. Rao P, Wu H, Jiang Y, et al. Development of an EvaGreen-based multiplex real-time PCR assay with melting curve analysis for simultaneous detection and differentiation of six viral pathogens of porcine reproductive and respiratory disorder. *J Virol Methods*. 2014;208(56-62)
- 53. Jiang Y, Guo Y, Wang P, et al. A novel diagnostic platform based on multiplex ligase detection-PCR and microarray for simultaneous detection of swine viruses. *J Virol Methods*. 2011;178(1-2):171-178.
- 54. Zhou Y, Chen L, Zhang L, et al. Simultaneous identification of 6 pathogens causing porcine reproductive failure by using multiplex ligation-dependent probe amplification. *Transbound Emerg Dis.* Nov 2020;67(6):2467-2474. doi:10.1111/tbed.13585
- 55. Wu X, Lin H, Chen S, et al. Development and application of a reverse transcriptase droplet digital PCR (RT-ddPCR) for sensitive and rapid detection of Japanese encephalitis virus. *J Virol Methods*. Oct 2017;248:166-171. doi:10.1016/j.jviromet.2017.06.015
- 56. Dhanze H, Bhilegaonkar K, Kumar G, et al. Comparative evaluation of nucleic acid-based assays for detection of Japanese encephalitis virus in swine blood samples. *Arch Virol.* 2015;160(5):7.
- 57. Tian C, Lin Z, He X, et al. Development of a fluorescent-intercalating-dye-based reverse transcription loop-mediated isothermal amplification assay for rapid detection of seasonal Japanese B encephalitis outbreaks in pigs. *Arch Virol.* 2012;157(8):1481-1488.
- 58. Liu H, Lu H, Liu Z, et al. Reverse transcription loop-mediated amplification for rapid detection of Japanese encephalitis virus in swine and mosquitoes. *J Anim Vet Adv.* 2012;11(21):8.
- 59. Zhang L, Tian G, Shi S. Development of a RT-PCR-RFLP assay for differentiation of Japanese encephalitis virus genotype I and genotype III. *Acta Vet et Zootech Sin.* 2014;45(9):1555-1560.
- 60. Zhang L, Cao S, Wu R, et al. Detection and differentiation of Japanese encephalitis virus genotype I and genotype III by reverse transcription loop-mediated isothermal amplification combined with restriction fragment length polymorphism. *Virus Genes*. Apr 2015;50(2):231-7. doi:10.1007/s11262-014-1158-5
- 61. Deng J, Pei J, Gou H, Ye Z, Liu C, Chen J. Rapid and simple detection of Japanese encephalitis virus by reverse transcription loop-mediated isothermal amplification combined with a lateral flow dipstick. *J Virol Methods*. 2015;213:7.
- 62. Mei L, Wu P, Ye J, et al. Development and application of an antigen capture ELISA assay for diagnosis of Japanese encephalitis virus in swine, human and mosquito. *Virol J*. 2012;9(4)

- 63. World Organization for Animal Health (OIE). Manual of Diagnostic Tests and Vaccines for Terrestrial Animals. Accessed March 9, 2014. http://www.oie.int/international-standard-setting/terrestrial-manual/access-online/
- 64. Cha GW, Lee EJ, Lim EJ, et al. A novel immunochromatographic test applied to a serological survey of Japanese encephalitis virus on pig farms in Korea. *PLoS One*. 2015;10(5):e0127313. doi:10.1371/journal.pone.0127313
- 65. Dhanze H, Bhilegaonkar KN, Rawat S, et al. Development of recombinant nonstructural 1 protein based indirect enzyme linked immunosorbent assay for sero-surveillance of Japanese encephalitis in swine. *J Virol Methods*. Oct 2019;272:113705. doi:10.1016/j.jviromet.2019.113705
- 66. Dhanze H, Kumar MS, Singh V, et al. Detection of recent infection of Japanese encephalitis virus in swine population using IgM ELISA: a suitable sentinel to predict infection in humans. *J Immunol Methods*. Nov 2020;486:112848. doi:10.1016/j.jim.2020.112848
- 67. Zhou D, Pei C, Yang K, et al. Development and application of a monoclonal-antibody-based blocking ELISA for detection of Japanese encephalitis virus NS1 antibodies in swine. *Arch Virol.* Jun 2019;164(6):1535-1542. doi:10.1007/s00705-019-04218-9
- 68. Litzba N, Klade CS, Lederer S, Niedrig M. Evaluation of serological diagnostic test systems assessing the immune response to Japanese encephalitis vaccination. *PLoS Negl Trop Dis.* Nov 16 2010;4(11):e883. doi:10.1371/journal.pntd.0000883
- 69. Wu Y, Wu X, Chen J, Hu J, Huang X, Zhou B. A novel protein chip for simultaneous detection of antibodies against four epidemic swine viruses in China. *BMC Vet Res.* May 26 2020;16(1):162. doi:10.1186/s12917-020-02375-7
- 70. Grace MR, Dhanze H, Pantwane P, Sivakumar M, Gulati BR, Kumar A. Latex agglutination test for rapid on-site serodiagnosis of Japanese encephalitis in pigs using recombinant NS1 antigen. J Vector Borne Dis. Apr-Jun 2019;56(2):105-110. doi:10.4103/0972-9062.263717
- 71. Grace MR, Chauhan J, Suman Kumar M, et al. Evaluation of carboxyl beads based latex agglutination test for rapid sero-diagnosis of Japanese encephalitis. *Biologicals*. Nov 2019;62:72-76. doi:10.1016/j.biologicals.2019.09.003
- 72. Dhanze H, Bhilegaonkar KN, Kumar C, Kumar MS, Singh P, Kumar A. Development and evaluation of lateral flow assay for sero-diagnosis of Japanese encephalitis in swine. *Anim Biotechnol*. Aug 2020;31(4):350-356. doi:10.1080/10495398.2019.1602539
- 73. Chauhan J, Dhanze H, Kumar HBC, Kumar MS, Bhilegaonkar KN. Development of dipstick enzyme linked immunosorbent assay for on-site sero-diagnosis of Japanese encephalitis in swine. *J Virol Methods*. Jun 2020;280:113876. doi:10.1016/j.jviromet.2020.113876
- 74. Lyons AC, Huang YS, Park SL, et al. Shedding of Japanese encephalitis virus in oral fluid of infected swine. *Vector Borne Zoonotic Dis.* Sep 2018;18(9):469-474. doi:10.1089/vbz.2018.2283
- 75. Katayama T, Saito S, Horiuchi S, et al. Nonsuppurative encephalomyelitis in a calf in Japan and isolation of Japanese encephalitis virus genotype 1 from the affected calf. *J Clin Microbiol*. Oct 2013;51(10):3448-53. doi:10.1128/jcm.00737-13
- 76. Kako N, Suzuki S, Sugie N, et al. Japanese encephalitis in a 114-month-old cow: pathological investigation of the affected cow and genetic characterization of Japanese encephalitis virus isolate. BMC Vet Res. Mar 11 2014;10:63. doi:10.1186/1746-6148-10-63
- 77. Schuh A, Ward M, Leigh Brown A, Barrett A. Dynamics of emergence and establishment of a newly dominant genotype of Japanese encephalitis virus throughout Asia. *J Virol* 2014;79(0):17-24.
- 78. Simon-Loriere E, Faye O, Prot M, et al. Autochthonous Japanese encephalitis with yellow fever coinfection in Africa. *N Engl J Med.* 04 2017;376(15):1483-1485. doi:10.1056/NEJMc1701600
- 79. Auerswald H, Maquart PO, Chevalier V, Boyer S. Mosquito vector competence for Japanese encephalitis virus. *Viruses*. Jun 2021;13(6). doi:10.3390/v13061154
- 80. Schuh A, Ward M, Brown A, Barrett A. Phylogeography of Japanese encephalitis virus: genotype is associated with climate. *PLoS Negl Trop Dis.* 2013;7(8)
- 81. Komiya T, Toriniwa H, Matsumura T, Takegami T, Nakayama T. Epidemiological study on Japanese encephalitis virus distribution in Ishikawa prefecture, Japan, by serological investigation using wild boar sera. *J Vet Med Sci.* Jun 21 2019;81(6):903-905. doi:10.1292/jvms.18-0613

- 82. Simmonds P, Becher P, Bukh J, et al. ICTV Virus Taxonomy Profile: Flaviviridae. *J Gen Virol*. 01 2017;98(1):2-3. doi:10.1099/jgv.0.000672
- 83. Chen YY, Fan YC, Tu WC, et al. Japanese encephalitis virus genotype replacement, Taiwan, 2009-2010. *Emerg Infect Dis.* Dec 2011;17(12):2354-6. doi:10.3201/eid1712.110914
- 84. Do LP, Bui TM, Hasebe F, Morita K, Phan NT. Molecular epidemiology of Japanese encephalitis in northern Vietnam, 1964-2011: genotype replacement. *Virol J*. Apr 1 2015;12:51. doi:10.1186/s12985-015-0278-4
- 85. Schuh A, Ward M, Leigh Brown A, Barrett A. Dynamics of emergence and establishment of a newly dominant genotype of Japanese encephalitis virus throughout Asia. *J Virol.* 2014;79(0):17-24.
- 86. Raut AA, Aasdev A, Mishra A, et al. Detection of coinfection of a divergent subgroup of genotype I Japanese encephalitis virus in multiple classical swine fever virus outbreaks in pigs of Assam, India. *Transbound Emerg Dis.* Oct 30 2020;doi:10.1111/tbed.13903
- 87. Kuwata R, Torii S, Shimoda H, et al. Distribution of Japanese encephalitis virus, Japan and Southeast Asia, 2016-2018. *Emerg Infect Dis.* Jan 2020;26(1):125-128. doi:10.3201/eid2601.190235
- 88. Wu R, Wang Q, Liu H, et al. Phylogenetic analysis reveals that Japanese encephalitis virus genotype III is still prevalent in swine herds in Sichuan province in China. *Arch Virol*. Jun 2016;161(6):1719-22. doi:10.1007/s00705-016-2814-y
- 89. Chai C, Wang Q, Cao S, et al. Serological and molecular epidemiology of Japanese encephalitis virus infections in swine herds in China, 2006-2012. *J Vet Sci*. Jan 31 2018;19(1):151-155. doi:10.4142/jvs.2018.19.1.151
- 90. Gao X, Liu H, Li M, Fu S, Liang G. Insights into the evolutionary history of Japanese encephalitis virus (JEV) based on whole-genome sequences comprising the five genotypes. *Virol J*. 2015;12(1):43.
- 91. Gangwar R, Shil P, Cherian S, Gore M. Delineation of an epitope on domain I of Japanese encephalitis virus envelope glycoprotein using monoclonal antibodies. *Virus Res.* 2011;158(1-2):179-187.
- 92. Conlan J, Vongxay K, Jarman R, et al. Serologic study of pig-associated viral zoonoses in Laos. *Am J Trop Med Hyg.* 2012;86(6):1077-1084.
- 93. Vannice KS, Hills SL, Schwartz LM, et al. The future of Japanese encephalitis vaccination: expert recommendations for achieving and maintaining optimal JE control. *NPJ Vaccines*. Jun 2021;6(1):82. doi:10.1038/s41541-021-00338-z
- 94. Wei J, Wang X, Zhang J, et al. Partial cross-protection between Japanese encephalitis virus genotype I and III in mice. *PLoS Negl Trop Dis*. Aug 2019;13(8):e0007601. doi:10.1371/journal.pntd.0007601
- 95. Ruget AS, Beck C, Gabassi A, et al. Japanese encephalitis circulation pattern in swine of northern Vietnam and consequences for swine's vaccination recommendations. *Transbound Emerg Dis*. Dec 2018;65(6):1485-1492. doi:10.1111/tbed.12885
- 96. Deng W, Guan C, Liu K, et al. Fine mapping of a linear epitope on EDIII of Japanese encephalitis virus using a novel neutralizing monoclonal antibody. *Virus Res.* 2014;179:133-139.
- 97. Kang BK, Hwang JM, Moon H, et al. Comparison of the antigenic relationship between Japanese encephalitis virus genotypes 1 and 3. *Clin Exp Vaccine Res.* Jan 2016;5(1):26-30. doi:10.7774/cevr.2016.5.1.26
- 98. Oliveira ARS, Cohnstaedt LW, Noronha LE, Mitzel D, McVey DS, Cernicchiaro N. Perspectives regarding the risk of introduction of the Japanese encephalitis virus (JEV) in the United States. *Front Vet Sci.* 2020;7:48. doi:10.3389/fvets.2020.00048
- 99. Ruiz-Fons F. A review of the current status of relevant zoonotic pathogens in wild swine (*Sus scrofa*) populations: changes modulating the risk of transmission to humans. *Transbound Emerg Dis.* 2015;64(1)
- 100. Diallo AOI, Chevalier V, Cappelle J, Duong V, Fontenille D, Duboz R. How much does direct transmission between pigs contribute to Japanese Encephalitis virus circulation? A modelling approach in Cambodia. *PLoS One*. 2018;13(8):e0201209. doi:10.1371/journal.pone.0201209