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Most recent publications:

1. **Saekhow P**, Kishizuka S, Sano N, Mitsui H, Akasaki H, Mawatari T, Ikeda H. Coincidental detection of genomes of porcine parvoviruses and porcine circovirus type 2 infecting pigs in Japan. *J Vet Med Sci*. 2016. 77(12):1581–6.
2. **Saekhow P**, Ikeda H. Prevalence and genomic characterization of porcine parvoviruses detected in Chiangmai area of Thailand in 2011. *Microbiol Immunol*. 2015. 59(2):82–8.
3. **Saekhow P**, Mawatari T, Ikeda H. Coexistence of multiple strains of porcine parvovirus 2 in pig farms. *Microbiol Immunol*. 2014. 58(7):382–7.
4. Thongprachum A, Khamrin P, **Saekhow P**, Pantip C, Peerakome S, Ushijima H, Maneekarn N. Analysis of the VP6 gene of human and porcine group A rotavirus strains with unusual subgroup specificities. *J Med Virol*. 2009. 81(1):183–91.
5. Chan-It W, Khamrin P, **Saekhow P**, Pantip C, Thongprachum A, Peerakome S, Ushijima H, Maneekarn N. Multiple combinations of P[13]-like genotype with G3, G4, and G5 in porcine rotaviruses. *J Clin Microbiol*. 2008. 46(4):1169–73.