

Supplementary Table 1. Identified Respiratory Pathogens in Patients with Human Bocavirus Pneumonia

Total (n = 185)	Upper Respiratory Infection (n = 75)	Pneumonia (n = 110)
Co-infection (n = 88, 47.6%)	n = 34, 45.3%	n = 54, 49.1%
Combined pathogens	Adenovirus (n = 1)	Adenovirus (n = 5)
	Coronavirus (n = 3)	Coronavirus (n = 3)
	Enterovirus (n = 2)	Human metapneumovirus (n = 2)
	Human metapneumovirus (n = 1)	Influenza A or B (n = 5)
	Influenza A or B (n = 4)	Parainfluenza virus 1, 3, 4 (n = 4)
	Parainfluenza virus 1, 3, 4 (n = 3)	Rhinovirus (n = 4)
	Rhinovirus (n = 8)	Acinetobacter baumannii (n = 10)
	Acinetobacter baumannii (n = 2)	Enterobacter cloacae (n = 1)
	Enterobacter aerogenes (n = 1)	Haemophilus influenzae (n = 1)
	Pseudomonas aeruginosas (n = 1)	Klebsiella pneumonia (n = 1)
	Staphylococcus aureus (n = 2)	Legionella pneumonia (n = 2)
	Streptococcus pneumonia (n = 1)	Mycoplasma pneumonia (n = 1)
	Stenotorophomonas maltophilia (n = 2)	Pseudomonas aeruginosas (n = 5)
	Aspergillus (n = 1)	Staphylococcus aureus (n = 9)
	TB/NTM (n = 4)	Streptococcus pneumonia (n = 2)
	Orientia tsutsugamushi (n = 1)	Stenotorophomonas maltophilia (n = 2)
		Streptococcus agalactiae (n = 1)
		Cytomegalovirus (n = 7)
		Pneumocystis jirovecii (n = 1)
		Aspergillus species (n = 2)
		Pulmonary TB/NTM (n = 5)

TB = tuberculosis, NTM = nontuberculous mycobacteria