

UNIVERSITY OF CENTRAL FLORIDA

HHS Select Agents and Toxins	
	Abrin
	Bacillus cereus Biovar anthracis
	Botulinum neurotoxins
	Clostridium botulinum and neurotoxin-producing species of Clostridia
	Conotoxins (Short, paralytic alpha conotoxins containing the following amino
	acid sequence X ₁ CCX ₂ PACGX ₃ X ₄ X ₅ X ₆ CX ₇)
	Coxiella burnetii
	Crimean-Congo hemorrhagic fever virus
	Diacetoxyscirpenol
	Eastern equine encephalitis virus
	Ebola virus
	Francisella tularensis
	Lassa fever virus
	Lujo virus
	Marburg virus
	Mpox virus Clade I
	1918-1919 H1N1 including reconstructed replication competent forms of the
	1918 pandemic influenza virus containing any portion of the coding regions of all
	eight gene segments (Reconstructed 1918 Influenza virus)
	Ricin
	Rickettsia prowazekii
	Severe acute respiratory coronavirus (SARS-CoV)
	SARS-CoV/SARS-CoV-2 chimeric viruses resulting from any deliberate
	manipulation of SARS-CoV-2 to incorporate nucleic acids coding for SARS-CoV
	virulence factors
	Saxitoxin
	Chapare virus
	Guanarito virus
	Junín virus
	Machupo virus
	Sabía virus
	Staphylococcal enterotoxins (subtypes A, B, C, D, E)
	T-2 toxin
	Tetrodotoxin
	Tick-borne encephalitis complex virus: Far Eastern subtype
	Tick-borne encephalitis complex virus: Siberian subtype
	Kyasanur Forest disease virus
	Omsk hemorrhagic fever virus
	Variola major virus (Smallpox virus)

This list of agents was obtained from the <u>Implementation Guidance</u> for the United States Government Policy for Oversight of Dual Use Research of Concern and Pathogens with Enhanced Pandemic Potential.



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	Variola minor virus (Alastrim)	
	Yersinia pestis	
Overla	Select Agents and Toxins	
	Bacillus anthracis	
	Bacillus anthracis Pasteur strain	
	Brucella abortus	
	Brucella melitensis	
	Brucella suis	
	Burkholderia mallei	
	Burkholderia pseudomallei	
	Hendra virus	
	Nipah virus	
	Rift Valley fever virus	
	Venezuelan equine encephalitis virus	
USDA Veterinary Services Select Agents and Toxins		
	African horse sickness virus	
	African swine fever virus	
	Avian influenza virus [this is included here as a veterinary select agent in 9 CFR	
	121.3. Low pathogenicity strains are excluded.]	
	Classical swine fever virus	
	Foot-and-mouth disease virus	
	Goat pox virus	
	Lumpy skin disease virus	
	Mycoplasma capricolum	
	Mycoplasma mycoides	
	Newcastle disease virus	
	Peste des petits ruminants virus	
	Rinderpest virus	
	Sheep pox virus	
	Swine vesicular disease virus	
USDA P	lant Select Agents and Toxins	
	Coniothyrium glycines	
	Peronosclerospora philippinensis (Peronosclerospora sacchari)	
	Ralstonia solanacearum	
	Rathayibacter toxicus	
	Sclerophthora rayssiae	
	Synchytrium endobioticum	
	Xanthomonas oryzae	
Other R	isk Group 4 Pathogens	

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	Tick-borne encephalitis virus complex including Absetterov, Central European	
	encephalitis, Hanzalova, Hypr, and Kumlinge	
	Herpesvirus simiae (herpes B or monkey B virus)	
	Hemorrhagic fever agents and viruses as yet undefined	
Other Risk Group 3 Pathogens		
	Bartonella	
	Brucella	
	Orientia tsutsugamushi	
	Pasteurella multocida type B -"buffalo" and other virulent strains	
	Rickettsia akari, R. australis, R. canada, R. conorii, R. rickettsii, R, siberica, R. typhi	
	(R. mooseri)	
	Chikungunya virus except the vaccine strain 181/25	
	Semliki Forest virus	
	St. Louis encephalitis virus	
	Flexal virus	
	Lymphocytic choriomeningitis virus (LCM) (neurotropic strains)	
	Hantaviruses, including Hantaan virus	
	Middle East respiratory syndrome coronavirus (MERS-CoV)	
	Severe acute respiratory coronavirus 2 (SARS-CoV-2)	
	Japanese encephalitis virus except strain SA 14-14-2	
	West Nile virus	
	Yellow fever virus	
	Human influenza A virus H2N2 (1957-1968)	
	Highly pathogenic avian influenza A virus H5Nx strains within the	
	Goose/Guangdong/96-like H5 lineage (e.g., H5N1, H5N6, H5N8 etc.)	
	Transmissible spongiform encephalopathy (TSE) agents (e.g., Creutzfeldt-Jacob	
	disease and kuru agents)	
Other		
	Any attenuated pathogen or vaccine strain that is currently excluded from the	
	Select Agent Regulations that exhibits the recovery of virulence at or near the	
	wild-type	
	Mpox virus clade I/II chimeric viruses resulting from any deliberate manipulation	
	of clade II to incorporate nucleic acids coding for clade I virulence factors	

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