

# Influenza A(H1N1)pdm09 <u>egg-derived</u><sup>1</sup> candidate vaccine viruses for development and production of vaccines for use in the 2021-2022 northern hemisphere influenza season

Antigenic and genetic analyses are performed by the WHO Collaborating Centres of the Global Influenza Surveillance and Response System (GISRS). Unless otherwise specified, all candidate vaccine viruses posted on this table have passed two-way haemagglutination inhibition (HI) test. National or Regional control authorities approve the composition and formulation of vaccines used in each country<sup>2</sup>

### 26 February 2021

### Candidate vaccine viruses (CVVs) (antigenically like A/Victoria/2570/2019 (Egg derived) - Accession number (GISAID): EPI\_ISL\_417210

Parent virus	Candidate vaccine virus	Type of virus or reassortant	Developing institute	Available from
	Wild type virus			WHO CCs and WHO ERLs
A/Indiana/02/2020	X-349	Classical	NYMC	NYMC, USA NIID, Japan NIBSC, UK
	X-349A	Classical	NYMC	NYMC, USA NIID, Japan NIBSC, UK
A/Victoria/2570/2019	Wild type virus			WHO CCs and WHO ERLs
	IVR-215	classical	Seqirus	WHO CCS and WHO ERL
A/Victoria/3/2020	Wild type virus			WHO CCs and WHO ERLs
	IVR-216	classical	Seqirus	WHO CCS and WHO EKE
A/Victoria/1/2020	Wild type virus			WILLO CC   WILLO EDI -
	IVR-217	classical	Seqirus	WHO CCs and WHO ERLS



## Influenza A(H3N2) <u>egg-derived</u> <sup>1</sup> candidate vaccine viruses for development and production of vaccines for use in the 2021-2022 northern hemisphere influenza season

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### 26 February 2021

Candidate vaccine viruses (antigenically like A/Cambodia/e0826360/2020 (egg derived) - Accession number (GISAID): EPI ISL 806547

Parent virus	Candidate vaccine virus	Type of virus or reassortant	Developing institute	Available from
	Wild type virus			WHO CCs
A/Cambodia/e0826360/2020	IVR-224	Classical	Seqirus	NIBSC, UK
	Wild type virus			WHO CCs
A/Tasmania/503/2020				NIBSC, UK
	IVR-221	Classical	Seqirus	WHO CCs
		Ciassical		NIBSC, UK

引用元:WHOウェブサイト



# Influenza B Victoria lineage <u>egg-derived</u><sup>1</sup> candidate vaccine viruses for development and production of vaccines for use in the 2021-2022 northern hemisphere influenza season

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### 26 February 2021

### Candidate vaccine viruses (antigenically like B/Washington/02/2019 (egg derived) - Accession number (GISAID): EPI\_ISL\_352076

Parent virus	Candidate vaccine virus	Type of virus or reassortant	Developing institute	Available from
	Wild type virus			WHO CCs
B/Hong Kong/574/2019	BX-87C	Classical	NYMC	NIBSC, UK NYMC, USA
	Wild type virus			WHO CCs NIBSC, UK
B/Washington/02/2019	BX-85C	Classical	NYMC	NIBSC, UK NIID, Japan NYMC, USA
B/Brisbane/35/2018	Wild type virus			NIBSC, UK
	Wild type virus			NIBSC, UK NIID, Japan
B/Victoria/705/2018	BVR-11	Classical	VIDRL	CCDC, China FCI, UK NIBSC, UK NIID, Japan VIDRL, Australia
	Wild type virus			WHO CCs
	BX-83A	Classical	NYMC	CCDC, China NYMC, USA
B/Sichuan-Gaoxin/531/2018	CNIC-1906	Classical	CCDC	CCDC, China FCI, UK NIBSC, UK VIDRL, Australia

引用元:WHOウェブサイト



# Influenza B Yamagata lineage <u>egg-derived</u><sup>1</sup> candidate vaccine viruses for development and production of vaccines for use in the 2021-2022 northern hemisphere influenza season

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### 26 February 2021

Candidate vaccine viruses (antigenically like B/Phuket/3073/2013 (egg derived) - Accession number (GISAID): EPI\_ISL\_168822

Parent virus	Candidate vaccine virus	Type of virus or reassortant	Developing institute	Available from
B/Phuket/3073/2013	Wild type virus			WHO CCs NIBSC, UK
,	BVR-1B	Classical	Segirus	VIDRL, Australia
	Wild type virus			CDC, USA
B/California/12/2015	BX-59A	Classical	NYMC	NIBSC, UK NYMC, USA
	BX-59B	Classical	NYMC	NIBSC,UK NYMC, USA
B/Brisbane/9/2014	Wild type virus			WHO CCs NIBSC, UK
B/Utah/09/2014	Wild type virus			CDC, USA NIBSC, UK
	Wild type virus			NIBSC, UK
B/Arizona/10/2015	BX-63	Classical	NYMC	NIBSC, UK NYMC, USA
	BX-63A	Classical	NYMC	NIBSC, UK NYMC, USA
B/Hong Kong/3417/2014	Wild type virus			NYMC, USA
	BX-57	Classical	NYMC	NIBSC, UK NYMC, USA

引用元:WHOウェブサイト