



Live virus vaccines based on a yellow fever vaccine backbone: Standardized template with key considerations for a risk/benefit assessment^{☆,☆☆}



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ABSTRACT

The Brighton Collaboration Viral Vector Vaccines Safety Working Group (V3SWG) was formed to evaluate the safety of live, recombinant viral vaccines incorporating genes from heterologous viruses inserted into the backbone of another virus (so-called “chimeric virus vaccines”). Many viral vector vaccines are in advanced clinical trials. The first such vaccine to be approved for marketing (to date in Australia, Thailand, Malaysia, and the Philippines) is a vaccine against the flavivirus, Japanese encephalitis (JE), which employs a licensed vaccine (yellow fever 17D) as a vector. In this vaccine, two envelope proteins (prM-E) of YF 17D virus were exchanged for the corresponding genes of JE virus, with additional attenuating mutations incorporated into the JE gene inserts. Similar vaccines have been constructed by inserting prM-E genes of dengue and West Nile into YF 17D virus and are in late stage clinical studies. The dengue vaccine is, however, more complex in that it requires a mixture of four live vectors each expressing one of the four dengue serotypes. This vaccine has been evaluated in multiple clinical trials. No significant safety concerns have been found. The Phase 3 trials met their endpoints in terms of overall reduction of confirmed dengue fever, and, most importantly a significant reduction in severe dengue and hospitalization due to dengue. However, based on results that have been published so far, efficacy in preventing serotype 2 infection is less than that for the other three serotypes. In the development of these chimeric vaccines, an important series of comparative studies of safety and efficacy were made using the parental YF 17D vaccine virus as a benchmark. In this paper, we use a standardized template describing the key characteristics of the novel flavivirus vaccine vectors, in comparison to the parental YF 17D vaccine. The template facilitates

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scientific discourse among key stakeholders by increasing the transparency and comparability of information. The Brighton Collaboration V3SWG template may also be useful as a guide to the evaluation of other recombinant viral vector vaccines.

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1. Preamble

1.1. Need for working group and development of a standardized template for collection of key information for risk/benefit assessment of viral vector vaccines

Recombinant viral vectors provide an effective means for heterologous antigen expression *in vivo* and thus represent promising platforms for developing novel vaccines against human pathogens such as Ebola, human immunodeficiency virus (HIV), tuberculosis, and malaria) [1–9]. Preclinical evaluation of such viral vector vaccines has indicated their potential for immunization and an increasing number of candidate vaccines are entering human clinical trials. Improving our ability to anticipate potential safety issues and meaningfully assess or interpret safety data from trials of such new viral vector vaccines will increase the likelihood of public acceptance should they be licensed [10–13].

The Brighton Collaboration (www.brightoncollaboration.org) was formed in 2000 as an international voluntary collaboration to enhance the science of vaccine safety research [e.g., via development of standardized case definitions of adverse events following immunizations (AEFI)] [14]. In recognition of these needs in this domain, the Brighton Collaboration created the Viral Vector Vaccines Safety Working Group (V3SWG) in October 2008. Analogous to the value embodied in standardized case definitions for AEFI, the V3SWG believes a standardized template describing the key characteristics of a novel vaccine vector, when completed and maintained with the latest research, will facilitate scientific discourse among key stakeholders by increasing the transparency and comparability of information. Fortunately, the International AIDS Vaccine Initiative (IAVI) had already developed an internal tool to assess the risk/benefit of different viral vectors under its sponsorship. The IAVI graciously shared this tool with the V3SWG for adaptation and broader use as a standardized template for collection of key information for risk/benefit assessment on any viral vector vaccine. This tool was aimed at identifying potential major hurdles or concerns that would need to be addressed during the development of a vectored vaccine. The template collects information on the characteristics of the wild type virus from which the vector was derived as well as known effects of the proposed vaccine vector in animals and humans, manufacturing features, toxicology and potency, nonclinical studies, and human use, with an overall adverse effect and risk assessment.

The V3SWG hopes that eventually all developers/researchers of viral vector vaccines (especially those likely to be used in humans in the near future) will complete this template and submit it to the V3SWG and Brighton Collaboration for peer review and eventual publication in Vaccine. Following this, to promote transparency, the template will be posted and maintained on the Brighton Collaboration website for use/reference by various stakeholders. Furthermore, recognizing the rapid pace of new scientific developments in this domain (relative to AEFI case definitions), we hope to maintain these completed templates “wiki-” style with the help of Brighton Collaboration and each vector vaccine “community.”

1.2. Need for risk/benefit assessment of live virus vaccines based upon a yellow fever vaccine backbone

Yellow fever (YF) is a mosquito-borne flavivirus disease that has long endangered persons in sub-Saharan Africa and

tropical areas in South America and is associated with a case fatality rate of 20–50% [15]. Since no effective anti-yellow fever virus medications are available and current mosquito-control measures are inadequate, vaccination remains paramount to YF prevention and control. Although appropriately controlled efficacy studies have never been carried out, the decline in YF cases following vaccination campaigns and the production in most studies of neutralizing antibodies in more than 95% of vaccinees are considered sufficient evidence that the 17D vaccine is effective. Regulations call for the vaccine to be administered every ten years; however, a recent recommendation by WHO's Strategic Advisory Group is that the vaccine need only be given once [16].

The live, attenuated YF 17D vaccine was previously deemed to be the world's safest and a model for the development of other live virus vaccines including polio, measles, mumps and varicella. Consequently, live vaccines against other flaviviruses, such as Japanese encephalitis virus, West Nile virus, and the four serotypes of dengue viruses, based on the YF 17D virus vaccine began to be developed. In 2001, however, severe rare reactions that were frequently fatal became recognized [17–19]. These reactions involved multiple organ systems and were named yellow fever vaccine-associated viscerotropic disease (YEL-AVD). As a consequence, in addition to live virus vaccines, inactivated vaccines, including one for YF, are being developed.

Risk groups for the development of YEL-AVD include elderly males as young as 56 years [20], women in their prime child-bearing years [21], and persons thymectomized as treatment for thymoma [22]. Guidelines for the definition of viscerotropic disease and for the association of YF vaccine with viscerotropic disease have been developed by a Brighton Collaboration working group [23]. In addition to YEL-AVD, other rare vaccine reactions include anaphylaxis and neurological disease called YF vaccine-associated neurological disease (YEL-AND). These reactions are rarely fatal or result in long term sequelae. Recognition of YEL-AVD has had a number of consequences including changes in recommendations for the vaccination of prospective travelers to and inhabitants in jungle (sylvatic) or savanna (intermediate) regions where they have a risk of exposure to YF virus-infected mosquitoes.

The feared complication of YF is spread of the virus to urban areas where the principal mosquito vector is *Aedes aegypti*, a mosquito that has become difficult if not impossible to eradicate in areas of huge tropical and subtropical municipalities with their large urban slums. Strategies for vaccination have also become complex as the number of cases of YEL-AVD in travelers have exceeded the number of cases of YF [15]. In South America, during periods of low virus activity, the risk to travelers of serious adverse events from the vaccine, particularly males over the age of 60, may be similar to that of developing yellow fever. In contrast, in Africa, the risk of acquiring yellow fever can be 600 times, and of death 700 times, the risk of vaccination.

An understanding of the makeup of the flavivirus genome is helpful in understanding the new vaccines being developed based upon the yellow fever vaccine virus as a vector. Flaviviruses are single stranded, positive sense RNA viruses. Their genomes encode three structural genes (capsid [C], pre-membrane [prM], envelope [E]) and at least seven non-structural genes (NS1, NS2a, NS2b, NS3, NS4a, NS4b, and NS5) in that order [24]. The coding region is flanked by non-coding regions at the 5' and 3' ends of the genome. The YF 17D vaccine was developed by serial passage of a

Table 1
Risk/benefit assessment for live virus vaccines based on a yellow fever vaccine backbone.

1. Basic information	Information		
1.1. Author(s)	Thomas P. Monath MD/modified by Stephen J. Seligman MD/moderated by Jim S. Robertson MD		
1.2. Date completed/updated	March 1, 2011 (JSR), updated September 25, 2014 (SJS,TPM)		
2. Vaccine vector information	Information		
2.1. Name of vaccines	Yellow fever 17D-204, Yellow fever 17DD, ChimeriVax-TDV (CYD-TDV), ChimeriVax-JE, Imojev®, ChimeriVax WN		
2.2. Class/subtype	Flaviviridae/Flavivirus single strand positive-sense RNA virus		
2.3. Proposed route of administration	Subcutaneously		
3. Characteristics of wild type agent and attenuated vaccine derived from them	Information	Comments/concerns	Reference(s)
3.1. Disease(s) caused by wild type, the strength of evidence, severity, and duration of disease for the following categories:	Wild-type yellow fever (YF) virus causes hepatitis and hemorrhagic fever	Mortality rates of 20–50%	[15]
3.1a. Disease(s) caused by attenuated live yellow fever virus vaccine			
Overall	Yellow fever vaccine associated neurotropic (YEL-AND) and viscerotropic (YEL-AVD) serious adverse effects	Reporting rate of YEL-AND is 0.8 per 100,000 overall. According to VAERS data, reporting rate of YEL-AVD is 0.4 per 100,000 overall. Other estimates vary widely. Risk for YEL-AVD increases in males with age ≥ 56 . In 2001, a series of reports appeared in Lancet describing severe, frequently fatal, viscerotropic reactions to yellow fever vaccine that stimulated surveillance of what had been considered the safest of live virus vaccines. Additional cases of YEL-AVD in prospective travelers and in inhabitants of South America were recognized. In S. America the incidence may be underestimated because much of the vaccine is administered to previously vaccinated individuals. In Peru the incidence of fatal reactions in a previously unimmunized population approached 1 in 10,000. In Africa surveillance for AEFI remains inadequate, although WHO has conducted follow-up for adverse events after mass YF vaccine campaigns, without definitive identification of YEL-AVD	[17–20,27–30]
In immunocompromized	Association of thymic disease with YEL-AVD. Immune deficiency a contraindication to YF 17D. Systemic lupus erythematosus on corticosteroids	Thymic disease is a contraindication to YF 17D. One report of YEL-AND (fatal) in a patient with HIV/AIDS 3 fatal cases. No safety issues after administration of YF17D in 102 HIV+ volunteers, although rate of serious adverse events of up to 3% cannot be excluded	[22,29,31–33]
In neonates, infants, children	Age < 6 months associated with risk of YEL-AND, and is a contraindication. Children aged 6–9 months only vaccinated under special circumstances and on the basis of current official advice		[15,34]
During pregnancy and in the unborn	Pregnancy a hypothetical contraindication. However congenital infections very rare and no reports of adverse effects on fetus	Two reports of neonates acquiring 17 D infection and YEL-AND via milk from recently vaccinated nursing mothers	[15,35]
Are there any other susceptible populations	Elderly have higher risk of YEL-AND and elderly males, YEL-AVD. Women of prime childbearing age	Reporting rate in persons >70 approx 2 per 100,000 for both YEL-AND and AVD	[20,21,27,36]
Animals	Wild-type YF causes similar disease in nonhuman primates (NHP), and (adapted strains) in hamsters, but is neurotropic (no hepatitis) in mice. YF 17D vaccine inoculated IC neurotropic/lethal in mice and causes self-limited encephalitis in monkeys	YF 17D vaccine is controlled by monkey neurovirulence test based on scoring inflammatory lesions in brain/spinal cord	[15]
3.2. Is there any known evidence of neurological or cardiac involvement of the wild type agent?	YEL-AND most often caused by neuroinvasion of CNS followed by acute meningoencephalitis, generally self-limited, rare sequelae. Rare except in infants <6 mos. Wild-type YF does not cause encephalitis (though the virus is neurovirulent in mice and monkeys after intracerebral inoculation). Wild-type (Wt) virus and YEL-AVD associated with myocarditis, but not a prominent feature		

Table 1 (Continued)

1. Basic information	Information		
3.3. What is known about the types of human cells infected and the receptors used in humans and animals?	In humans and NHP, wild-type YF infects multiple types of lymphoid cells, including DCs and Kupffer cells, then spreads to hepatocytes. YF 17D infects skin at site of inoculation, regional nodes, then reticulo-endothelial system (RES) and bone marrow. Wt dengue virus infects a large variety of cell types, including monocytes, macrophages, DCs and possibly endothelial cells. Chimeric YF/DEN vaccines infect DCs in vitro		[37–39]
3.4. Does the agent replicate in the nucleus?	No		
3.5. What is the risk of integration into the human genome?	None		
3.6. Does the agent establish a latent or persistent infection?	Persistent YF 17D virus infection of experimentally infected monkeys reported. No evidence in wild-type yellow fever of humans, but not specifically studied. One report of RNA genomes of YF 17D in urine of humans approx. 6 mos. after vaccination. Chronic infection with West Nile, Japanese encephalitis, and TBE have been described in animal models and in humans	IgM antibody lasting up to 18 months following 17D vaccination in one study suggested the possibility of persistent infection	[40–43]
3.7. How does the wild type agent normally transmit?	By agency of blood feeding <i>Aedes</i> and <i>Haemagogus</i> spp. mosquitoes	YF 17D vaccine incapable of infecting mosquitoes. Recombinants produced by the insertion of dengue 4, JE and WN prM-E genes into wild-type YF showed a significant decrease in infectivity for mosquito vectors compared to wild-type YF	[15,44–49]
3.8. What is known about the mechanisms of immunity to the wild type agent?	Both wild-type and vaccine induced immunity against future or repeated infection principally via neutralizing antibodies. Innate immune responses play an important role in early defense. CTLs responsible for clearing infection and recovery	Defects in innate immunity (interferon pathways) may underlie susceptibility of rare individuals to severe vaccine associated SAEs (YEL-AVD). Interferon receptor k/o mice develop viscerotropic disease	[50–52]
3.9. Is there treatment required and readily available for the disease caused by the wild type agent?	No. see cited review		[53]
4. Characteristics of proposed chimeric vaccines	Information	Comments/concerns	Reference(s)
4.1. What is the basis of attenuation/inactivation?	Chimerization (replacement of structural membrane and envelope genes (prM-E) with genes from another flavivirus). Evidence that chimerization itself contributes to attenuation also comes from experiments showing that insertion of prM-E genes from wt DENV 4 in a wt YF Asibi backbone results in decreased virulence. Multiple mutations in the YF 17D non-structural genes that occurred during empirical passage to develop the attenuated 17D vaccine. In some vaccine constructs, mutations were also inserted in the prM-E genes of the gene donor virus to decrease neurovirulence (WN vaccine). For constructing the chimeric JE vaccine, prM-E genes from the attenuated SA1414.2 vaccine were used, already containing some mutations as compared to wt JE virus		[39,54–58]
4.2. What is the risk of reversion to virulence or recombination with wild type or other agents?	Negligible due to replacement of entire PrM-E gene (chimerization), redundant mutations. See discussion of recombination in reference cited	Attenuated YF 17D and 17DD vaccines differ from wild-type YF in the genes encoding 20 amino acids. In addition there are 4 nT changes in the 3'UTR that also might affect attenuation. The prM-E sequences in the YF vaccine result in 8 amino acid changes. Accordingly, the chimeric vaccines lack the mutations in the prM-E genes of 17D virus. Evaluation of chimeric vaccine candidates versus parental YF 17D with respect to markers of attenuation in vitro and in vivo have consistently shown that the chimeric viruses are more attenuated than the parent. More than 29,000 humans subjects have received ChimeriVax vaccines without significant safety issues. Empirical studies have been performed to assess the biological consequences of non-homologous recombination of the chimeric vaccine viruses with prM-E or backbone sequences from virulent flaviviruses. These studies have shown that such recombinational events do not result in restoration of a virulent phenotype or a virus that has increased potential to disseminate via mosquito bite	[44,56,59–63]

Table 1 (Continued)

1. Basic information	Information		
4.3 Are yellow fever virus and chimeric viruses derived from them genetically stable?	YF 17D vaccine is an uncloned mixture (genetic swarm) as shown for all other flaviviruses. The 17D vaccine contains multiple diverse plaque populations and subpopulations of virions that react with wild-type specific YF monoclonal antibodies. In one fatal case of encephalitis caused by YF 17D vaccine, the virus isolated from brain contained two mutations that were associated with increased neurovirulence. In contrast, 17D isolates from cases of YEL-AVD have not shown mutations or selection of a specific subpopulation or variant associated with these adverse events. Overall, sequencing studies show that yellow fever 17D vaccines as manufactured using the seed lot system, and the chimeric constructs derived from them, have demonstrated low rates of mutation during <i>in vitro</i> and <i>in vivo</i> replication. Furthermore, YF appears to be more genetically stable than other RNA viruses. This may be due in part to the high fidelity of its polymerase. However, 27% of the yellow fever fusion peptide sequences deposited in GenBank show changes in this highly conserved region in contrast to 5% of other pathogenic flaviviruses suggesting that yellow fever virus is not more genetically stable than other flaviviruses at least in this small region	[64–72]	
4.3a. Are the chimeric vaccine candidates genetically stable during multiple passages?	Yes	All chimeric vaccine candidates have been assessed for genetic stability through serial passages. YF polymerase appears to be quite stable compared to other RNA viruses. ChimeriVax vaccines have all been made by molecular cloning and transfection of RNA into acceptable cell substrates to produce vaccine seed. The resulting seed stock is then biologically cloned (by plaque purification). Molecular and biological cloning promote genetic homogeneity of the vaccine virus. The actual vaccines are passed only a few times in cell cultures during manufacture. They are assessed for genetic stability on serial passage; after these viruses accumulate a few mutations related to adaptation to Vero cells, they have been quite stable on serial passages. Quality control procedures (sequencing) is used to monitor genetic stability of virus seeds and vaccine lots. Specifications for the consensus sequence are used for vaccine release	[39,57,68,70]
4.4. What is known about the genetic stability during <i>in vivo</i> replication?	Less empirical information, but no indication of instability, reversion to virulence	Serial <i>in vivo</i> passages in mice showed no reversion	[69]
4.5. Will a replication competent agent be formed?	The chimeric vaccines are replication competent		
4.6. What is the potential for shedding and transmission?	Considered unlikely, except for possible secretion in breast milk. The presence of virus titers as high as 6.2×10^9 genome equivalents/g in individuals with YEL-AVD increases the possibility of transmission	Chimeric YF/West Nile vaccine studied in horses; no evidence for shedding but vaccines intended for other species may need studies. Limited persistence and bio-distribution in monkeys after vaccination with Chimerivax-WN	[35,38,73,74]
4.7. Will the agent survive in the environment?	No		
4.8. Is there a non-human 'reservoir'?	Not for the attenuated vaccine virus		
4.9. Is there any evidence for or against safety during pregnancy?	See discussion of YF17D vaccine		
4.10. Can the vector accommodate multigenic inserts or will several vectors be required for multigenic vaccines?	See comments	Tetravalent YF/dengue vaccine required making 4 separate vectors and mixing in a formulation	[39,54]
4.11. What is known about the effect of pre-existing immunity on 'take', safety or efficacy in animal models?	No antivector immunity because prM-E (containing neutralizing epitopes) of YF replaced by corresponding genes of target vaccine virus. Prior YF immunity does not interfere. T cell responses to YF 17D non-structural proteins do not preclude effective immunization or re-use of vectors	Positive effect on chimeric dengue vaccine immunogenicity due to prior 17D immunity	See above reviews

Table 1 (Continued)

1. Basic information	Information		
		Comments/Concerns	Reference(s)
5. Manufacturing	Information		
5.1. Describe the source (e.g., isolation, synthesis)	Infectious cDNA clone of YF 17D vaccine virus Sanofi Pasteur – Swiftwater PA		
5.2. Describe the provenance of the vector including passage history and exposure to animal products	FDA licensed YF 17D vaccine	No fetal bovine serum (FBS) or animal products used in manufacturing process	
5.3. Can the vector be produced in an acceptable cell substrate?	Yes	Vero cells, WHO-87 cells, FDA bank deposited at ATCC	
5.4. Describe the proposed production process	prM-E of vector replaced with corresponding genes of virus against which immunity is desired	Chimeric RNA transfected into Vero cells to produce chimeric live attenuated vaccine	See above reviews
5.5. What are some Purity/Potential contaminants?	Virus purified from supernatant fluid by depth filtration, ultrafiltration, diafiltration		
Is there a large scale manufacturing feasibility?	Vaccines already made at commercial scale		
Are there any IP issues and is there free use of the vector?	Proprietary, vector system is covered by multiple patents	IP licensed by Acambis from St Louis University and NIH, and multiple new patents filed by Acambis. Acambis acquired by sanofi Pasteur in 2008	
6. Toxicology and potency (Pharmacology)	Information	Comments/Concerns	Reference(s)
6.1. What is known about the replication, transmission and pathogenicity in animals?	Multiple Good Laboratory Practice (GLP) toxicity studies performed in nonhuman primates (NHP) for chimeric vaccines against Japanese encephalitis, dengue, West Nile. The live chimeric viruses replicate in mice, hamsters, monkeys, cause transient viremia. Neurovirulence tests performed in infant mice and NHP, using licensed YF 17D vaccine as a reference control	In rodents and NHP, the chimeric vaccines were significantly less virulent than the licensed YF 17D vaccine	[55,57,75–77]
6.2. For replicating vectors, has a comparative virulence and viral kinetic study been conducted in permissive and susceptible species? (yes/no) If not, what species would be used for such a study? Is it feasible to conduct such a study?	Many such studies, see references		See above [54]
6.3. Does an animal model relevant to assess attenuation exist?	Reduced viremia in NHP infected with ChimeriVax-DEN versus Wt DEN viruses. For the encephalitides, e.g., JE and WN, rodents and NHP are excellent models in which to assess attenuation of neurovirulence. YF 17D is neurovirulent (and in infant mice neuroinvasive), whereas the chimeric vaccines are significantly attenuated. Specifications for release of seed viruses include demonstration of reduced neurovirulence in a GLP test in NHP	One of the difficulties in developing a dengue vaccine has been the absence of a convenient animal model. In nonhuman primates wild-type dengue causes a transient viremia, but no disease. A mouse model for dengue has been described, but since it involves immunocompromised mice, it can no be used to evaluate vaccine efficacy	See above [54,57,75,78]
6.4. Does an animal model for safety including immuno-compromised animals exist?	Unpublished studies in hamsters treated with cyclophosphamide showed no increase in virulence except for prolonged subclinical viremia. Chimeric JE vaccine was fully attenuated in type I/II IFN receptor KO mice (A129 or AG129)		[57,79]
6.5. Does an animal model for reproductive toxicity exist?	Yes, but not tested	Ongoing for chimeric dengue vaccine	
6.6. Does an animal model for immunogenicity and efficacy exist?	Hamsters, mice, horses, monkeys	See references and reviews	[80]
6.7. What is known about biodistribution?	Replicates in skin at site of inoculation, then draining nodes, then RES and bone marrow	Virus cleared after adaptive immunity established	[38]
6.8. Have neurovirulence studies been conducted?	Multiple	All lots of vaccine tested by sensitive neurovirulence test in infant mice. GLP neurovirulence studies in NHP for all new vaccine constructs, using licensed YF 17D as reference control	[75]
6.9. What is the evidence that the vaccines will generate a beneficial immune response in Rodent?	Rodents (mice, hamsters) immunized with single inoculation develop neutralizing antibodies and protected against challenge with wild-type virus corresponding to the inserted prM-E gene	Examples include Japanese encephalitis and West Nile	See above [54,57]

Table 1 (Continued)

1. Basic information	Information		
Non rodent? NHP?	Horses immunized and protected against West Nile Multiple studies showing immunogenicity and protection against challenge with dengue, JE, WN	Licensed vaccine (Prevenile®, Intervet) for horses	[77,79,81–84] [54,77,81,83]
Human?	The ChimeriVax-JE vaccine (Imojev®) was approved in Australia, Thailand and other Asian countries based on Phase 1–3 studies showing non-inferiority of neutralizing antibody responses to approved JE vaccine. The ChimeriVax-WN vaccine has been shown to elicit protective levels of neutralizing antibodies as well as strong T cell responses in Phase 1–2 clinical trials. A placebo-controlled phase 2b study of tetravalent ChimeriVax-DEN in Thai children conducted in a single site demonstrated protection against types 1, 3 and 4 but, despite the production of neutralizing antibodies, there was no observed protection from disease with type 2 dengue, the most prevalent circulating serotype. A subsequent phase 3 study in Asian children conducted in 5 countries and 11 sites found 56% overall protection against dengue fever (which met the proscribed study end-point), however efficacy against dengue 2 was lower than for other serotype. 35% protection against type 2 dengue was observed, compared to 50 to 78% protection against the other serotypes. Vaccine efficacy of against dengue hemorrhagic fever (DHF) was 88.5% (per protocol) and 67.2% against hospitalization (intent to treat). A phase 3 study in Latin America further confirmed the results obtained in Asia, with a 61% overall protection against dengue fever, with an efficacy of 50% against serotype 1, 42% against serotype 2, 74% against serotype 3 and 77% against serotype 4. A significant protection was also observed against severe disease and hospitalization. Multiple previous Phase 2 studies also suggested a similar level of efficacy	The Thai study of ChimeriVax-DEN with confirmation from an Asian and Latin American study, raises the possibility that protective efficacy is lower against type 2 dengue. However, protection against hospitalization and severe dengue involved all 4 serotypes. In the phase 3 studies there were no deaths caused by dengue in the control or vaccine groups. Accordingly the efficacy of the vaccine in preventing mortality remains to be studied	[38,39,45,57,85–89] Sanofi Pasteur, unpublished data, [90]
6.10. Have challenge or efficacy studies been conducted with HIV?	See 3.1a in immunocompromised		
Other diseases?	JE, dengue, WN (see refs)		
7. Previous human use	Please type one of the following: yes, no, Unknown, N/A (non-applicable)	Comments	Reference(s)
7.1. Has the vector already been used for targeting the disease of vector origin (measles, BCG, rabies)?	Yes, licensed vaccine against YF		
7.2. What is known about the replication, transmission and pathogenicity of the vector in: healthy people? Immunocompromised? neonates, infants, children? pregnancy and in the unborn? gene therapy experiments? any other susceptible populations?	See first section		
7.3. Is there any previous human experience with a similar vector including in HIV+ (safety and immunogenicity records)?	non-applicable		
7.4. Is there any previous human experience with present vector including HIV+ (safety and immunogenicity records)?	see 3.1a		
7.5. What is known about the effect of pre-existing immunity on 'take', safety or efficacy in any human studies with this or different insert?	prior administration of 17D does not interfere with the chimeric vaccines		

Table 1 (Continued)

1. Basic information		Information		
7.6. Name some other non-HIV vaccines using same vector and describe some of the public health considerations.	none			
8. Overall Risk Assessment	Describe the toxicities	Please rate the risk as one of the following: none, minimal, low, moderate, high, or unknown	Comments	Reference(s)
8.1. What is the potential for causing serious unwanted effects and toxicities in: Healthy people?	Multiple trials of chimeric YF with JE, dengue, WN gene inserts have not shown any significant safety issues	The chimeric JE vaccine (Imojev®) is licensed, the chimeric dengue vaccine is in Phase III and the WN vaccine is in Phase II. A veterinary vaccine against WN is licensed		See previous and [91]
Immunocompromised?	No data			
Neonates, infants, children?	Trials of the chimeric JE and dengue vaccines have been conducted in children 2 years and older, and have shown them to be safe and well tolerated.			
Pregnancy and in the unborn?	No data. Parental YF 17D (and therefore ChimeriVax) not be administered during pregnancy unless clearly required, based on a high risk exists of natural infection			
Other susceptible populations?				
8.2. What is the risk of neurotoxicity/neuroinvasion or cardiac effects?	The vaccines are significantly less neurovirulent than the licensed YF 17D vaccine			[75]
8.3. What is the potential for shedding and transmission in risk groups?	YF 17D rarely transmitted to infants via breast milk Viremia titers as high as 6.2×10^9 genome equivalents/g in individuals dying with YEL-AVD raise the possibility of transmission by blood but likelihood of adverse event low since the virus in such individuals has few/no mutations	However, multiple studies have shown that the chimeric vaccines are not infectious for mosquitoes		[45–49,74]
8.4. What is the risk of adventitious agent (including TSE) contamination?	Minimal. Seed viruses and each vaccine lot are tested for adventitious viruses. No animal products used in manufacturing.			
8.5. Can the vector be manufactured at scale in an acceptable process?	Yes, the JE vaccine is at commercial scale now; Sanofi Pasteur has constructed a commercial factory for the dengue vaccine	10,000–50,000 doses per L—very efficient process		
8.6. Can virulence, attenuation and toxicity be adequately assessed in preclinical models?	Yes			
8.7. Rate the evidence that a beneficial response will be obtained in humans.	Already known—neutralizing antibodies in >95% of human subjects. T cell responses to the E protein conserved in humans	The dengue vaccine contains a mixture of 4 separate viruses each representing 1 serotype. Interference between the individual viruses is observed, so that 2–3 doses are required for complete immunization. In contrast the JE and WN vaccines are monovalent and a single dose provides durable immunity and T cell memory. The proposed correlate of protection for chimeric vaccines against encephalitic viruses (JE, WN) is the level of neutralizing antibodies (PRNT50 titer), and a level of >10 is considered protective; the chimeric vaccines elicit neutralizing antibody titers in great excess over this minimum protective level. However, an immune correlate is not established yet for dengue vaccines. Cellular immunity may also contribute to protection, and it has been shown that chimeric vaccines are able to induce significant responses in this respect.		[39,57,91–93]
9. Adverse Effect Assessment	Describe the adverse effects	Please rate the risk as one of the following: none, minimal, low, moderate, high, or unknown	Comments	Reference(s)
9.1. Describe the adverse effects observed				
Mild local reactions	Erythema, pain	Minimal		
Mild systematic reactions	Similar to placebo			
Moderate local reactions	Erythema, pain	Minimal		
Moderate systematic reactions	Similar to placebo			
Severe local reactions		None		
Severe systematic reactions		None		
10. Administration Assessment	Information	Comments/Concerns		Reference(s)

Table 1 (Continued)

1. Basic information	Information
10.1. What is the average Tissue Culture Infections Dose per millimeter (TCID/ml)?	Human dose is 3–5 log ₁₀ plaque forming units (PFU).
10.2. What is the highest TCID/ml that can be used before cell toxicity?	The viruses cause CPE in vitro at low multiplicities of infection (MOI).
10.3. Are different demographics affected differently?	No information
References	Information

wild-type virus in chicken embryo cultures resulting in the acquisition of nucleotide mutations some of which do not cause changes in the translated amino acids (synonymous) and some of which do (non-synonymous). In the YF-17D vectored vaccines that are being developed for Japanese encephalitis, West Nile and the four serotypes of dengue viruses, the coding regions of the prM-E proteins of the corresponding viruses are substituted for those in the YF 17D virus vaccine, resulting in a chimeric virus. The YF-17D viral genome excluding the prM-E genes is referred to as the “backbone”. Although the production of such chimeric viruses alone may decrease their virulence, preclinical studies have demonstrated that the YF backbone has sufficient mutations by itself to maintain vaccine attenuation of chimeric vaccines bearing the dengue surface antigens [22]. Another aspect of the chimeric flavivirus technology that is distinguished from other viral vectors, is that anti-vector immunity is not a significant problem for flavivirus vaccine development. This feature is due to the fact that the prM-E region is solely responsible for generating epitopes recognized by neutralizing antibodies. Thus, the chimeric vector contains the only neutralizing antigens of the intended target for immunization. Previous immunization with YF17D and T cell responses to the YF 17D backbone are insufficient to prevent effective immunization with a chimeric vector expressing a heterologous flavivirus prM-E transgene.

The efforts of the V3SWG were focused initially on the above flavivirus vaccines. Not addressed are additional possible vaccines in which nucleotide sequences encoding for epitopes of other microorganisms might be added into the complete YF virus vaccine yielding live virus vaccines with additional vaccination potential. Vaccines employing other methods of attenuating flaviviruses through mechanisms such as nucleotide deletions and which do not involve a viral vector are not considered in this project [25].

The chimeric vaccines indicated for the prevention of dengue, Japanese encephalitis, and West Nile will be considered new entities from the regulatory perspective, and will need to be independently assessed for safety and efficacy. However, in the development of these new vaccines, the parental YF 17D vaccine virus has provided an important comparator and benchmark in all preclinical and many clinical trials. For example, the monkey neurovirulence test is an important measure of safety of YF 17D vaccines, and YF 17D was used as the reference strain in many studies, which showed that the chimeric vaccines were more attenuated than parental YF 17D. An important regulatory question thus arises as to whether the age range for vaccination, precautions and contraindications in labelling for use of YF 17D vaccines should apply to the new, chimeric vaccines. This question is particularly important since data on very rare adverse events will likely not be available at the time the new vaccines are approved. The template supplied in this paper contains information that can potentially be useful in considering how new chimeric vaccines should be described in reference to parental YF 17D vaccines.

1.3. Methods for developing, completing and reviewing the standardized template

Following the process described in the accompanying overview paper [26] as well as on the Brighton Collaboration website (<http://cms.brightoncollaboration.org:8080/public/what-we-do/setting-standards/case-definitions/process.html>), the Brighton Collaboration V3SWG was formed in October 2008 and includes ~15 members with clinical, academic, public health, regulatory and industry backgrounds with appropriate expertise and interest. The composition of the working and reference group as well as results of the web-based survey completed by the reference group with subsequent discussions in the working group can be viewed at <http://www.brightoncollaboration.org/internet/en/index/working-groups.html>. The workgroup meets via emails and monthly conference calls coordinated by a secretariat [currently at CDC's Division of HIV/AIDS Prevention].

The V3SWG invited a flavivirus expert, Thomas P. Monath (TPM), who has been intimately associated with the development of flavivirus vaccines based on the YF virus vaccine backbone to complete the template in 2011. The first draft was then critiqued by a member of the working group knowledgeable about flaviviruses, Stephen J. Seligman (SJS), moderated by another member, James S. Robertson (JSR), discussed by the V3SWG as a whole, and then peer reviewed by reference groups (e.g., American Society of Virology, American Society of Tropical Medicine and Hygiene) and the Brighton Collaboration membership. Bruno Guy updated the template with new information as of October, 2014. Sections 8 (overall risk assessment) and 9 (adverse effect assessment) of the template seeks to rate the risk of the viral vector in various situations as: none, minimal, low, moderate, high, or unknown. An initial assessment was made by TPM and then reviewed by others, based largely on the anticipated frequency and severity of the vaccine adverse event versus the expected frequency and severity of the target vaccine preventable disease [15]. Depending on the season, the risk of the yellow fever vaccine approaches the risk of YF in S. America. In Africa the risk of yellow fever is usually much greater than the risk of the vaccine. The V3SWG may develop more explicit criteria for standardizing the rating of these risks in the future with further experience.

The resulting template is submitted as a guideline for evaluating the current issues in development of vaccines based on the yellow fever virus vaccine backbone.

2. Standardized template (Table 1)

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Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at <http://dx.doi.org/10.1016/j.vaccine.2014.10.004>.

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