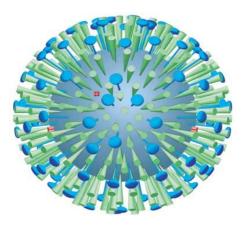
Nanoparticulate Vaccine Design: The VesiVax[®] System



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Influenza

- Each year up to 20% of the world's population contracts influenza
- 250,000 to 500,000 people die annually from influenza-associated complications
- Pacific Bridge Life Sciences estimates the associated cost of influenza at \$5.2B in 2004
- "Avian influenza" mortality rates greater than 50%



Influenza Virus Type A

Enveloped RNA Virus Genome encodes 10 proteins Major Viral Envelope Proteins Neuraminidase - Hemagglutinin (H) - Neuraminidase (N) Hemagglutinin **RNA** M2 protein



Survival Strategies Employed by the Influenza Virus

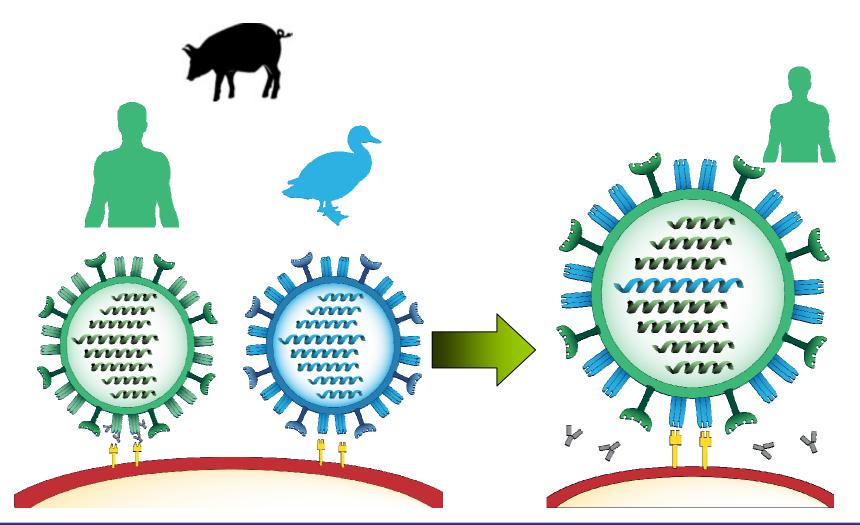
Antigenic Drift

- High mutation rates
- RNA viruses lack proofreading capabilities
 - Often one mutation per genome copy
 - Evolutionary advantage
 - Active response to changes in environment and drug regimen
- Antigenic Shift
 - Shuffling of viral genes gives rise to "reassortants"
 - Recombination of H and N creates new strains
- Infective for multiple hosts
 - Humans, pigs, birds, horses, dogs, mice
- Hardy

 Able to survive and retain virulence for up to 48 hours on hard non-porous surfaces



Antigenic Shift





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Influenza Vaccines

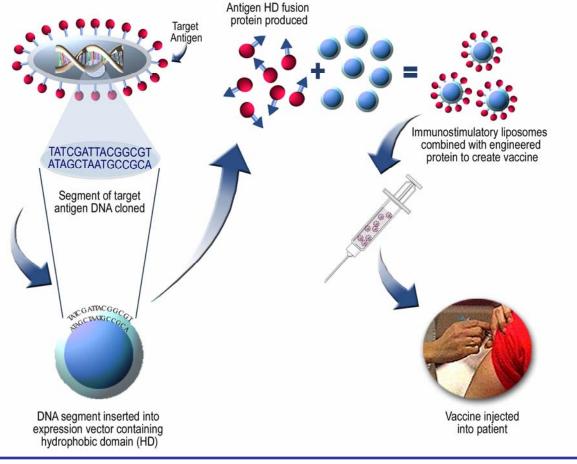
Time-intensive production process

- Generates inefficiencies that gives the virus an advantage
- Specific strains and virulence must be forecasted and produced well in advance of each flu season
- There is no feedback loop in this process, when the forecasts are inaccurate, supply chain is already committed, and course corrections can not be made
- Because the influenza virus changes constantly, clinical evaluation of new vaccines is impractical



The VesiVax[®] System

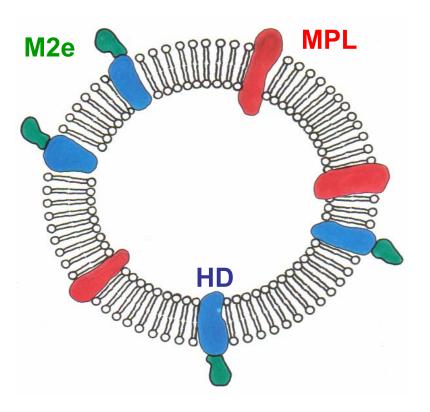
Designed to facilitate rapid vaccine development -





VesiVax[®] influenza vaccine targets the highly conserved M2 ectodomain segment (M2e)

- HD = Hydrophobic domain
- M2e = Antigen (~100/Liposome)
- **MPL** = Adjuvant (~2500/Liposome)





Vaccine Design

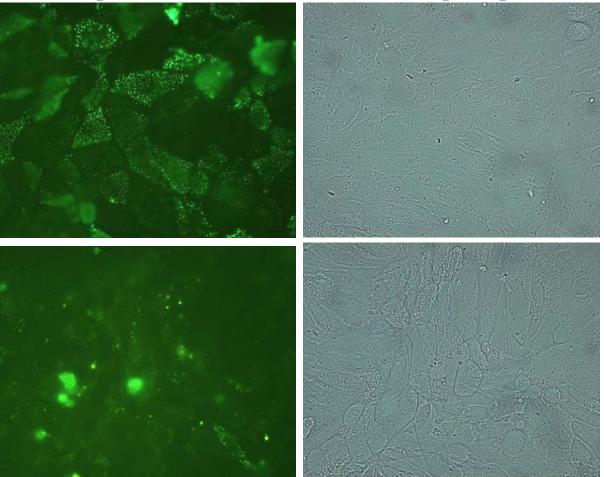
Influenza type A M2e sequences

Subtype	M2eA Sequence	Year
H1N1	MSLLTEVETPTRNEWGCRCNDSSD	1918 – "Spanish″
H1N1	MSLLTEVETPIRNEWGCRCNGSSD	1934 - PR/8
H2N2	MSLLTEVETPIRNEWGCRCNDSSD	1957 – "Asian"
H3N2	MSLLTEVETPIRNEWGCRCNDSSD	1968 – "Hong Kong"
H5N1	MSLLTEVETLTRNGWECKCRDSSD	1997 – "Avian"
H9N2	MSLLTEVETPTRNGWECKCNDSSD	1999 – "Avian"
H5N1 H6N2	MSLLTEVETPTRNEWECRCSDSSD MSLLTEVETPIRNEWGCRCNDSSD	2004 – "Avian" X-88

H5N1- First evidence of influenza virus transmitted from birds to humans. It is important to note that the majority of Influenza type A strains have high sequence homology for M2e. The conserved nature of M2e allows for the potential to create a vaccine that is effective against all strains.

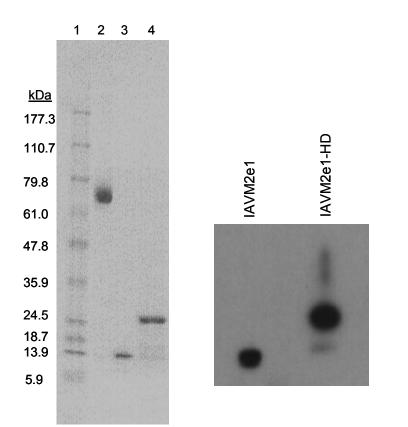


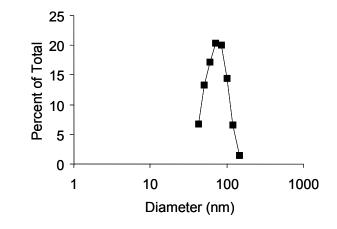
VesiVax[®] Influenza Vaccine Virally Infected Cells Display M2e

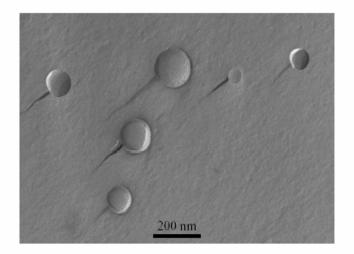




Preparation of L-M2e-HD

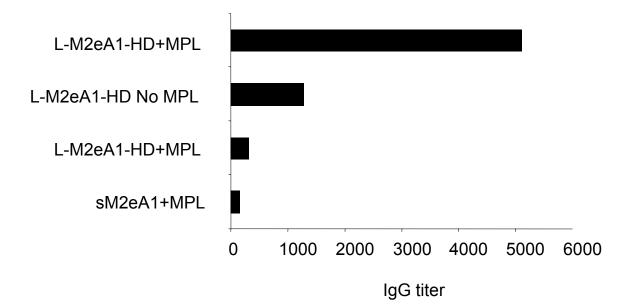








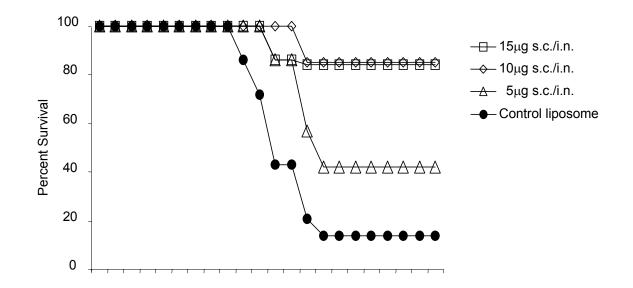
Preliminary Formulation Screen



BALB/c mice (n=5) Immunized twice (SubQ/IN)



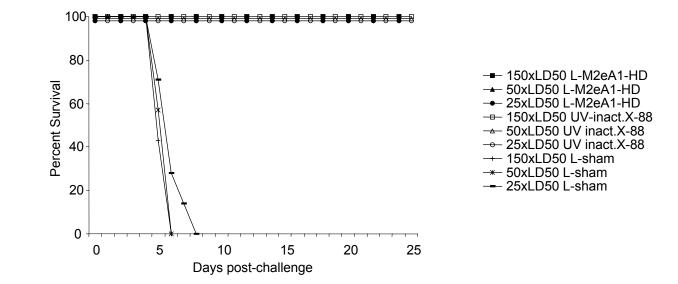
Dose Ranging Study



BALB/c mice (n=7) Immunized twice (SubQ/IN) Challenged with 10X LD50



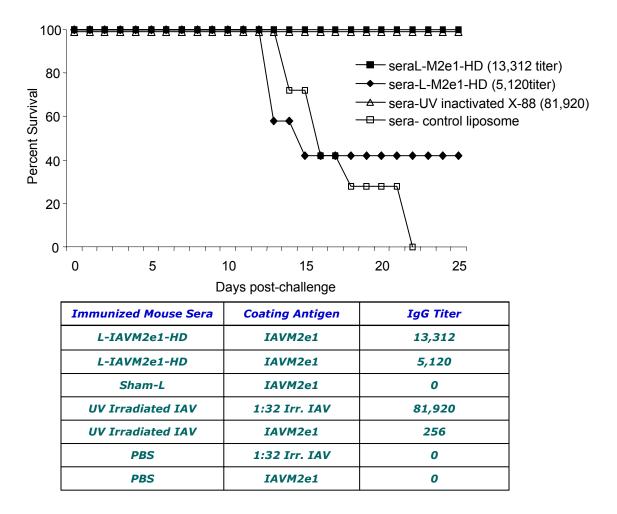
Maximal Viral Challenge



BALB/c mice (n=7) Immunized twice (SubQ/IN) Challenged with X-88

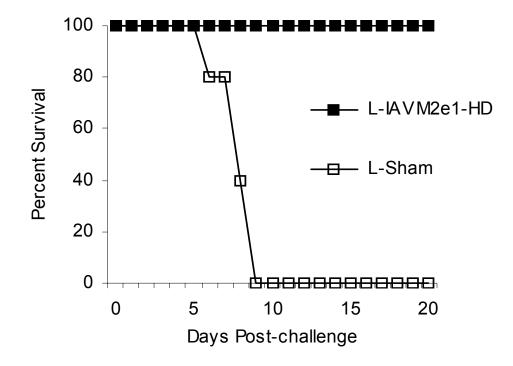


VesiVax[®] Influenza Vaccine Passive Transfer of Immunity





Cross-Protection of M2e





Reduction of Viral Burden

Immunization	M2e1	M2e2	M2e3
Challenge	H1N1	H5N1	H9N2
N-fold Reduction	>300	>10	>50



M2 as a target for vaccine development

- Evolutionarily conserved
- Not as susceptible to the high mutation and reassortment rates observed with the H and N epitopes
- Present on the surface of viruses and infected cells

Is M2 a good flu vaccine target?

- Significant protection observed for epidemic and pandemic strains of influenza
- Cross protection against strains with the same M2 demonstrated
- The data suggests that M2 mediated immunity is antibody dependent
- M2 specific antibodies recognize M2 on the surface of infected cells and on the virus



VesiVax[®] Vaccination Studies

Protection from severe infection observed

Against viral and bacterial pathogens In different species and strains of animals In both sexes In short and long term studies In adults and young animals Through different routes of vaccination

Assays of immunological response parameters demonstrate

Antibody titers increase >30x over placebo Antigen specific proliferation of immune cells increase >10x Key cytokine levels increase by >10x over placebo

No significant side effects observed



The VesiVax[®] Advantage

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VesiVax[®] Influenza Vaccine

- Recombinant DNA system allows "cut & paste" design of M2e antigens
- Flexible design facilitates rapid engineering of new influenza vaccines
- Routine scale-up procedure
- Production simplified
- Minimal biohazard (BL1)
- Selective antigen display (M2e)
- Reduced possibility of side effects
- No risk of infection

Influenza Virus Vaccine

- Pathogen-based vaccines are not amenable to rapid development
- Time and labor intensive manufacturing process
- Complex production procedures (eggs)
- Biohazard requirements (BL2-BL4)
- Non-selective antigen display
- Inflammation at the site of injection
- Increased possibility of clinical complications



Implications

VesiVax® technology

- Represents a leap forward in vaccine development and production
- Demonstrated efficacy with Influenza
- Demonstrated efficacy with HSV2
 - Significantly shortens time of vaccine production
 - Can potentially respond to new pathogens in weeks, not months



Acknowledgements

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