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- Genetic analysis is a powerful approach toward understanding the structure and function <u>of the viral genome</u>, its <u>gene</u> <u>products</u>, and <u>their roles</u> in infection and disease.
- Variation in viral properties is of *great importance for human medicine*:
- Viruses that have stable antigens on their surfaces (poliovirus, measles virus) can be controlled by vaccination,
- Other viruses that exist as many antigenic types (rhinoviruses),
- Or change frequently (influenza virus A) are difficult to control by vaccination.
- Viral genetics analysis may help to develop more effective vaccines and *antiviral therapy*.



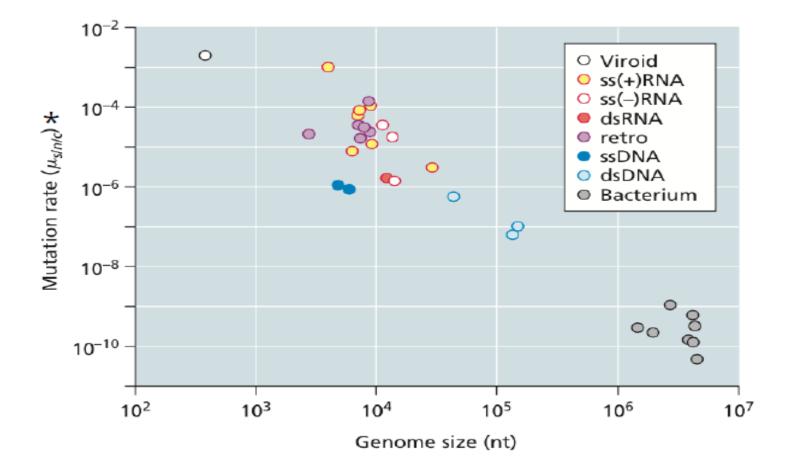
- The following terms are basic to a discussion of genetics:
- ✓ The **genome** is the sum of the genes of an organism.
- ✓ **Genotype** refers to the genetic constitution of an organism.
- Phenotype refers to the observable properties of an organism, which are produced by the genotype in cooperation with the environment and other factors such as epigenetic factors
- Wild-type virus denotes the original virus from which mutants are derived and with which the mutants are compared.
- ✓ A **mutation** is a heritable change in the genotype.
- Defective Viruses A defective virus is one that lacks one or more functional genes required for viral replication.





- The study of viral genetics falls into two general areas:
- (1) mutations and their effect on replication and pathogenesis
- (2) the interaction of two genetically dis-
- tinct viruses that infect the same cell.

## Genome size and mutation Rate



## **Types of Virus Mutants**

Some markers commonly used FOR **DETECTION OF MUTATIONS** include: plaque morphology, antibody escape or resistance to neutralizing antisera, loss of a virus protein, drug resistance, host range, and inability to grow at low or high temperatures (Conditional-lethal mutants). • Defective Viruses

## **Types of Virus Mutants**

- Mutations that inactivate essential genes are termed lethal mutations. These mutants are difficult to isolate because the virus cannot replicate.
- A **deletion mutant** results from loss or selective removal of a portion of the genome and the function it encodes.
- Other mutations may produce **plaque mutants**, which differ from the **wild type** in the size or appearance of the infected cells;
- **host range mutants, which differ in the tissue type or species of target** cell that can be infected;
- **attenuated mutants, which** are variants that cause less serious disease in animals or humans.
- Conditional mutants, such as temperature-sensitive (ts) or cold-sensitive mutants,
- have a mutation in a gene for an essential protein that allows virus production only at certain temperatures. Whereas ts mutants generally grow well or relatively better at 30° C to 35° C, the encoded protein is inactive at elevated temperatures of 38° C to 40° C, preventing virus production.
- Live virus vaccines are often conditional or host range mutants and attenuated

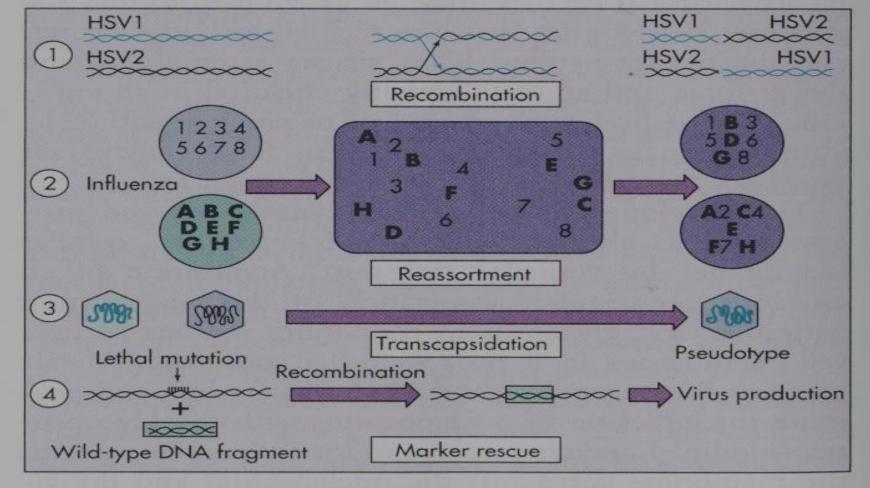
for human disease.



## INTERACTIONS

• When two genetically distinct viruses infect a cell, different phenomena can ensue. (1) Recombination (2) Genetic Reactivation - Multiplicity reactivation - Marker rescue (3) Re assortment (4) Phenotypic mixing (5) Complementation

(6) Interference



**FIGURE 6–15.** Genetic exchange between viral particles can give rise to new viral types, as illustrated. Representative viruses include the following: 1, intertypic recombination of herpes simplex virus type 1 (HSV1) and type 2 (HSV2); 2, reassortment of two strains of influenza virus; 3, rescue of a papovavirus defective in assembly by a complementary defective virus (transcapsidation); and 4, marker rescue of a lethal or conditional mutation.

#### Interactions Among Viruses:

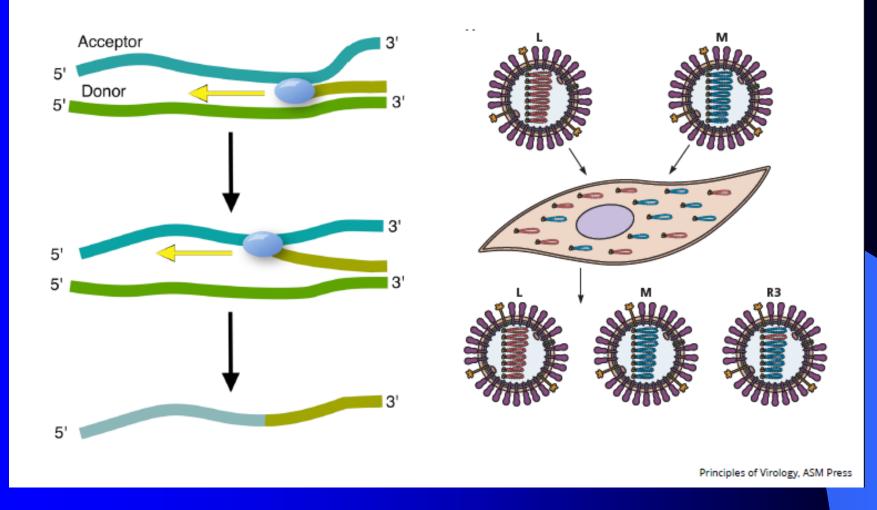
When two or more virus particles infect the same host cell, they may interact in a variety of ways. They must be sufficiently closely related, usually within *the same viral family*, for most types of interactions to occur:

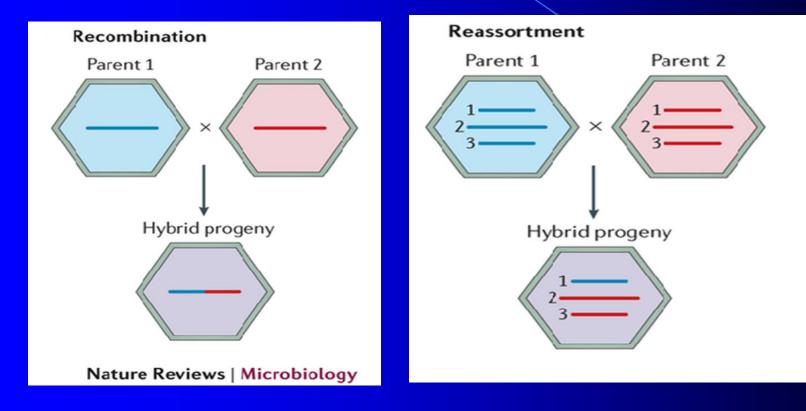
- <u>Genetic interaction</u> results in some progeny that are heritably (genetically) different from either parent.
- *A) Recombination:* Recombination results in the production of progeny virus (**recombinant**) that carries traits not found together in either parent.

B) **Reassortment**: In the case of viruses with *segmented genomes*, eg, influenza virus, the formation of recombinants is due to reassortment.



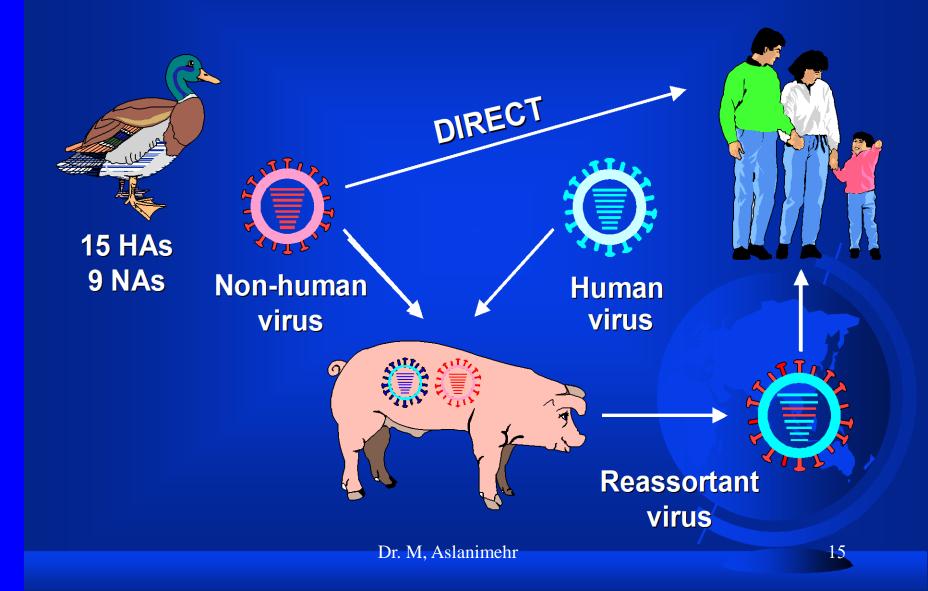
### Variation further generated by recombination and reassortment



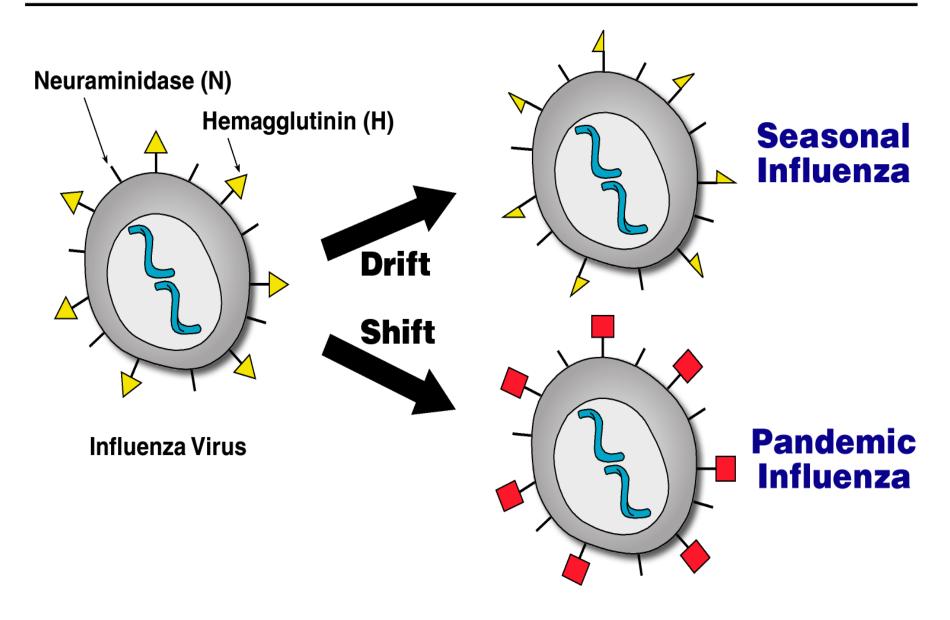


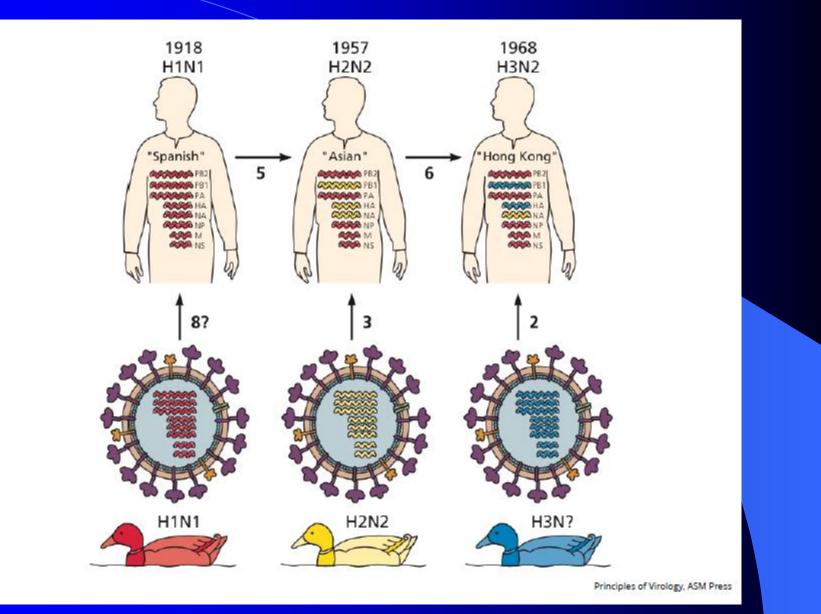


## **Mechanism of antigenic shift**



## **Influenza: Antigenic Drift and Shift**





- C. Complementation: This refers to the interaction of viral gene products in cells infected with two viruses, *one or both* of which *may be defective*.
- It results in the replication of one or both under conditions in which replication would not ordinarily occur. The basis for complementation is that one virus provides a gene product in which the second is defective, allowing the second virus to grow.
- D. Phenotypic Mixing: This occurs when the genome of one virus becomes randomly incorporated within capsid proteins specified by a different virus or a capsid consisting of components of both viruses.
- If the genome is encased in a *completely heterologous protein coat*, this extreme example of phenotypic mixing may be called "*phenotypic masking*" or "**transcapsidation**."



*E) Interference:* Infection of either cell cultures or whole animals with two viruses often leads to an inhibition of multiplication of one of the viruses, an effect called interference.

Several *mechanisms* have been elucidated as causes of interference:

- (1) One virus may inhibit the ability of the second to adsorb to the cell, either **by blocking its receptors** (retroviruses, enteroviruses) or by **destroying its receptors** (orthomyxoviruses).
- (2) One virus may **compet**e with the second for components of the replication apparatus (eg, polymerase, translation initiation factor).
- (3) The first virus may cause the infected cell to produce an inhibitor (**interferon**) that prevents replication of the second virus.

