JOURNAL OF MICROBIOLOGY & BIOLOGY EDUCATION, May 2016, p. 300-301 DOI: http://dx.doi.org/10.1128/jmbe.v17i2.1096



# **Supplemental Materials**

for

## **Modeling Influenza Antigenic Shift and Drift with LEGO Bricks**

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### **Table of Contents**

(Total pages 4)

Appendix 1: List of materials and instructor's notes

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### Appendix 1: List of materials and instructor's notes.

#### List of Materials

	Item	Number of pieces	Source
1	Celebrate it <sup>TM</sup> Gift box	1	Michaels arts and crafts store http://www.michaels.com/search?q=346385
2	1X2/red LEGO bricks	4	
3	1X4/red LEGO bricks	4	
4	1X6/red LEGO bricks	4	Online LEGO® shop
5	1X2/blue LEGO bricks	1	http://shop.lego.com/en-US/Pick-A-Brick-ByTheme
6	1X4/blue LEGO bricks	1	
7	1X6/blue LEGO bricks	1	

#### Modeling Antigenic Shift with Lego® Bricks

Instructor's notes



Figure 1. Antigenic Shift Exercise: Manipulatives and Outcomes.

Antigenic shift is modeled using i) a non-transparent box (panel A) as an infected host cell and ii) two sets of Lego® bricks with different color (panel B) representing the genomes of distinct Influenza strains replicating with the same rate. Single-row LEGO® bricks (to emphasize ssRNA genome) with two, four and six studs work the best. It is straightforward to purchase the needed pieces using the "Pick a Brick by Theme" option of the online LEGO® store. Total cost of 1 genome is under 1\$. The box shown in panel A was purchased from Michael's crafts store, also for under one dollar. Students are asked to package 10 viruses by picking Lego® bricks based on their size without peeking in the box. The genetic makeup of each packaged virus is recorded as non-recombinant (NR) if all selected Lego® bricks are the same color or recombinant (R), if the colors are mixed (panel C). Lego® bricks are returned in the box before the next virus is assembled. Class results are tallied in a table and analyzed.

#### Modeling Antigenic Drift with Lego® Bricks

Instructor's notes



Figure 2. Antigenic Drift Exercise: Manipulatives and Outcomes.

Each pair of students is given three model Influenza genomes of the same color (panel A) and a marker (not shown). Students are asked to keep one genome as a reference for the genetic makeup of the virus that initially infected the cell (panel B–top) and to generate two genome copies taking into account the error prone nature of the RNA polymerase copying RNA. It is assumed that the enzyme introduces on average 3 mutations per every10 nucleotides copied. Mutations are visualized by coloring studs of the Lego® bricks (panel B). Class results are displayed together to demonstrate the vast diversity of Influenza viruses that can arise.