

Video Article

Production of Pseudotyped Particles to Study Highly Pathogenic Coronaviruses in a Biosafety Level 2 Setting

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URL: <https://www.jove.com/video/59010>

DOI: [doi:10.3791/59010](https://doi.org/10.3791/59010)

Keywords: Pseudotyped particle, pseudovirion, coronavirus, CoV, spike protein, severe acute respiratory syndrome coronavirus, SARS-CoV, Middle East respiratory syndrome coronavirus, MERS-CoV, murine leukemia virus, MLV

Date Published: 10/31/2018

Citation: Millet, J.K., Tang, T., Nathan, L., Jaimes, J.A., Hsu, H.L., Daniel, S., Whittaker, G.R. Production of Pseudotyped Particles to Study Highly Pathogenic Coronaviruses in a Biosafety Level 2 Setting. *J. Vis. Exp.* (), e59010, doi:10.3791/59010 (2018).

Abstract

The protocol aims to generate coronavirus (CoV) spike (S) fusion protein pseudotyped particles with a murine leukemia virus (MLV) core and luciferase reporter, using a simple transfection procedure of the widely available HEK-293T cell line. Once formed and released from producer cells, these pseudovirions incorporate a luciferase reporter gene. Since they only contain the heterologous coronavirus spike protein on their surface, the particles behave like their native coronavirus counterparts for entry steps. As such, they are the excellent surrogates of native virions for studying viral entry into host cells. Upon successful entry and infection into target cells, the luciferase reporter gets integrated into the host cell genome and is expressed. Using a simple luciferase assay, transduced cells can be easily quantified. An important advantage of the procedure is that it can be performed in biosafety level 2 (BSL-2) facilities instead of BSL-3 facilities required for work with highly pathogenic coronaviruses such as Middle East respiratory syndrome coronavirus (MERS-CoV) and severe acute respiratory syndrome coronavirus (SARS-CoV). Another benefit comes from its versatility as it can be applied to envelope proteins belonging to all three classes of viral fusion proteins, such as the class I influenza hemagglutinin (HA) and Ebola virus glycoprotein (GP), the class II Semliki forest virus E1 protein, or the class III vesicular stomatitis virus G glycoprotein. A limitation of the methodology is that it can only recapitulate virus entry steps mediated by the envelope protein being investigated. For studying other viral life cycle steps, other methods are required. Examples of the many applications these pseudotype particles can be used in include investigation of host cell susceptibility and tropism and testing the effects of virus entry inhibitors to dissect viral entry pathways used.

Introduction

Host cell entry constitutes the initial steps of the viral infectious life cycle. For enveloped viruses, this involves binding to a single host cell receptor or several receptors, followed by fusion of viral and cellular membranes. These essential functions are carried out by viral envelope glycoproteins^{1,2}. The coronavirus envelope glycoprotein is called the spike (S) protein and is a member of the class I viral fusion proteins^{2,3,4,5,6}. Studying viral envelope glycoproteins is critical for understanding many important characteristics of a given virus, such as: lifecycle initiation, its host and cellular tropism, interspecies transmission, viral pathogenesis, as well as host cell entry pathways. Viral pseudotyped particles, also named pseudovirions, are powerful tools that enable us to easily study the function of viral fusion proteins. Pseudotyped particles or pseudovirions are chimeric virions that consist of a surrogate viral core with a heterologous viral envelope protein at their surface. The protocol's main purpose is to show how to obtain coronavirus spike pseudotyped particles that are based on a murine leukemia virus (MLV) core and contain a luciferase reporter gene. As examples, the method to produce pseudotyped particles with the spike proteins of the highly pathogenic severe acute respiratory syndrome (SARS) and Middle East respiratory syndrome (MERS) coronaviruses are presented. The protocol describes the transfection procedure involved, how to infect susceptible target cells, and infectivity quantification by luciferase assay.

Since the entry steps of the pseudovirions are governed by the coronavirus S at their surface, they enter cells in a similar fashion to native counterparts. As such, they are excellent surrogates of functional infectivity assays. Pseudotyped particles are usually derived from parental model viruses such as retroviruses (MLV^{7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22}) and the lentivirus human immunodeficiency virus - HIV^{23,24,25,26,27,28,29,30,31,32,33,34,35}) or rhabdoviruses (vesicular stomatitis virus - VSV^{36,39,40,41,42,43,44,45,46,47}). When used in pseudotyping, the parental viruses' genomes are modified to remove essential genes, rendering them defective for accomplishing a complete replication cycle. This feature allows them to be used in intermediate biosafety level facilities (BSL-2) and is an important advantage over using highly pathogenic native viruses that require higher biosafety facilities (BSL-3, BSL-4 which are not as readily available) when conducting virus entry studies. Here, the S proteins of risk group 3 pathogens SARS-CoV and MERS-CoV are used as examples of viral envelope proteins being incorporated into MLV pseudotyped particles, generating SARS-CoV S and MERS-CoV S pseudovirions (SARS-Spp and MERS-Spp, respectively). These pseudovirions have been successfully used in studies focusing on entry events of these viruses^{48,49,50,51}. Another advantage is that the technique described here is not limited to pseudotyping coronavirus S proteins: it is very flexible and can be used to incorporate representatives of all three classes of viral fusion proteins. Examples include influenza hemagglutinin (HA, class I)⁵², Ebola virus glycoprotein (GP class I), E1 protein of the

alphavirus Semliki Forest virus (SFV, class II), and VSV glycoprotein (G, class III)⁵³. In addition, more than one kind of viral glycoprotein can be co-incorporated into a pseudotyped particle, as in the case of influenza HA- and NA- pseudotyped particles⁵¹.

Based on the work performed by Bartosch et al.²⁰, this protocol describes the generation of MLV pseudotyped particles with a three-plasmid co-transfection strategy using the widely available and highly transfection-competent HEK-293T cell line⁵⁴. The first plasmid encodes the MLV core genes *gag* and *pol* but lacks the MLV envelope *env* gene. The second plasmid is a transfer vector that encodes a firefly luciferase reporter gene, an MLV Ψ -RNA packaging signal, along with 5'- and 3'-flanking MLV long terminal repeat (LTR) regions. The third plasmid encodes the fusion protein of interest, in this case either the SARS-CoV S or MERS-CoV S protein. Upon co-transfection of the three plasmids using a transfection reagent, viral RNA and proteins get expressed within transfected cells allowing generation of pseudotyped particles. Since MLV is used as pseudovirion backbone, this occurs at the plasma membrane: the RNAs containing the luciferase gene reporter and packaging signal get encapsulated into nascent particles that also incorporate plasma membrane-expressed coronavirus spike proteins. The particles that bud out from cells contain the coronavirus S protein at their surface and are harvested for use in infectivity assays. Because pseudotyped particles harbor the coronavirus S protein and not the MLV envelope protein, when used for infecting cells, they behave like their native coronavirus counterparts for entry steps. The viral RNA containing the luciferase reporter and flanking LTRs is then released within the cell and the retroviral polymerase activities enable its reverse transcription into DNA and integration into the host cell genome. Quantification of the infectivity of viral pseudotyped particles in infected cells is then performed with a simple luciferase activity assay. Because the DNA sequence that gets integrated into the host cell genome only contains the luciferase gene and none of the MLV or coronavirus protein-encoding genes, they are inherently safer to use than replication-competent native viruses.

In addition to being safer surrogates and highly adaptable to allow incorporation of various kinds of envelope glycoproteins, the pseudotyped particles described here are also highly versatile and can be used to study many aspects of virus entry. This includes but is not limited to: testing host cell susceptibility to virus infection, analyzing the cellular entry pathways an enveloped virus uses, studying the effects of pharmacological inhibitors and drug screenings, conducting neutralization antibody assays, characterizing host cell entry of enveloped viruses that cannot be cultured, and generating viral vectors for gene delivery, stable cellular expression of genes of interest, or gene silencing.

Protocol

1. Cell seeding for pseudotyped particle production

NOTE: Perform this step in the biosafety cabinet.

- By standard cell culture techniques, obtain an 80-90% confluent 75 cm² flask of HEK-293T/17 cells passaged in complete Dulbecco's Modified Eagle's Medium (DMEM-C) containing 10% (vol/vol) fetal bovine serum (FBS), 10 mM 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid (HEPES), 100 IU/mL penicillin, and 100 µg/mL streptomycin. Prepare DMEM-T medium for transfections (its composition is the same as DMEM-C but without antibiotics).
- Wash cells with 10 mL of pre-warmed (37 °C) Dulbecco's Phosphate Buffered Saline (DPBS) twice.
NOTE: Handle HEK293T/17 cells with care as they easily detach.
- Aspirate the supernatant and detach cells with 1 mL of 0.25% trypsin solution pre-warmed at 37 °C. Place the flask of cells at 37 °C, 5% CO₂ incubator for 3–5 min or until cells start detaching.
NOTE: Avoid incubating cells with trypsin for more than 5 minutes as this typically leads to cell clumping.
- Deactivate trypsin by adding 4 mL of DMEM-C medium and count cells using a cell counting slide and light microscope.
NOTE: To avoid having to count too many cells, an additional dilution step may be required beforehand. Remember to factor in this dilution when calculating the actual cell density of trypsinized cells.
- Dilute cells to 5 × 10⁵ cells/mL with DMEM-C.
- Seed 6-well tissue culture plate with 2 mL of cell solution per well and gently move the plate back and forth and side to side to evenly distribute cells, avoiding circular motion.
NOTE: This is a key step. Evenly distributed cells will ensure that cells do not clump at the center of wells. In turn, this will ensure good transfection efficiencies and pseudotyped particle production.
- Incubate the plate overnight (16-18 h) in a 37 °C, 5% CO₂ cell culture incubator.

2. Three-plasmid co-transfection

NOTE: Perform this step in the biosafety cabinet.

- Observe cells under an inverted light microscope to check for the cell morphology and density.
NOTE: Ideally, cell density should be in the 40-60% confluency range. It is critical that cells are neither too confluent (80–90% confluent) nor too sparsely distributed (20–30% confluent) in each well. A cell density of 40–60% confluency will ensure good pseudotyped particles production.
- Plasmids mix**
 - Calculate the plasmid mix for each envelope glycoprotein following the quantities for one well of a 6-well plate shown in **Table 1**. Multiply quantities if transfecting several wells and include an extra well to avoid running out of mix.
NOTE: Along with the SARS-CoV S and MERS-CoV S encoding plasmids, include empty vector control for the generation of negative control particles which lack viral envelope glycoproteins (Δ env particles) along with a positive control glycoprotein such as vesicular stomatitis virus (VSV) G glycoprotein that is known to robustly infect a very wide range of cells (VSV-Gpp). Plasmids are available upon request.
 - Mix calculated volumes of plasmids and reduced serum cell culture medium (see **Table of Materials**) in a microcentrifuge tube.
- Lipid-based transfection reagent mix** (see **Table of Materials**)

1. Calculate the volumes for the transfection reagent mix from the quantities shown in **Table 2** for one well (1:3 transfection ratio, multiply quantities as needed). Include extra wells to avoid running out of transfection reagent mix.
2. Mix calculated volumes of lipid-based transfection reagent (3 μ L per well) and reduced serum cell culture medium (47 μ L per well) in a microcentrifuge tube, making sure to add the transfection reagent into the reduced serum cell culture medium and not the other way around.
4. Incubate both mixes (for one well: 50 μ L of plasmids mix and 50 μ L of lipid-based transfection reagent mix) separately for 5 min at room temperature.
5. Add the contents of the transfection reagent mix to the plasmids mix at a 1:1 ratio (for 1 well: 50 μ L of each mix)
6. Perform thorough up-down pipetting of the resulting mix.
7. Incubate the mix for at least 20 min at room temperature.
8. Aspirate the spent medium of cells.
9. Add gently 1 mL of pre-warmed (37 °C) reduced serum cell culture medium per well.
10. Add dropwise 100 μ L of transfection mix to each well.
NOTE: Exercise care when adding the transfection mix to wells of HEK-293T as they detach easily.
11. Incubate cells in a 37 °C, 5% CO₂ cell culture incubator for 4-6 h.
12. Add 1 mL per well of pre-warmed (37 °C) DMEM-T medium, which does not contain antibiotics
NOTE: This is a key step. Transfection reagents increase cell permeability and increase sensitivity to antibiotics. To ensure good transfection efficiency and pseudotyped particle production, it is important to avoid using cell culture medium containing antibiotics
13. Incubate cells in a 37 °C, 5% CO₂ cell culture incubator for 48 h.

3. Pseudotyped particles collection

NOTE: Perform this step in the biosafety cabinet.

1. Observe cells under inverted light microscope to check for cell morphology and general condition. Also check the color of the medium which should be light pink/slightly orange.
NOTE: This is an important step. If there is too much cell death associated with the transfection or the medium color turned orange/yellow (acidic pH), this will typically be associated with lower yields in infectious pseudotyped particles
2. Transfer supernatants of transfected cells to 50 mL conical centrifuge tubes.
3. Centrifuge tubes at 290 x g for 7 min to remove cell debris.
4. Filter clarified supernatants through a sterile 0.45 μ m pore-sized filter.
5. Make small volume aliquots (e.g., 500 μ L or 1 mL) of pseudotyped virus solution in cryovials.
6. Store at -80 °C.
NOTE: The protocol can be paused here. Pseudotyped particles are stable at -80 °C for many months but once thawed, avoid re-freezing them as they will lose infectivity

4. Pseudotyped particle infection of susceptible cells

NOTE: Perform this step in the biosafety cabinet.

1. **Cell seeding of susceptible cells in 24-well plate**
 1. Obtain by standard cell culture techniques 80–90% confluent 75 cm² flask of susceptible cells: Vero-E6 cells for SARS-CoV pseudotyped particles (SARS-Spp) and Huh-7 cells for MERS-CoV S pseudotyped particles (MERS-Spp).
NOTE: To confirm whether the pseudotyped particles that have been produced are infectious, it is important to carefully choose an appropriate susceptible cell line for pseudovirion infectivity assays. Using poorly permissive cells will lead to low infectivity.
 2. Wash cells twice with 10 mL of pre-warmed (37 °C) DPBS.
 3. Aspirate the supernatant and detach cells with 1 mL of 0.25% trypsin solution pre-warmed at 37 °C. Place the flask of cells at 37 °C, 5% CO₂ incubator for 3–5 minutes or until cells start detaching.
NOTE: Avoid incubating cells with trypsin for more than 5 minutes as this typically leads to cell clumping. Huh-7 cells are especially sensitive to this effect.
 4. Deactivate trypsin by adding DMEM-C medium and count cells using a cell counting slide and light microscope.
 5. Dilute cells to 5 x 10⁵ cells/mL with DMEM-C.
 6. Seed wells of a 24-well plate with 0.5 mL of cell solution per well and gently move the plate back and forth and side to side to evenly distribute cells, avoiding circular motion
NOTE: This is a key step. Evenly distributed cells will ensure that cells do not clump at the center of wells. In turn, this will ensure good infectivity assays. For each pseudotyped particle (SARS-Spp, MERS-Spp) and condition, prepare three wells for three experimental replicates. Include wells for the non-infected (N.I.), empty vector Δ env particles and positive control particles such as VSV-Gpp
 7. Incubate the plate overnight (16–18 h) in a 37 °C, 5% CO₂ cell culture incubator
2. **Pseudotyped particle infection**
 1. Observe cells under light microscope and visually confirm that there is a confluent carpet of cells.
 2. Bring cryovials of pseudotyped virus to thaw on ice.
 3. Wash cells three times with 0.5 mL pre-warmed (37 °C) DPBS
NOTE: This is a key step. Cells that are not properly rinsed prior to infection typically lead to poor infectivity readouts
 4. Aspirate the supernatants of cells.
 5. Inoculate cells with 200 μ L of thawed pseudotyped particle solution.
 6. Incubate cells in a 37 °C, 5% CO₂ cell culture incubator for 1–2 h.
 7. Add 300 μ L of pre-warmed (37 °C) DMEM-C medium.

8. Incubate cells in a 37 °C, 5% CO₂ cell culture incubator for 72 h.

5. Infectivity quantification by luciferase assay readout

NOTE: Perform initial steps in the biosafety cabinet.

1. Thaw luciferin substrate (stored at –80 °C) and 5x luciferase assay lysis buffer (stored at –20 °C) until they reach room temperature.
2. Dilute luciferase assay lysis buffer to 1x with sterile water.
3. Aspirate supernatants of cells infected with pseudotyped particles.
4. Add 100 µL of 1x luciferase assay lysis buffer to each well.
5. Place the plate on a rocker and incubate for 15 min with rocking at room temperature (from this point onwards the plate can be handled outside of a biosafety cabinet).
6. Prepare microcentrifuge tubes for each well by adding 20 µL of luciferin substrate in each tube.
7. Turn on the luminometer.
8. Perform luciferase activity measurement one well at a time by transferring 10 µL of lysate to one tube containing 20 µL of luciferin substrate.
9. Flick the tube gently to mix contents, but avoid displacing the liquid on walls of tube.
10. Place the tube in device and close lid.
11. Measure the luminescence value of the tube by using the luminometer.
12. Record the relative light unit's measurement.
13. Repeat steps 5.8-5.12 until all wells are analyzed.

NOTE: With the appropriate equipment such as a plate reading luminometer, this process can be performed automatically. The assay will need to be scaled to the plate format (e.g., 96-well plate format).

6. Data analysis

1. Calculation and plotting of relative luciferase units' averages and standard deviations

1. Use a graph plotting software to calculate luciferase assay measurement averages and standard deviations of experimental and biological replicates.
2. Plot data as bar chart with standard deviations.

NOTE: When performing statistical analyses on data, make sure to include at least three biological replicates in data sets.

Representative Results

Representative results of infectivity assays of SARS-CoV S and MERS-CoV S pseudotyped particles are shown in **Figure 1**. As expected, for both **Figure 1A and 1B**, the VSV G pseudotyped positive control particles (VSV-Gpp) gave very high average infectivity in the 10^6 to 10^7 relative luciferase units (RLU) range respectively. For SARS-CoV S pseudotyped particles (**Figure 1A**) infection of susceptible Vero-E6 cells, a strong average infectivity was measured at around 9.8×10^4 RLU. This value is almost 3 orders of magnitude higher than the values measured for the non-infected control (1.1×10^2 RLU), or the Δ env particles (1.5×10^2 RLU), which do not harbor any viral envelope glycoproteins at their surface. Similarly, for MERS-CoV S pseudotyped particles (**Figure 1B**) infection of Huh-7 cells, a high average infectivity was measured at around 1.0×10^6 RLU. This is almost 4 orders of magnitude higher than the values measured for the non-infected control (0.8×10^2 RLU), or the Δ env particles (2.0×10^2 RLU). An additional infectivity assay was performed in which the SARS-Spp and MERS-Spp were serially diluted and used to infect Vero-E6 cells (**Figure 2A**). This assay confirms that the luciferase activity measured is dependent on the concentration of the particles used to infect cells. To confirm the role of angiotensin converting enzyme 2 (ACE2) and dipeptidyl peptidase 4 (DPP4), the receptors of SARS-CoV and MERS-CoV, respectively, in mediating attachment and entry of the pseudotyped particles, we used poorly-permissive HEK-293T cells and transfected them to express either ACE2 or DPP4 (**Figure 2B**). The transfected cells were then used for an infectivity assay. This analysis demonstrates that upon overexpression of ACE2 and DPP4, there is a ~4-log and ~2-log increase for SARS-Spp and MERS-Spp infectivity, respectively, confirming that receptor usage of pseudovirions is similar to that of native viruses.

The examples shown here demonstrate the importance of including negative (non-infected, Δ env particles) as well as positive control (VSV-Gpp) conditions when producing pseudotyped particles. Indeed, the positive control VSV-Gpp particles allow us to assess whether a particular batch of pseudotyped particles was successful in yielding functional and infective pseudovirions. Expected results of typical infection by VSV-Gpp particles in most mammalian cell lines are in the 10^6 – 10^7 RLU range. Problems with HEK-293T/17 producer cells (high passage number, issues with cell densities) or poor transfection efficiencies can impact overall pseudotyped particle production and infectivity. Furthermore, the negative control conditions are also important as they allow us to assess the baselines of RLU measurements in a particular cell line (non-infected condition) and non-specific internalization of particles (Δ env infection) which is not mediated by a viral envelope protein. Ideally, for a given type of particle pseudotyped with a viral envelope protein of interest, it is recommended to obtain values that are a few orders of magnitude higher than the negative control values, as shown here in **Figure 1A,B**. However, if for a given cell type a pseudotyped virus infection gives very little infectivity (i.e., close to negative controls such as in **Figure 2B** in non-transfected N.T. conditions) it does not necessarily mean that the pseudotyped particle production was faulty. It could well be that the particular cell line used is not or poorly permissive to infection. It is recommended to check whether a given cell type is expected to be susceptible to infection by the virus being investigated. Transfecting poorly permissive cells with plasmid(s) expressing the viral receptor(s) can allow more efficient viral entry and infection to occur, as shown in **Figure 2B**, where upon transfection of ACE2 and DPP4 receptors in target HEK-293T cells, there is a ~4- and ~2-log increase in infectivity, respectively.

Plasmid/reagent	Quantity
pCMV-MLVgagpol MLV gag and pol encoding plasmid	300 ng
pTG-Luc transfer vector with luciferase reporter	400 ng
pcDNA-SARS-S, pcDNA-MERS-S or empty vector	300 ng
Reduced serum cell culture medium	To 50 μ L

Table 1. Quantities of plasmids and reduced serum cell culture medium required to transfect one well of a 6-well plate of HEK-293T/17 cells for pseudotyped particle production. From the concentration of plasmid preparations, calculate the required volume to reach the required amount of plasmid DNA. If more than one well is transfected with the same plasmids, multiply volumes by required number of wells to transfect, and include an extra well in calculations to avoid running out of mix in later steps. The total amount of DNA being transfected is 1 μ g/well. Plasmids are available upon request to authors.

Reagent	Quantity
Transfection reagent	3 μ L
Reduced serum cell culture medium	47 μ L

Table 2. Quantities of transfection reagent and reduced serum cell culture medium required to transfect one well of a 6-well plate of HEK-293T/17 cells for pseudotyped particle production. Multiply volumes by required number of wells to transfect and include extra wells in calculations to avoid running out of mix in later steps. The transfection reagent: plasmid DNA ratio used is 3:1.

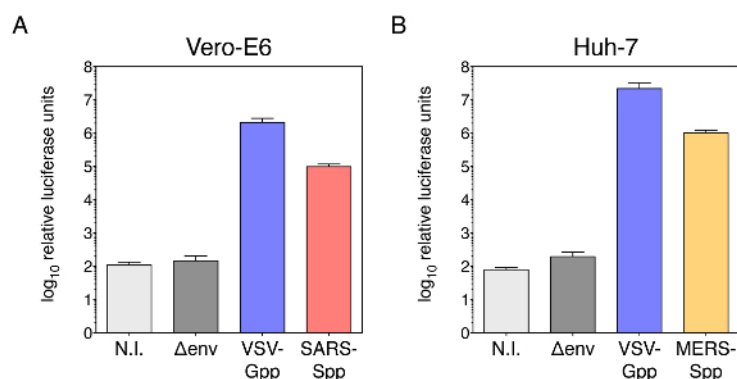


Figure 1. Coronavirus S-pseudotyped particle infectivity assays in susceptible host cells using a murine leukemia virus (MLV) backbone and luciferase reporter gene. (A) SARS-CoV S pseudotyped particle infectivity assay in Vero-E6 cells. (B) MERS-CoV S pseudotyped particle infectivity assay in Huh-7 cells. For both (A) and (B), plotted data corresponds to the average relative luciferase units from three independent experiments, with error bars corresponding to standard deviation (s.d.). Data plotted in log₁₀ scale on y-axis. N.I.: non-infected control; Δenv: infection with pseudotyped particles lacking viral envelope glycoproteins and VSV-Gpp: infection with pseudotyped particles bearing positive control VSV G envelope glycoprotein. Other abbreviation used, SARS-Spp: infection with SARS-CoV S pseudotyped particles, MERS-Spp: infection with MERS-CoV S pseudotyped particles. [Please click here to view a larger version of this figure.](#)

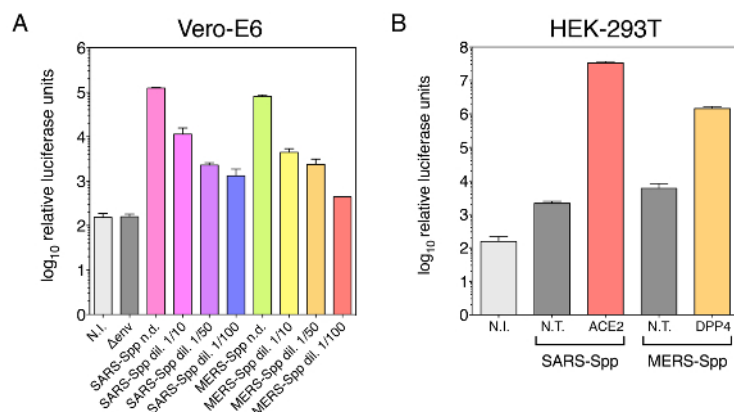


Figure 2. Concentration-dependence of CoV S-pseudovirion infectivity and role of ACE2 and DPP4 receptors in SARS-Spp and MERS-Spp entry. (A) Concentration-dependence of SARS-Spp and MERS-Spp infectivity measured by luciferase activity assay. Pseudovirions were serially diluted and used to infect Vero-E6 cells. (B) Infectivity assay of SARS-Spp and MERS-Spp in poorly permissive HEK-293T target cells transfected to express ACE2, and DPP4 receptors, respectively. Data plotted in log₁₀ scale on y-axis as in Figure 1 from duplicate experiments, with error bars corresponding to standard deviation (s.d.). N.T.: non-transfected control HEK-293T cells; ACE2: angiotensin converting enzyme 2 (SARS-CoV receptor) and DPP4: dipeptidyl peptidase 4 (MERS-CoV receptor). Other abbreviations used are the same as in Figure 1. [Please click here to view a larger version of this figure.](#)

Discussion

This protocol describes a method to efficiently produce pseudotyped particles bearing the S protein of risk group 3 coronaviruses, SARS-CoV and MERS-CoV, in a BSL-2 setting. These particles, which incorporate a luciferase reporter gene, enable us to easily quantify coronavirus S-mediated entry events by a relatively simple luciferase assay^{48,49,50,51}. In infectivity assays using permissive cells, we confirm that the luciferase activity measured is dependent on the concentration of particles. In addition, ACE2 and DPP4 receptor transfection allows for more efficient entry in poorly permissive cells lines, such as HEK-293T cells. The method is highly adaptable to other viral envelope glycoproteins and has been used extensively^{48,49,50,51,52,53,55,56,57,58,59}, often to complement other assays like biochemical analyses or native virus infections.

The protocol we describe here is based on the retrovirus MLV that incorporates a luciferase reporter. However, it is important to emphasize that there is a very wide array of other pseudotyping systems that have been successfully developed for packaging coronavirus S^{12,13,25,26,30,31,32} and other viral envelope glycoproteins^{10,11,14,16,17,23,24,29,33,38,40,42,44,46}. Some of these other systems are based on the commonly used MLV retroviral core^{7,8,9,10,11,12,13,14,15,16,17,18,19,20}, or based on the widely used lentiviral HIV-1 pseudotyping system using different strategies^{23,24,25,26,27,28,29,30,31,32,33,34,35}, or with the rhabdovirus vesicular stomatitis virus (VSV) as core, which allows to incorporate a wide variety of envelope glycoproteins, and again with various strategies employed^{37,38,39,40,41,42,43,44,45,46,47}. In addition, other reporters such as fluorescent proteins like GFP^{11,13} and RFP³⁶, or enzymes other than luciferase like β-galactosidase^{16,17} and secreted alkaline phosphatase (SEAP)⁴² have been successfully employed for measurements. Furthermore, in the assay presented in this protocol, a transient transfection was used to express the MLV and CoV S genes and proteins. However, there are other strategies for expression, such as generation of stable cell lines for production of pseudotyped viruses^{7,14}. As each of these systems have their advantages and disadvantages, it is important to consider the following important parameters when deciding which pseudotyping system best suits an investigator's needs: pseudovirion core (MLV, HIV-1, VSV or others), how selective a particular pseudotyping core is in incorporating a specific viral envelope glycoprotein, reporter for assay readout (GFP, luciferase, SEAP or other), and the transfection strategy (number of plasmids involved in co-transfection, transient transfection or generation of stable cell lines).

There are a number of critical steps in the method that are important to emphasize. Cell density, particularly of the HEK-293T/17 producer cell line is a critical factor in ensuring successful transfection. A cell density in the range of 40-60% confluency was found to be optimal. Higher densities typically result in low transfection efficiencies and low particle production. Also, it is important to keep in mind that HEK-293T/17 cells are less adherent than other cell lines. Care should be exercised when handling them to avoid detaching them unnecessarily. One option is to treat cell culture plastic surfaces with poly-D-lysine to enhance adherence. Furthermore, higher cell passage often results in poor transfection rates. After adding the transfection reagent to HEK-293T/17 cells, it is also important to remember that cell permeability increases. This is why at this point it is best to avoid using medium containing antibiotics as they may increase cytotoxicity. Before collecting pseudotyped particles, check the color of transfected HEK-293T/17 cell supernatants. Typically, after 48 h of transfection, the cell culture medium color takes an orange-pink tinge. Yellow-colored medium usually translates to poor pseudotyped particles yields and is often a result of issues with cell seeding density or high passage number.

In this protocol, pseudotyped particle production is performed in a 6-well plate format. To increase volume of produced particles, several wells of a 6-well plate can be transfected with the same plasmids mix and the supernatants can be pooled together. The pool can then be clarified, filtered and aliquoted. Alternatively, to scale production up, other kinds of vessels (e.g., 25, or 75 cm² flasks) can be used. In this case, transfection conditions should be scaled up accordingly. In this protocol, the infectivity assay is performed using a 24-well plate format and a luminometer that only allows measurements one tube at a time. For high throughput screenings, other formats are also possible, such as 96-well plate format and a plate reader luminometer. Volumes and reagents for the luciferase assay need to be adapted accordingly. Storage of pseudotyped particles in cryovials at -80 °C maintains their stability for several months without noticeable decrease in infectivity. It is not recommended to subject them to freeze-thaw cycles as this will decrease their infectivity over time. Thus, it is best to store them in small aliquots such as 0.5-1 mL and thaw them before an infection.

The method presented here has several limitations. An important one is the fact that pseudotyped particles recapitulate only viral entry events. To analyze other steps in the infectious life cycle, other assays are required. Furthermore, as MLV particles bud at the plasma membrane, it is important to bear in mind that the envelope glycoprotein being studied needs to also traffic to the plasma membrane for incorporation into pseudovirions during production. As such, it is important to know where in the cell a particular viral envelope glycoprotein is expressed in transfection conditions, such as by visualizing subcellular localization with an immunofluorescence assay, and/or by checking for retention signals within the protein. Also, while the protocol describes steps to produce and test infectivity, it does not detail how to measure incorporation of viral envelope glycoproteins into pseudotyped particles. One method is to perform western blot assays on concentrated solutions of particles, as previously described^{50,51} for MERS-CoV S incorporation. In these assays, the S envelope glycoprotein of MERS-CoV is probed along with the capsid (p30) protein of MLV, which allow us to normalize incorporation of the S protein into particles. Other examples of such assays analyzing viral envelope glycoprotein incorporation into pseudovirions have been performed for SARS-CoV S incorporation in an HIV-1 lentiviral pseudovirion system³², Ebola glycoprotein (GP) in another MLV pseudotyped particle system¹⁷, and influenza hemagglutinin (HA) and neuraminidase (NA) in VSV pseudovirions³⁸. A recent development in characterizing pseudotyped particle production is the use of innovative imaging devices such as Nanosight: it enables us to directly visualize, quantify, and size viral particles⁵⁰. The device provides detailed information on overall particle production; however, it is important to keep in mind that it does not provide information on envelope glycoprotein incorporation. A future direction for the application of these versatile pseudovirion particles is to analyze individual viral fusion events using single particle tracking, microfluidics and total internal reflection fluorescence microscopy^{60,61,62}. Such approaches were successfully applied to influenza virus and feline coronavirus particles as well as influenza HA- and NA-pseudotyped VSV-based pseudovirions⁶³. The deployment of such techniques applied to coronavirus S-pseudotyped MLV-based particles is currently being developed.

Disclosures

The authors have nothing to disclose.

Acknowledgements

We wish to thank all members of the Whittaker and Daniel labs for helpful comments. Research funding was provided by the NIH grants R21 AI111085 and R01 AI135270. T.T. acknowledges support from the Presidential Life Science Fellowship in Cornell University and National Science Foundation Graduate Research Fellowship Program under Grant No. DGE-1650441. L.N. acknowledges support from a Samuel C. Fleming Family Graduate Fellowship and National Science Foundation Graduate Research Fellowship Program under Grant No. DGE-1650441. This work is also supported by National Science Foundation 1504846 (to S.D. and G.R.W.).

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