

## **APPENDICES**

**APPENDIX A**  
**OPTIMIZED DENV-2 prM-E GENE**

5'-

GGATCCGCCACCATGAAATGTCTGCTGTACCTGGCCTTCCTGTTTCATCGGC  
GTGAATTGTTTCCACCTGACCACCAGGAACGGCGAGCCCCACATGATCGTG  
AGCAGACAGGAGAAGGGCAAGAGCCTGCTGTTCAAGACCGAGGACGGCG  
TGAACATGTGTACCCTGATGGCCATGGACCTGGGCGAGCTGTGCGAGGAC  
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CACTTCCAGAGAGCCCTGATCTTCATCCTGCTGACCGCCGTGGCCCCCAGC  
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CCCCAGAGCAGCATCACAGAGGCCGAGCTGACCGGCTACGGCACAGTGA  
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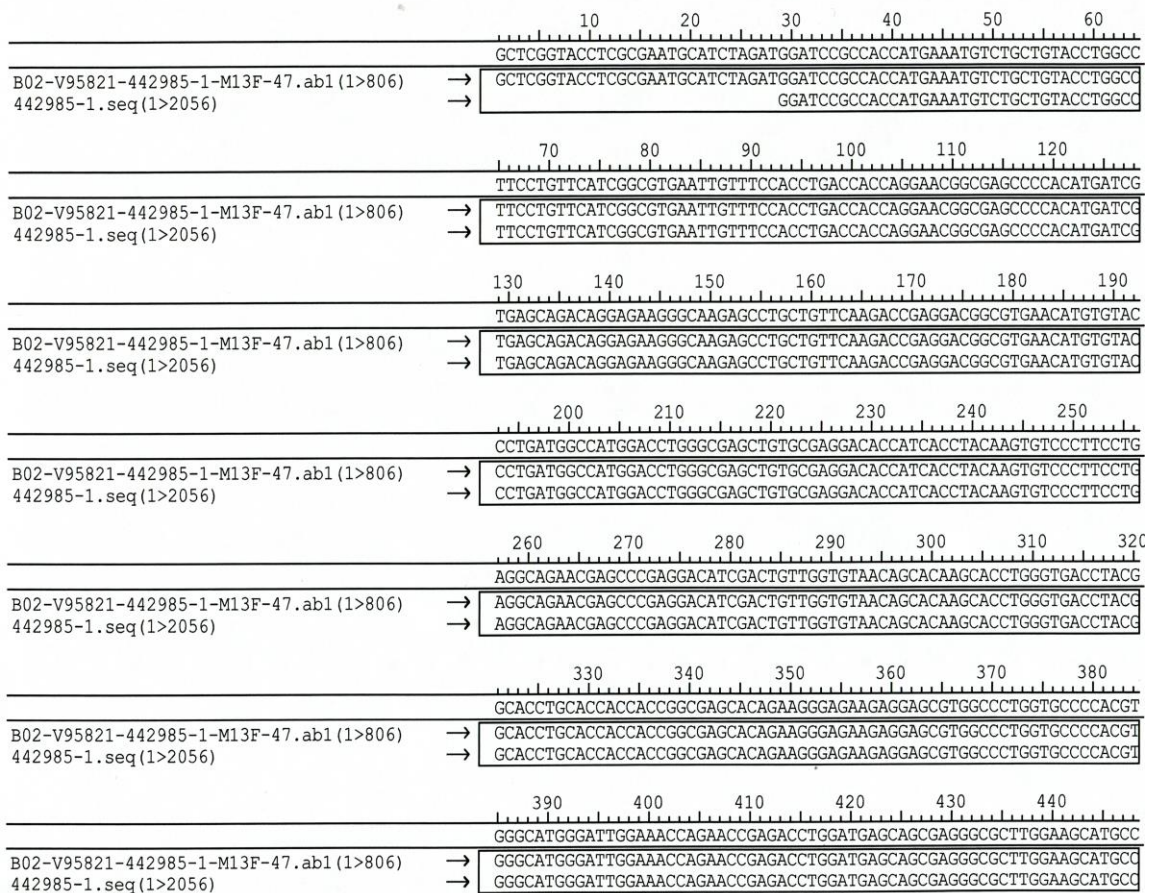
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ACAGCATGTGTACAGGCAAGTTCAAGGTGGTGAAGGAGATCGCCGAGACC  
CAGCACGGCACCATCGTGATCAGAGTGCAGTACGAGGGCGATGGCAGCCC  
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GCAGACTGATCACCGTGAACCCCATCGTGACCGAGAAGGATAGCCCCGTG  
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GAGCCCGGCCAGCTGAAGCTGAACTGGTTCAAGAAGGGCAGCAGCATCGG  
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GCGTGTCCTGGACCATGAAGATCCTGATCGGCGTGATCATCACCTGGATCG  
GCATGAATTCCAGAAGCACCAGCCTGAGCGTGTCCCTGGTGTGCTGGTCGGA  
GTGGTGACCCTGTACCTGGGCGTGATGGTGCAGGCCTGATGACTCGAG-3



### APPENDIX C

## NUCLEOTIDE SEQUENCE ALIGNMENT BETWEEN pUC57-D2opt.prM-E CONSTRUCT AND TARGETED INSERTED SEQUENCE

QC Items	Specifications	Results
Sequencing Alignment	Sequencing results are consistent with the targeted insert sequence.	<input checked="" type="checkbox"/> Pass <input type="checkbox"/> Fail



		450	460	470	480	490	500	510
		CAGAGAATCGAGACCTGGATTCTGAGACACCCCGGCTTCACCATCATGGCCGCCATCCTGGCCT						
B02-V95821-442985-1-M13F-47.ab1 (1>806)	→	CAGAGAATCGAGACCTGGATTCTGAGACACCCCGGCTTCACCATCATGGCCGCCATCCTGGCCT						
442985-1.seq (1>2056)	→	CAGAGAATCGAGACCTGGATTCTGAGACACCCCGGCTTCACCATCATGGCCGCCATCCTGGCCT						
		520	530	540	550	560	570	
		ACACCATCGGCACCACCCACTTCCAGAGAGCCCTGATCTTCATCCTGCTGACCGCCGTGGCCCC						
B02-V95821-442985-1-M13F-47.ab1 (1>806)	→	ACACCATCGGCACCACCCACTTCCAGAGAGCCCTGATCTTCATCCTGCTGACCGCCGTGGCCCC						
442985-1.seq (1>2056)	→	ACACCATCGGCACCACCCACTTCCAGAGAGCCCTGATCTTCATCCTGCTGACCGCCGTGGCCCC						
		580	590	600	610	620	630	640
		CAGCATGACCATGAGATGCATCGGCATCAGCAACAGAGACTTCGTGGAGGGCGTGAGCGCGGGC						
B02-V95821-442985-1-M13F-47.ab1 (1>806)	→	CAGCATGACCATGAGATGCATCGGCATCAGCAACAGAGACTTCGTGGAGGGCGTGAGCGCGGGC						
442985-1.seq (1>2056)	→	CAGCATGACCATGAGATGCATCGGCATCAGCAACAGAGACTTCGTGGAGGGCGTGAGCGCGGGC						
		650	660	670	680	690	700	
		AGCTGGGTGGACATCGTGTGGAGCACGGCAGCTGTGTGACCACCATGGCCAAGAACAAGCCCA						
B02-V95821-442985-1-M13F-47.ab1 (1>806)	→	AGCTGGGTGGACATCGTGTGGAGCACGGCAGCTGTGTGACCACCATGGCCAAGAACAAGCCCA						
442985-1.seq (1>2056)	→	AGCTGGGTGGACATCGTGTGGAGCACGGCAGCTGTGTGACCACCATGGCCAAGAACAAGCCCA						
B04-V95821-442985-1-442985-1seq.ab1 (1>814)	→	TCGTGCTGGAGCACGGCAGCTGTGTGACCACCATGGCCAAGAACAAGCCCA						
		710	720	730	740	750	760	
		CACTGGACTTCGAGCTGATCAAGACCGAGGCCAAGCAGCCCGCCACCCTGAGAAAGTACTGTAT						
B02-V95821-442985-1-M13F-47.ab1 (1>806)	→	CACTGGACTTCGAGCTGATCAAGACCGAGGCCAAGCAGCCCGCCACCCTGAGAAAGTACTGTAT						
442985-1.seq (1>2056)	→	CACTGGACTTCGAGCTGATCAAGACCGAGGCCAAGCAGCCCGCCACCCTGAGAAAGTACTGTAT						
B04-V95821-442985-1-442985-1seq.ab1 (1>814)	→	CACTGGACTTCGAGCTGATCAAGACCGAGGCCAAGCAGCCCGCCACCCTGAGAAAGTACTGTAT						
		770	780	790	800	810	820	830
		CGAGGCCAAGCTGACCAACACCACCACCGACAGCAGATGCCCCACCCAGGGCGAGCCAGCCTC						
B02-V95821-442985-1-M13F-47.ab1 (1>806)	→	CGAGGCCAAGCTGACCAACACCACCACCGACAGCAGATGCCCCACCCAGGGCGAGCCAGCCTC						
442985-1.seq (1>2056)	→	CGAGGCCAAGCTGACCAACACCACCACCGACAGCAGATGCCCCACCCAGGGCGAGCCAGCCTC						
B04-V95821-442985-1-442985-1seq.ab1 (1>814)	→	CGAGGCCAAGCTGACCAACACCACCACCGACAGCAGATGCCCCACCCAGGGCGAGCCAGCCTC						
		840	850	860	870	880	890	
		AATGAGGAGCAGGACAAGAGATTTCGTGTGTAAGCACAGCATGGTGGACAGAGGCTGGGGCAACG						
442985-1.seq (1>2056)	→	AATGAGGAGCAGGACAAGAGATTTCGTGTGTAAGCACAGCATGGTGGACAGAGGCTGGGGCAACG						



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      840      850      860      870      880      890
      |-----|-----|-----|-----|-----|
AATGAGGAGCAGGACAAGAGATTCTGTGTGTAAGCACAGCATGGTGGACAGAGGCTGGGGCAACG
B04-V95821-442985-1-442985-1seq.ab1 (1>814) → AATGAGGAGCAGGACAAGAGATTCTGTGTGTAAGCACAGCATGGTGGACAGAGGCTGGGGCAACG

      900      910      920      930      940      950      960
      |-----|-----|-----|-----|-----|
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B04-V95821-442985-1-442985-1seq.ab1 (1>814) → GCTGTGGCCTGTTCCGCAAGGGCGGCATCGTGACCTGTGCCATGTTACATGCAAGAAGAACAT

      970      980      990      1000     1010     1020
      |-----|-----|-----|-----|-----|
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B04-V95821-442985-1-442985-1seq.ab1 (1>814) → GAAGGGCAAGTGGTGCAGCCTGAGAACCTGGAGTACACCATCGTGATCACCCCTCACTCTGGC

      1030     1040     1050     1060     1070     1080
      |-----|-----|-----|-----|-----|
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442985-1.seq (1>2056) → GAGGAGCATGCCGTGGGCAACGACACCGGCAAGCAGGCAAGGAGATCAAGATCACCCCCAGA
B04-V95821-442985-1-442985-1seq.ab1 (1>814) → GAGGAGCATGCCGTGGGCAACGACACCGGCAAGCAGGCAAGGAGATCAAGATCACCCCCAGA

      1090     1100     1110     1120     1130     1140     1150
      |-----|-----|-----|-----|-----|
GCAGCATCACAGAGCCGAGCTGACCGGCTACGGCACAGTGACCATGGAGTGTAGCCCTAGAAC
442985-1.seq (1>2056) → GCAGCATCACAGAGCCGAGCTGACCGGCTACGGCACAGTGACCATGGAGTGTAGCCCTAGAAC
B04-V95821-442985-1-442985-1seq.ab1 (1>814) → GCAGCATCACAGAGCCGAGCTGACCGGCTACGGCACAGTGACCATGGAGTGTAGCCCTAGAAC

      1160     1170     1180     1190     1200     1210
      |-----|-----|-----|-----|-----|
CGGCCTGGATTTC AACGAGATGGTGTGCTGCTGCAAATGGAGAACAAGGCCTGGCTGGTGCACAGA
442985-1.seq (1>2056) → CGGCCTGGATTTC AACGAGATGGTGTGCTGCTGCAAATGGAGAACAAGGCCTGGCTGGTGCACAGA
B04-V95821-442985-1-442985-1seq.ab1 (1>814) → CGGCCTGGATTTC AACGAGATGGTGTGCTGCTGCAAATGGAGAACAAGGCCTGGCTGGTGCACAGA

      1220     1230     1240     1250     1260     1270     1280
      |-----|-----|-----|-----|-----|
CAATGGTTCCTGGATCTGCCCTCTGCCCTGGCTGCCTGGCGCCGACACCCAGGGAAGCAACTGGA
442985-1.seq (1>2056) → CAATGGTTCCTGGATCTGCCCTCTGCCCTGGCTGCCTGGCGCCGACACCCAGGGAAGCAACTGGA
B04-V95821-442985-1-442985-1seq.ab1 (1>814) → CAATGGTTCCTGGATCTGCCCTCTGCCCTGGCTGCCTGGCGCCGACACCCAGGGAAGCAACTGGA
    
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		1290	1300	1310	1320	1330	1340	
		TTCAGAAGGAGACCTGGTGACCTTCAAGAACCCCCACGCCAAGAAGCAGGACGTGGTGGTGCT						
442985-1.seq (1>2056)	→	TTCAGAAGGAGACCTGGTGACCTTCAAGAACCCCCACGCCAAGAAGCAGGACGTGGTGGTGCT						
B04-V95821-442985-1-442985-1seq.ab1 (1>814)	→	TTCAGAAGGAGACCTGGTGACCTTCAAGAACCCCCACGCCAAGAAGCAGGACGTGGTGGTGCT						
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	ACCCCCACGCCAAGAAGCAGGACGTGGTGGTGCT						
		1350	1360	1370	1380	1390	1400	
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B04-V95821-442985-1-442985-1seq.ab1 (1>814)	→	GGGCAGCCAGGAGGGGCCATGCACACCGCCCTGACAGGCGCCACCGAGATCCAGATGAGCAGC						
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	GGGCAGCCAGGAGGGGCCATGCACACCGCCCTGACAGGCGCCACCGAGATCCAGATGAGCAGC						
		1410	1420	1430	1440	1450	1460	1470
		GGCAACCTGCTGTTACCGGCCATTTGAAATGTAGACTGAGAATGGATAAGCTGCAGCTGAAGG						
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B04-V95821-442985-1-442985-1seq.ab1 (1>814)	→	GGCAACCTGCTGTTACCGGCCATTTGAAATGTAGACTGAGAATGGATAAGCTGCAGCTGAAGG						
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	GGCAACCTGCTGTTACCGGCCATTTGAAATGTAGACTGAGAATGGATAAGCTGCAGCTGAAGG						
		1480	1490	1500	1510	1520	1530	
		GCATGTCTTACAGCATGTGTACAGGCAAGTTC AAGTGGTGAAGGAGATCGCCGAGACCCAGCA						
442985-1.seq (1>2056)	→	GCATGTCTTACAGCATGTGTACAGGCAAGTTC AAGTGGTGAAGGAGATCGCCGAGACCCAGCA						
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	GCATGTCTTACAGCATGTGTACAGGCAAGTTC AAGTGGTGAAGGAGATCGCCGAGACCCAGCA						
		1540	1550	1560	1570	1580	1590	1600
		CGGCACCATCGTGATCAGAGTGCAGTACGAGGCGGATGGCAGCCCTGTAAGATCCCTTCGAG						
442985-1.seq (1>2056)	→	CGGCACCATCGTGATCAGAGTGCAGTACGAGGCGGATGGCAGCCCTGTAAGATCCCTTCGAG						
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	CGGCACCATCGTGATCAGAGTGCAGTACGAGGCGGATGGCAGCCCTGTAAGATCCCTTCGAG						
		1610	1620	1630	1640	1650	1660	
		ATCATGGATTTGGAGAAGAGACAGTGTGGGAGACTGATCACCGTGAACCCATCGTGACCG						
442985-1.seq (1>2056)	→	ATCATGGATTTGGAGAAGAGACAGTGTGGGAGACTGATCACCGTGAACCCATCGTGACCG						
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	ATCATGGATTTGGAGAAGAGACAGTGTGGGAGACTGATCACCGTGAACCCATCGTGACCG						
		1670	1680	1690	1700	1710	1720	
		AGAAGGATAGCCCCGTGAACATCGAGGCCGAGCCCCCTTTCGGCGACAGCTACATCATCATCGG						
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		1670	1680	1690	1700	1710	1720		
		----- ----- ----- ----- ----- -----							
		AGAAGGATAGCCCCGTGAACATCGAGGCCGAGCCCCCTTCGGCGACAGCTACATCATCATCGG							
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	AGAAGGATAGCCCCGTGAACATCGAGGCCGAGCCCCCTTCGGCGACAGCTACATCATCATCGG							
		1730	1740	1750	1760	1770	1780	1790	
		----- ----- ----- ----- ----- -----							
		CGTGGAGCCCGCCAGCTGAAGCTGAACTGGTTCAAGAAGGGCAGCAGCATCGGCCAGATGATC							
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B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	CGTGGAGCCCGCCAGCTGAAGCTGAACTGGTTCAAGAAGGGCAGCAGCATCGGCCAGATGATC							
		1800	1810	1820	1830	1840	1850		
		----- ----- ----- ----- ----- -----							
		GAGACCACCATGAGAGGAGCCAAGCGGATGGCCATCCTGGGCGACACCGCCTGGGACTTCGGCT							
442985-1.seq (1>2056)	→	GAGACCACCATGAGAGGAGCCAAGCGGATGGCCATCCTGGGCGACACCGCCTGGGACTTCGGCT							
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	GAGACCACCATGAGAGGAGCCAAGCGGATGGCCATCCTGGGCGACACCGCCTGGGACTTCGGCT							
		1860	1870	1880	1890	1900	1910	1920	
		----- ----- ----- ----- ----- -----							
		CTCTGGGCGGCGTGTTCACCTCCATCGGCAAGGCCCTGCACCAAGTGTTCGGCGCCATCAGCG							
442985-1.seq (1>2056)	→	CTCTGGGCGGCGTGTTCACCTCCATCGGCAAGGCCCTGCACCAAGTGTTCGGCGCCATCAGCG							
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	CTCTGGGCGGCGTGTTCACCTCCATCGGCAAGGCCCTGCACCAAGTGTTCGGCGCCATCAGCG							
		1930	1940	1950	1960	1970	1980		
		----- ----- ----- ----- ----- -----							
		CGCCGCCTTCTCCGGCGTGTCTTGACCATGAAGATCCTGATCGGCGTATCATCACCTGGATC							
442985-1.seq (1>2056)	→	CGCCGCCTTCTCCGGCGTGTCTTGACCATGAAGATCCTGATCGGCGTATCATCACCTGGATC							
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	CGCCGCCTTCTCCGGCGTGTCTTGACCATGAAGATCCTGATCGGCGTATCATCACCTGGATC							
		1990	2000	2010	2020	2030	2040		
		----- ----- ----- ----- ----- -----							
		GGCATGAATTCCAGAAGCACCAGCCTGAGCGTGTCCCTGGTGTGGTGGGAGTGGTGACCCTGT							
442985-1.seq (1>2056)	→	GGCATGAATTCCAGAAGCACCAGCCTGAGCGTGTCCCTGGTGTGGTGGGAGTGGTGACCCTGT							
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	GGCATGAATTCCAGAAGCACCAGCCTGAGCGTGTCCCTGGTGTGGTGGGAGTGGTGACCCTGT							
		2050	2060	2070	2080	2090	2100	2110	
		----- ----- ----- ----- ----- -----							
		ACCTGGGCGTGTGGTGCAGGCTGATGACTCGAGATCGGATCCCGGGCCCGTGCAGTGCAGAG							
442985-1.seq (1>2056)	→	ACCTGGGCGTGTGGTGCAGGCTGATGACTCGAG							
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	ACCTGGGCGTGTGGTGCAGGCTGATGACTCGAGATCGGATCCCGGGCCCGTGCAGTGCAGAG							
		2120							
		-----							
		GCCTGCATGC							
B03-V95821-442985-1-M13R-48.ab1 (1>812)	←	GCCTGCATGC							

## APPENDIX D

### SOLUTION PREPARATION

#### 1. Solutions for agarose gel electrophoresis

<b>1.1 0.5M EDTA, pH 8.0</b>	<b>100 ml</b>
Na <sub>2</sub> EDTA (MW 372.2)	18.61 g
Distilled H <sub>2</sub> O	60 ml
Adjust pH with concentrated NaOH to	8.0
Adjust volume with distilled H <sub>2</sub> O up to	100 ml
store at room temperature	
<b>1.2 5X TBE</b>	<b>1 L</b>
Tris base	53.89 g
Boric acid	27.51 g
0.5 M EDTA, pH 8.0	20 ml
Adjust volume with distilled H <sub>2</sub> O to	1 L
Store at room temperature	
<b>1.3 0.5X TBE</b>	<b>1 L</b>
5X TBE	1 L
Distilled H <sub>2</sub> O	100 ml
Store at room temperature	
<b>1.4 1% agarose in 0.5X TBE buffer</b>	<b>100 ml</b>
Agarose	1 g
0.5X TBE buffer	100 ml
Boil until solution is clear	

## 2. Solution for bacterial culture and cloning

<b>2.1 LB both</b>	<b>1</b>	<b>L</b>
NaCl	10	g
Tryptone	10	g
Yeast extract	5	g
Distilled H <sub>2</sub> O to	1	L

Sterilize by autoclave at 121 °C, 15 psi for 20 min and store at 4 °C

Add antibiotics as design (Ampicillin 50 or 100 µg/ml, Kanamycin 50 µg/ml final concentration)

<b>2.2 LB agar</b>	<b>1</b>	<b>L</b>
NaCl	10	g
Tryptone	10	g
Yeast extract	5	g
European bacteriological agar	20	g

Sterilize by autoclave at 121 °C, 15 psi for 20 min and store at 4 °C

To make agar plate, melt LB agar and cool it down to 55-60 °C. Add antibiotics as design (Ampicillin 25, 50 or 100 µg/ml, Kanamycin 50 µg/ml final concentration) and pour into petridishes (approximately 25 ml/plate)

<b>2.3 100 mg/ml Ampicillin (1,000X)</b>	<b>10</b>	<b>ml</b>
Ampicillin sodium salt	1	g
Sterile MilliQ H <sub>2</sub> O	10	ml
Filtrate through 0.22 µm filter and store at -20 °C		

<b>2.4 50 mg/ml Kanamycin (1,000X)</b>	<b>10</b>	<b>ml</b>
Kanamycin	0.5	g
Sterile MilliQ H <sub>2</sub> O	10	ml
Filtrate through 0.22 µm filter and store at -20 °C		



### 3. Cell culture media and reagents

<b>3.1 1X DMEM</b>	<b>1</b>	<b>L</b>
DMEM powder	1	pack
NaHCO <sub>3</sub>	3.7	g
Sterile MilliQ H <sub>2</sub> O to	900	ml
Adjust pH with concentrated HCl to	7.2	
Adjust volume with sterile MilliQ H <sub>2</sub> O to	1	L
Filtrate through 0.22 µm filter and store at 4 °C		
<b>3.2 Growth medium for HEK 293 and HEK 393T cell</b>	<b>500</b>	<b>ml</b>
1X DMEM	450	ml
Heat-inactivated FBS	45	ml
store at 4 °C		
Add antibiotics as design (G418 500 – 2000 µg/ml)		
<b>3.3 1X PBS for cell culture</b>	<b>4</b>	<b>L</b>
NaCl	32.03	g
KCl	0.81	g
Na <sub>2</sub> HPO <sub>4</sub>	2.44	g
KH <sub>2</sub> PO <sub>4</sub>	0.76	g
MilliQ H <sub>2</sub> O to	4	L
Sterilize by autoclave at 121 °C, 15 psi for 20 min and store at room temperature		
<b>3.4 0.5M EDTA pH 8.0</b>	<b>300</b>	<b>ml</b>
EDTA	55.836	g
Distilled H <sub>2</sub> O to	200	ml
Adjust pH with concentrated HCl to	8.0	
Adjust volume with distilled H <sub>2</sub> O to	300	ml
Sterilize by autoclave at 121 °C, 15 psi for 20 min and store at room temperature		



<b>3.5 0.25 (w/v) Trypsin/EDTA</b>	<b>1</b>	<b>L</b>
Trypsin	2.5	g
0.5M EDTA pH 8.0	2	ml
Adjust volume with 1X PBS to	1	L
Filtrate through 0.22 $\mu$ m filter and store at -20 °C		

<b>3.6 0.4% (w/v) Tryphan blue</b>	<b>100</b>	<b>ml</b>
Tryphan blue	0.4	g
1X PBS	100	ml
Store at room temperature		

#### **4. Solutions for SDS-PADE and Western blot analysis**

<b>4.1 10% (w/v) SDS</b>	<b>100</b>	<b>ml</b>
Sodium dodecyl sulfate (SDS)	10	g
Distilled H <sub>2</sub> O to	100	ml
Store at room temperature		

<b>4.2 1M Tris-HCl pH 6.8</b>	<b>100</b>	<b>ml</b>
Tris base	12.1	g
distilled H <sub>2</sub> O to	70	ml
Adjust pH with concentrated HCl to	6.8	
Adjust volume with distilled H <sub>2</sub> O to	100	ml
Store at room temperature		

<b>4.3 1.5M Tris-HCl pH 8.8</b>	<b>150</b>	<b>ml</b>
Tris base	27.26	g
distilled H <sub>2</sub> O to	70	ml
Adjust pH with concentrated HCl to	8.8	
Adjust volume with distilled H <sub>2</sub> O to	100	ml
Store at room temperature		



<b>4.4 1M DTT</b>	<b>32.4 ml</b>
DTT	5 g
distilled H <sub>2</sub> O to	32.4 ml
store at -20 °C	
<b>4.5 5X SDS loading buffer</b>	<b>50 ml</b>
1M Tris-HCl pH 6.8	3.125 ml
10% (w/v) SDS	20 ml
87% glycerol	11.5 ml
Bromophenol blue	7.5 mg
distilled H <sub>2</sub> O to	10.375 ml
store at room temperature	
Add 10 µl of 1M DTT before use	
<b>4.6 10X running buffer (stock solution)</b>	<b>1 L</b>
Tris-base	30.2 g
Glycine	141.4 g
SDS	10 g
Distilled H <sub>2</sub> O to	1 L
<b>4.7 1X running buffer (working solution)</b>	<b>1 L</b>
10X running buffer	100 ml
Distilled H <sub>2</sub> O to	900 ml
Freshly preparation before used	
<b>4.8 10% (w/v) APS</b>	<b>1 L</b>
APS	0.1 g
Distilled H <sub>2</sub> O to	1 ml
Store at 4 °C and use within 1 week	



**4.9 Polyacrylamide gel****Reagents for 10% separating gel**

40% acrylamide/bis-acrylamide	2.5	ml
1.5M Tris-HCl pH 8.8	2.5	ml
Distilled H <sub>2</sub> O to	4.8	ml
10% (w/v) SDS	0.1	ml
10% (w/v) APS	0.15	ml
TEMED	0.0125	ml

**Reagents for 5% stacking gel**

40% acrylamide/bis-acrylamide	0.375	ml
1.5M Tris-HCl pH 6.8	0.38	ml
Distilled H <sub>2</sub> O to	2.225	ml
10% (w/v) SDS	0.030	ml
10% (w/v) APS	0.045	ml
TEMED	0.0125	ml

**4.10 1X transfer buffer**

	<b>1</b>	<b>L</b>
Tris base	3	g
Glycine	14	g
100% methanol	200	ml
Distilled H <sub>2</sub> O to	1	L
Freshly preparation and store at -20 °C before used		

**4.11 10X TBS (Tris-buffered saline) pH 7.6**

	<b>1</b>	<b>L</b>
Tris base	24.2	g
NaCl	80	g
Distilled H <sub>2</sub> O to	800	ml
Adjust pH with concentrated HCl to	7.6	
Adjust volume with distilled H <sub>2</sub> O to	1	L
Store at room temperature		



<b>4.12 1X TBS, 0.1% Tween-20</b>	<b>1</b>	<b>L</b>
10X TBS	100	ml
Tween-20	1	ml
Distilled H <sub>2</sub> O to	1000	ml
Store at room temperature		
<b>4.13 0.2% (w/v) Ponceau S / 1% (v/v) acetic acid</b>	<b>500</b>	<b>ml</b>
Ponceau S	1	g
Distilled H <sub>2</sub> O to	495	ml
Acetic acid	5	ml
Store at room temperature		
<b>4.14 5% (w/v) Skimmed milk in 1X TBS with 0.1% Tween-20</b>	<b>100</b>	<b>ml</b>
Skimmed milk	5	g
1X TBS with 0.1% Tween-20	100	ml
store at 4 °C and stir before use		

## 5. Solution for immunofluorescent assay (IFA)

### 5.1 1M PBS (10X PBS) for IFA

NaCl	90	g
Na <sub>2</sub> HPO <sub>4</sub>	71	g
NaH <sub>2</sub> PO <sub>4</sub>	60	g
Distilled H <sub>2</sub> O to	800	ml
Adjust pH to	7.4	
Adjust volume with distilled H <sub>2</sub> O to	1	L
store at room temperature		

### 5.2 0.1M PBS (1X PBS)

1M PBS (10X PBS)	100	ml
Distilled H <sub>2</sub> O to	900	ml
store at room temperature		



<b>5.3 0.3% (v/v) Triton X-100 / 1X PBS</b>	<b>100</b>	<b>ml</b>
Triton X-100	0.3	ml
1X PBS to	100	ml
store at room temperature		
<b>5.4 0.03% (v/v) Triton X-100 / 1X PBS</b>	<b>100</b>	<b>ml</b>
Triton X-100	0.03	ml
1X PBS to	100	ml
store at room temperature		