

October

October 7th

Experiment: Lysin induction

Responsables: Pablo Delgado

Protocol code: InEc

Protocol modifications or specifications: The culture was prepared with 10 mL of preculture, 90 mL of LB medium and 100 mg/mL of ampicillin. At the induction step, 1 mM IPTG was added. It was incubated for 10 hours.

Results: SDS-PAGE from october 8th.

October 7th

Experiment: Lysin extraction

Responsables: María José

Protocol code: ExEc

Protocol modifications or specifications: total volume from induction was centrifuged and pellet was resuspended in 10 mL of Lysis Buffer. Then, 80 uL of PMSF, 0.5 uL of DNase, 200 uL of MgCl₂ and 0.6 uL of mercaptoethanol were added.

Results: SDS-PAGE from October 8th.

October 7th

Experiment: Lysin purification

Responsables: María José

Protocol code: LysP

Protocol modifications or specifications: Supernatant from protein extraction was incubated overnight with 100 uL of resin, previously washed. Resin was resuspended in 10 mM Imidazole.

Results: SDS-PAGE from october 8th.

October 8th

Experiment: Lysin purification

Responsables: María José

Protocol code: LysP

Protocol modifications or specifications: After incubation, the protein was purified by affinity chromatography. Non Bound, Wash 1 with 10 mM Imidazole, Wash 2 with 20 mM Imidazole and three Elutions with 500 mM Imidazole were obtained.

Results: SDS-PAGE from october 8th.

October 8th

Experiment: SDS-PAGE

Responsables: María José

Protocol code: SDSPAGE

Protocol modifications or specifications: SDS-PAGE was run with samples of lysin induction, extraction and purification from October 7th. The gel was run for 30 min at 90 V and 1 hour and a half at 120 V.

Results:



October 9th

Experiment: Lysin induction

Responsables: Pablo Delgado

Protocol code: InEc

Protocol modifications or specifications: The culture was prepared with 25 mL of preculture, 375 mL of LB medium and 100 mg/mL of ampicillin. At the induction step, 1 mM IPTG was added. It was incubated for 6 hours.

Results: SDS-PAGE from October 10th.

October 9th

Experiment: Lysin extraction

Responsables: María José

Protocol code: ExEc

Protocol modifications or specifications: total volume from induction was centrifuge and then, the eight pellets were resuspended in 5 mL of Lysis Buffer. Resuspended pellets were joined in two falcons (20 mL each one) and then 200 uL of PMSF, 1 uL of DNase, 400 uL of MgCl₂ and 1.7 uL of mercaptoethanol were added.

Results: SDS-PAGE from October 10th.

October 10th

Experiment: Lysin purification

Responsables: María José

Protocol code: LysP

Protocol modifications or specifications: Supernatant from protein extraction was incubated overnight with 400 uL of resin, previously washed. Resin was resuspended in 10 mM Imidazole. After incubation, the protein was purified by affinity chromatography. Non Bound, Wash 1 with 10 mM Imidazole, Wash 2 with 20 mM Imidazole and three Elutions with 500 mM Imidazole were obtained.

Results: SDS-PAGE from October 10th.

October 10th

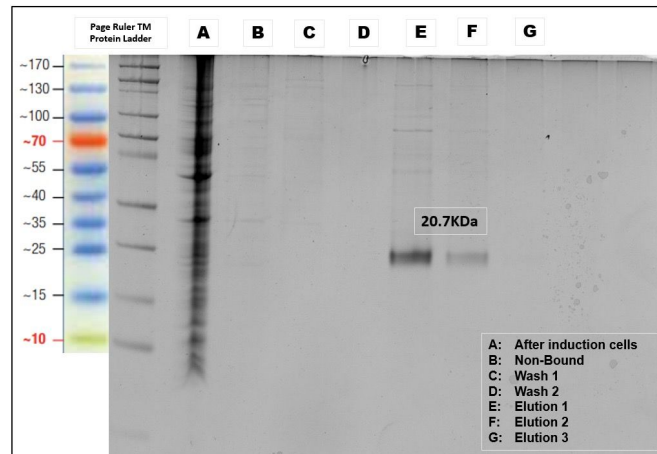
Experiment: SDS-PAGE

Responsables: María José

Protocol code: SDSPAGE

Protocol modifications or specifications: SDS-PAGE was run with samples of lysin induction, extraction and purification from October 7th. The gel was run 30 min at 90 V and 1 hour and a half at 120 V.

Results:



October 11th

Experiment: Plasmid Isolation

Responsables: Pablo Delgado

Protocol code: ThermoScientific Kit, K0503

Protocol modifications or specifications: At the final step, plasmidial DNA was eluted using 40μL of nuclease free water.

Results:

Name	Resistance	Concentration (ng/uL)	Absorbance
ILC1	C	231.5	1.85
ILC2	C	218	1.85
ILC3	C	204.6	1.84
ILC C+	C	172.5	1.85
ILC C-	C	99.6	1.82
Lisina E. coli	A	415	1.87
K2235000	A	430	1.86

October 11th

Experiment: Lysin dialysis and concentration

Responsables: María José

Protocol code: Protein_DC

Protocol modifications or specifications: Lysin was dialysed and concentrated with buffer 50 mM Tris pH 7.45 in a 3K Amicon. It was centrifuged at 5000 g for approximately 1 hour/round.

Results:

Concentration	
Lysin*	0.12 mg/mL
Imidazole	0.59 mM

*Protein correction factor was 0.659

October 11th

Experiment: *Clostridium difficile* NTCC 13307 culture

Responsables: Anthony Mora

Protocol code: a cryopreserved culture was stroke in Blood agar.

Protocol modifications or specifications: None

Results: Not needed

October 12th

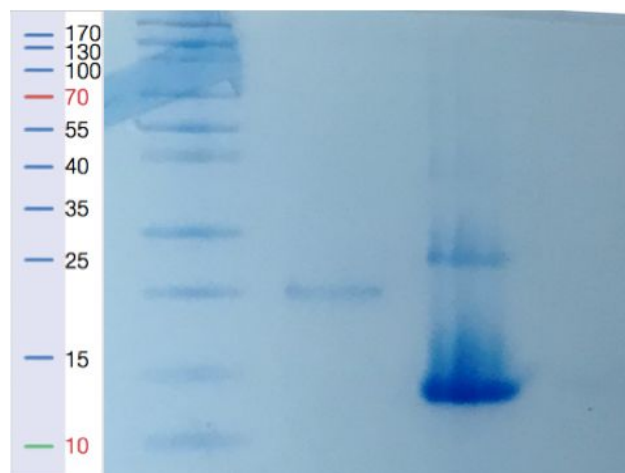
Experiment: SDS-PAGE

Responsables: María José

Protocol code: SDSPAGE

Protocol modifications or specifications: SDS-PAGE was run with samples of lysin dialysis and concentration from October 11th. The gel was run 30 min at 90 V and 1 hour and a half at 120 V.

Results:



(1) Molecular weight marker. (2) Lysin dialysed and concentrated. (3) Lysozyme 0.5 mg/mL.

October 12th

Experiment: AIP induction

Responsables: María José Durán

Protocol code: InEc

Protocol modifications or specifications: The culture was prepared with 40 mL of preculture, 360 mL of LB medium and 100 mg/mL of ampicillin. At the induction step, 1 mM IPTG was added. It was incubated O/N.

Results: SDS-PAGE from October 15th.

October 13th

Experiment: AIP extraction

Responsables: María José

Protocol code: ExEc

Protocol modifications or specifications: total volume from induction was centrifuge and then, the eight pellets were resuspended in 5 mL of Lysis Buffer. Resuspended pellets were joined in two falcons (20 mL each one) and then 200 uL of PMSF, 1 uL of DNase, 400 uL of MgCl₂ and 1.7 uL of mercaptoethanol were added.

Results: SDS-PAGE from October 15th.

October 13th

Experiment: AIP concentration

Responsables: María José

Protocol code: Protein_DC

Protocol modifications or specifications: AIP was dialysed and concentrated with buffer 50 mM Tris pH 7.45 in a 3K Amicon. It was centrifuged at 5000 g for approximately 1 hour/round.

Results: SDS-PAGE from October 15th.

October 13th

Experiment: Lysin induction

Responsables: María José

Protocol code: InEc

Protocol modifications or specifications: The culture was prepared with 40 mL of preculture, 360 mL of LB medium and 100 mg/mL of ampicillin. At the induction step, 1 mM IPTG was added. It was incubated for 6 hours.

Results: SDS-PAGE from October 13th.

October 13th

Experiment: Lysin extraction

Responsables: María José

Protocol code: ExEc

Protocol modifications or specifications: total volume from induction was centrifuged and then, the eight pellets were resuspended in 5 mL of Lysis Buffer. Resuspended pellets were joined in two falcons (20 mL each one) and then 200 uL of PMSF, 1 uL of DNase, 400 uL of MgCl₂ and 1.7 uL of mercaptoethanol were added.

Results: SDS-PAGE from October 13th.

October 13th

Experiment: Lysin purification

Responsables: María José

Protocol code: LysP

Protocol modifications or specifications: Supernatant from protein extraction was incubated overnight with 400 uL of resin, previously washed. Resin was resuspended in 10 mM Imidazole. After

incubation, the protein was purified by affinity chromatography. Non Bound, Wash 1 with 10 mM Imidazole, Wash 2 with 20 mM Imidazole and three Elutions with 500 mM Imidazole were obtained.

Results: SDS-PAGE from October 13th.

October 13th

Experiment: Lysin dialysis and concentration

Responsables: María José

Protocol code: Protein_DC

Protocol modifications or specifications: Lysin was dialysed and concentrated with buffer 50 mM Tris pH 7.45 in a 3K Amicon. It was centrifuged at 5000 g for approximately 1 hour/round.

Results:

Concentration	
Lysin*	-
Imidazole	11.9 mM

*Protein correction factor was 0.659

October 14th

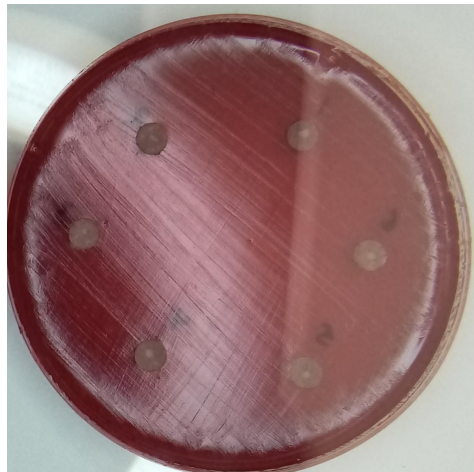
Experiment: Lysine Susceptibility Testing on *Clostridium difficile* NTCC 13307

Responsables: Anthony Mora

Protocol code: Lys_Sus

Protocol modifications or specifications: PBS was used as a negative control

Results: There was no protein activity, no inhibition halos were observed



Columbia agar with *C. difficile* NTCC 13307 inhibition zones

October 15th

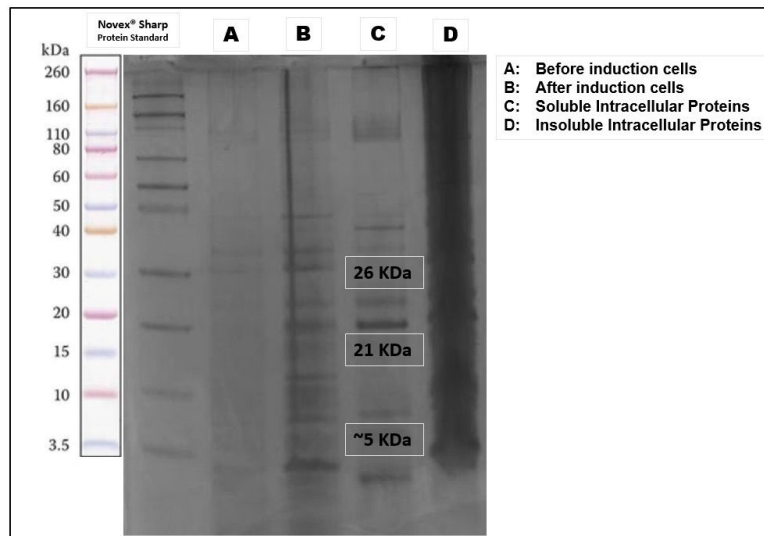
Experiment: SDS-PAGE

Responsables: María José

Protocol code: SDSPAGE

Protocol modifications or specifications: SDS-PAGE was run with the dialysed AIP supernatant extract from October 13th. The gel was run for 30 min at 90 V and 1 hour and a half at 120 V.

Results:



October 16th

Experiment: AIP induction

Responsables: María José Durán

Protocol code: InEc

Protocol modifications or specifications: The culture was prepared with 7.5 mL of preculture, 67.5 mL of LB medium and 100 mg/mL of ampicillin. At the induction step, 1 mM IPTG was added. It was incubated for 9 hours. After centrifugation, supernatant was stored at 4°C for a later lyophilization.

Results: SDS-PAGE from October 19th.

October 17th

Experiment: AIP extraction

Responsables: María José

Protocol code: ExEc

Protocol modifications or specifications: total volume from induction was centrifuged and then, the eight pellets were resuspended in 5 mL of Lysis Buffer. Resuspended pellets were joined in two falcons (20 mL each one) and then 200 uL of PMSF, 1 uL of DNase, 400 uL of MgCl₂ and 1.7 uL of mercaptoethanol were added.

Results: SDS-PAGE from October 19th.

October 17th

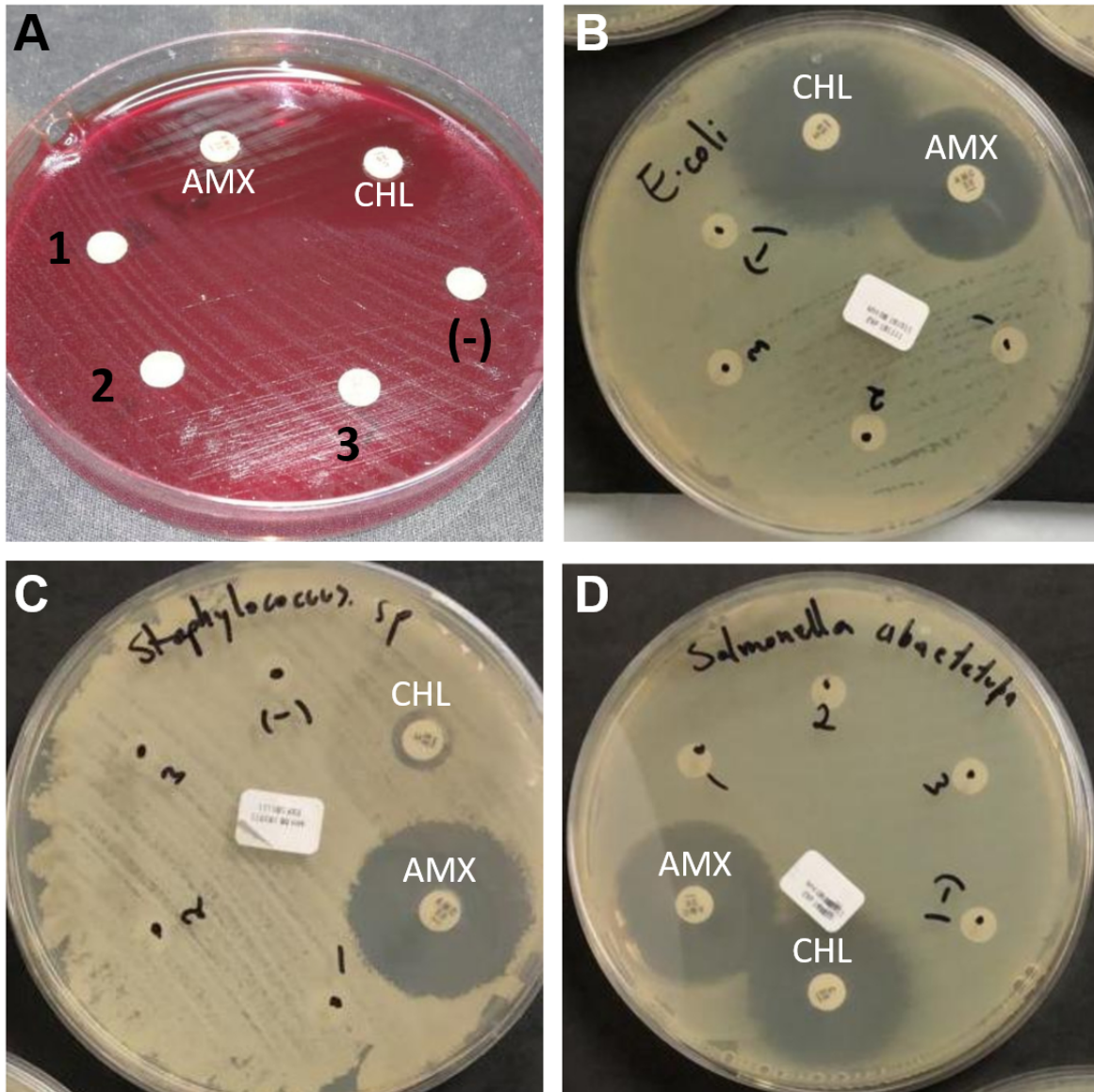
Experiment: Lysin Susceptibility Testing on *Clostridium difficile* NTCC 13307

Responsables: Anthony Mora

Protocol code: Lys_Sus

Protocol modifications or specifications: *Salmonella abaeetuba*, *Escherichia coli* and *Staphylococcus aureus* were tested in Mueller-Hinton agar with amoxicillin and chloramphenicol as positive controls and PBS as negative control. The inoculum was standardized to McFarland 0.5.

Results: There was no protein activity, no inhibitory halos were observed around the disks with protein (1, 2 and 3)



Lysis assay of endolysin CD27L1-179 on: A) *Clostridium difficile*, B) *Escherichia coli*, C) *Staphylococcus* sp. and D) *Salmonella abacetuba*. Amoxicillin (AMX) and chloramphenicol (CHL) were tested as positive control agents and PBS as negative control (-); both antibiotics were used by recommendation of Andino-Molina and colleagues (2019). Lysin was assayed with three different concentrations (1: 120µg/mL, 2: 60µg/mL and 3: 30µg/mL) to evaluate its growth inhibition capacity.

October 19th

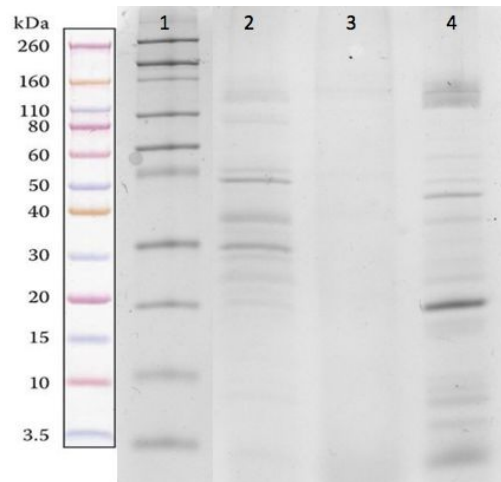
Experiment: SDS-PAGE

Responsables: María José

Protocol code: SDSPAGE

Protocol modifications or specifications: SDS-PAGE was run with the dialysed AIP supernatant from induction and extraction, and lyophilized induction supernatant from October 13th. The gel was run for 30 min at 90 V and 1 hour and a half at 120 V.

Results:



(1) Molecular Marker, (2) Extracellular proteins after dialyzation, (3) Extracellular proteins after lyophilization, (4) Soluble Intracellular proteins.

October 20th

Experiment: Quantification of bacterial fluorescence using independent calibrants for promotor K225300 characterization.

Responsables: Anthony Mora, Paula Thiel, Pablo Delgado and Diego Rojas

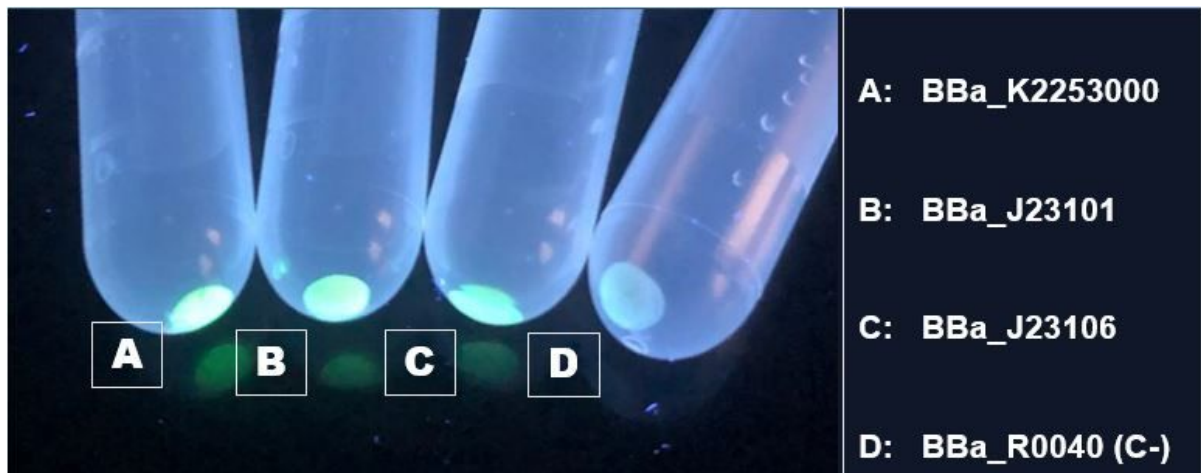
Protocol code: Quan_fluo

Protocol modifications or specifications: The device 3: J23117.B0034.E0040.B0015 in pSB1C3 was not tested in the experiment

Results:

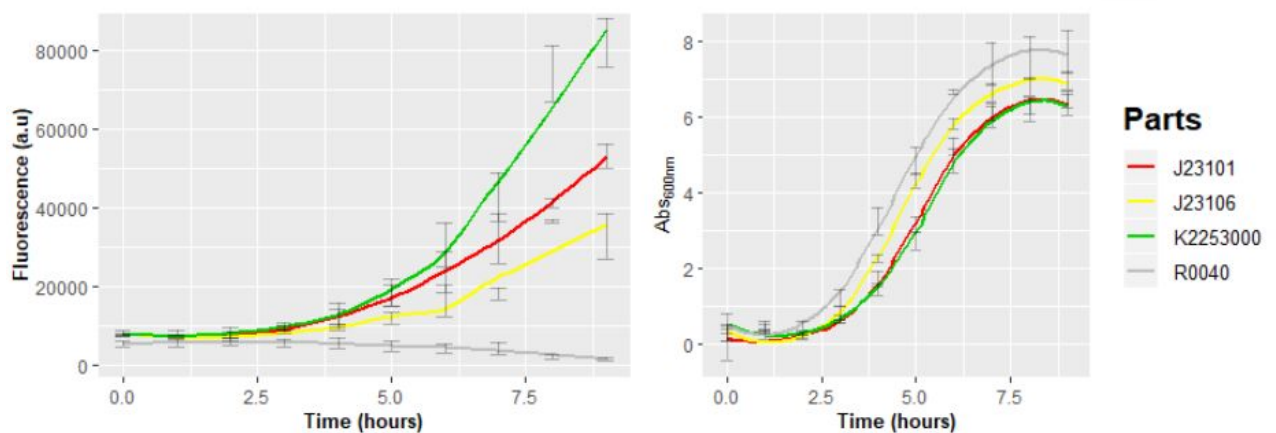
In order to measure the expression strength of this part from *Lactococcus* in *E. coli* we placed a GFP and a terminator downstream of this sequence. The final construct designed for this characterization consisted on the following iGEM parts: K2253000+E0040+B0015.

Therefore, we used two other GFP expression cassettes (provided by iGEM 2016 InterLab), which were regulated under a well characterized Anderson Promoter and an Elowitz RBS (B0034), as references for comparison. The strong expression cassette contained the J23101 promoter and the medium expression cassette the J23106 promoter. Both of these cassettes were cloned in the same iGEM backbone as the K2253000 characterization construct.



Qualitative differences between this part and standard biobricks for InterLab iGEM2016

The curve was started with an OD_{600} 0.05. This was done in triplicate. It was incubated for 9 hours at 37°C and 200 rpm. Samples for GFP (485nm, 528nm) and OD_{600} quantification were taken every hour and measured in a Tecan Biotek plate reader. Non parametric statistics analysis were implemented with Minitab 19 Statistical Software (2019).



Growth curve and fluorescence measurement of K2253000 promoter with GFP

October 21st

Experiment: iGEM 2019 Plate Reader Abs600 (OD) Calibration

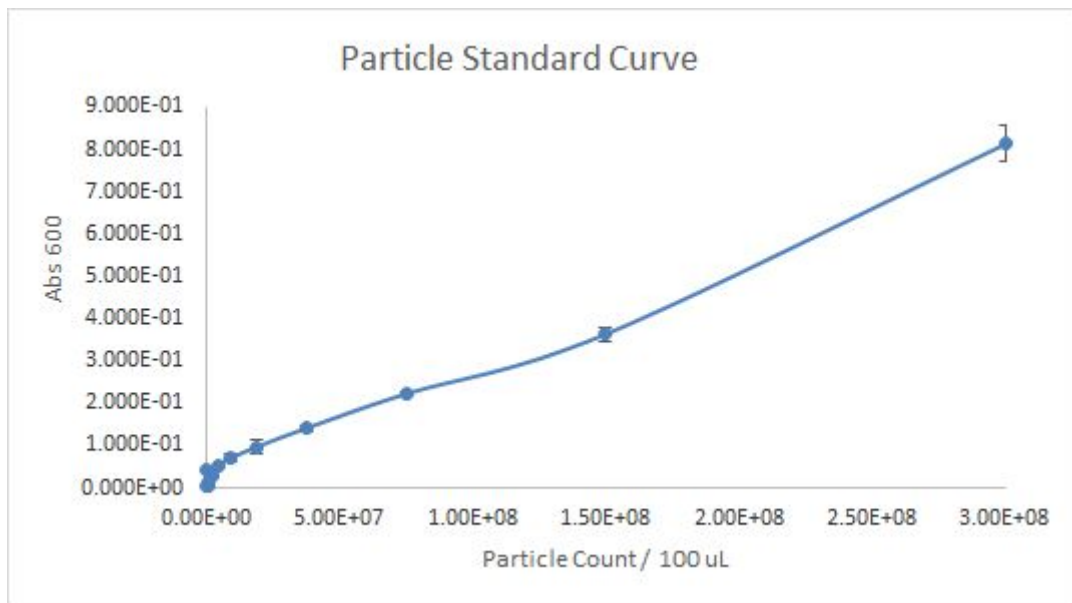
Responsables: Diego Rojas

Protocol code: none

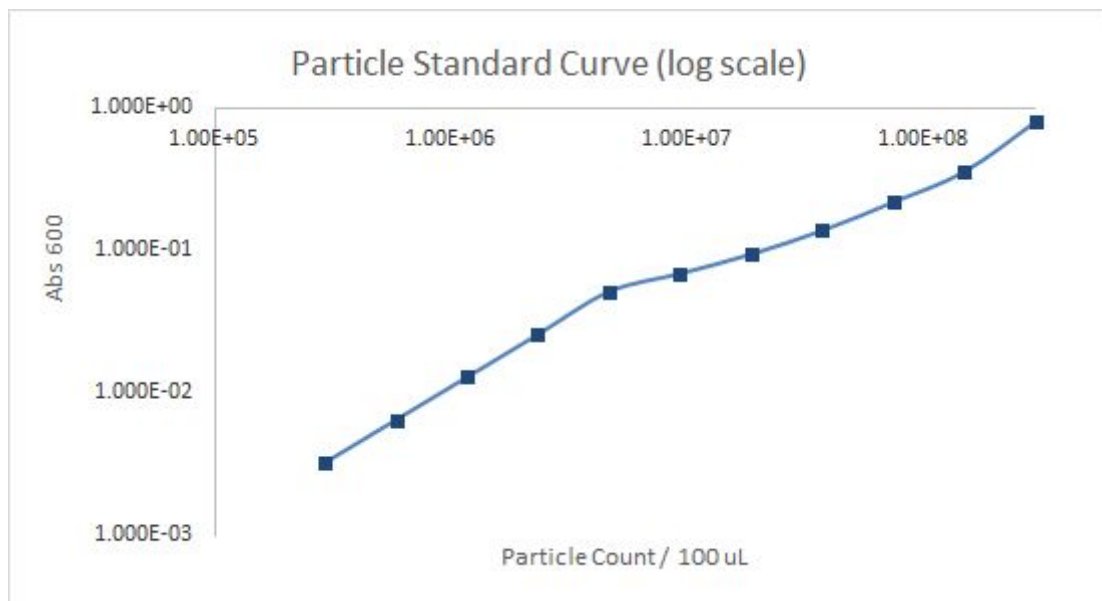
Protocol modifications or specifications: none

Results:

After following the iGEM 2019 calibration protocol we got a Mean particles/Abs600 value of 3.90×10^8 for Mean of med-high levels to standardized our experiments and OD_{600} measurements with a R^2 correlation value of 0.9909 in our Biotek Synergy HTX Multi-mode microplate reader.



Silica Beads Particles Standard Curve Calibration



Silica Beads Particles Standard Curve Calibration at log scale

October 21st

Experiment: iGEM 2019 Plate Reader Fluorescence Calibration

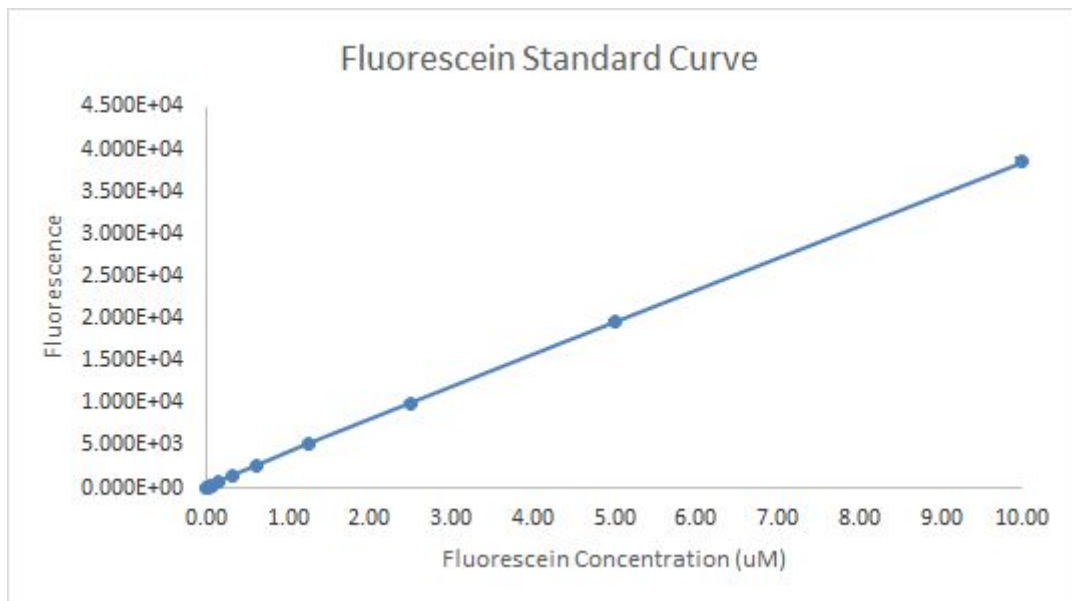
Responsables: Diego Rojas

Protocol code: none

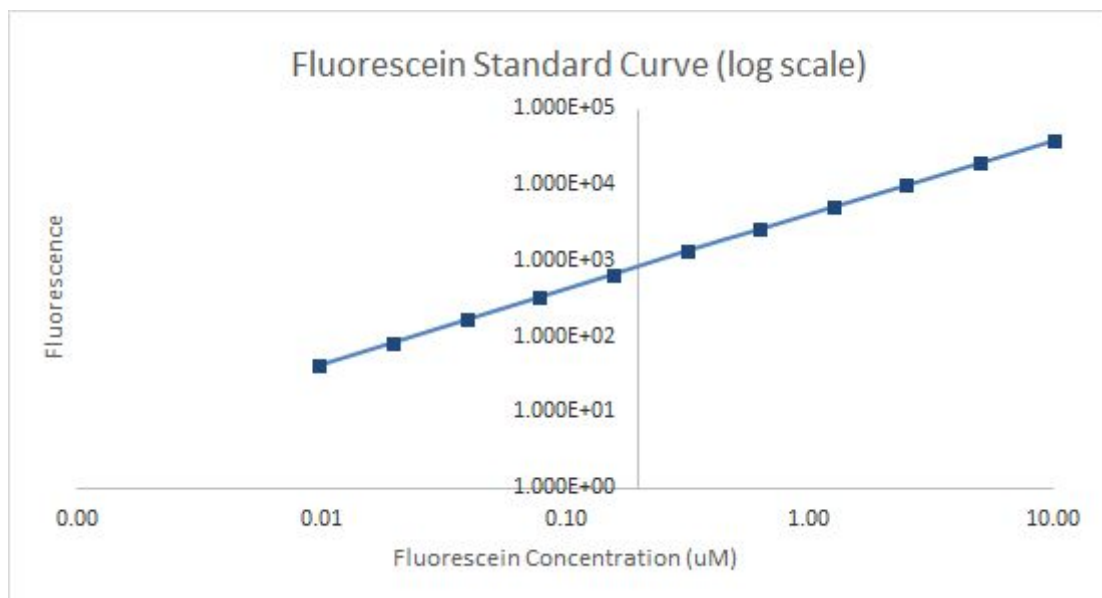
Protocol modifications or specifications: none

Results:

After following the iGEM 2019 calibration protocol we got a MEFL/a.u. value of 1.47×10^{10} to standardized our experiments and fluorescence measurements with a R^2 correlation value of 0.9999 in our Biotek Synergy HTX Multi-mode microplate reader.



Fluorescein Standard Curve Calibration



Fluorescein Standard Curve Calibration at log scale