

Detecting and Quantifying Bacteriophage M13 on Human Skin

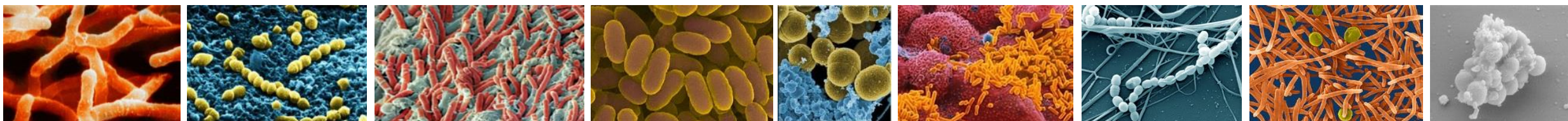
Young Hun (Colin) Kim

Baoqing Zhou

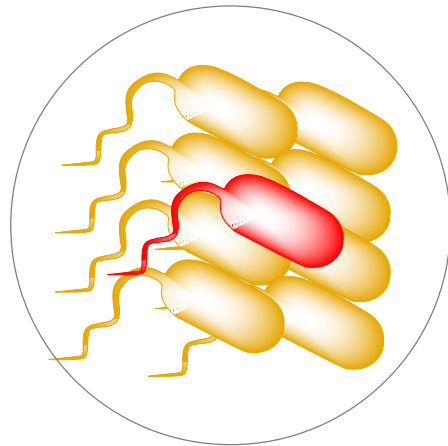
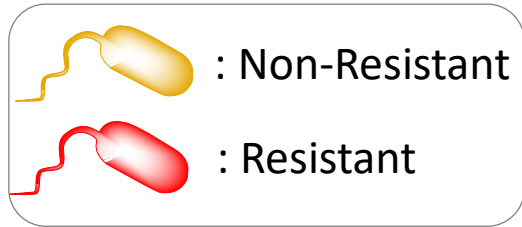
Biochemistry, College of Creative Studies

Chen Lab

Department of Chemistry and Biochemistry, UCSB

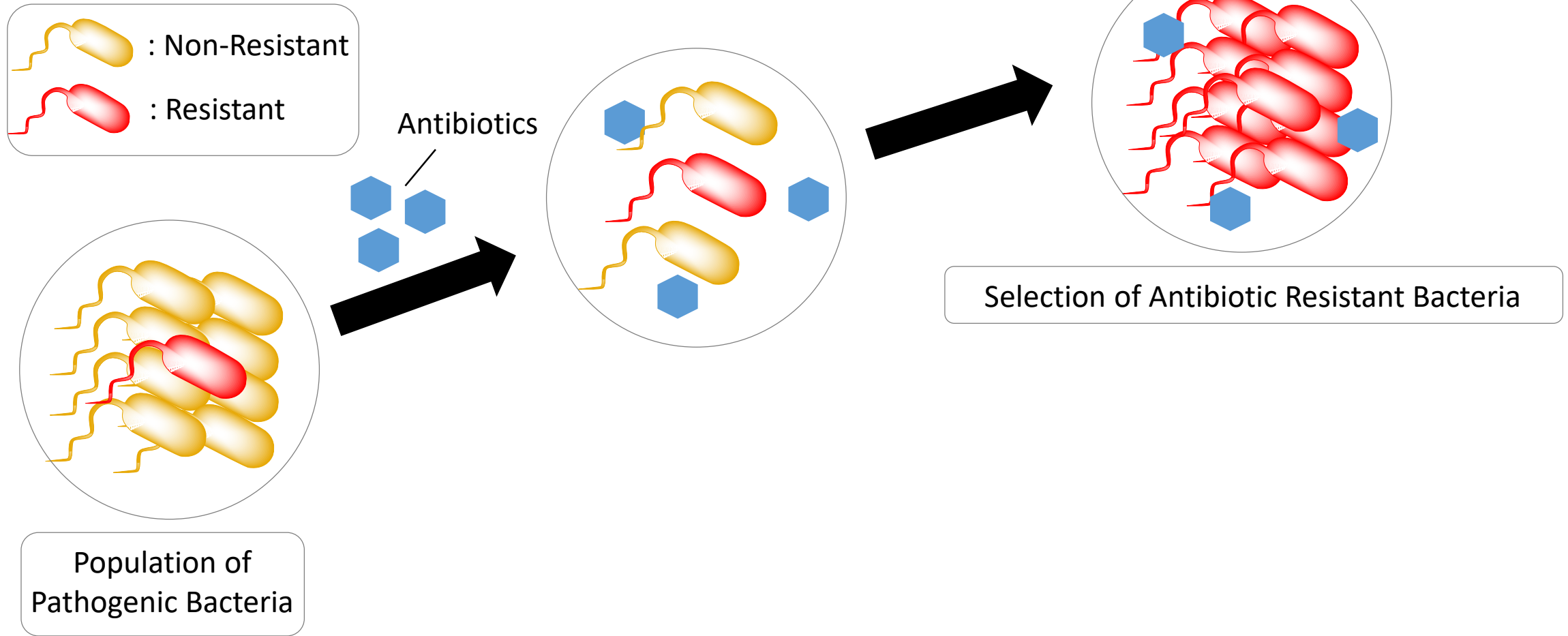


Rapid evolution of antibiotic resistant bacteria

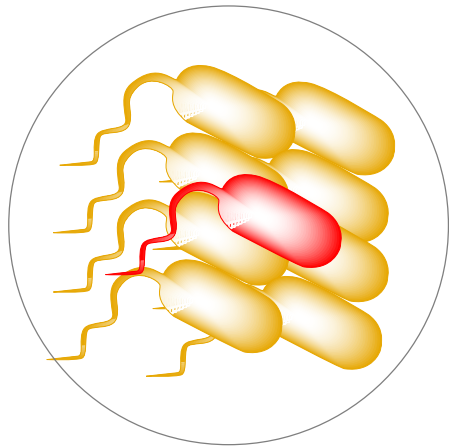
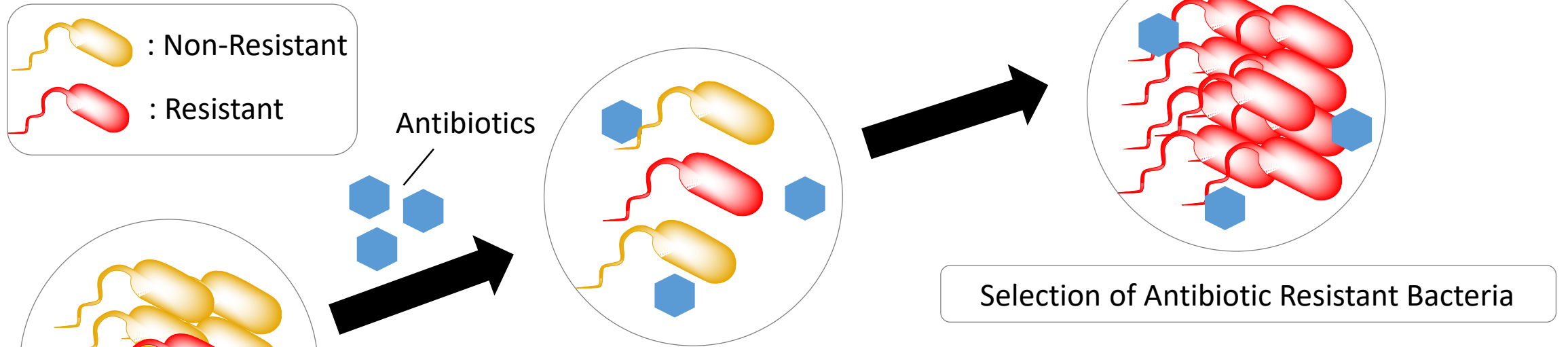


Population of
Pathogenic Bacteria

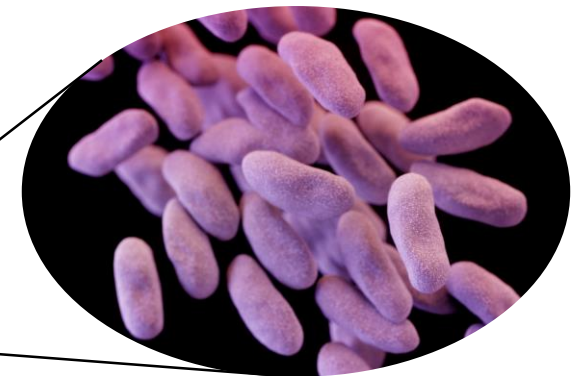
Rapid evolution of antibiotic resistant bacteria



Rapid evolution of antibiotic resistant bacteria




Population of Pathogenic Bacteria

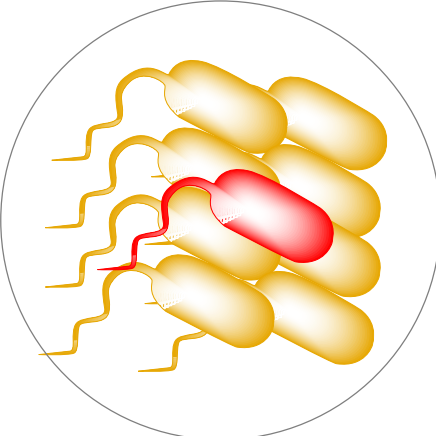


Carbapenem-Resistant Enterobacteriaceae

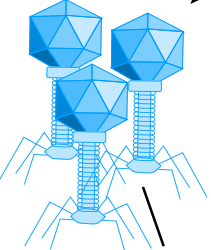
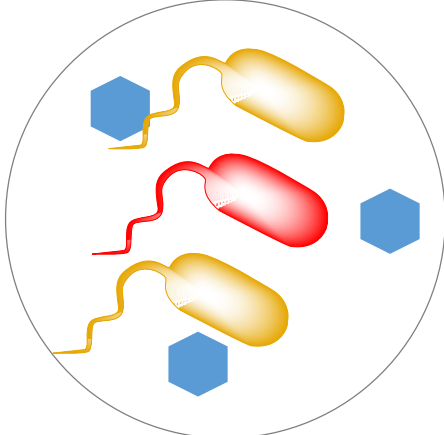
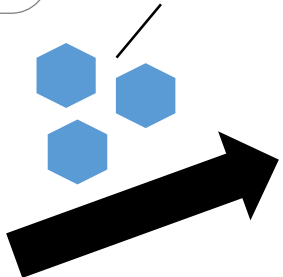
Bacteriophages: natural viruses that infect bacteria

 : Non-Resistant
 : Resistant

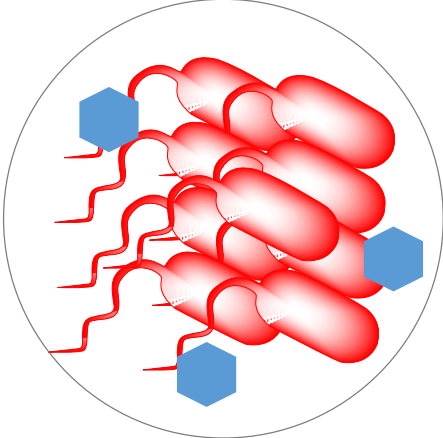
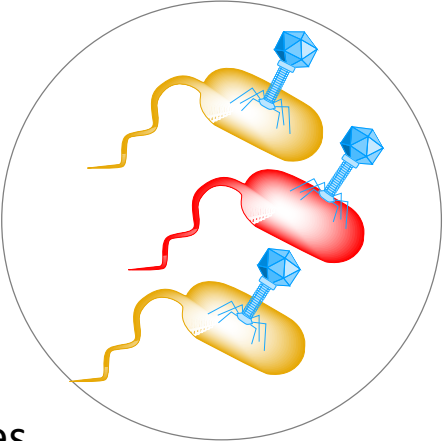
Antibiotics



Population of Pathogenic Bacteria




Bacteriophages

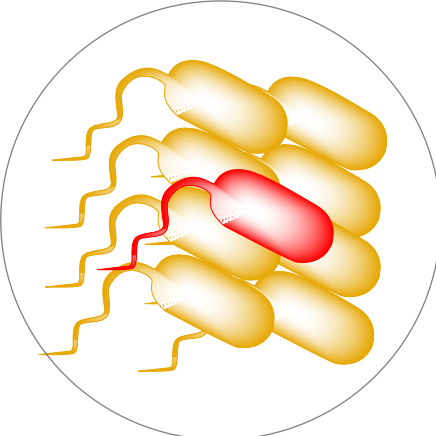


Selection of Antibiotic Resistant Bacteria

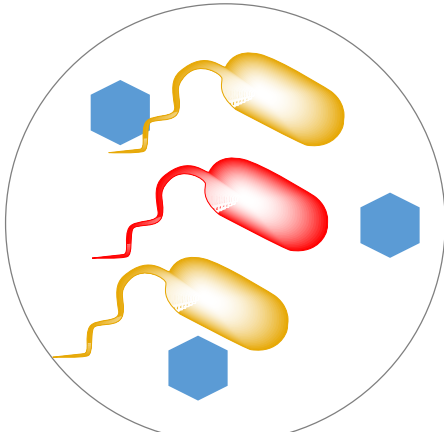
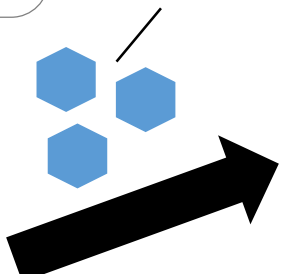
Bacteriophages: natural viruses that infect bacteria

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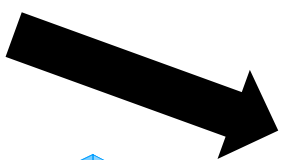
Antibiotics



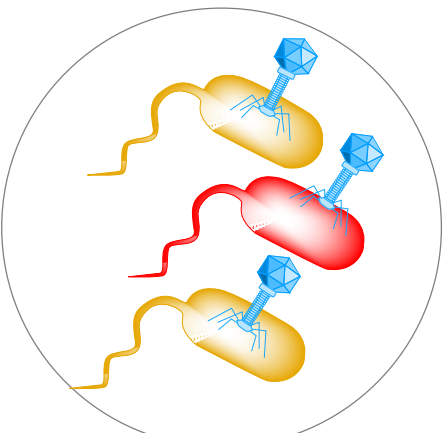
Population of Pathogenic Bacteria



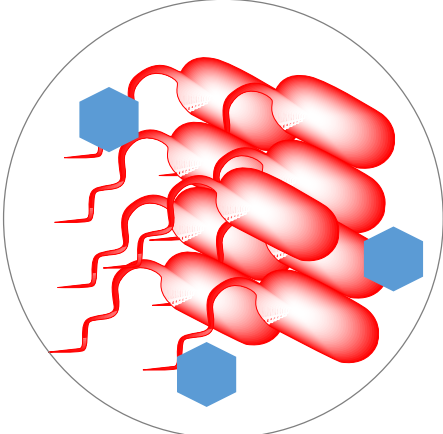
Selection of Antibiotic Resistant Bacteria



Bacteriophages



Successful Treatment of Pathogenic Bacteria



Characterizing the microbial community of wounds



Characterizing the microbial community of wounds

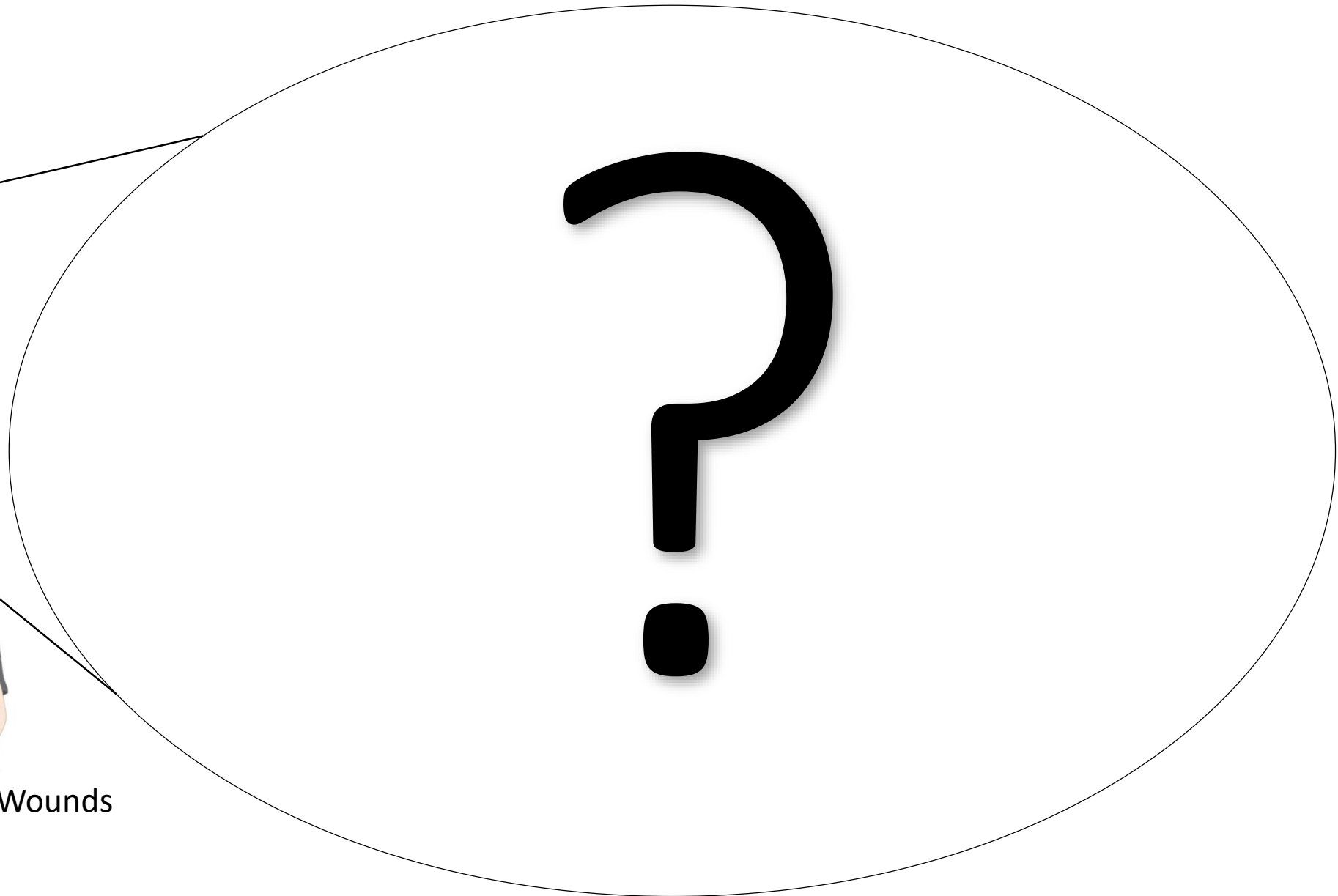
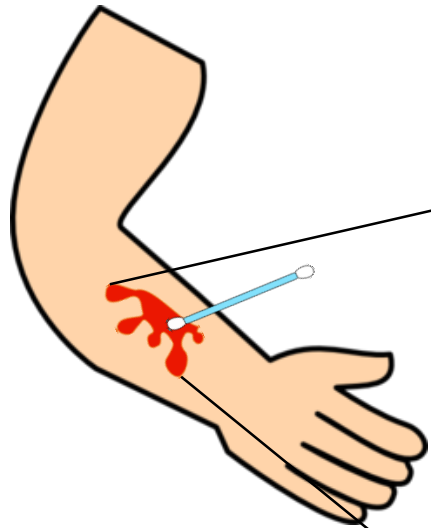


Diabetic Foot Ulcers



Surgical Wounds

Characterizing the microbial community of wounds

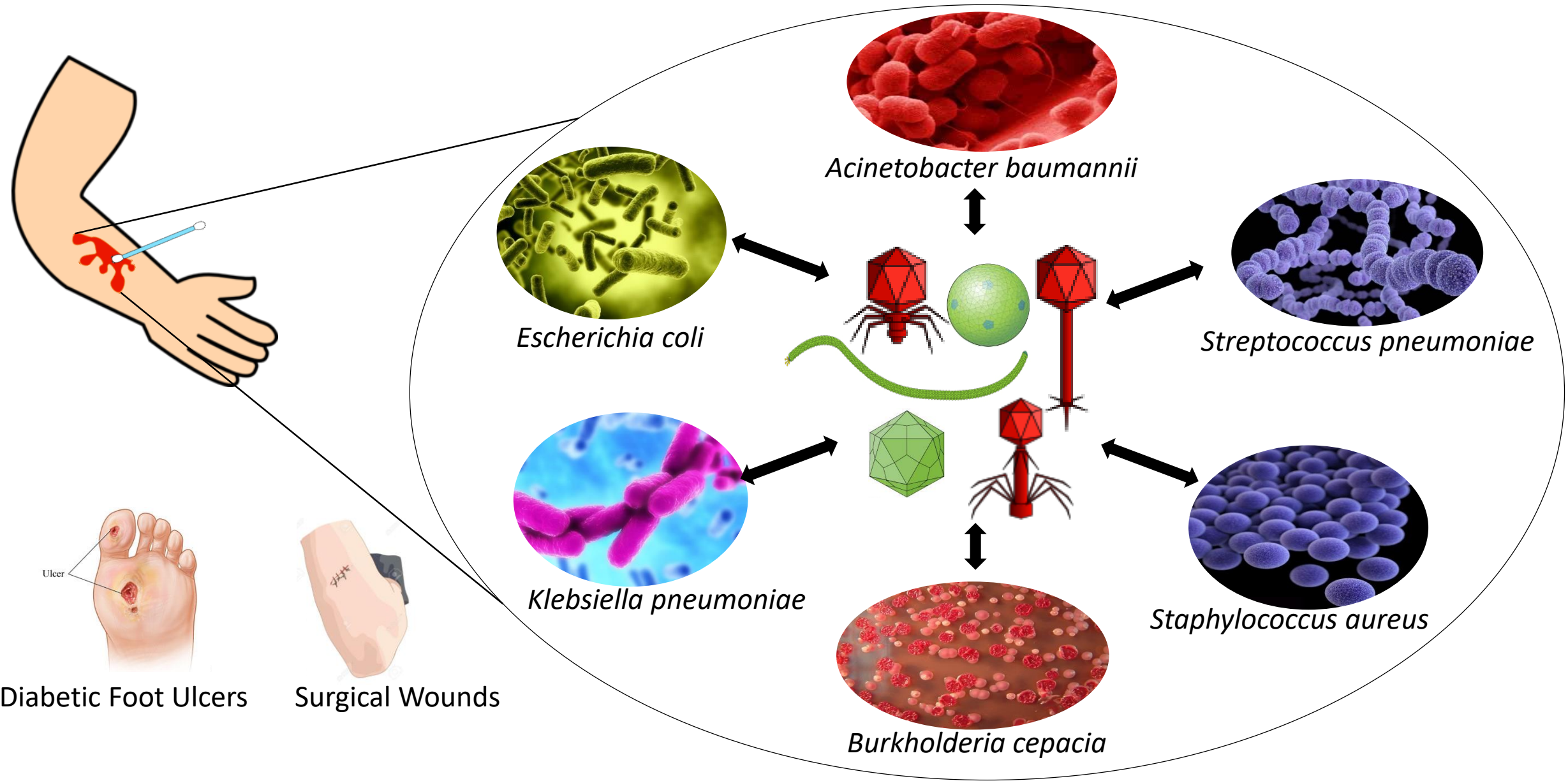


Diabetic Foot Ulcers



Surgical Wounds

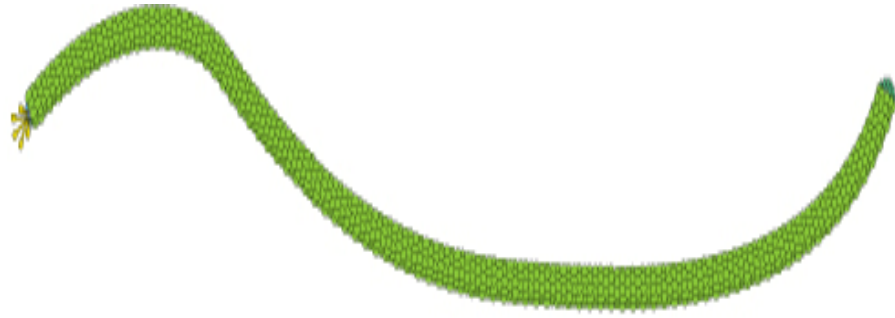
Characterizing the microbial community of wounds



Diabetic Foot Ulcers

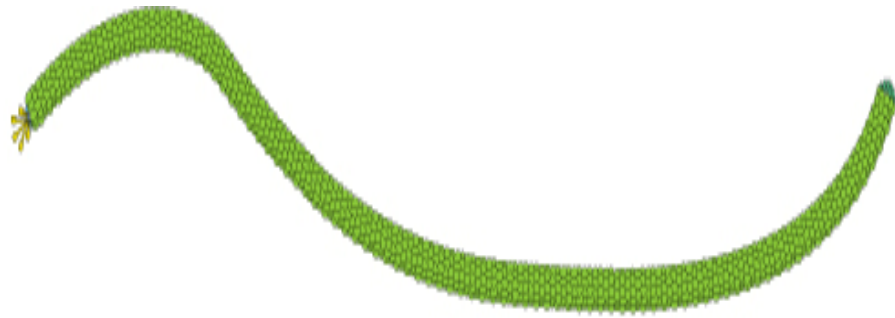
Surgical Wounds

Using phage M13 to develop the skin swab protocol



Bacteriophage M13

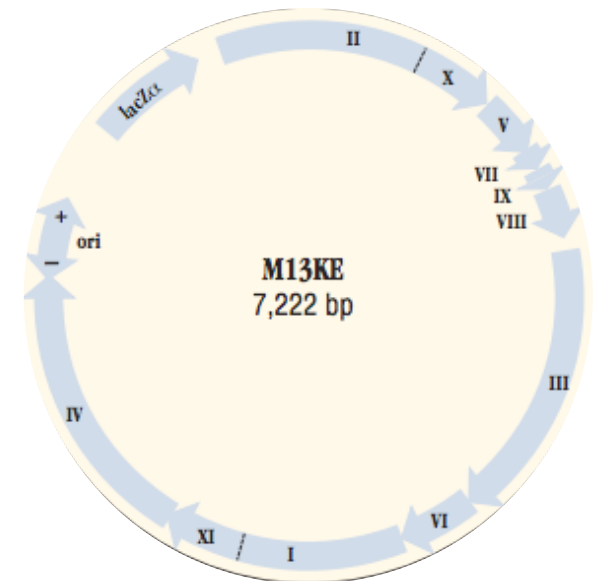
Using phage M13 to develop the skin swab protocol



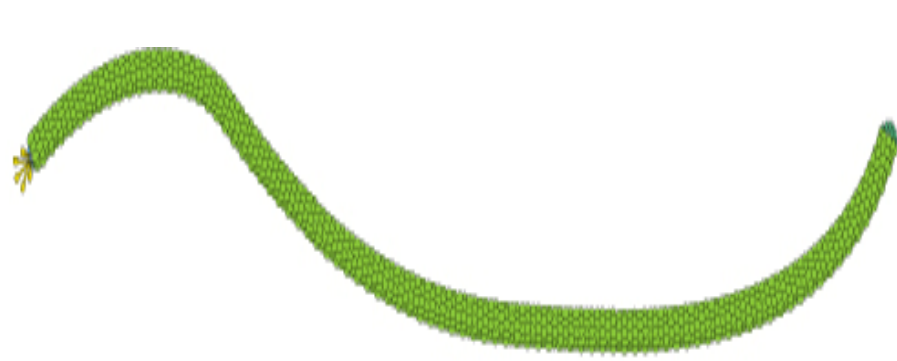
Bacteriophage M13



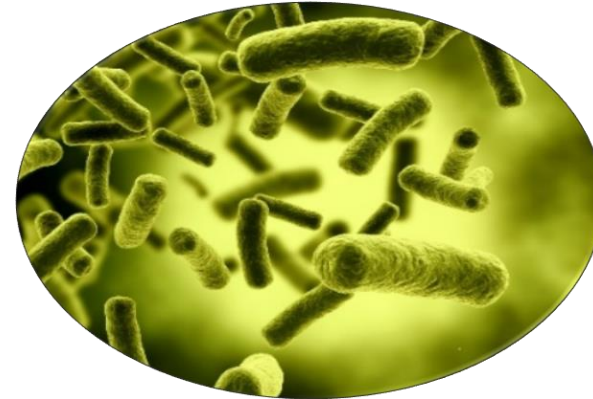
Escherichia coli



Using phage M13 to develop the skin swab protocol

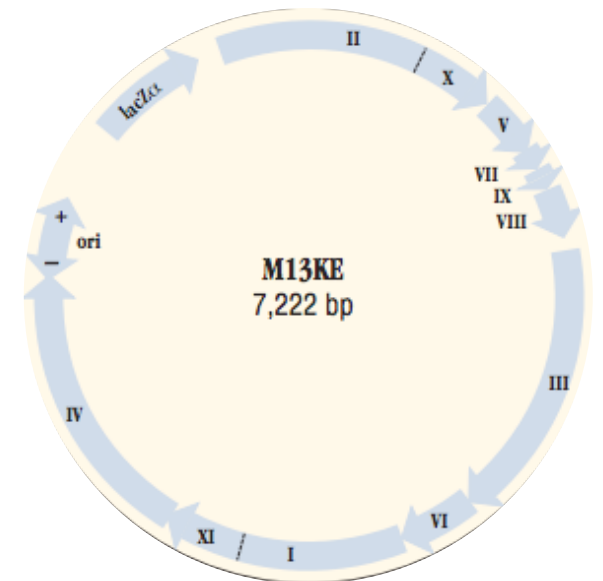
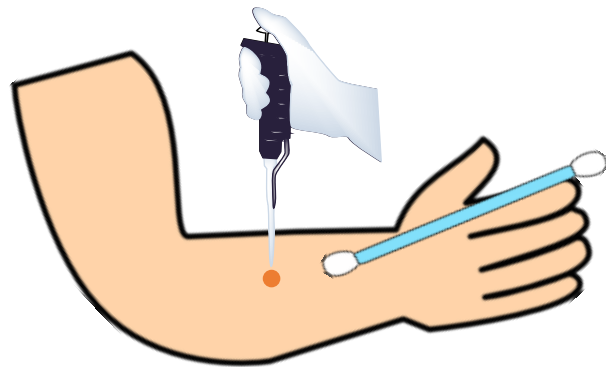


Bacteriophage M13



Escherichia coli

How much phage can we recover by swabbing a human skin?



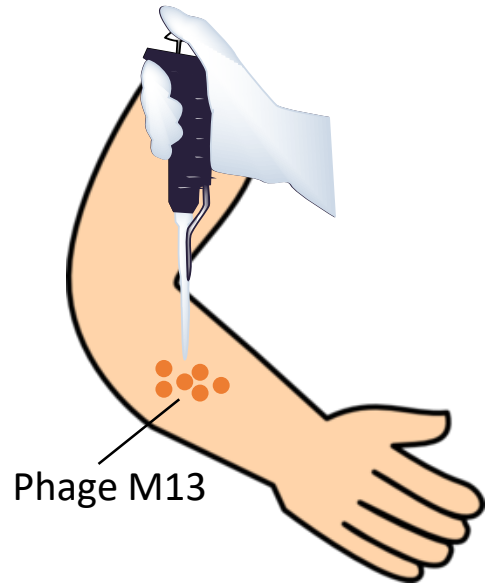
Recovering phage M13 from human skin swab

1. Skin Swab Sample Collection



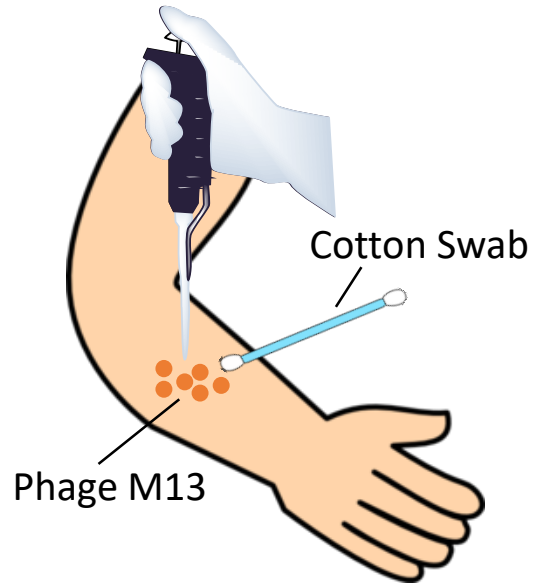
Recovering phage M13 from human skin swab

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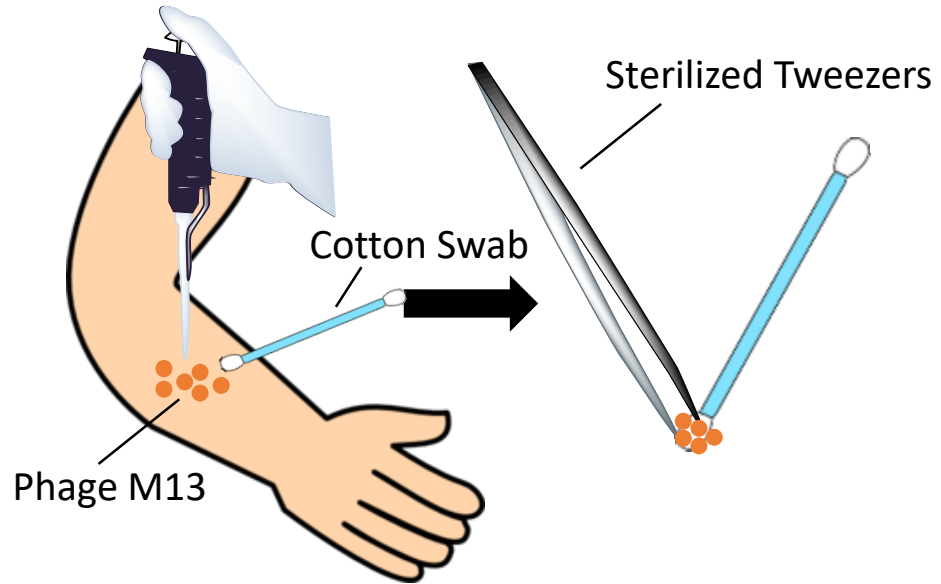
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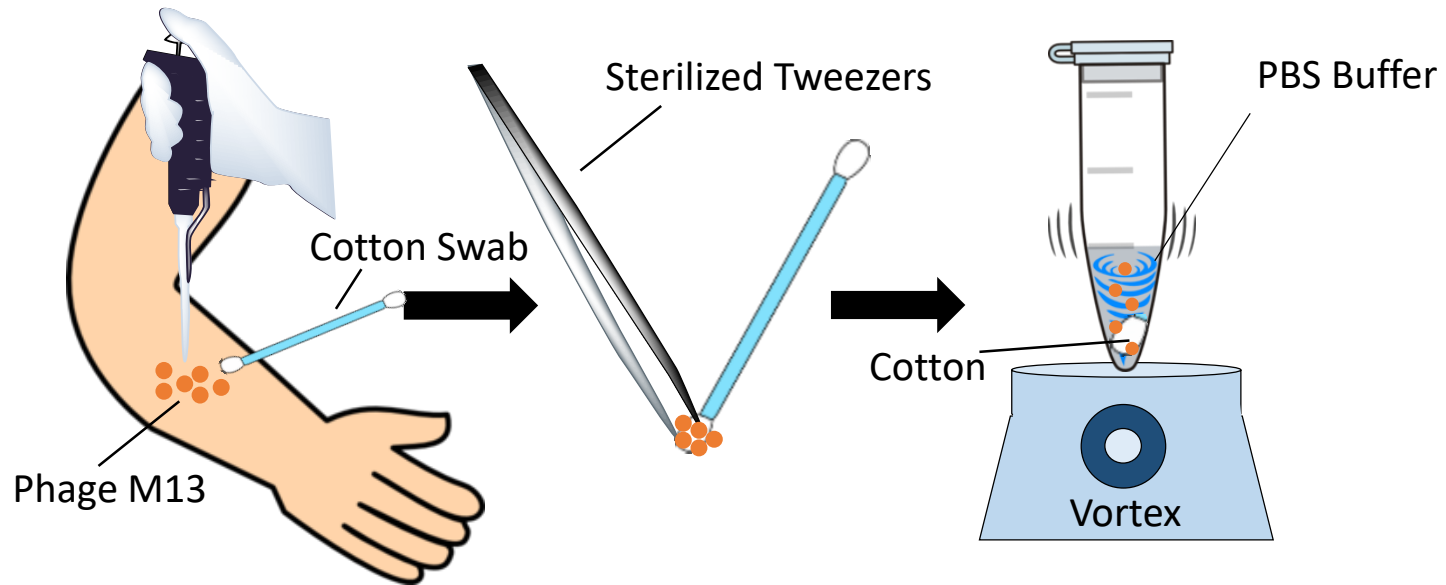
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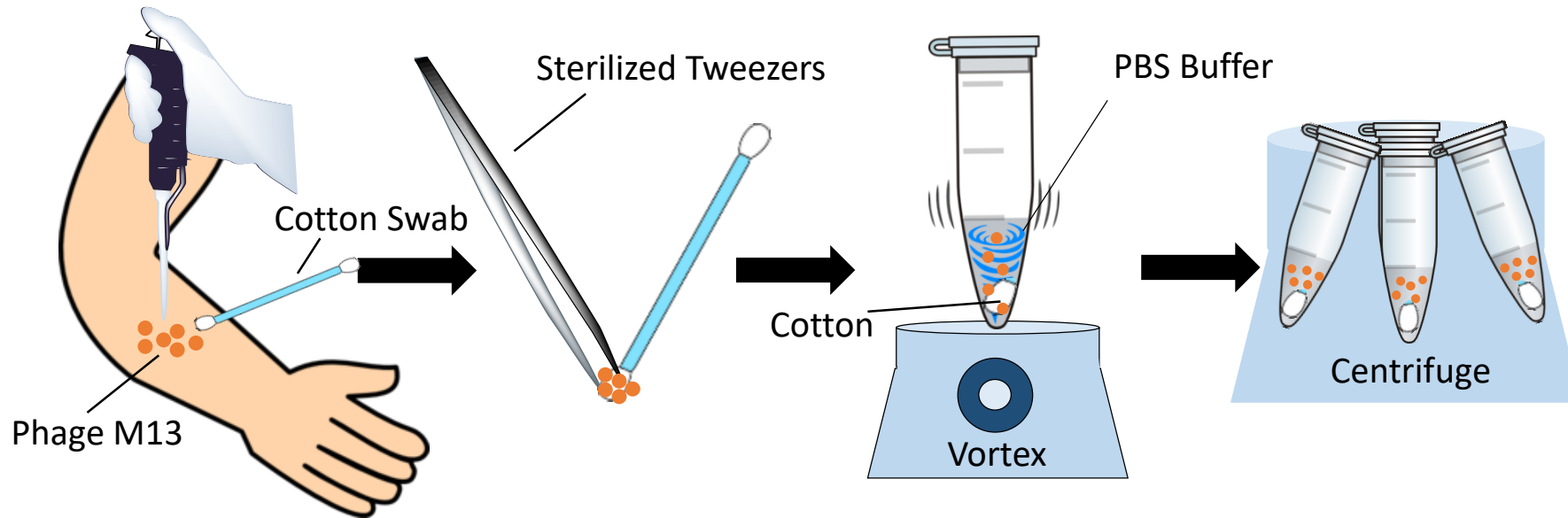
Recovering phage M13 from human skin swab

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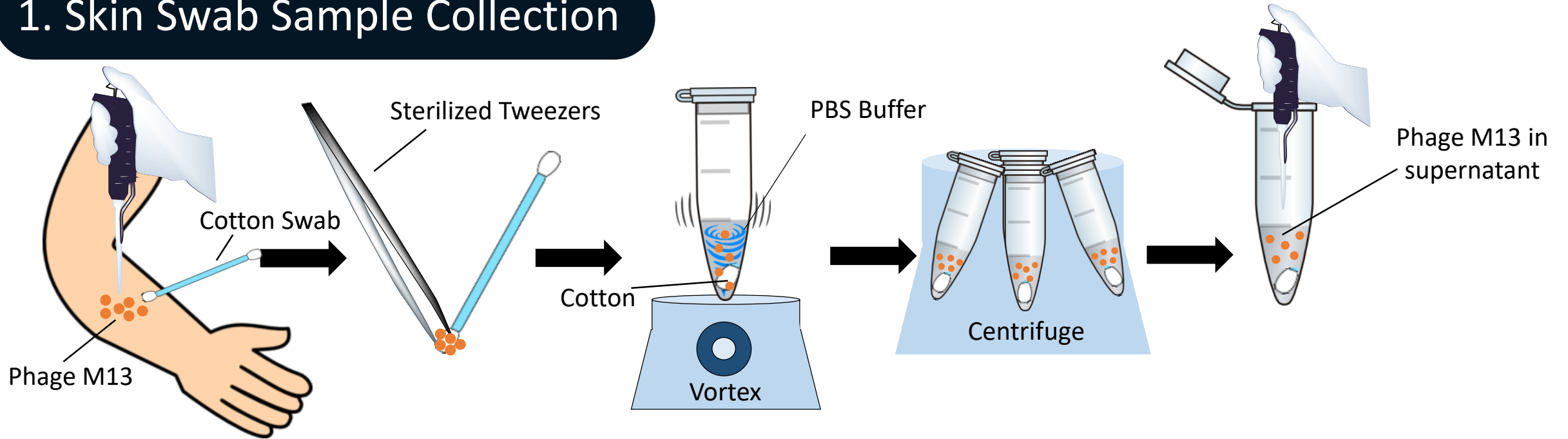
Recovering phage M13 from human skin swab

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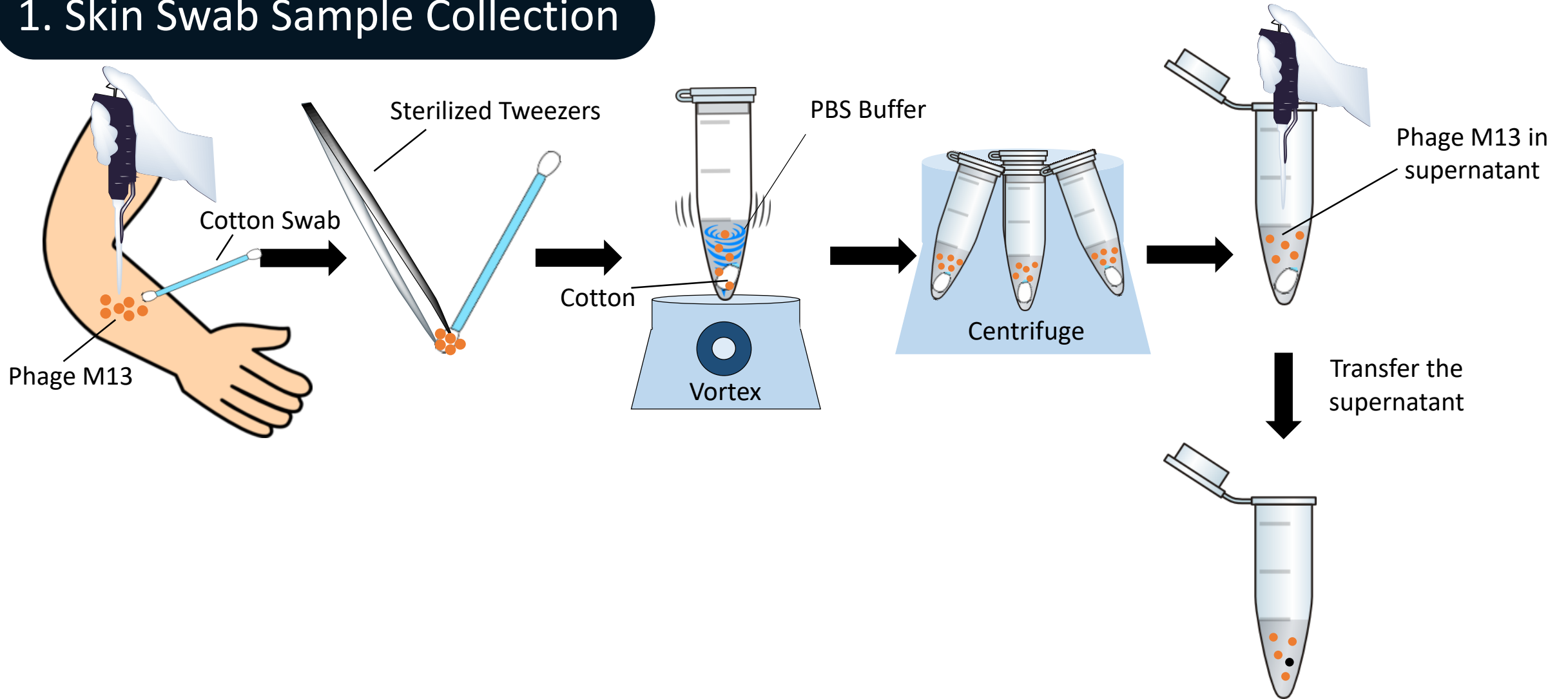
Recovering phage M13 from human skin swab

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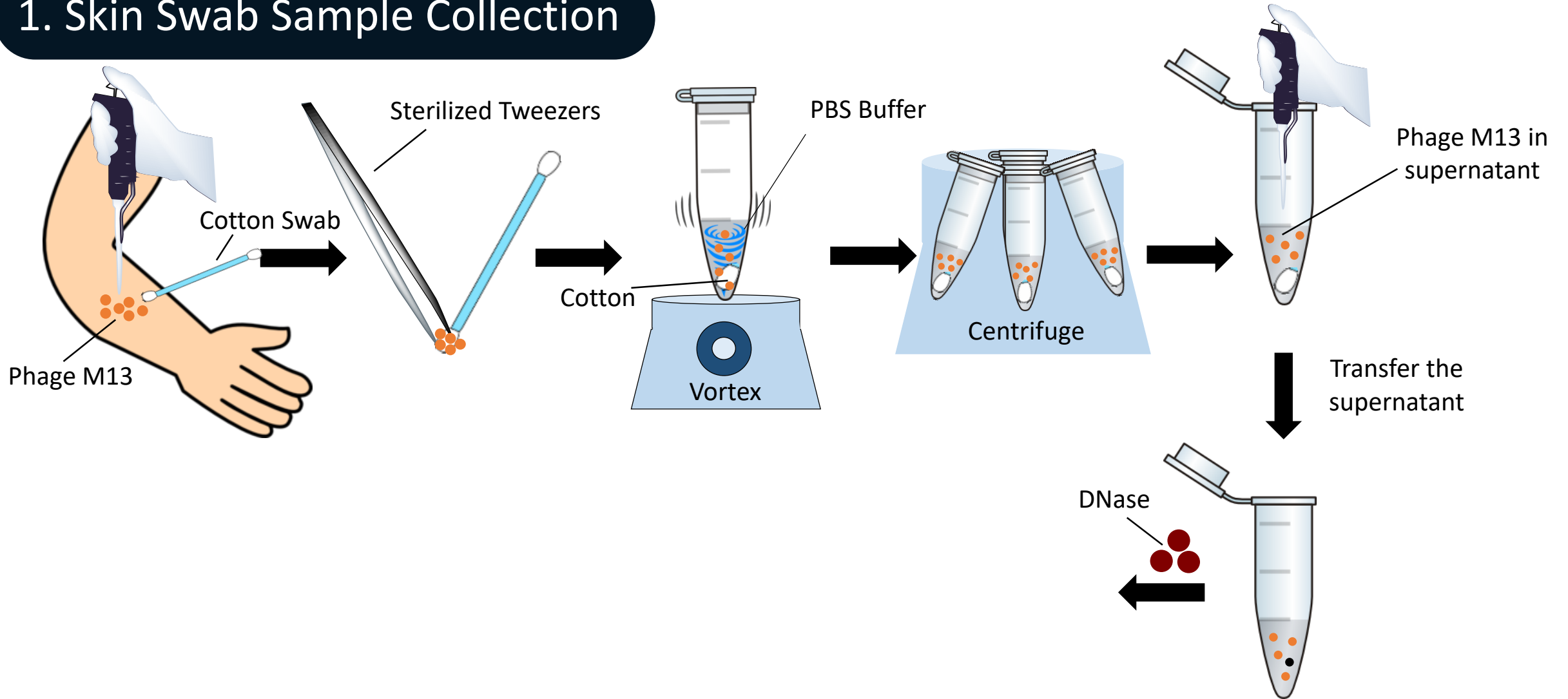
Recovering phage M13 from human skin swab

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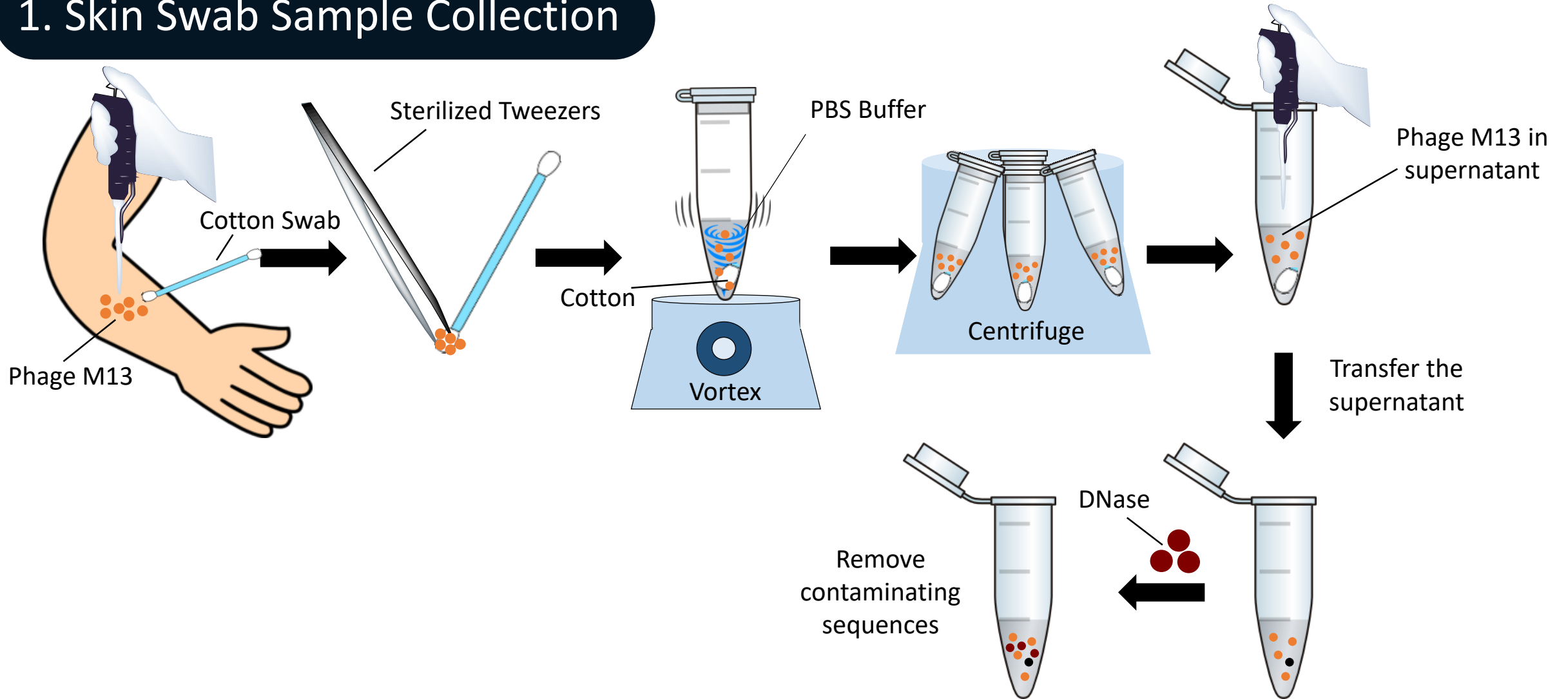
Recovering phage M13 from human skin swab

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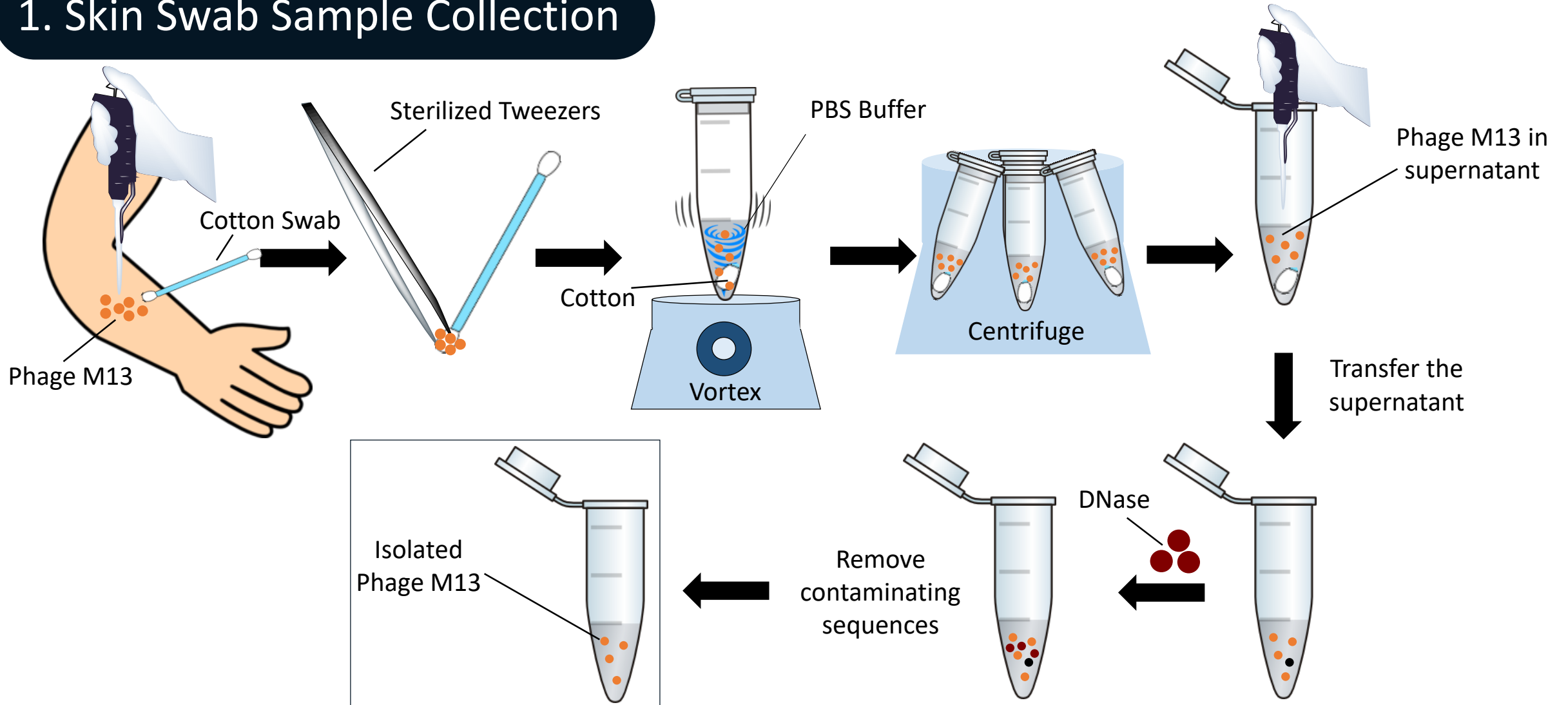
Recovering phage M13 from human skin swab

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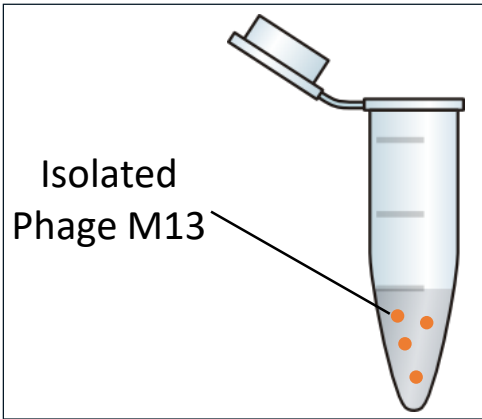
Recovering phage M13 from human skin swab

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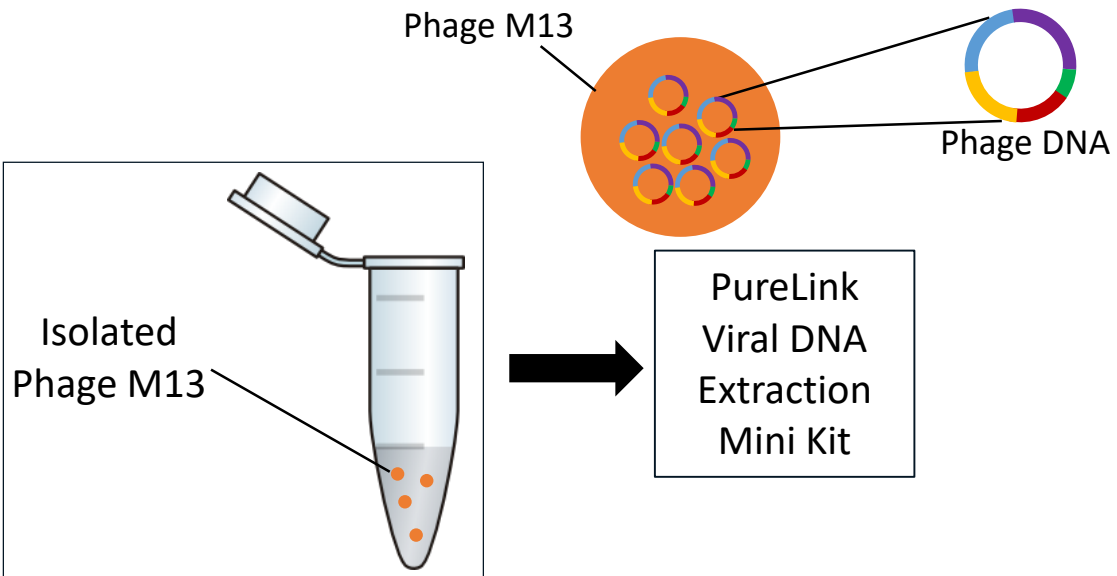
Recovering phage M13 from human skin swab

2. Phage DNA Extraction



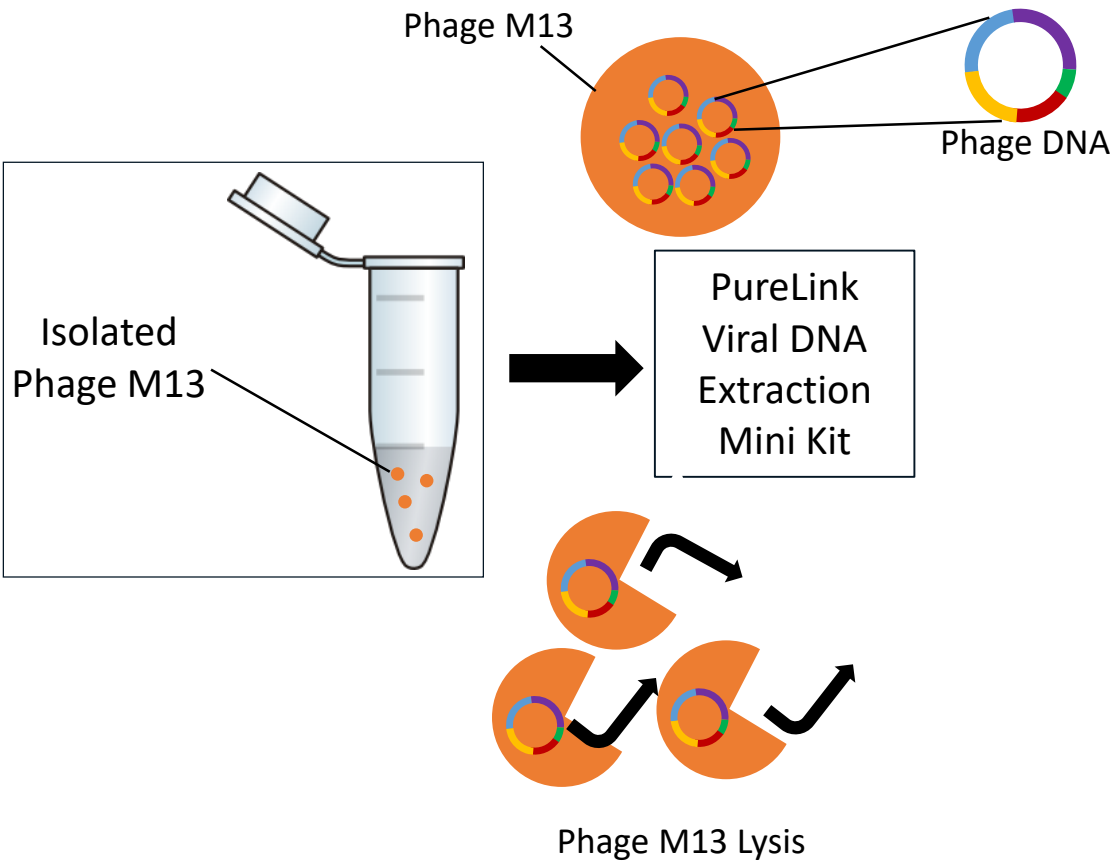
Recovering phage M13 from human skin swab

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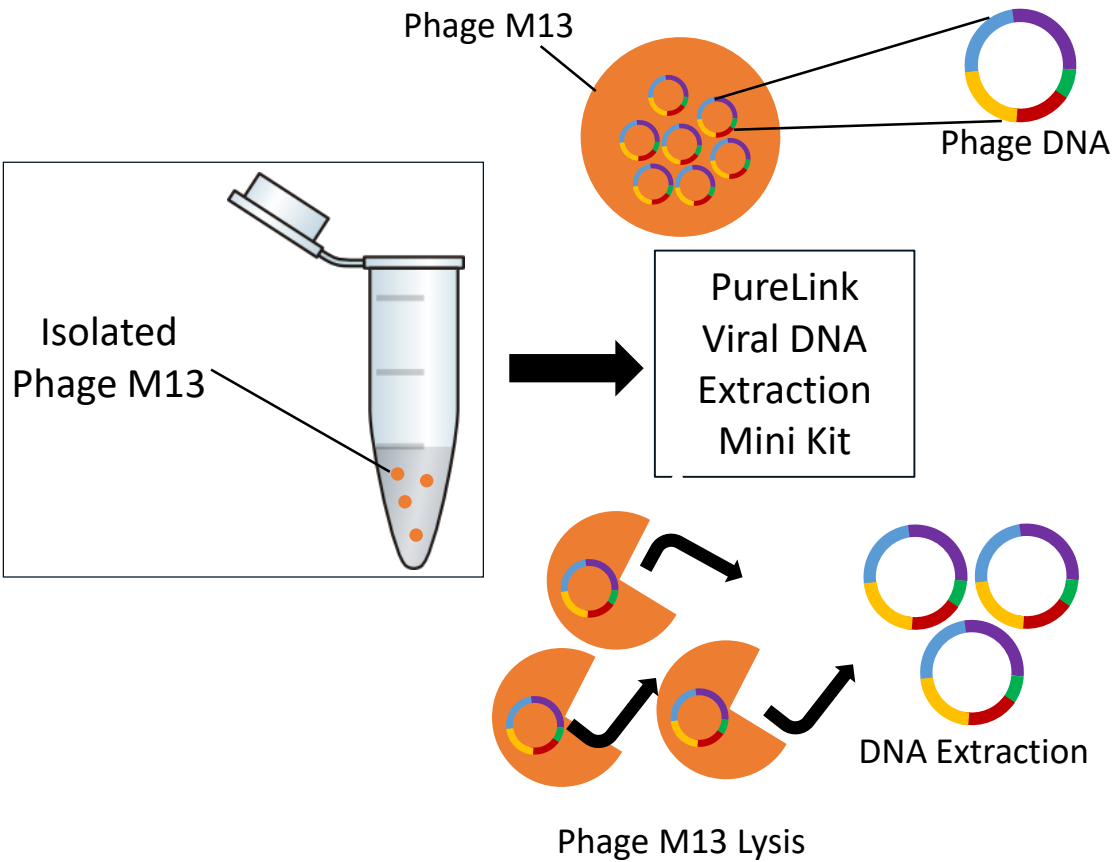
Recovering phage M13 from human skin swab

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Recovering phage M13 from human skin swab

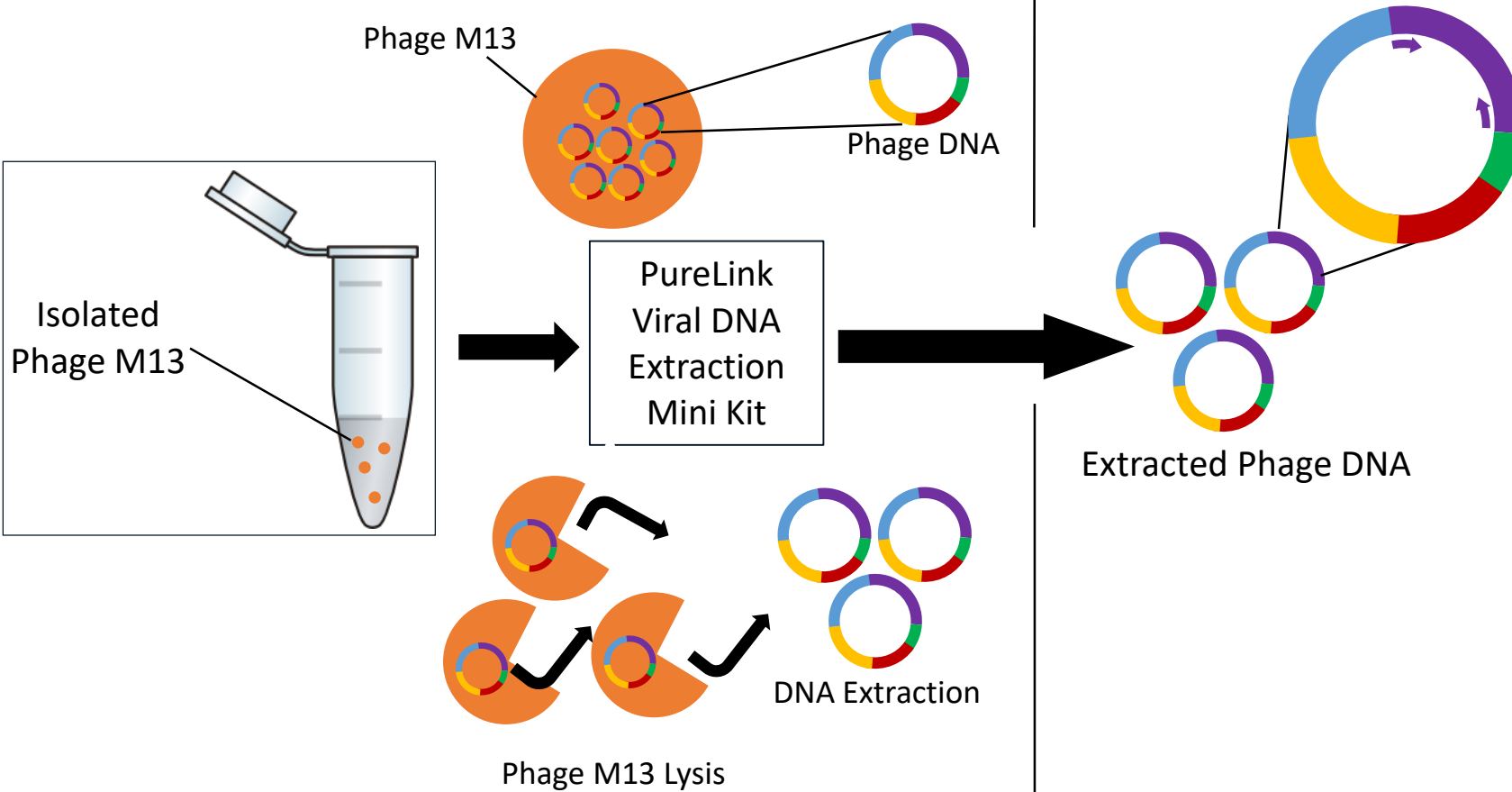
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Recovering phage M13 from human skin swab

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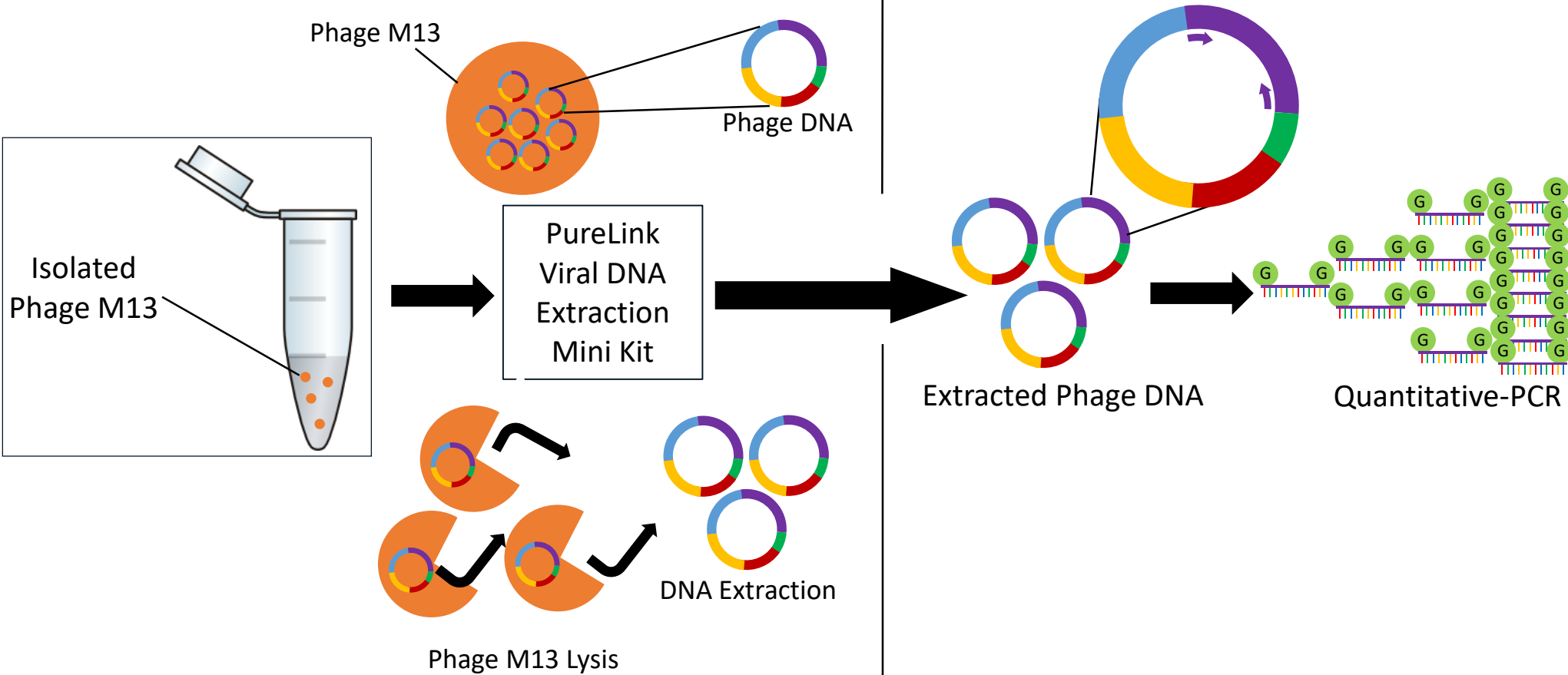
3. Quantification of Phage DNA



Recovering phage M13 from human skin swab

2. Phage DNA Extraction

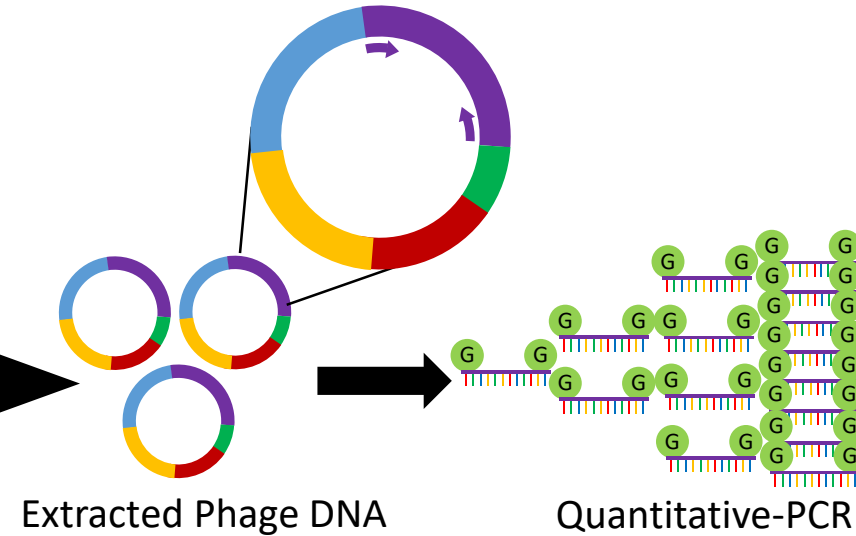
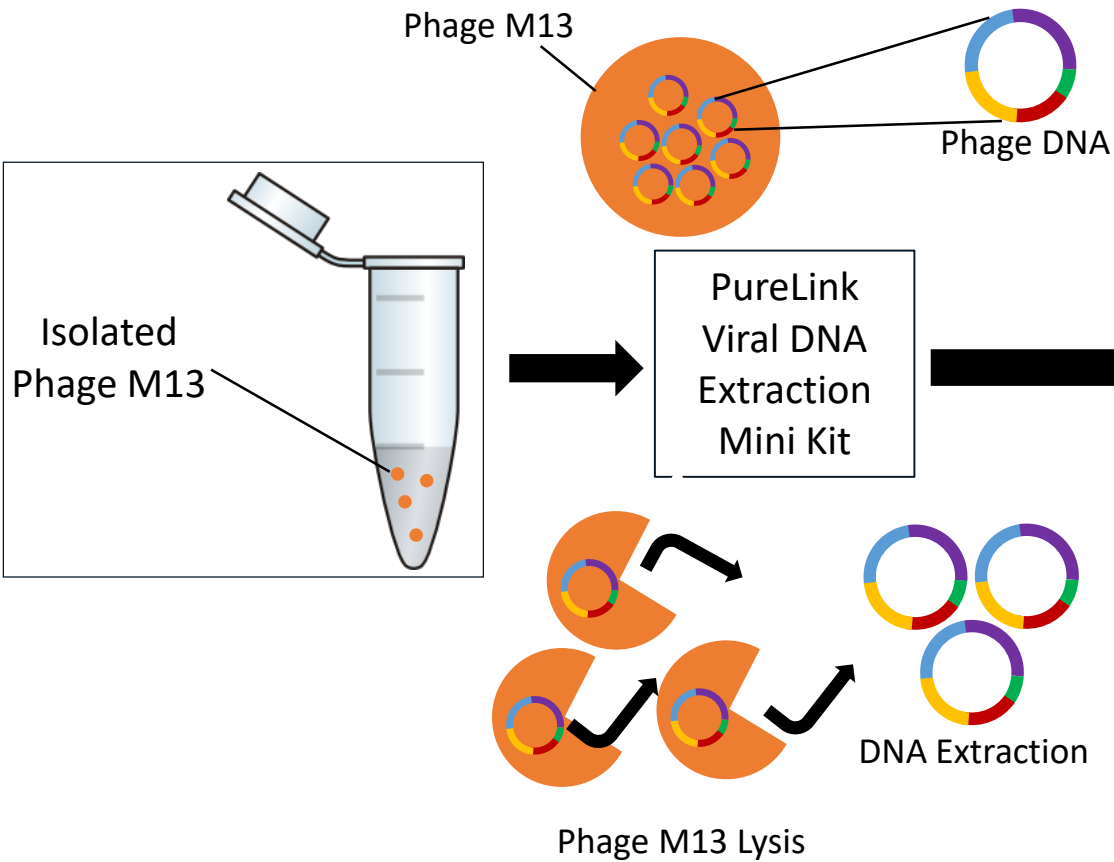
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Recovering phage M13 from human skin swab

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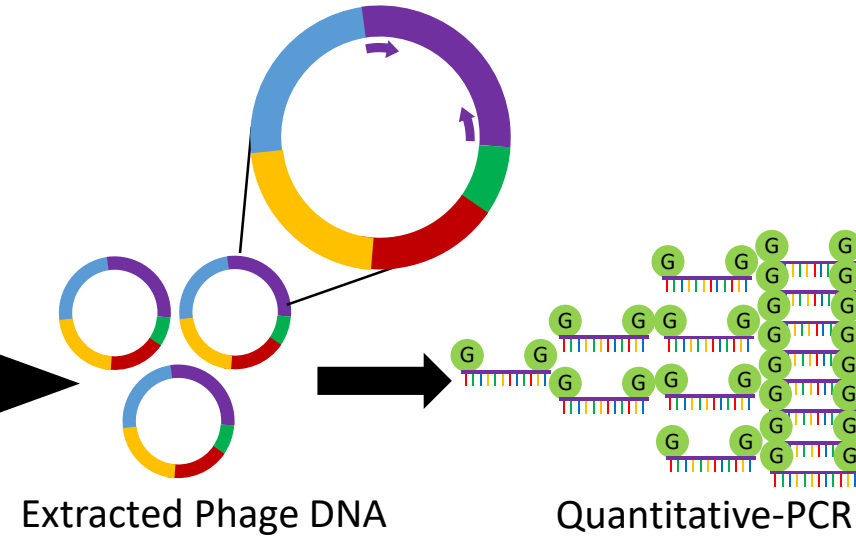
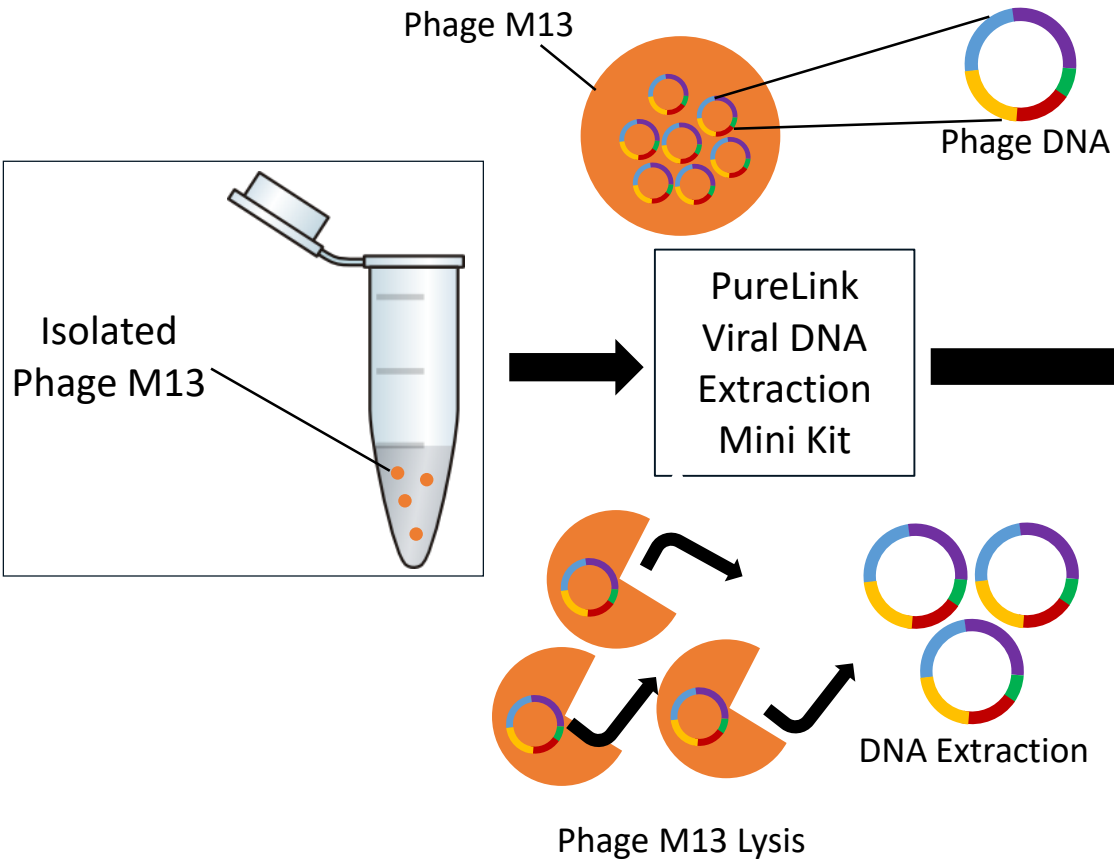
3. Quantification of Phage DNA



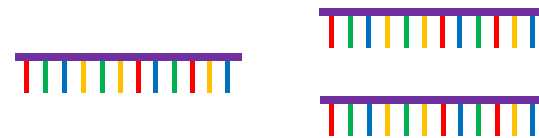
Recovering phage M13 from human skin swab

2. Phage DNA Extraction

3. Quantification of Phage DNA



qPCR Principles

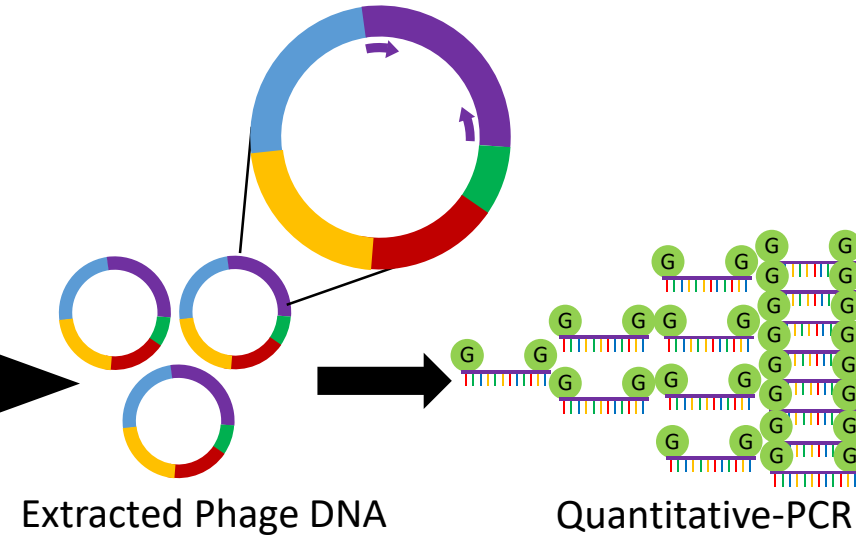
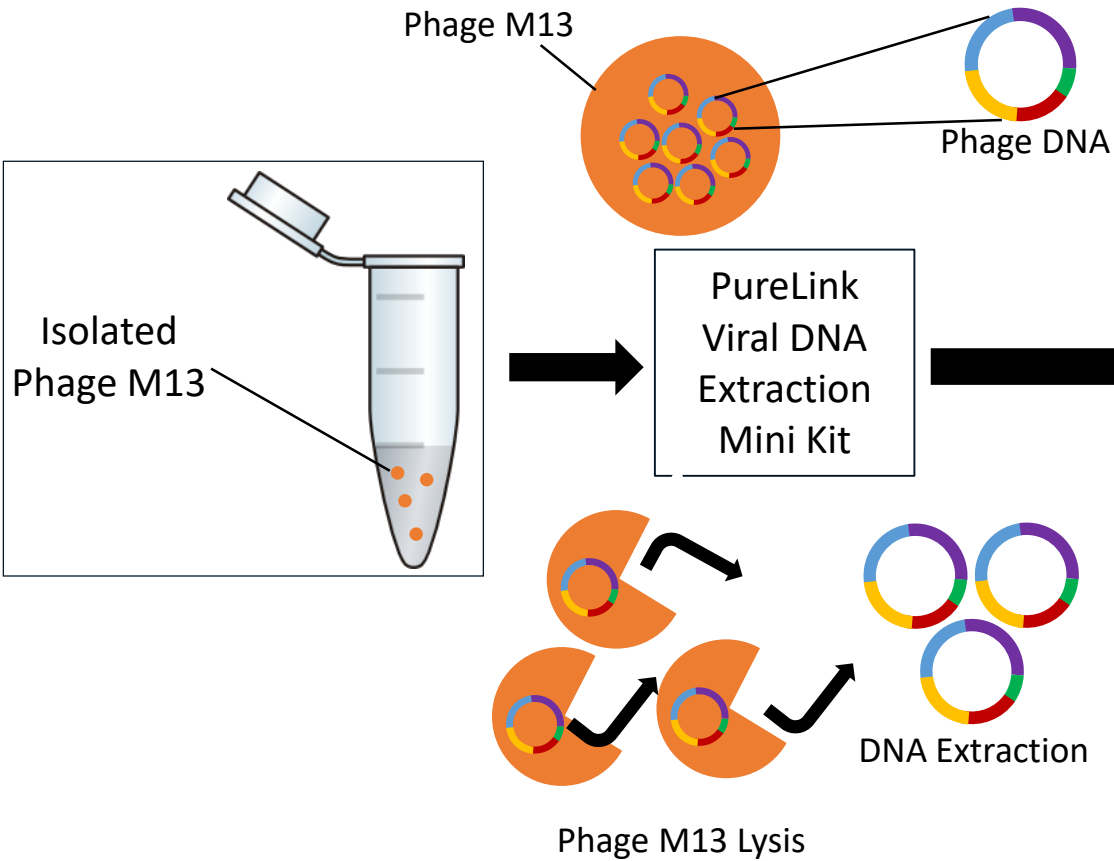


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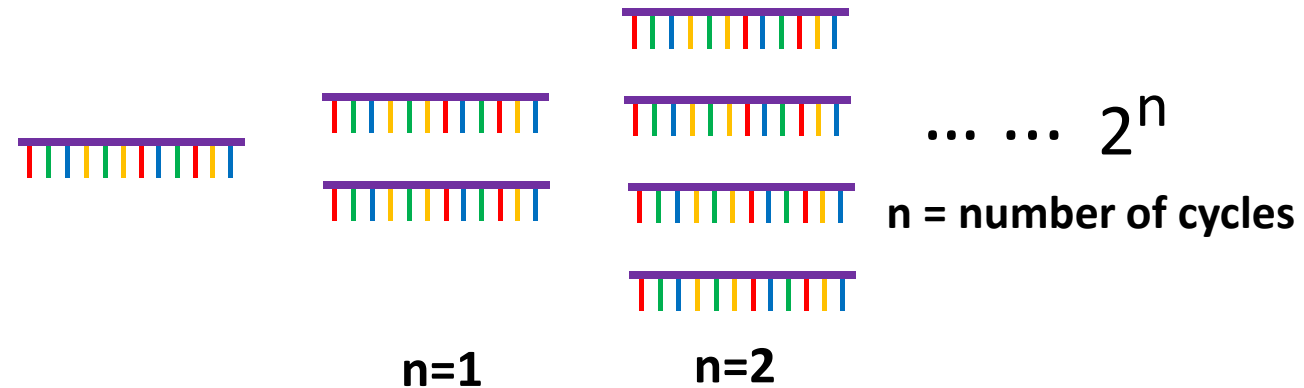
Recovering phage M13 from human skin swab

2. Phage DNA Extraction

3. Quantification of Phage DNA



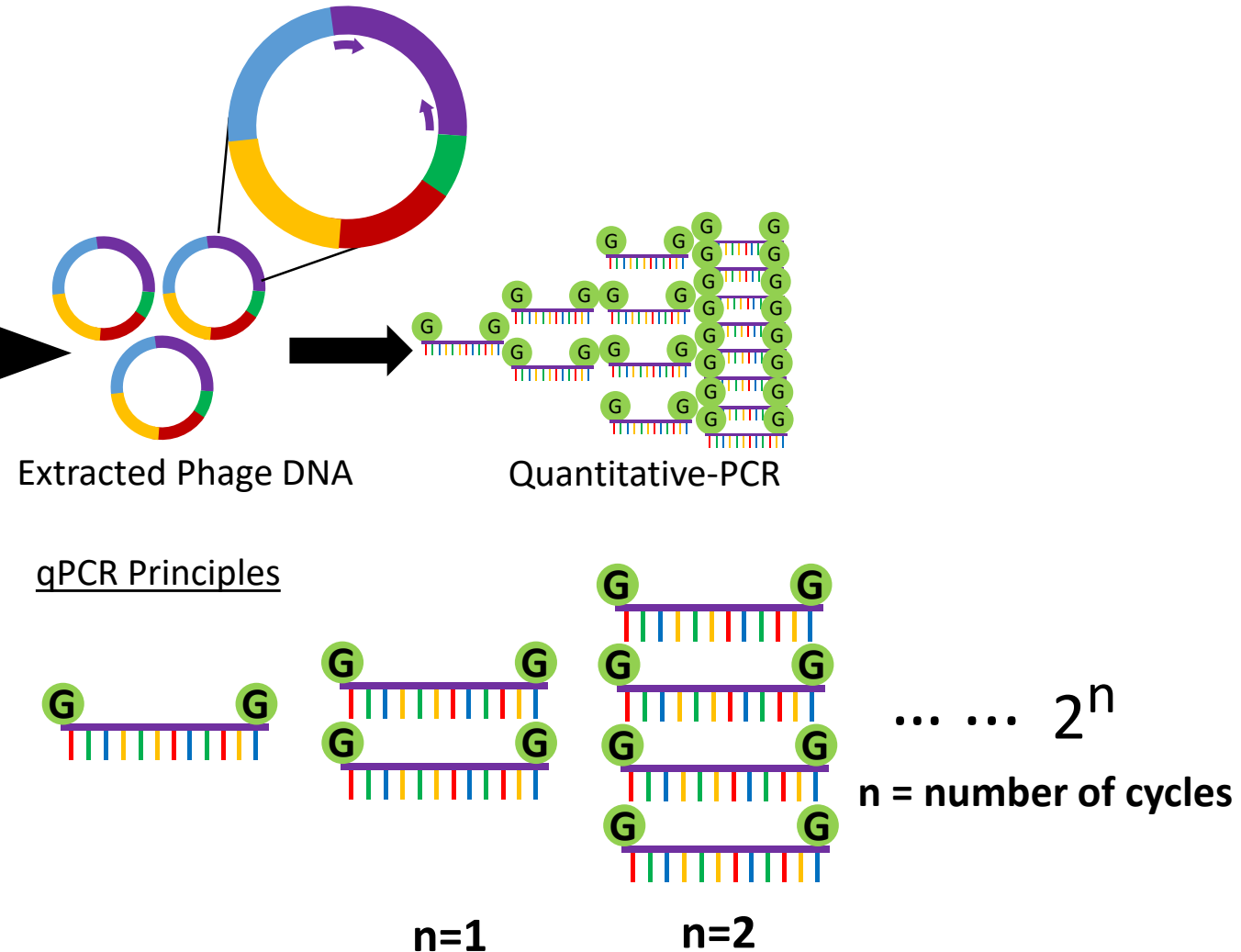
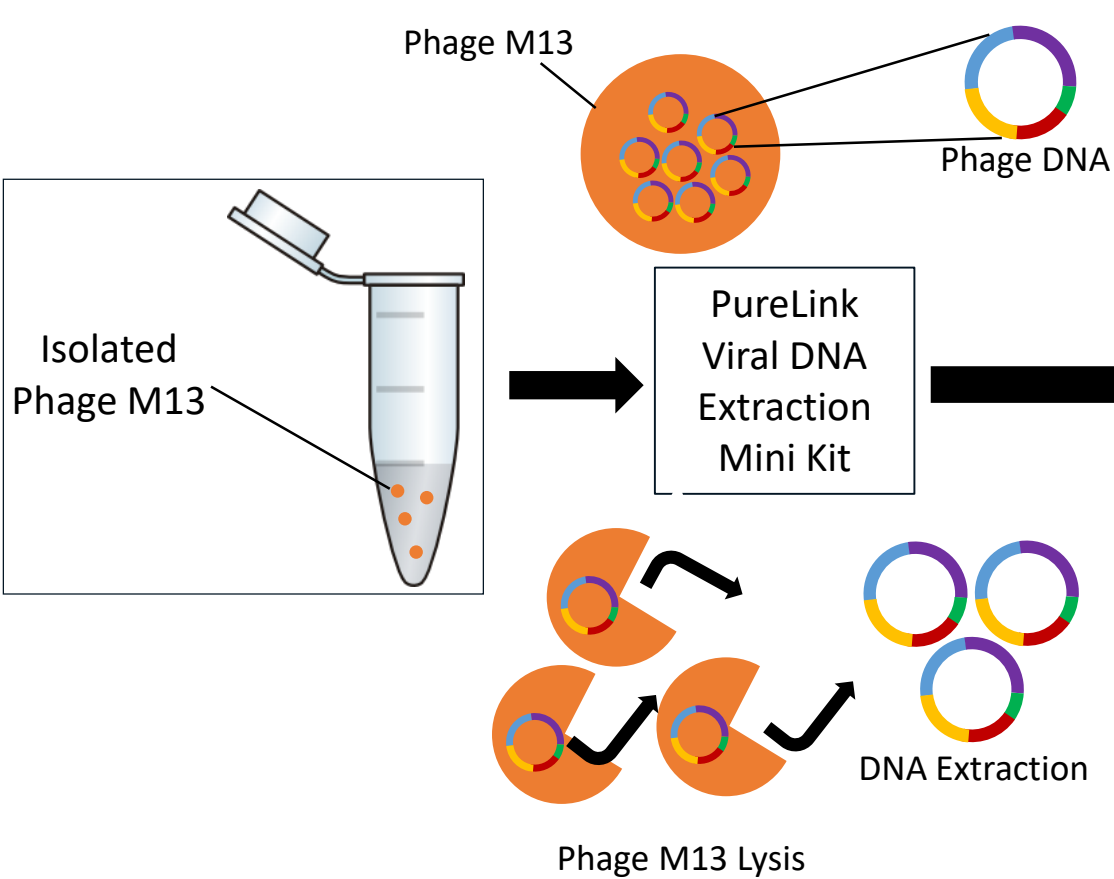
qPCR Principles



Recovering phage M13 from human skin swab

2. Phage DNA Extraction

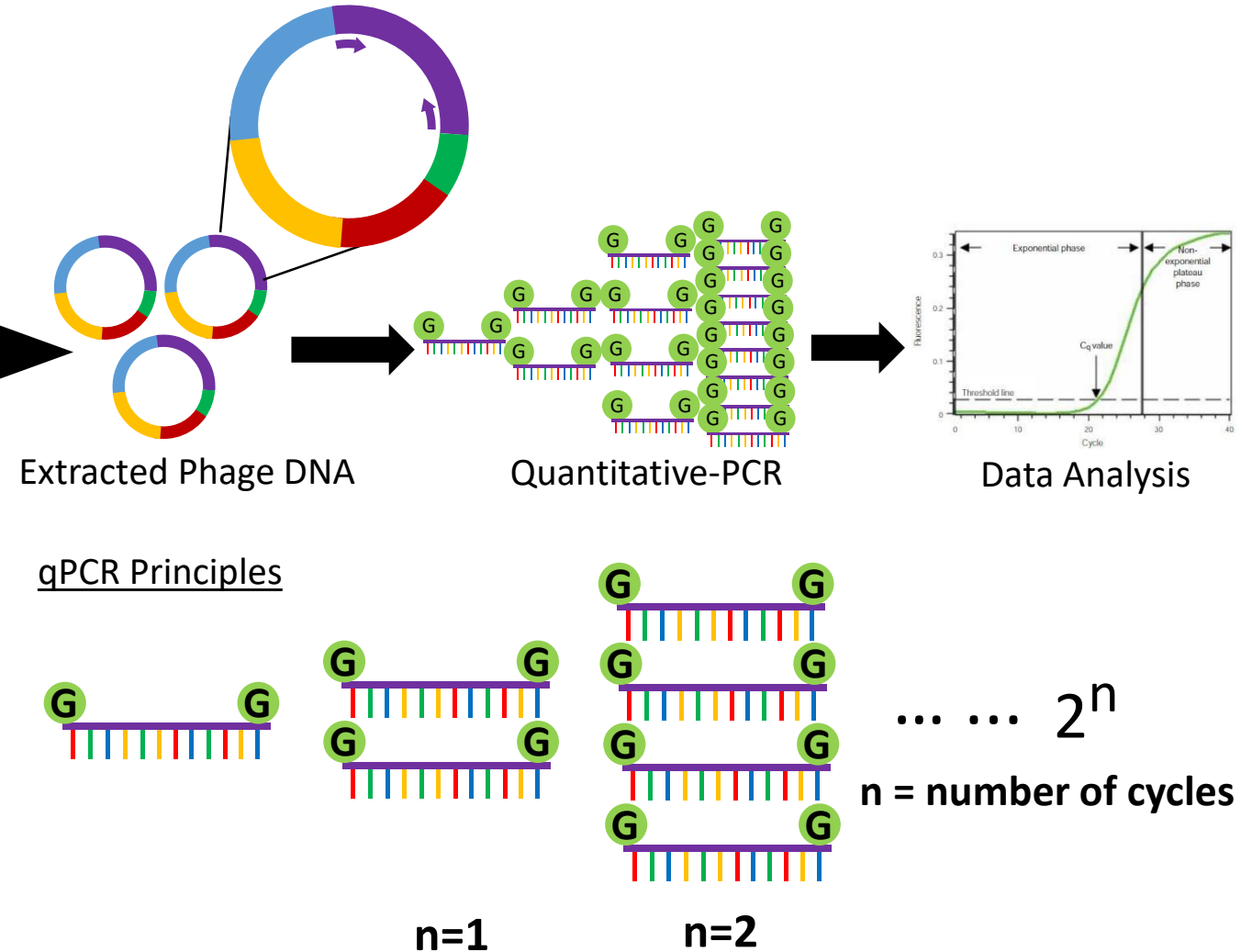
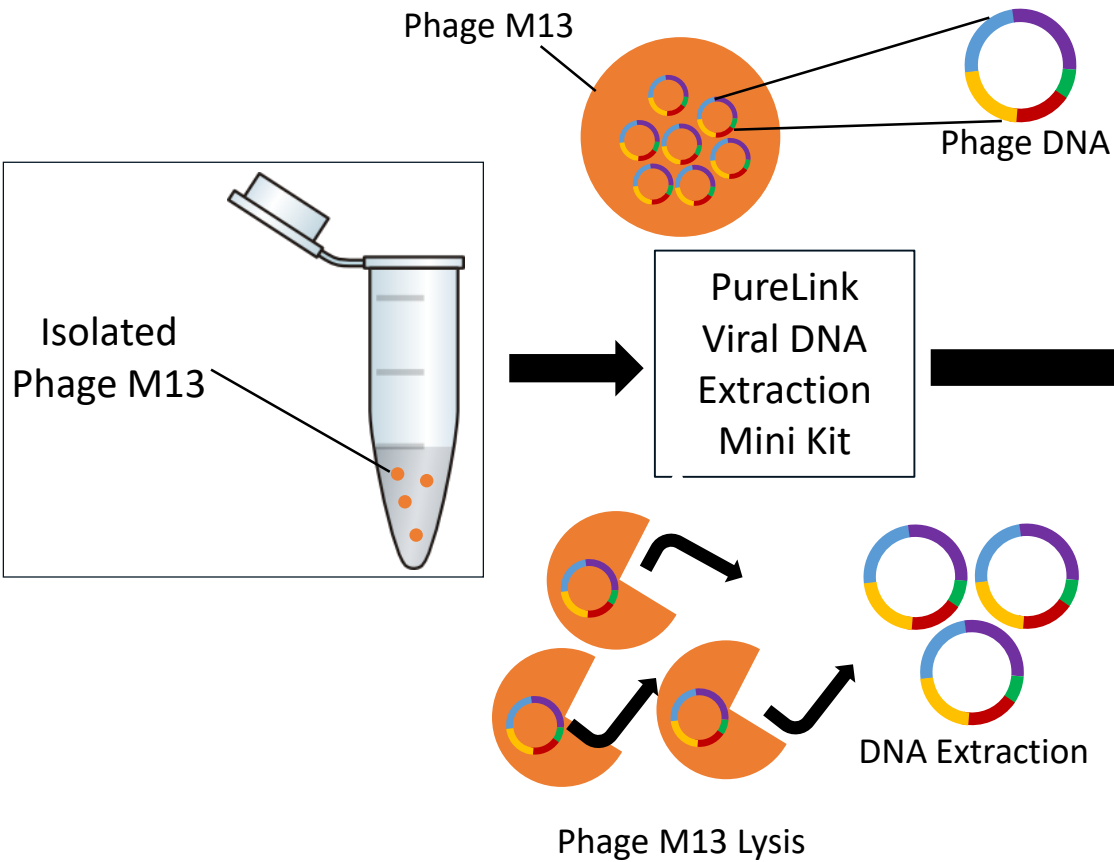
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Recovering phage M13 from human skin swab

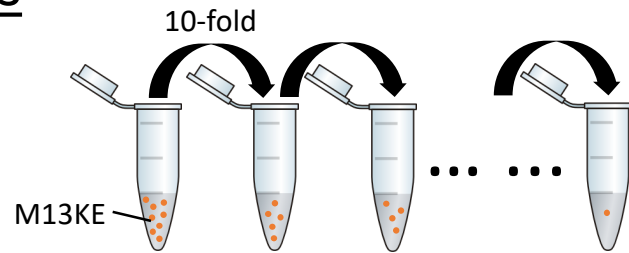
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Identifying qPCR limit of detection for phage M13 DNA

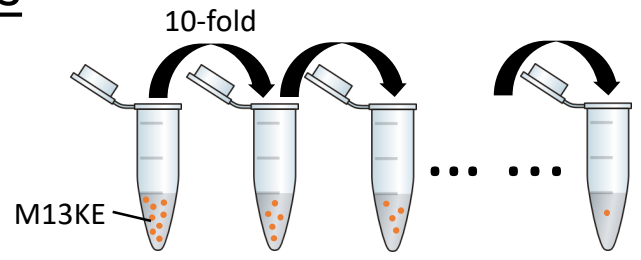
Samples



Identifying qPCR limit of detection for phage M13 DNA

Samples

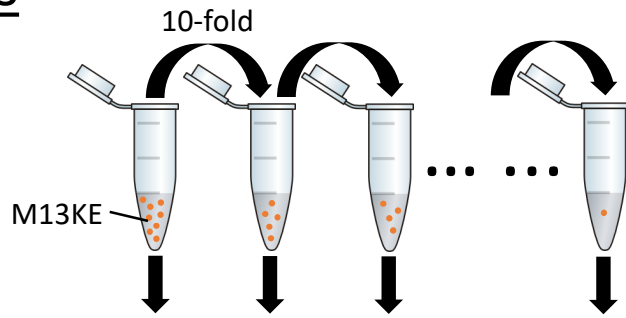
Control



Identifying qPCR limit of detection for phage M13 DNA

Samples

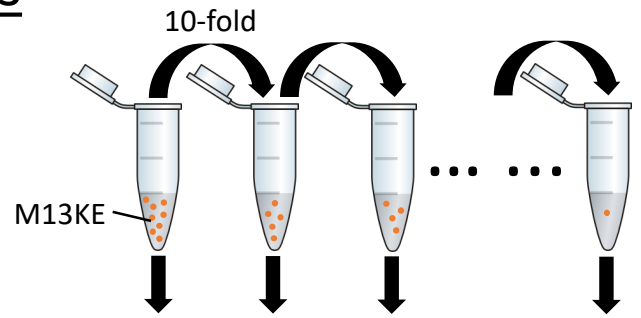
Control



Identifying qPCR limit of detection for phage M13 DNA

Samples

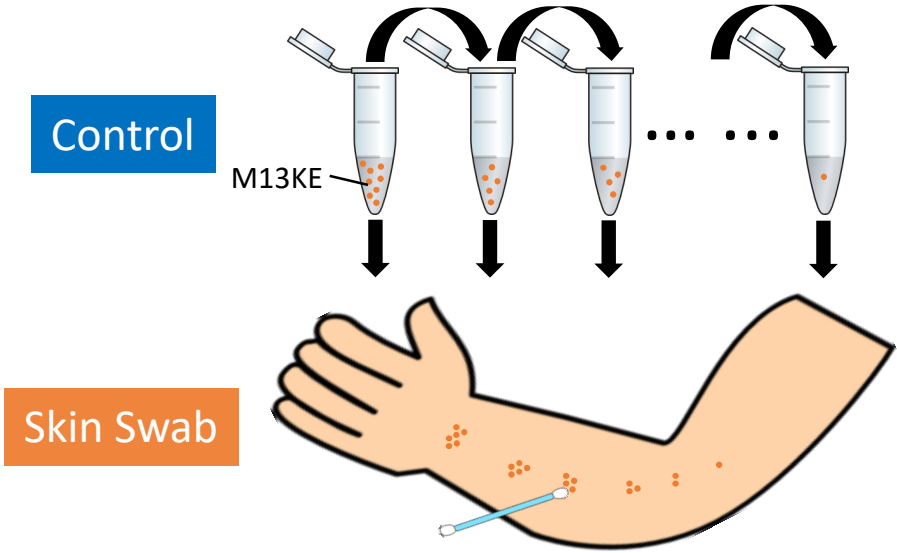
Control



Skin Swab

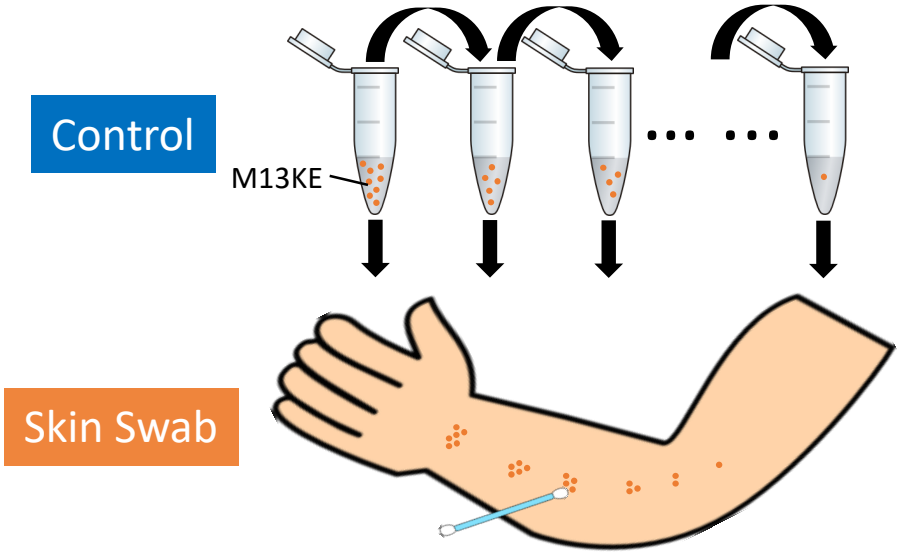
Identifying qPCR limit of detection for phage M13 DNA

Samples



Identifying qPCR limit of detection for phage M13 DNA

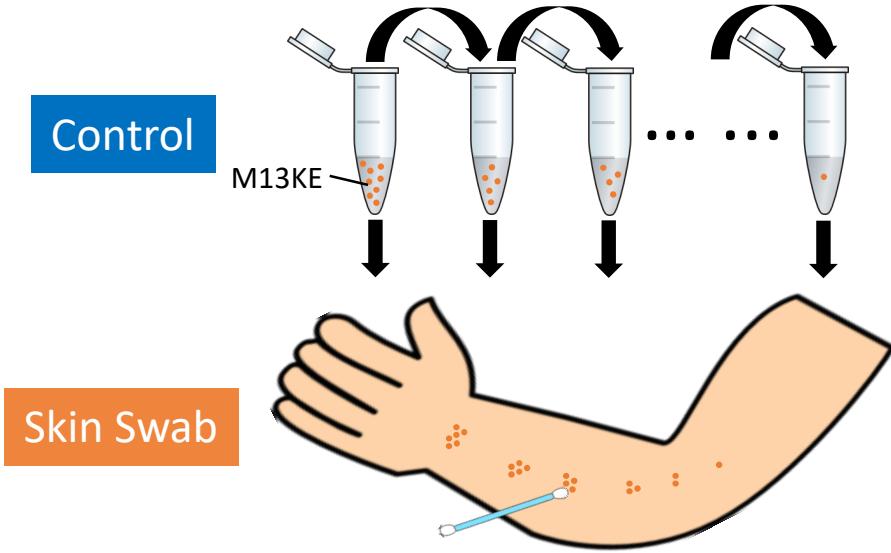
Samples



Samples were assayed in triplicate

Identifying qPCR limit of detection for phage M13 DNA

Samples



Control

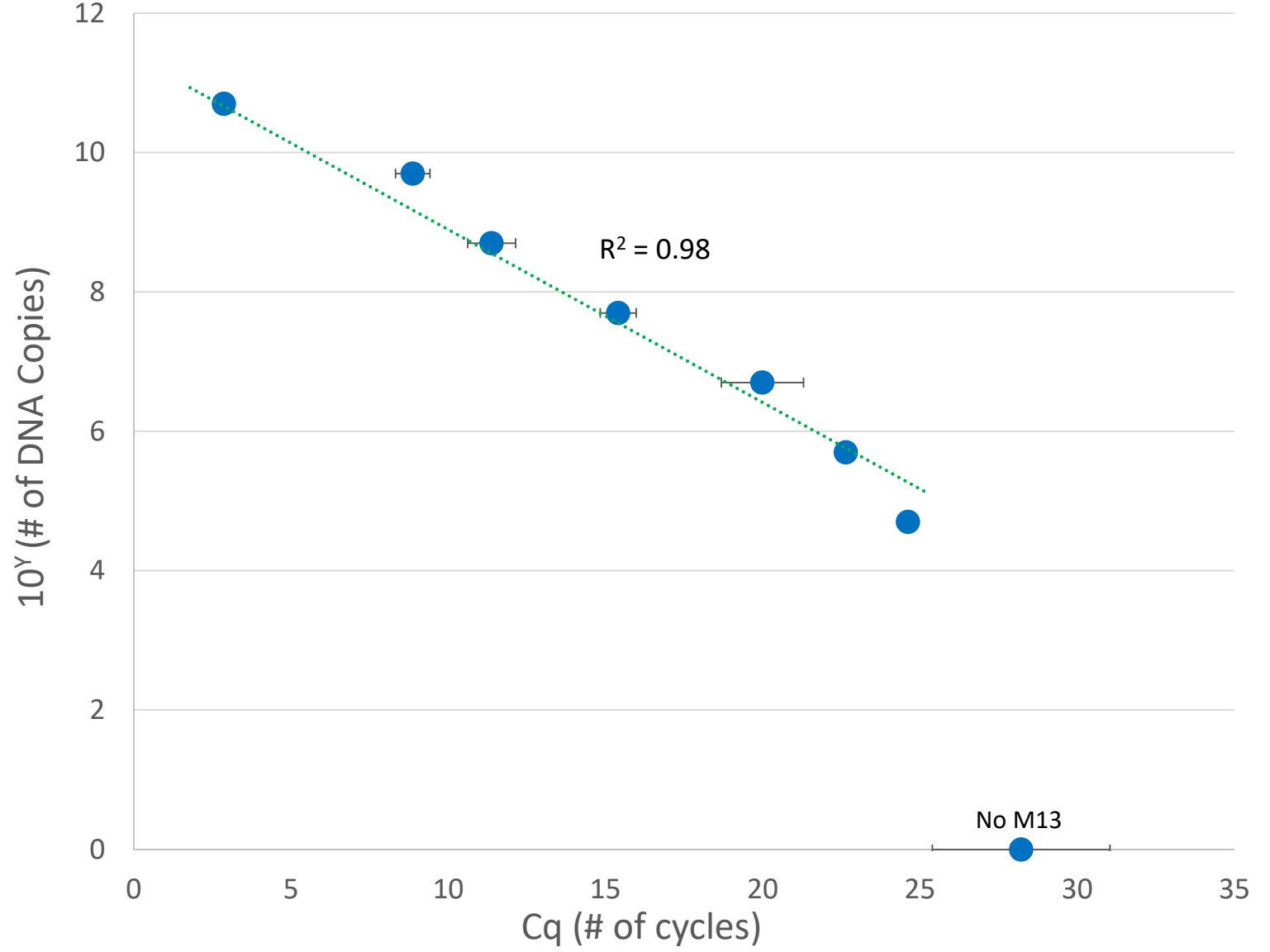
M13KE

10-fold

Skin Swab

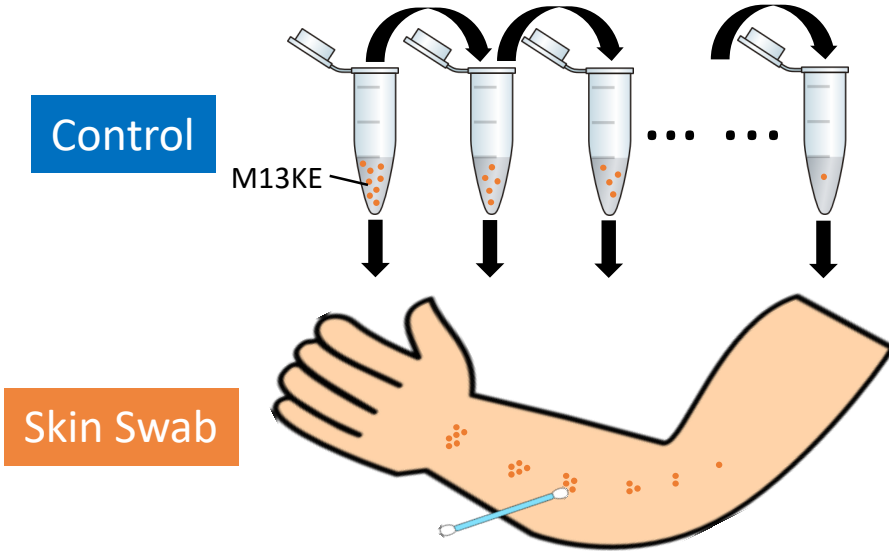
Samples were assayed in triplicate

Absolute Quantification of M13KE Skin Swab Samples

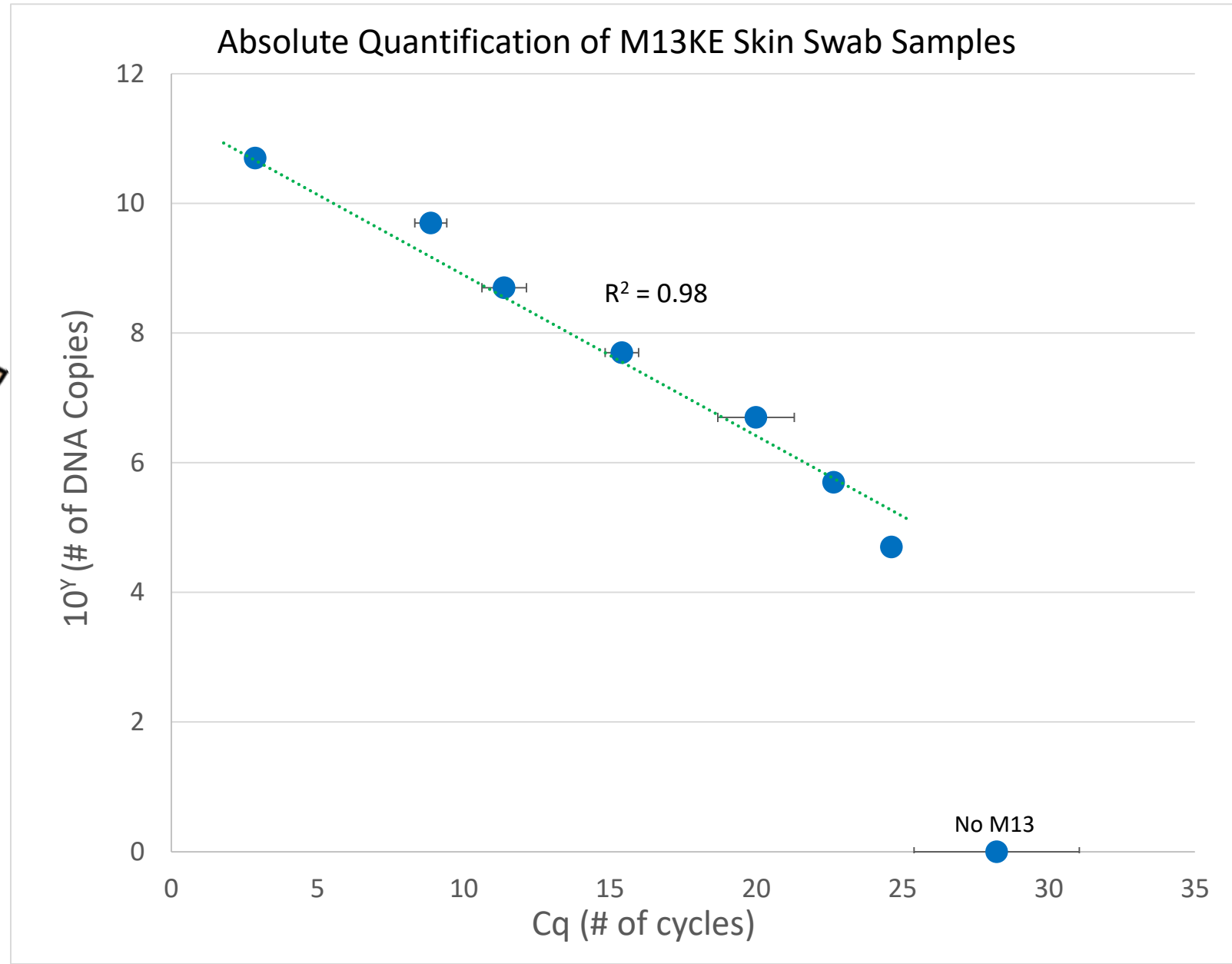


Identifying qPCR limit of detection for phage M13 DNA

Samples

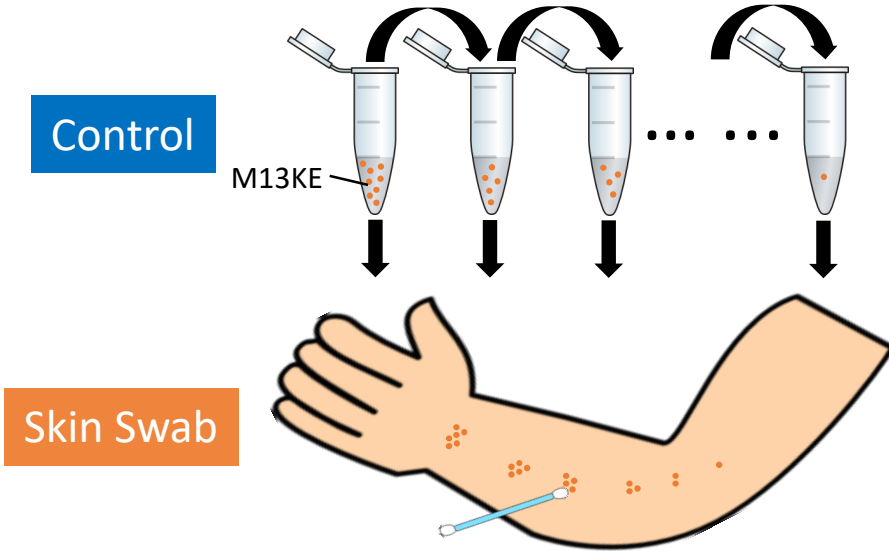


- qPCR baseline of detection = 2.36×10^4 copies

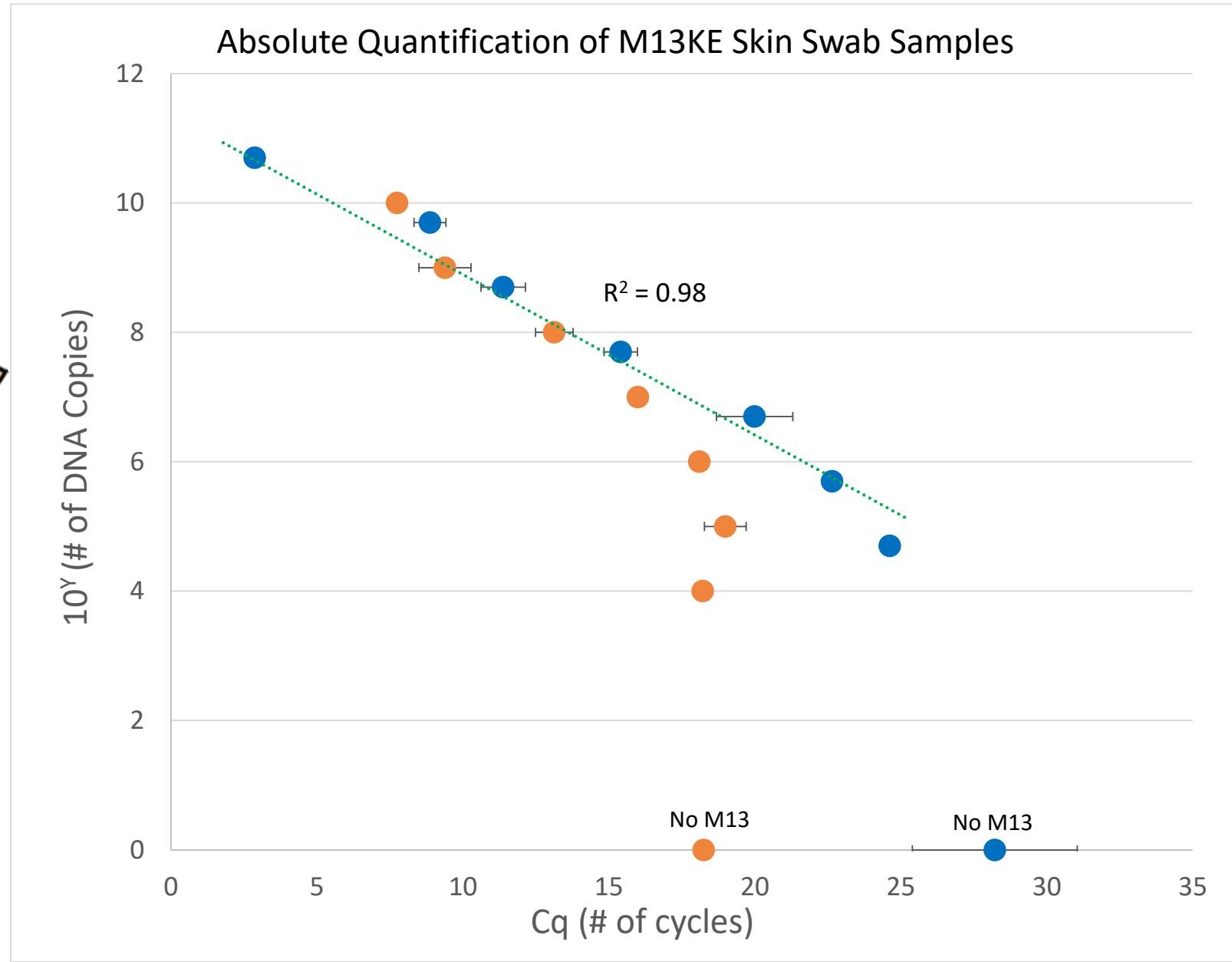


Identifying qPCR limit of detection for phage M13 DNA

Samples

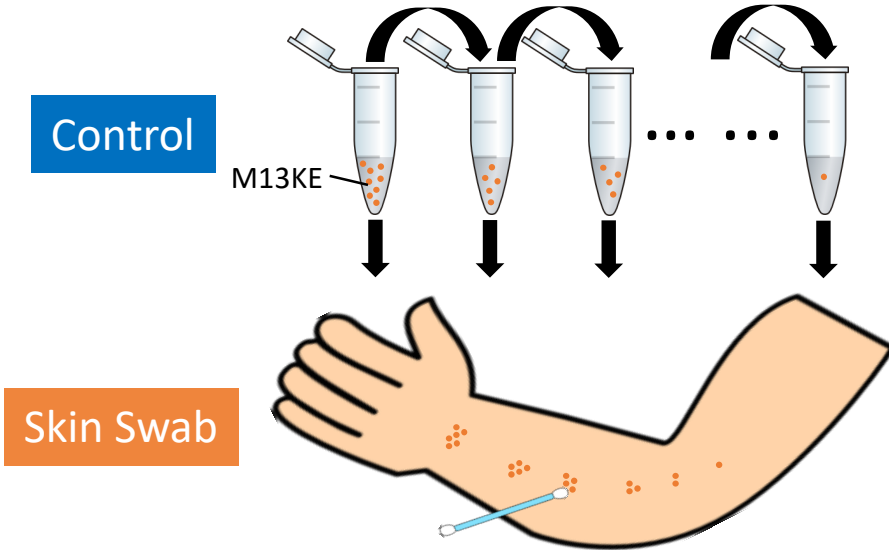


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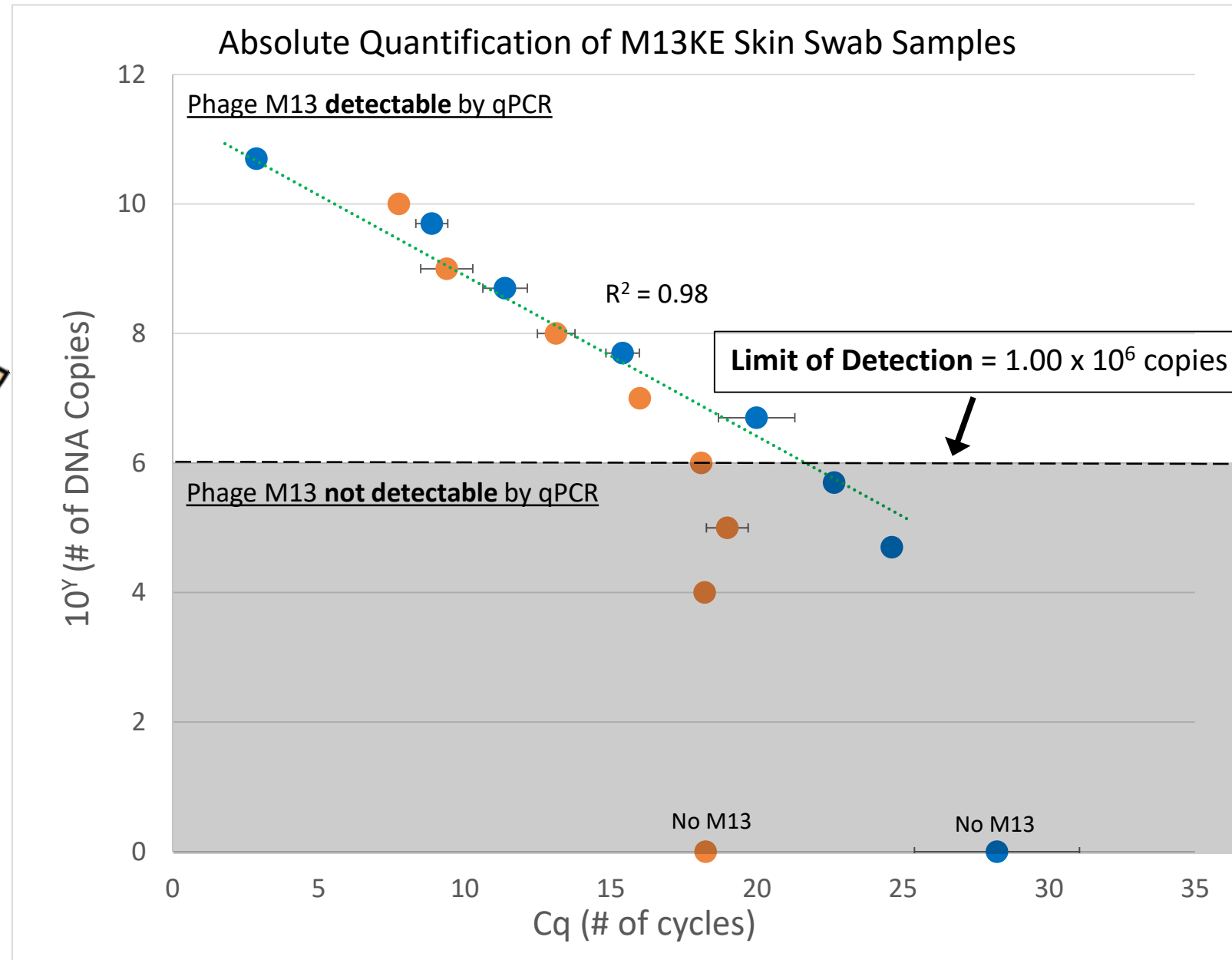


Identifying qPCR limit of detection for phage M13 DNA

Samples

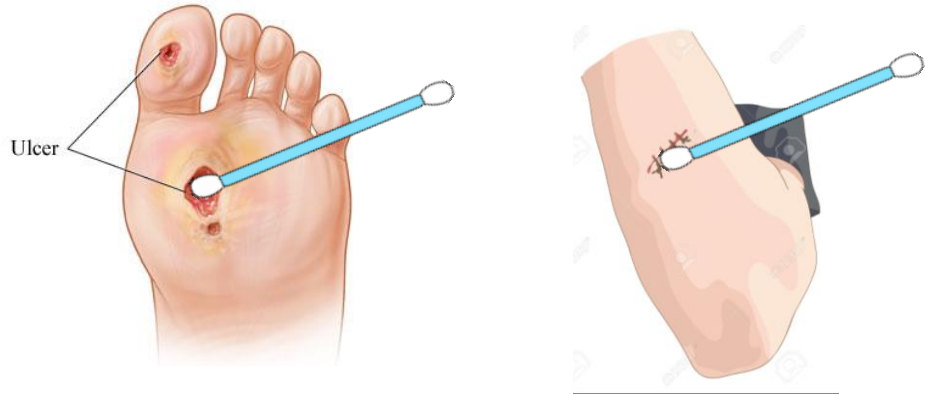


- qPCR baseline of detection = 2.36×10^4 copies
- Limit of detection for human skin swab = 1.00×10^6 copies
- Contaminating sequences on human skin



Establishing a protocol to quantify phages from wounds

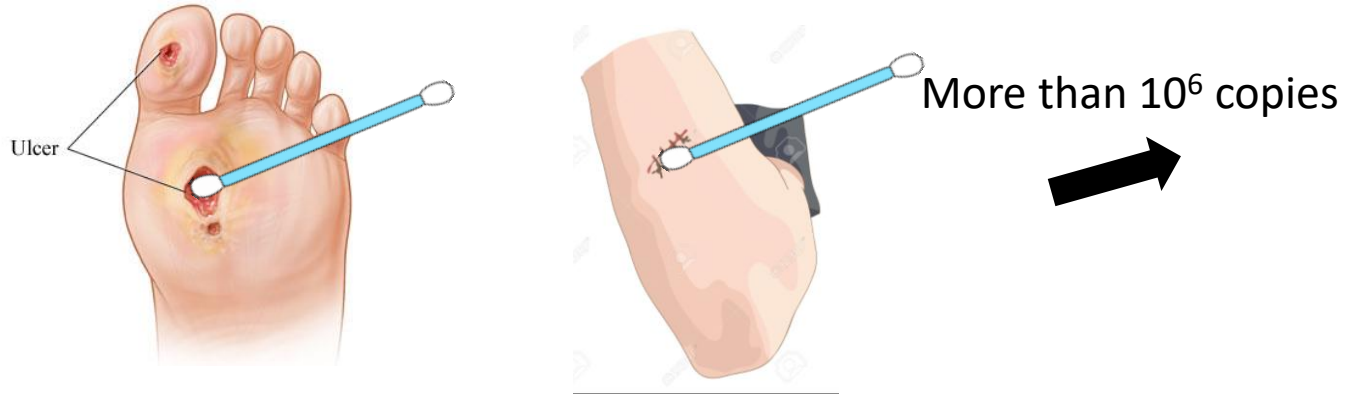
qPCR limit of detection for human skin swab = 1.00×10^6 copies



Human skin wounds

Establishing a protocol to quantify phages from wounds

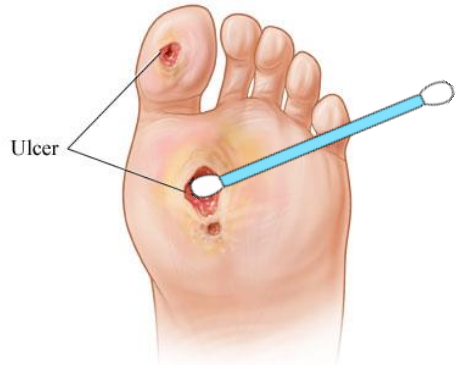
qPCR limit of detection for human skin swab = 1.00×10^6 copies



Human skin wounds

Establishing a protocol to quantify phages from wounds

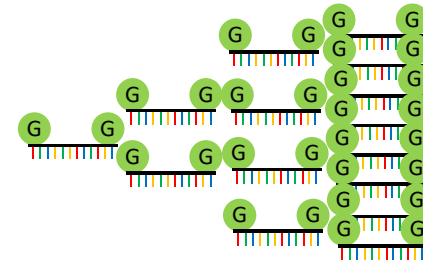
qPCR limit of detection for human skin swab = 1.00×10^6 copies



Ulcer



More than 10^6 copies

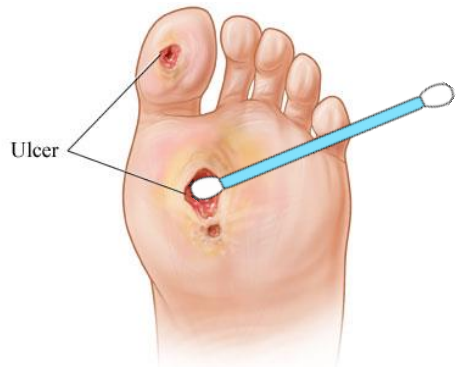


Can quantify using qPCR

Human skin wounds

Establishing a protocol to quantify phages from wounds

qPCR limit of detection for human skin swab = 1.00×10^6 copies



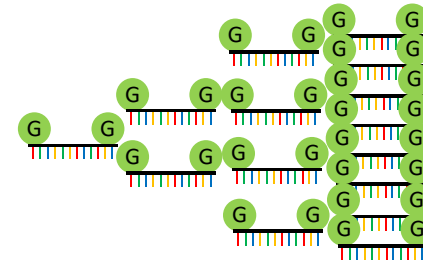
Human skin wounds



More than 10^6 copies



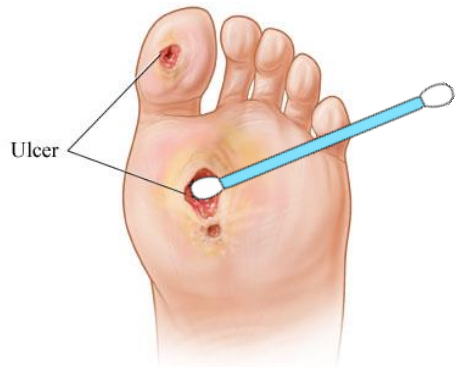
Less than 10^6 copies



Can quantify using qPCR

Establishing a protocol to quantify phages from wounds

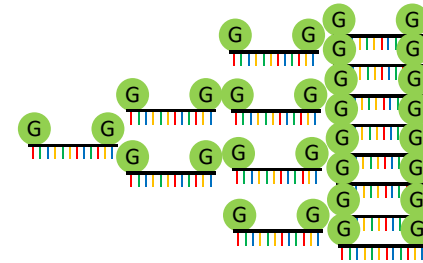
qPCR limit of detection for human skin swab = 1.00×10^6 copies



Human skin wounds



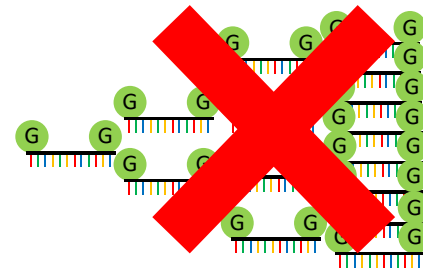
More than 10^6 copies



Can quantify using qPCR



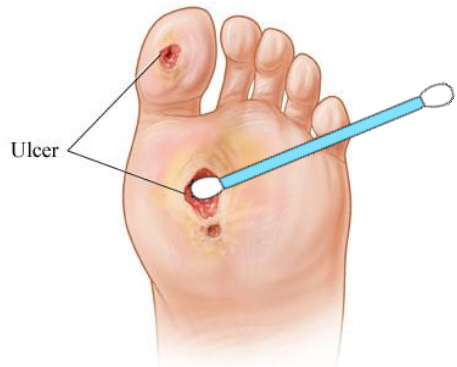
Less than 10^6 copies



Cannot quantify using qPCR

Establishing a protocol to quantify phages from wounds

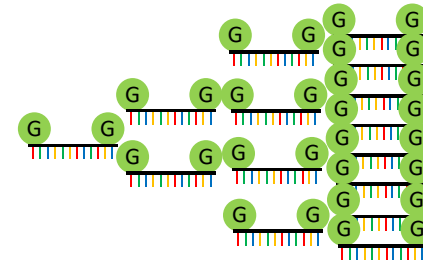
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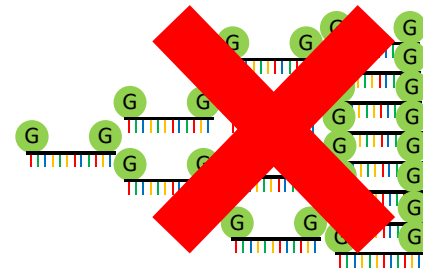
More than 10^6 copies



Can quantify using qPCR



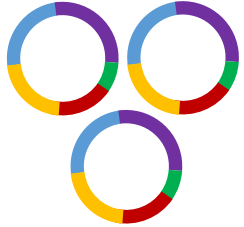
Less than 10^6 copies



Cannot quantify using qPCR

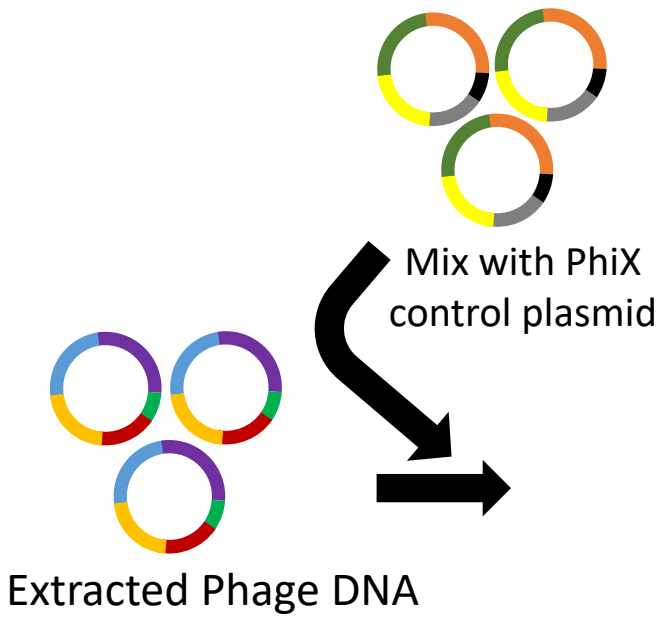


Sequencing phage M13 genome for quantification

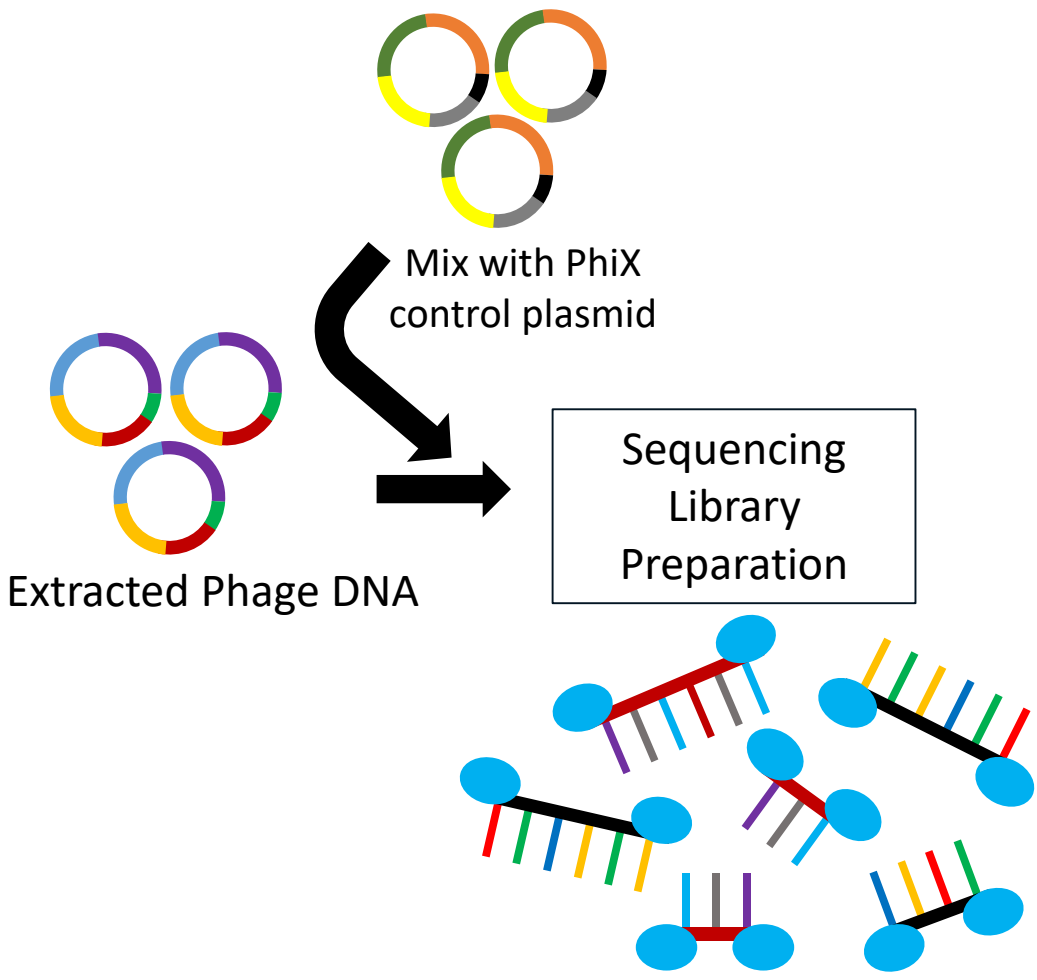


Extracted Phage DNA

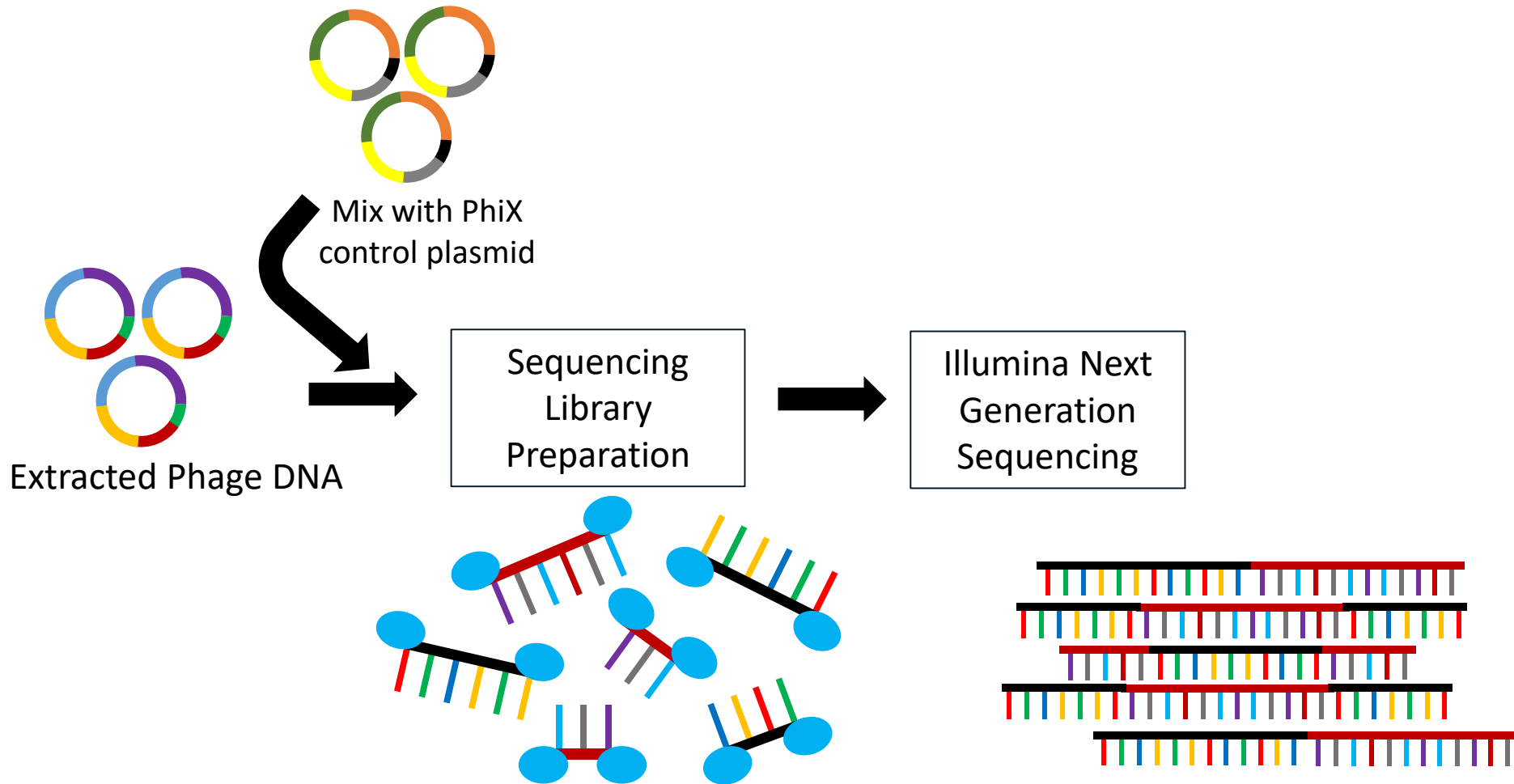
Sequencing phage M13 genome for quantification



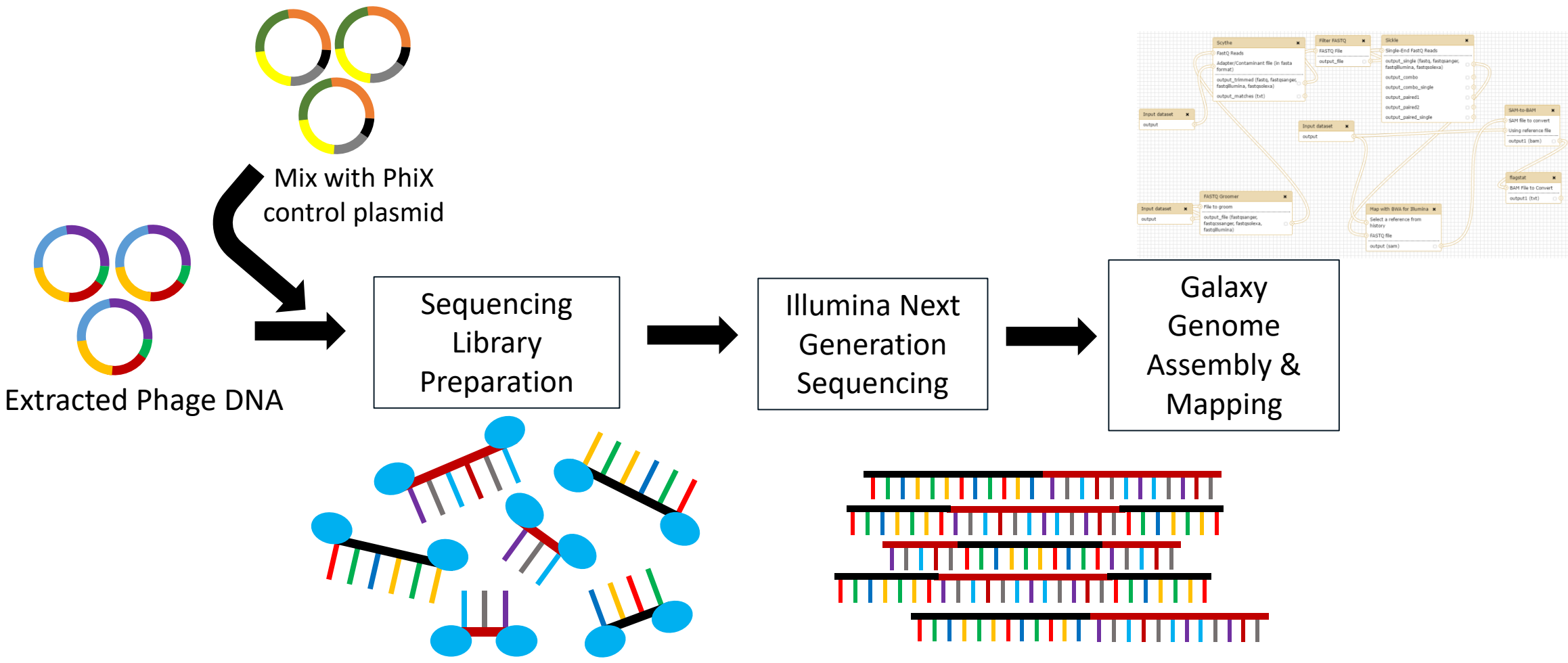
Sequencing phage M13 genome for quantification



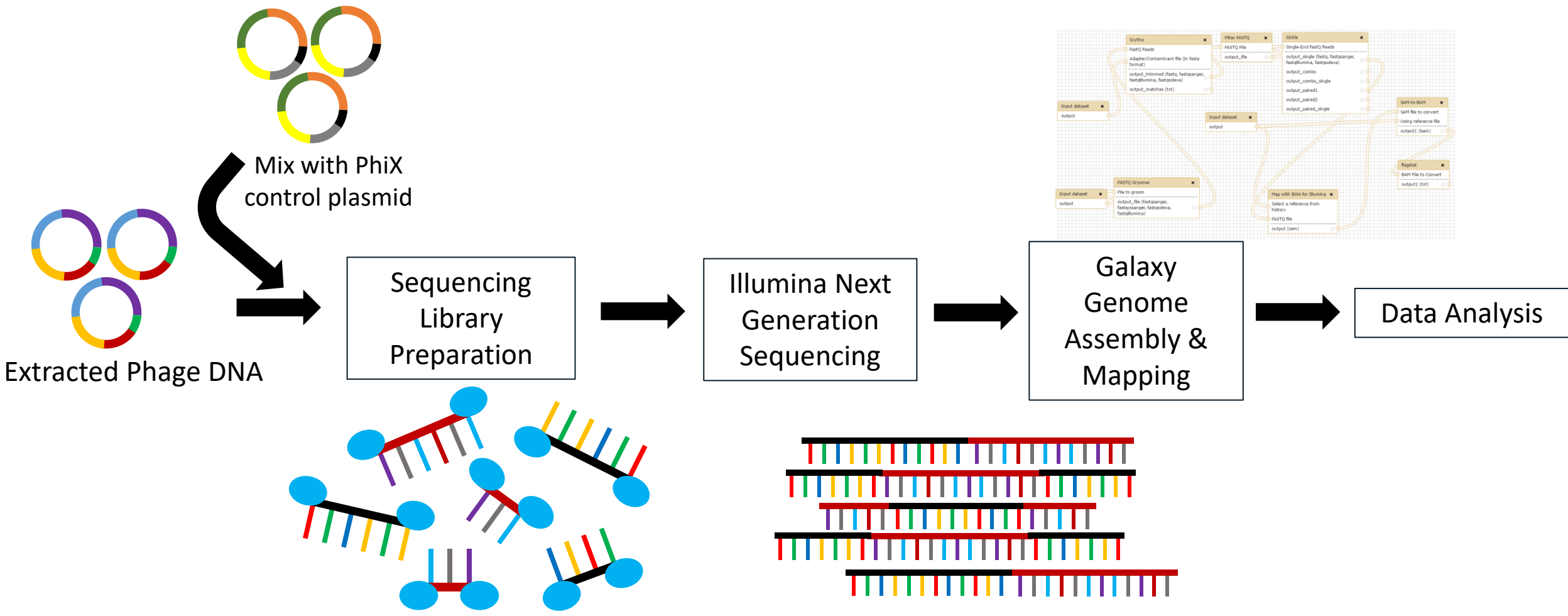
Sequencing phage M13 genome for quantification



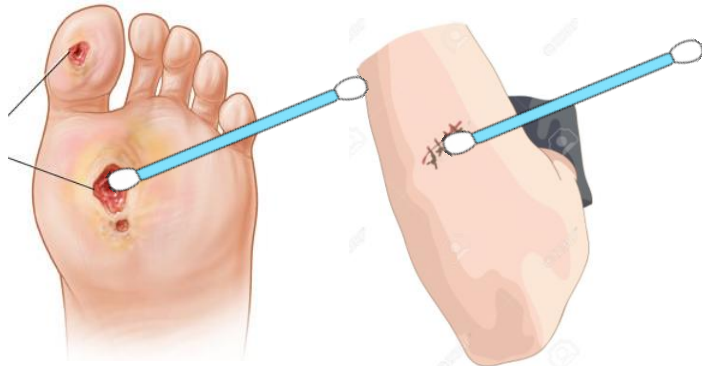
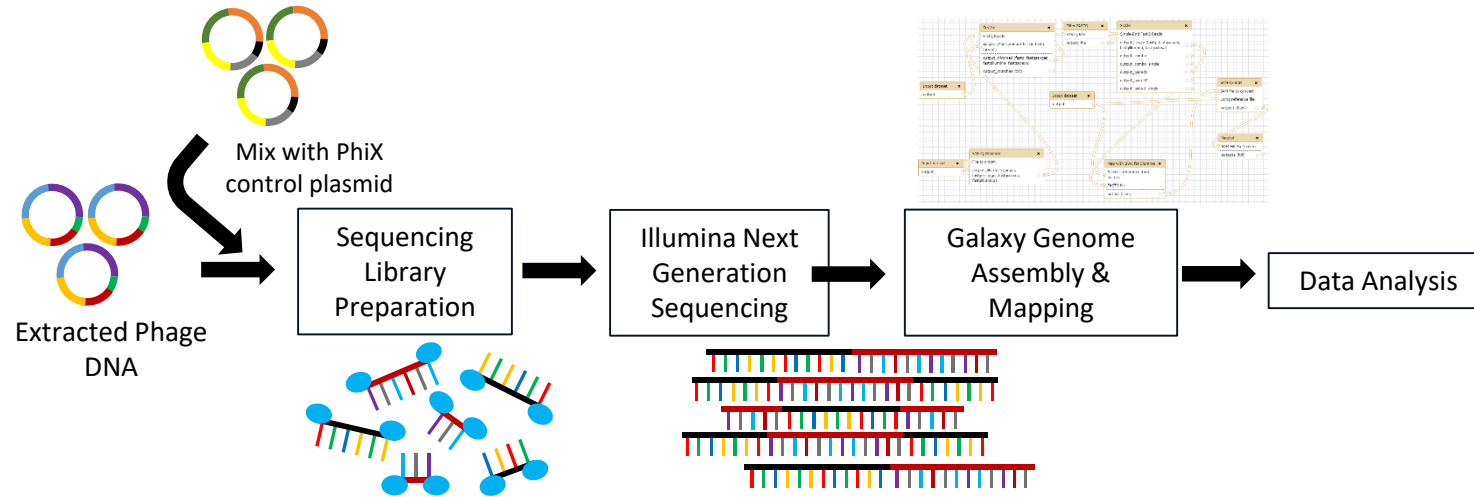
Sequencing phage M13 genome for quantification



Sequencing phage M13 genome for quantification

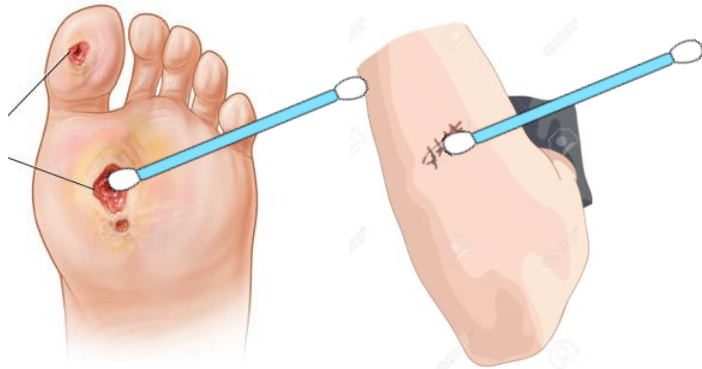
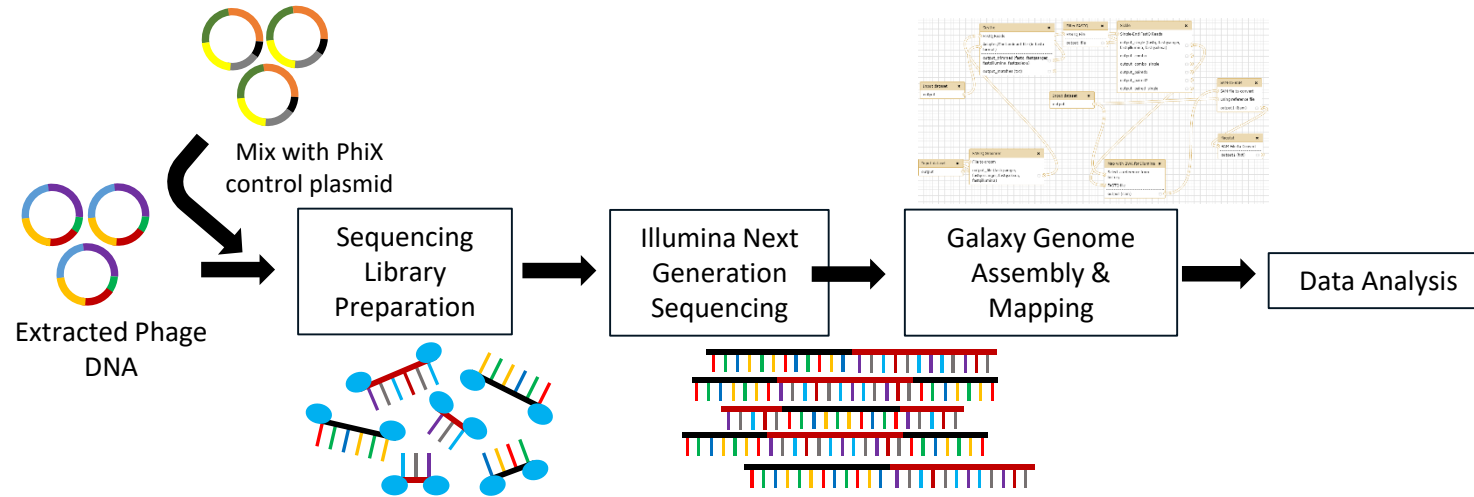


Applying the protocol to analyze human skin wounds

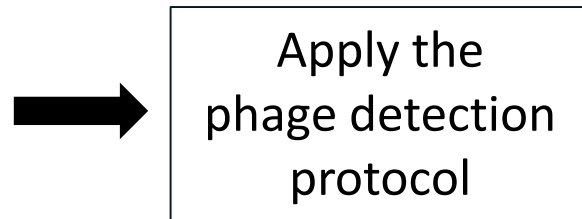


Human skin wounds

Applying the protocol to analyze human skin wounds

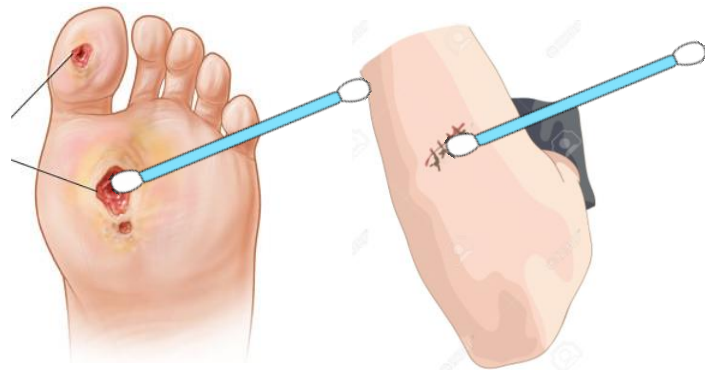
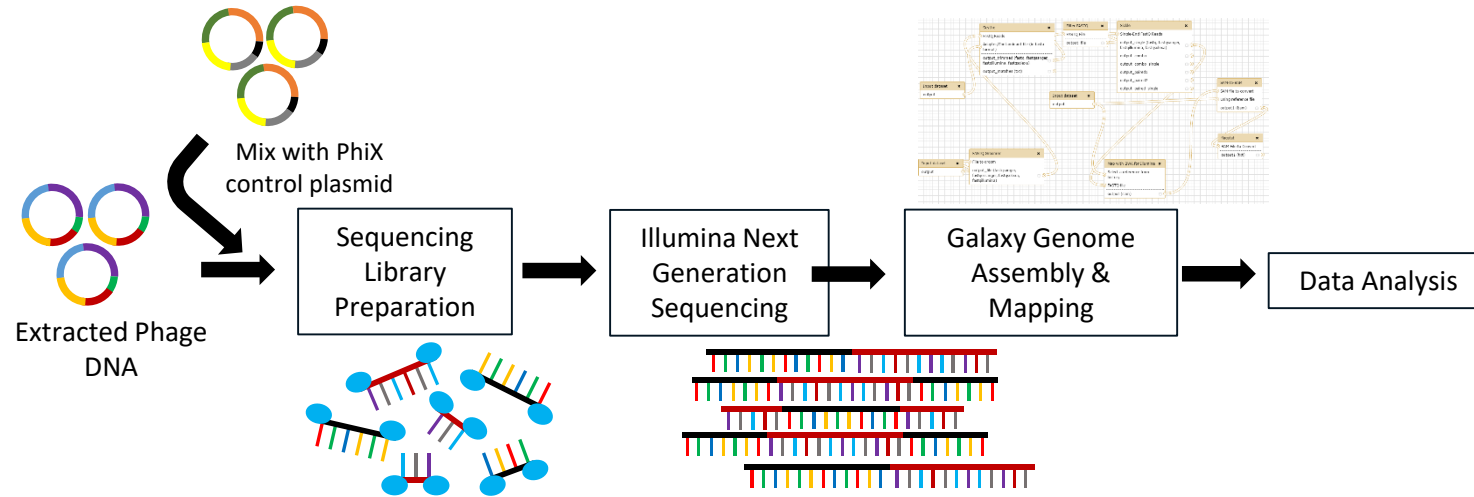


Human skin wounds

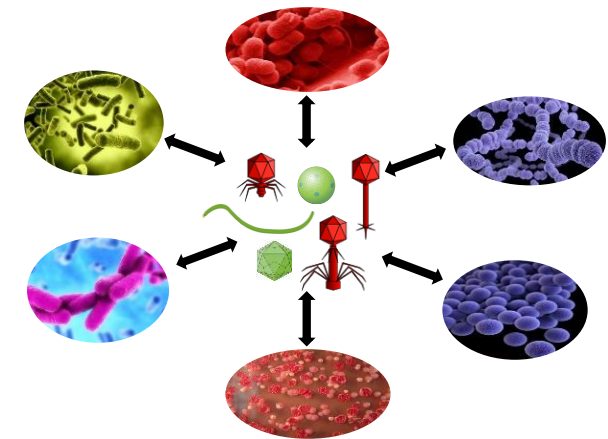
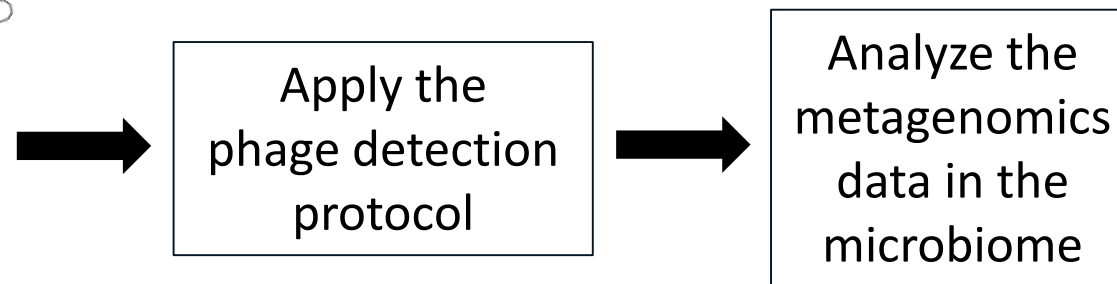


Apply the
phage detection
protocol

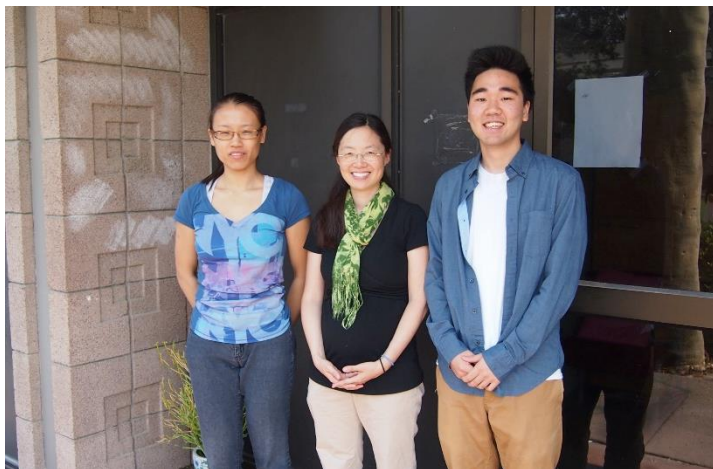
Applying the protocol to analyze human skin wounds



Human skin wounds



Acknowledgments



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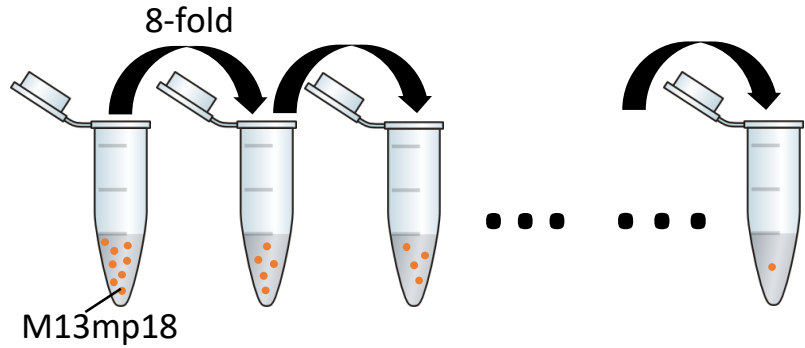
Arica Lubin
Stephanie Mendes
Joel Rothman



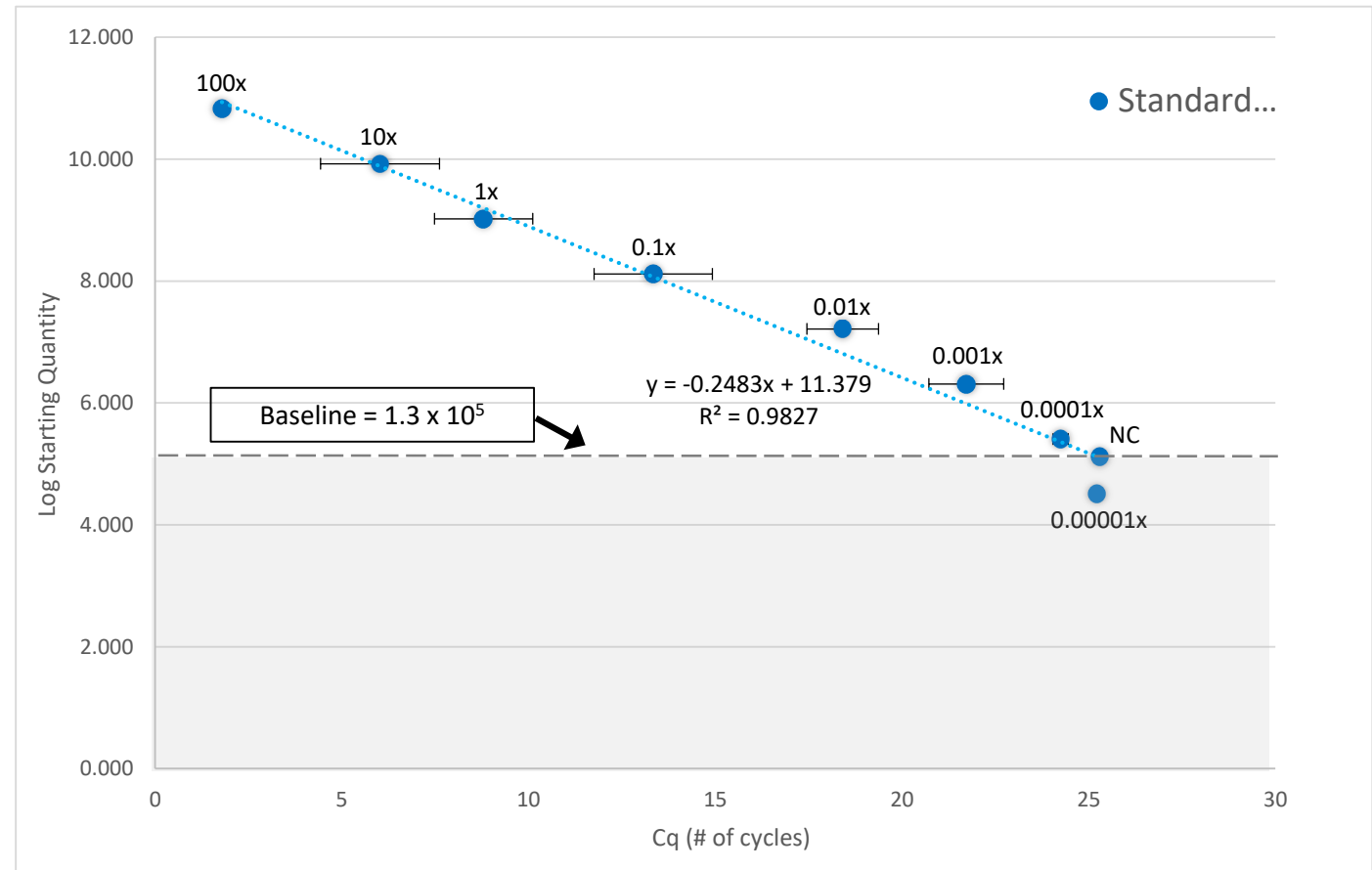
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Quantifying phage M13 DNA by qPCR – Standard Curve

Serial dilution of M13mp18



| Sample | Expected | | Observed | |
|--------------------------------------|----------------------|----------------------|----------|----------------|
| | Total # of DNA | Total # of DNA | Mean Cq | Standard Error |
| 100x | 6.7×10^{10} | 6.7×10^{10} | 1.79 | 0 |
| 10x | 8.4×10^9 | 8.4×10^9 | 6.02 | 1.59 |
| 1x | 1.0×10^9 | 1.0×10^9 | 8.80 | 1.32 |
| 0.1x | 1.3×10^8 | 1.3×10^8 | 13.33 | 1.59 |
| 0.01x | 1.6×10^7 | 1.6×10^7 | 18.41 | 0.96 |
| 0.001x | 2.0×10^6 | 2.0×10^6 | 21.72 | 1.00 |
| 0.0001x | 2.6×10^5 | 2.6×10^5 | 24.25 | 0.20 |
| 0.00001x | 3.2×10^4 | 3.2×10^4 | 25.22 | 0.05 |
| Negative Control (dH ₂ O) | 0 | 1.3×10^5 | 25.29 | 0.155 |

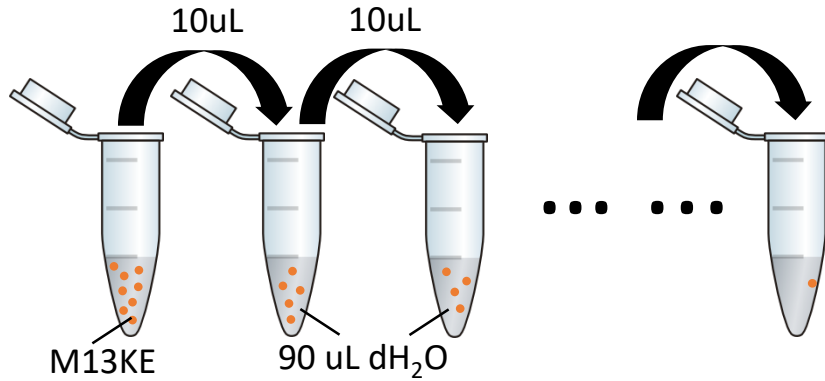


- Baseline of detection = 1.3×10^5 copies/uL
- Linear equation, $R^2 = 0.98271$

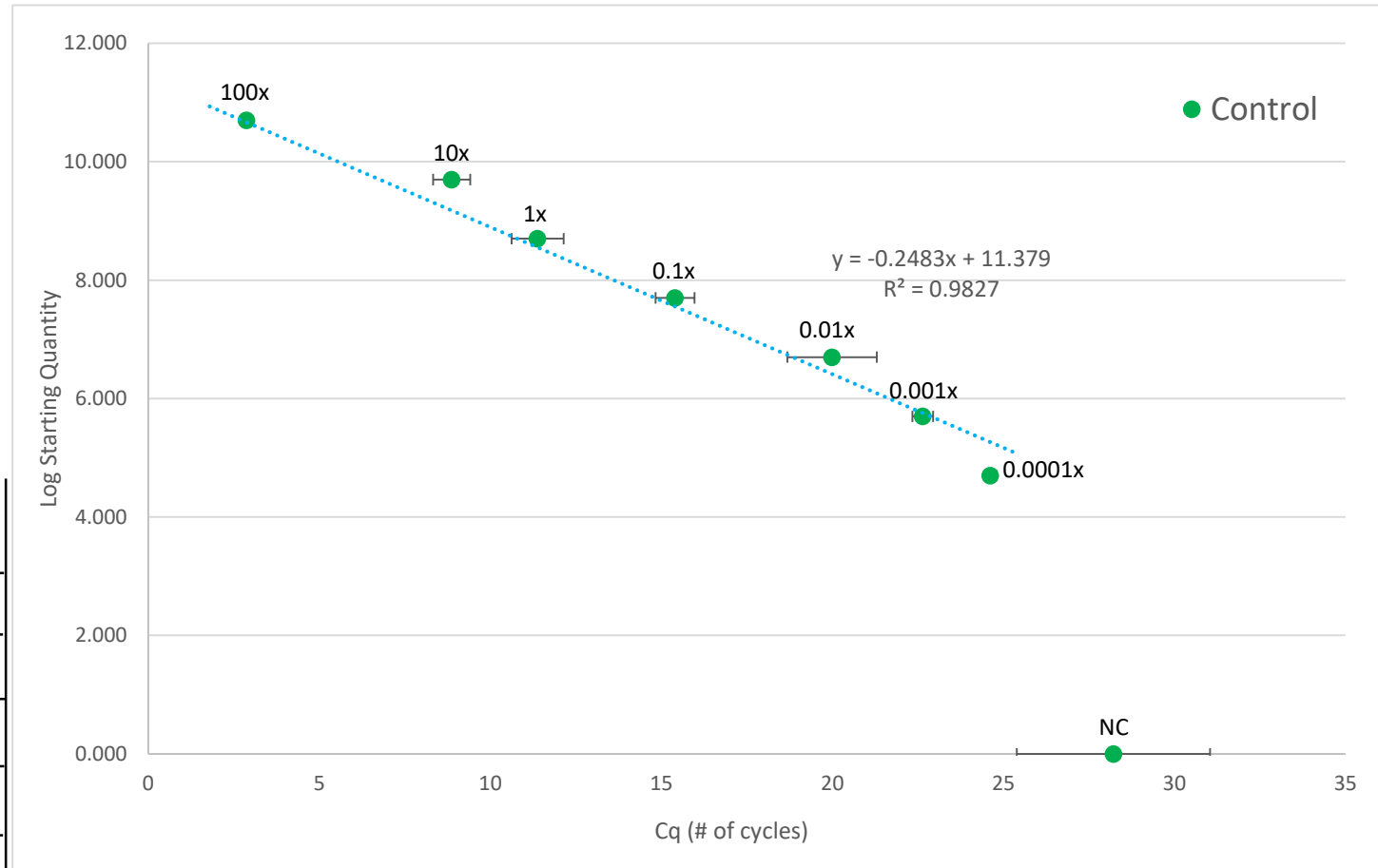
This standard curve was used in the absolute quantification of our samples.

Quantifying phage M13 DNA by qPCR – Control Samples

10-fold serial dilution of M13KE



| Sample | <i>Expected</i> | | <i>Observed</i> | |
|------------------------|----------------------|-----------------------|-----------------|----------------|
| | Total # of DNA | Total # of DNA | Mean Cq | Standard Error |
| 100x | 5.0×10^{10} | 4.63×10^{10} | 2.87 | 0 |
| 10x | 5.0×10^9 | 1.50×10^9 | 8.87 | 0.54 |
| 1x | 5.0×10^8 | 3.57×10^8 | 11.38 | 0.76 |
| 0.1x | 5.0×10^7 | 3.58×10^7 | 15.41 | 0.57 |
| 0.01x | 5.0×10^6 | 2.60×10^6 | 19.99 | 1.31 |
| 0.001x | 5.0×10^5 | 5.70×10^5 | 22.65 | 0.30 |
| 0.0001x | 5.0×10^4 | 1.84×10^5 | 24.62 | 0.17 |
| Negative Control (PBS) | 0 | 2.36×10^4 | 28.22 | 2.82 |

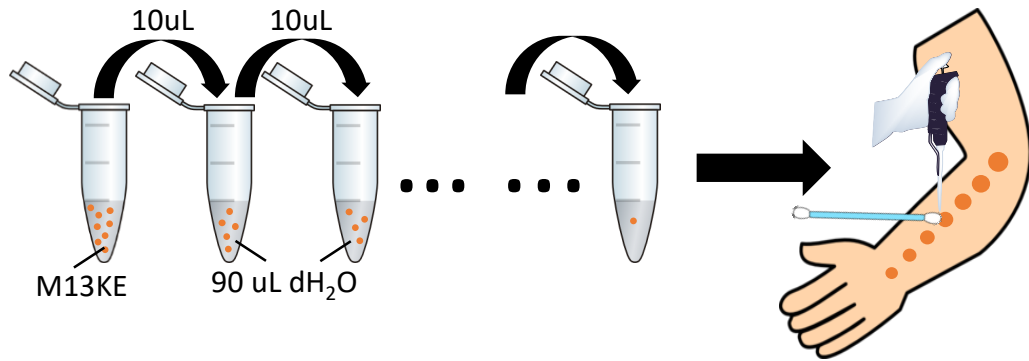


- Baseline of detection for M13KE = 2.36×10^4 copies

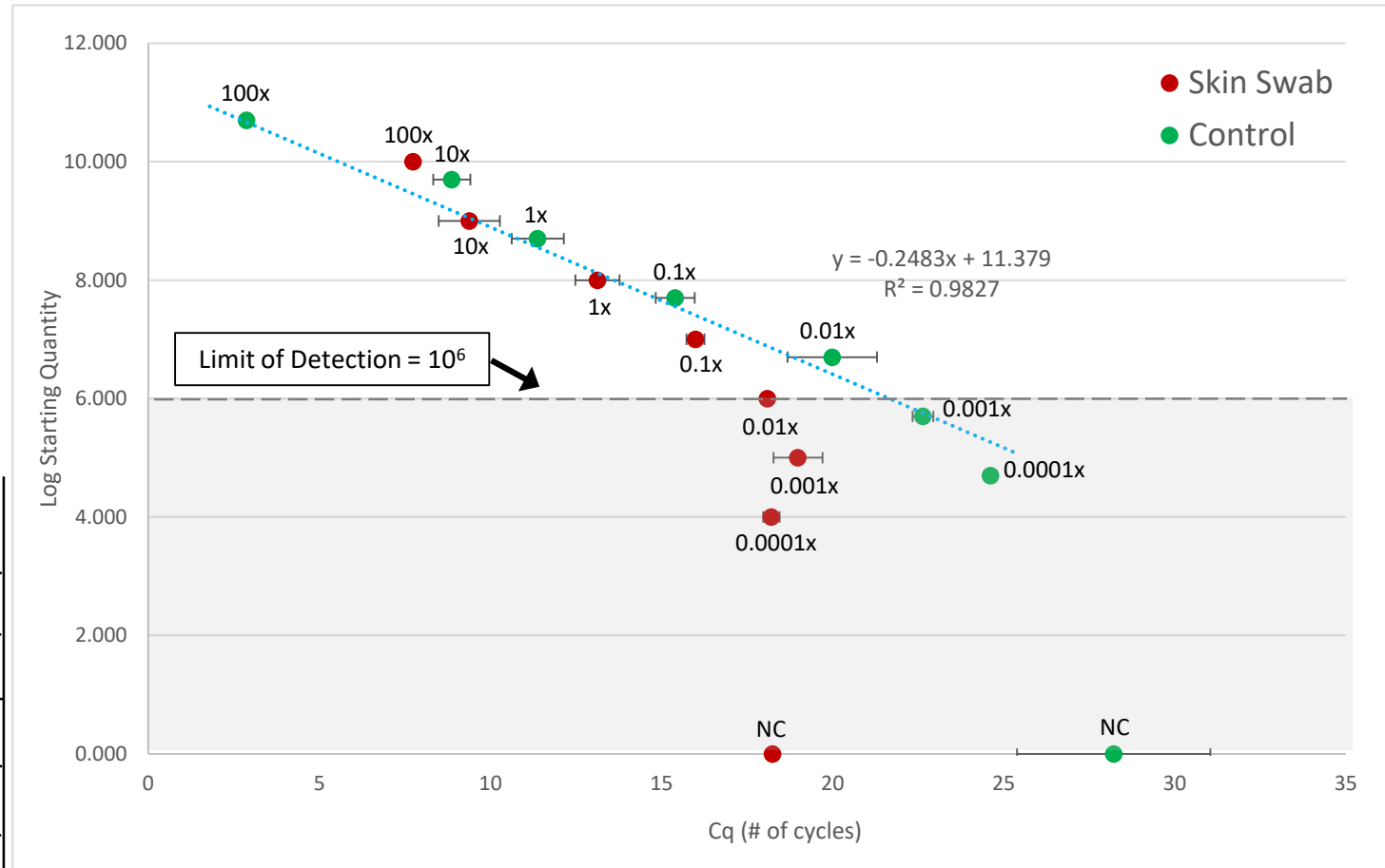
Control samples were compared to skin swab samples

Quantifying phage M13 DNA by qPCR – Swab Samples

10-fold serial dilution of M13KE



| Sample | <i>Expected</i> | <i>Observed</i> | | Standard Error |
|------------------------|----------------------|-----------------------|---------|----------------|
| | Total # of DNA | Total # of DNA | Mean Cq | |
| 100x | 1.0×10^{10} | 2.86×10^{10} | 7.75 | 0 |
| 10x | 1.0×10^9 | 1.12×10^9 | 9.38 | 0.89 |
| 1x | 1.0×10^8 | 1.31×10^8 | 13.13 | 0.65 |
| 0.1x | 1.0×10^7 | 2.56×10^7 | 15.99 | 0.26 |
| 0.01x | 1.0×10^6 | 7.68×10^6 | 18.09 | 0.13 |
| 0.001x | 1.0×10^5 | 4.61×10^6 | 18.99 | 0.71 |
| 0.0001x | 1.0×10^4 | 7.20×10^6 | 18.21 | 0.24 |
| Negative Control (PBS) | 0 | 7.04×10^6 | 18.24 | 0.13 |



- Limit of detection for human skin swab = 1.0×10^6 copies
- Contaminating sequences on human skin