Happy holiday! Remember to take care of yourself and your loved ones!

```
- using Pkg

- using Interact

url =
"https://raw.githubusercontent.com/CSSEGISandData/COVID-19/master/csse_covid_19_data
- url ="https://raw.githubusercontent.com/CSSEGISandData/COVID-
19/master/csse_covid_19_data/csse_covid_19_time_series/time_series_covid19_confirmed_global.csv"

"covid_data.csv"
- download(url, "covid_data.csv")

- using CSV
- using DataFrames
```

	Province/State	Country/Region	Lat	Long	1/2	
1	missing	"Afghanistan"	33.9391	67.71	0	
2	missing	"Albania"	41.1533	20.1683	0	
3	missing	"Algeria"	28.0339	1.6596	0	
4	missing	"Andorra"	42.5063	1.5218	0	
5	missing	"Angola"	-11.2027	17.8739	0	
6	missing	"Antigua and Barbuda"	17.0608	-61.7964	0	
7	missing	"Argentina"	-38.4161	-63.6167	0	
8	missing	"Armenia"	40.0691	45.0382	0	
9	"Australian Capital Territory"	"Australia"	-35.4735	149.012	0	
10	"New South Wales"	"Australia"	-33.8688	151.209	0	
mo	ore					
280	missing	"Zimbabwe"	-19.0154	29.1549	0	

CSV.read("covid\_data.csv", DataFrame)

data = CSV.read("covid\_data.csv", DataFrame);

	Province/State	Country/Region	Lat	Long	1/2	
1	missing	"Afghanistan"	33.9391	67.71	0	
2	missing	"Albania"	41.1533	20.1683	0	
3	missing	"Algeria"	28.0339	1.6596	0	
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8	missing	"Armenia"	40.0691	45.0382	0	
9	"Australian Capital Territory"	"Australia"	-35.4735	149.012	0	
10	"New South Wales"	"Australia"	-33.8688	151.209	0	
mo	ore					
280	missing	"Zimbabwe"	-19.0154	29.1549	0	

### data

# countries =

["Afghanistan", "Albania", "Algeria", "Andorra", "Angola", "Antigua and Barbuda", "A

countries = collect(data[:,2])

## unique\_countries =

["Afghanistan", "Albania", "Algeria", "Andorra", "Angola", "Antigua and Barbuda", "A

unique\_countries = unique(countries)

### using WebIO

(::Base.Fix2{typeof(startswith), String}) (generic function with 1 method)

startswith("I")

I\_countries = [startswith(country,"I") for country in countries];

	Province/State	Country/Region	Lat	Long	1/22/20	1/23/20	1/24/20	1/25
1	missing	"Iceland"	64.9631	-19.0208	0	0	0	0
2	missing	"India"	20.5937	78.9629	0	0	0	0
3	missing	"Indonesia"	-0.7893	113.921	0	0	0	0
4	missing	"Iran"	32.4279	53.688	0	0	0	0
5	missing	"Iraq"	33.2232	43.6793	0	0	0	0
6	missing	"Ireland"	53.1424	-7.6921	0	0	0	0
7	missing	"Israel"	31.0461	34.8516	0	0	0	0
8	missing	"Italy"	41.8719	12.5674	0	0	0	0

# data[I\_countries, :]

BitVector: [false, false, fals

countries .== "India"

### $IND\_row = 148$

IND\_row = findfirst(countries .=="India")

### IND\_data\_row =

DataFrameRow (719 columns)

	Province/State	Country/Region	Lat	Long	1/22/20	1/23/20	1/24/20	1/25
	String63	String63	Float64?	Float64?	Int64	Int64	Int64	Int
148	missing	India	20.5937	78.9629	0	0	0	0

IND\_data\_row = data[IND\_row, :]

### IND\_data =

[0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, 2, 3, 3, 3, 3, 3, 3, 3, 3, 3, more ,34799691, 3480888

IND\_data = collect(IND\_data\_row[5:end])

using Plots

plot(IND\_data);

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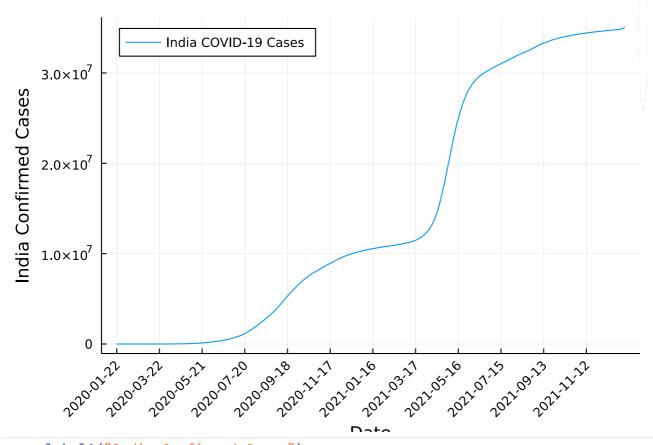
```
date_strings =
```

["1/22/20", "1/23/20", "1/24/20", "1/25/20", "1/26/20", "1/27/20", "1/28/20", "1/29/

- date\_strings = String.(names(data))[5:end]
- using Dates

format = dateformat"m/d/Y"

- format = Dates.DateFormat("m/d/Y")
- dates = parse.(Date, date\_strings, format) + Year(2000);
- plot(dates,IND\_data, xticks=dates[1:60:end], xrotation=45, leg=:topleft,
  label="India COVID-19 Cases");
- xlabel!("Date");



ylabel!("India Confirmed Cases")

# Simple Model of infection

- md"
- Simple Model of infection
- . "

### UndefVarError: @cI\_n\_cmd not defined

- 1. top-level scope @ :0
- 2. #macroexpand#51 @ expr.jl:115 [inlined]
- 3. macroexpand @ expr.jl:114 [inlined]
- 4. try\_macroexpand(::Module, ::Base.UUID, ::Expr) @ PlutoRunner.jl:253
- 5. var"#run\_expression#25"(::Bool, ::typeof(Main.PlutoRunner.run\_expression), ::Module, ::Expr, ::Base.UUID, ::Nothing, ::Nothing) @ PlutoRunner.jl:482
- 6. top-level scope @ none:1

$$I_{n+1} = I_n + (cI_n)$$

$$I0 = 1$$

 $I_0 = 1$ 

$$I _0 = 1$$

$$I_0 = 1$$

$$I_0 = 1$$

$$c = 0.01$$

c = 0.01 # average no. of people that each individual infects on each day

$$\lambda = 1 + \underline{c};$$

### $I_1 = 1.01$

$$I_1 = \underline{\lambda} * \underline{I_0}$$

### $I_2 = 1.0201$

$$I_2 = \underline{\lambda} * \underline{I_1}$$

### $I_3 = 1.030301$

$$I_3 = \lambda * I_2$$

I =

$$I = zeros(\underline{T})$$

1

$$I[1] = \underline{I}_{\underline{0}}$$

$$T = 20$$

T=20

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```
• for n in 1:T-1
        I[n+1] = \underline{\lambda} * \underline{I}[n]
        @show n, I[n]
 end
 [1.0, 1.01, 1.0201, 1.0303, 1.0406, 1.05101, 1.06152, 1.07214, 1.08286, 1.09369, 1.10
   plot(I, m=:o, label="I[n]", legend=:topleft);
p = 0.02
   p = 0.02
\alpha = 0.01
 \alpha = 0.01
N = 1000
   N = 1000
run_infection (generic function with 2 methods)
   function run_infection(T=20)
        I = zeros(T)
       I[1] = I_0
       for n in 1:T-1
            I[n+1] = \underline{\lambda} * I[n]
        end
        return I
   end
# 2 methods for generic function run_infection:

    run_infection() in Main.workspace#134 at

     /Users/virajvaidya/Downloads/MITMATH2.jl#==#afef1536-fd7e-47cc-8b6b-
     a9d4e716f882:1
   • run infection(T) in Main.workspace#134 at
     /Users/virajvaidya/Downloads/MITMATH2.jl#==#afef1536-fd7e-47cc-8b6b-
```

```
methods(<u>run_infection</u>)
```

```
[1.0, 1.01, 1.0201, 1.0303, 1.0406, 1.05101, 1.06152, 1.07214, 1.08286, 1.09369]
run_infection(10)
```

a9d4e716f882:1

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### Multiple definitions for <u>I\_result</u>

```
I_result = run_infection(10);
```

UndefVarError: I\_result not defined

1. top-level scope @ | Local: 1

I\_result

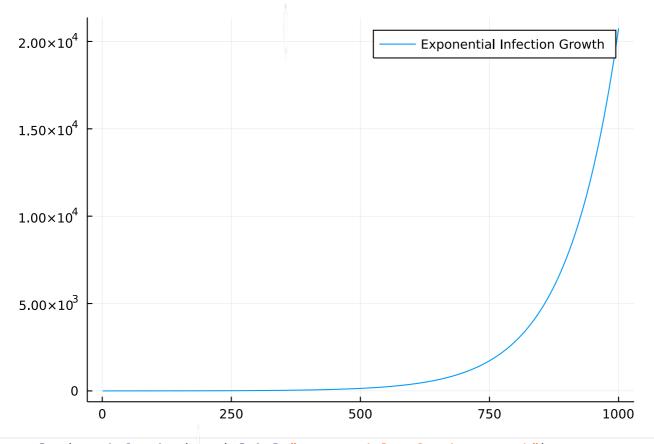
# UndefVarError: I\_result not defined

1. top-level scope @ [Local: 1]

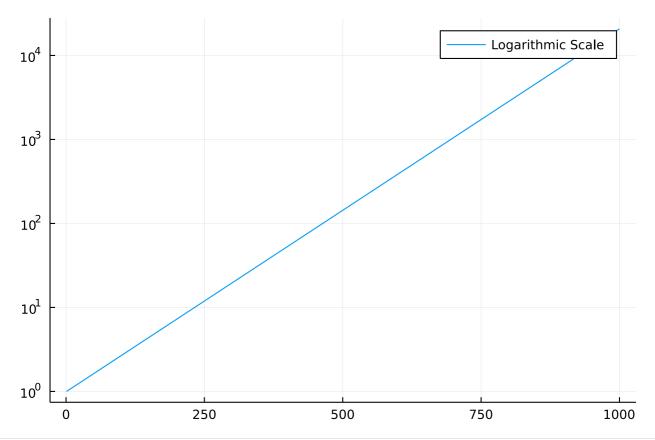
```
plot(I_result, m=:0)

end_T = 1000

end_T = 1000
```



plot(run\_infection(1000), label="Exponential Infection Growth")



plot(run\_infection(1000), yscale=:log10, label="Logarithmic Scale")

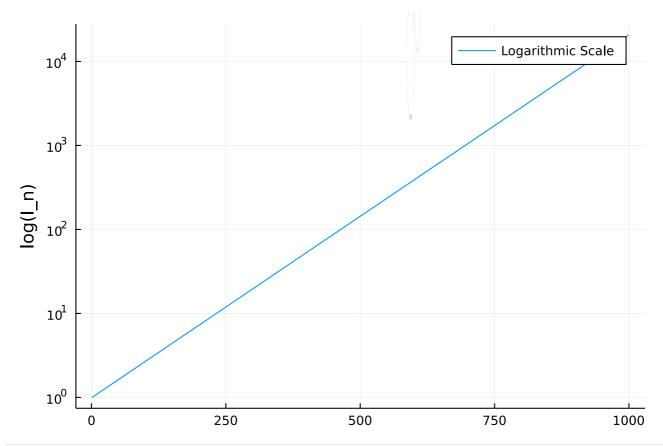
# Multiple definitions for $\underline{I}$ result

I\_result = run\_infection(1000)

# UndefVarError: I\_result not defined

1. top-level scope @ Local: 1

plot(log10.(I\_result))



ylabel!("log(I\_n)")

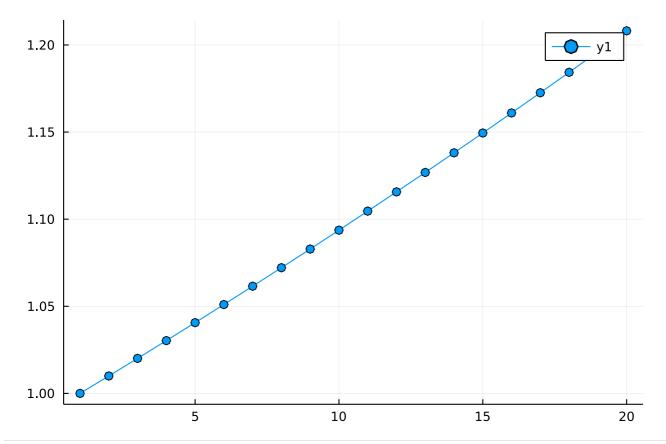
β (generic function with 1 method)

$$\beta(\mathbf{I}, S) = p * \alpha * S$$

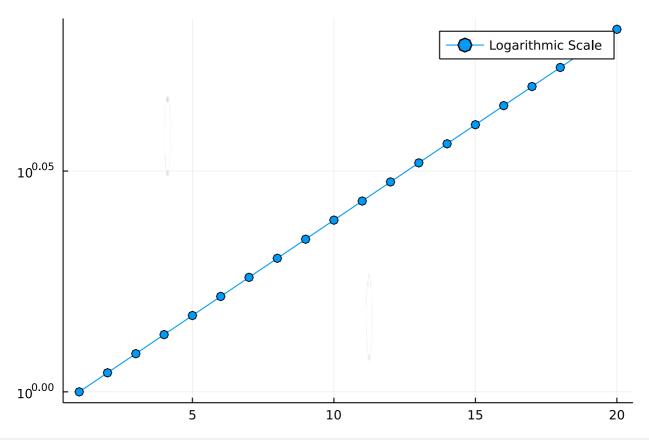
I\_20 =

[1.0, 1.01, 1.0201, 1.0303, 1.0406, 1.05101, 1.06152, 1.07214, 1.08286, 1.09369, 1.10

 $I_20 = run_infection(20)$ 



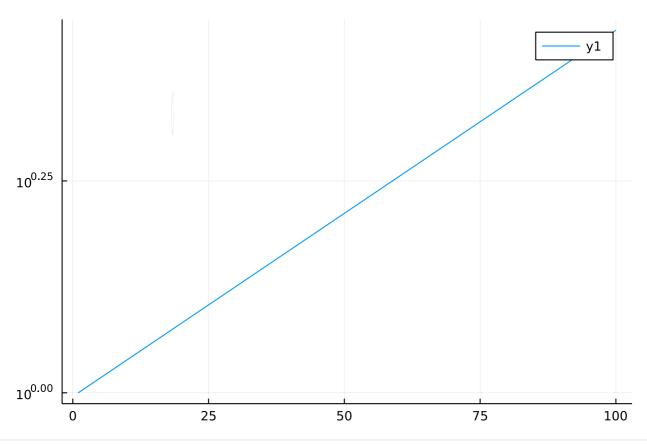
plot(<u>I\_20</u>, m=:o)



plot(<u>I\_20</u>, m=:o, yscale=:log10, label="Logarithmic Scale")

<sup>2</sup> ,2.44863, 2.47312, 2.49785, 2.52283, 2.54806, 2.57354, 2.59927, 2.62527, 2.65152, 2.67

I\_100= run\_infection(100)



plot(I\_100, yscale=:log10)

c\_average = 1.1

c\_average = 1.1

cs =

[0.984051, 1.01137, 1.1932, 1.21093, 1.07294, 1.02978, 1.04538, 1.1157, 1.08298, 0.94

cs = [c\_average + 0.1\*randn() for i in 1:100]

