

# Expresiones regulares

Bioinformática  
Facultad de Ciencias e Ingeniería

# Expresión regular (regex)

- Cadena de caracteres que define un patrón que se encuentra en un bloque de texto.
- Se puede construir una expresión regular para permitir múltiples coincidencias posibles, mientras se restringen otras posibilidades.

A la clase de profundización en bioinformática del periodo 2025-1 asisten 10 estudiantes.

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Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)  
reference genome

SARS-CoV-2 genome

LOCUS MT324064 67 bp DNA linear INV 03-OCT-2021  
DEFINITION mitochondrial cytochrome oxidase subunit 1 (COI) gene, partial cds; mitochondrial  
ACCESSION MT324064.1  
VERSION calc MT324064.1  
KEYWORDS lese, coi, positive  
SOURCE /mgi/mitochondrion *Mythimna separata* (*Pseudaletia separata*)  
ORGANISM *Mythimna separata*  
https://eutils.ncbi.nlm.nih.gov/entrez/eutils/efetch.fcgi?db=Eukaryota; Metazoa; Ecdysozoa; Arthropoda; Hexapoda; Insecta;  
/home/paul/Pterygota; Neoptera; Endopterygota; Lepidoptera; Glossata;  
Ditrysia; Noctuoidea; Noctuidae; Hadeninae; Mythimna.  
REFERENCE 1 (bases 1 to 667)  
AUTHORS Mahat,K., Mitchell,A. and Zangpo,T.  
TITLE An updated global COI barcode reference data set for Fall Armyworm (*Spodoptera frugiperda*) and first record of this species in Bhutan  
JOURNAL J Asia Pac Entomol 24 (1), 105-109 (2021)  
REFERENCE 2 (bases 1 to 667)  
AUTHORS Mitchell,A. and Mahat,K.  
TITLE Direct Submission  
JOURNAL Submitted (09-APR-2020) Entomology, Australian Museum, 1 William  
Dawes Street, DARLINGHURST, New South Wales 2010, Australia  
FEATURES Location/qualifiers  
source 1..667  
/organism="Mythimna separata"  
/organelle="mitochondrion"  
/mol\_type="genomic DNA"  
/specimen\_voucher="am11901"  
/db\_xref="BOLD:BPI001-19.COI-5P"  
/country="Bhutan: Punakha, Shengana"  
/collection\_date="10-May-2013"  
/collected\_by="Kiran Mahat"  
/PCR\_primers="fwd\_seq:  
gtaaaacgacggccagttcwcacwaaycayaarrwtatyygg, rev\_seq:  
caggaaacacgactatgacaaaatrtawacytcddggrtgnc"  
gene  
CDS  
gene  
<1..>667  
/gene="COI"  
<1..>667  
/gene="COI"  
/codon\_start=2  
/transl\_table=5  
/product="cytochrome oxidase subunit 1"  
/protein\_id="UBQ033860.1"  
/translation="TLYIFGIWAGMVGTSLSLLIRAEGLTPGSILGDDQIYNTIVTA  
HAFIMIFFMVMPIMIGGFGNWLVPMLGAPDMAPRMNNMSFWLPPSLTLISSIV  
ENGAGTGTVYPPLSSIAHGSSVLDALIFSLHLAGISIISLGAINFTTIINMRNSL  
SFDMQPLFIHAVGITAFLLLSLPVLAGITMLLTDRLNNTSFDPAGGGDPILYQHL  
ORIGIN  
1 aacattatat ttatTTTTG gaatttggc tggatagt ggaacttcat taagattact  
61 aattcgact gaatttagaa cccctggatc ttatTTGGA gatgaccaaa ttataatcac  
121 tattgttaca gtcatgtc ttatataat ttttttata ttataccata ttataattgg  
181 aggattttgtt aattttatgt tacccatata attagagct cttatccatcg

```
>MT324064_Bhutan
Lo es menor que 1 no hay selección positiva. La
idea es calcular ese radio para cada rama del arbol identificar
1 5 |aacattatat tttatTTTg gaatttgagc tggtagatTT ggaacttcat taagattact
61 aattcgagct gaatttaggaa cccctggatc tttaattgga gatgaccaaa tttataatac
121 tattgttaca gctcatgctt ttattataat ttttttataa gttataccta ttataattgg
181 aggatttgtt aattgattag taccttaat attaggagct cctgatatag catttcctcg
241 tataaataat ataagttttt gattacttcc cccatcttaa actttactaa tttcaagtag
301 aattgttagaa aatggagcag gaacaggatg aacagtttat cccccacttt catcaaataat
361 tgctcatgga ggttagatctg tagatTTAGC tatttttctt ttacatttag ctggaaatttc
421 ctctatTTA ggtgctatta attttattac tacaattatc aatacgtat taaatagttt
481 atcatttgat caaatacctt tatTTatTTg agctgttggg attactgcatttttactatt
541 attatctta cctgtattag caggagctat tactataactt ttaacagatc gaaatcttaa
601 tacatctttt tttgatcctg ctggaggagg tgatccaatt ttatataaac atttattttg
661 atttttt
```

hay de todas las especies de  
Mythimna de NCBI para hacer  
arbolitos.

La idea es armar arbol con  
secuencias de Indonesia, EU

## Wildcards

\w	Letters, numbers and _
.	Any character except \n \r
\d	Numerical digits
\t	Tab
\r	Return character. Also used as the generic end-of-line character in TextWrangler
\n	Line-feed character. Also used as the generic end-of-line character in Notepad++
\s	Space, tab, or end of line
[A-Z]	A single character of the ranges indicated in square brackets
[^A-Z]	A single character including all characters not in the brackets. Note that this will include \n unless otherwise specified, and may cause you to match across lines
\	Used to escape punctuation characters so they are searched for as themselves, not interpreted as wildcards or special symbols
\\	The \ symbol itself, escaped

## Boundaries

^	Match the start of the line, i.e., the position before the first character
\$	Match the last position before the end-of-line character

## Quantifiers, used in combination with characters and wildcards

+	Look for the longest possible match of one or more occurrences of the character, wildcard, or bracketed character range immediately preceding. The match will extend as far as it can while still allowing the entire expression to match.
*	As above, matches as many of the previous character to occur, but allows for the character not to occur at all if the match still succeeds
?	Modifies greediness of + or * to match the shortest possible match instead of longest
{}	Specify a range of numbers to repeat the match of the previous character. For example: \d{2,4} matches between 2 and 4 digits in a row [AC]{4,} matches 4 or more of the letter A or C in a row

## Capturing and replacing

( )	Capture the search results between the parentheses for use in the replacement term
\1 \$1	Substitute the contents of the matched into the replacement term, in numerical order. Syntax depends on the text editor or language that you are using.