

This week's highlights

- Evaluate the truth value of a compound proposition given truth values of its constituent variables.
- Prove propositional equivalences using truth tables
- Prove propositional equivalences using other known equivalences, e.g.
 - DeMorgan's laws
 - Double negation laws
 - Distributive laws, etc.
- Compute the CNF and DNF of a given compound proposition.
- Define a predicate over a finite domain using a table of values and as properties
- Determine the truth value of the proposition resulting from evaluating a predicate
- Describe the set of domain elements that make a predicate with one input evaluate to true.
- Evaluate universal and existential statements about finite domains (with no quantifier alternations).
- Counterexample and witness-based arguments for predicates with infinite domains
- Practice combinations of \wedge , \rightarrow in conjunction with universal and existential quantifiers
- State and apply DeMorgan's law for quantified statements.
- Translate sentences from English to propositional logic using appropriate propositional variables and boolean operators.

- Decide and justify whether or not a collection of propositions is consistent.
- Use predicates with set of tuples as their domain to relate values to one another
- Evaluate nested quantifiers: both alternating and not.

Lecture videos

Week 4 Day 1 YouTube playlist

Week 4 Day 2 YouTube playlist

Week 4 Day 3 YouTube playlist

Monday January 25

Consider the following algorithm to introduce redundancy in a string of 0s and 1s.

Create redundancy by repeating each bit three times

```

1 procedure redun3( $a_{k-1} \cdots a_0$ : a binary string)
2 for  $i := 0$  to  $k-1$ 
3    $c_{3i} := a_i$ 
4    $c_{3i+1} := a_i$ 
5    $c_{3i+2} := a_i$ 
6 return  $c_{3k-1} \cdots c_0$ 

```

Decode sequence of bits using majority rule on consecutive three bit sequences

```

1 procedure decode3( $c_{3k-1} \cdots c_0$ : a binary string whose length is an integer multiple of 3)
2 for  $i := 0$  to  $k-1$ 
3   if exactly two or three of  $c_{3i}, c_{3i+1}, c_{3i+2}$  are set to 1
4      $a_i := 1$ 
5   else
6      $a_i := 0$ 
7 return  $a_{k-1} \cdots a_0$ 

```

Give a recursive definition of the set of outputs of the *redun3* procedure, *Out*, **Basis step:** _____

Recursive step: _____

Consider the message $m = 0001$ so that the sender calculates $\text{redun3}(m) = \text{redun3}(0001) = 000000000111$.

Introduce ____ errors into the message so that the signal received by the receiver is _____ but the receiver is still able to decode the original message.

Challenge: what is the biggest number of errors you can introduce?

Building a circuit for line 3 in *decode* procedure: given three input bits, we need to determine whether the majority is a 0 or a 1.

c_{3i}	c_{3i+1}	c_{3i+2}	a_i	Circuit
1	1	1		
1	1	0		
1	0	1		
1	0	0		
0	1	1		
0	1	0		
0	0	1		
0	0	0		

Wednesday January 27

Definition: A **predicate** is a function from a given set (domain) to $\{T, F\}$.

A predicate can be applied, or **evaluated** at, an element of the domain.

Two predicates over the same domain are **equivalent** means they evaluate to the same truth values for all possible assignments of domain elements to the input.

Input	Output		
x	$P(x)$ $[x]_{2c,3} > 0$	$N(x)$ $[x]_{2c,3} < 0$	$Mystery(x)$
000	F		T
001	T		T
010	T		T
011	T		F
100	F		F
101	F		T
110	F		F
111	F		T

The domain for each of the predicates $P(x)$, $N(x)$, $Mystery(x)$ is _____

Definition: The **truth set** of a predicate is the collection of all elements in its domain where the predicate evaluates to T .

The truth set for the predicate $P(x)$ is _____.

The truth set for the predicate $N(x)$ is _____.

The truth set for the predicate $Mystery(x)$ is _____.

Definition 15 (Universal quantification) The **universal quantification** of $P(x)$ is the statement “ $P(x)$ for all values of x in the domain” and is written $\forall xP(x)$. An element for which $P(x) = F$ is called a **counterexample** of $\forall xP(x)$.

The **existential quantification** of $P(x)$ is the statement “There exists an element x in the domain such that $P(x)$ ” and is written $\exists xP(x)$. An element for which $P(x) = T$ is called a **witness** of $\exists xP(x)$.

Example: _____ is a true existential quantification.

Statements involving predicates and quantifiers are **logically equivalent** means they have the same truth value no matter which predicates (domains and functions) are substituted in.

Quantifier version of De Morgan’s laws: $\neg\forall xP(x) \equiv \exists x(\neg P(x))$

$$\neg\exists xQ(x) \equiv \forall x(\neg Q(x))$$

Example: _____ is a false universal quantification. It is logically equivalent to _____

Recall: Each RNA strand is a string whose symbols are elements of the set $B = \{\mathbf{A}, \mathbf{C}, \mathbf{G}, \mathbf{U}\}$. The **set of all RNA strands** is called S . The function *rlen* that computes the length of RNA strands in S is:

$$\begin{array}{lll} \text{Basis Step:} & \text{If } b \in B \text{ then} & \begin{array}{ll} rlen : S & \rightarrow \mathbb{Z}^+ \\ rlen(b) & = 1 \end{array} \\ \text{Recursive Step:} & \text{If } s \in S \text{ and } b \in B, \text{ then} & rlen(sb) = 1 + rlen(s) \end{array}$$

Example predicates on S

$H(s) = T$	Truth set of H is _____
$L_3(s) = \begin{cases} T & \text{if } rlen(s) = 3 \\ F & \text{otherwise} \end{cases}$	Strand where L_3 evaluates to T is e.g. _____ Strand where L_3 evaluates to F is e.g. _____
F_A is defined recursively by: Basis step: $F_A(\mathbf{A}) = T, F_A(\mathbf{C}) = F_A(\mathbf{G}) = F_A(\mathbf{U}) = F$ Recursive step: If $s \in S$ and $b \in B$, then $F_A(sb) = F_A(s)$	Strand where F_A evaluates to T is e.g. _____ Strand where F_A evaluates to F is e.g. _____
P_{AUC} is defined as the predicate whose truth set is the collection of RNA strands where the string AUC is a sub-string (appears inside s , in order and consecutively)	Strand where P_{AUC} evaluates to T is e.g. _____ Strand where P_{AUC} evaluates to F is e.g. _____

Friday January 29

Definition (Rosen p123): The **Cartesian product** of the sets A and B , $A \times B$, is the set of all ordered pairs (a, b) , where $a \in A$ and $b \in B$. That is: $A \times B = \{(a, b) \mid (a \in A) \wedge (b \in B)\}$. The Cartesian product of the sets A_1, A_2, \dots, A_n , denoted by $A_1 \times A_2 \times \dots \times A_n$, is the set of ordered n -tuples (a_1, a_2, \dots, a_n) , where a_i belongs to A_i for $i = 1, 2, \dots, n$. That is, $A_1 \times A_2 \times \dots \times A_n = \{(a_1, a_2, \dots, a_n) \mid a_i \in A_i \text{ for } i = 1, 2, \dots, n\}$

Recall: Each RNA strand is a string whose symbols are elements of the set $B = \{A, C, G, U\}$. The **set of all RNA strands** is called S . The function $rnalen$ that computes the length of RNA strands in S is:

$$\begin{array}{lll} & & rnalen : S \rightarrow \mathbb{Z}^+ \\ \text{Basis Step:} & \text{If } b \in B \text{ then} & rnalen(b) = 1 \\ \text{Recursive Step:} & \text{If } s \in S \text{ and } b \in B, \text{ then} & rnalen(sb) = 1 + rnalen(s) \end{array}$$

A function $basecount$ that computes the number of a given base b appearing in a RNA strand s is:

$$\begin{array}{lll} & & basecount : S \times B \rightarrow \mathbb{N} \\ \text{Basis Step:} & \text{If } b_1 \in B, b_2 \in B & basecount(b_1, b_2) = \begin{cases} 1 & \text{when } b_1 = b_2 \\ 0 & \text{when } b_1 \neq b_2 \end{cases} \\ \text{Recursive Step:} & \text{If } s \in S, b_1 \in B, b_2 \in B & basecount(sb_1, b_2) = \begin{cases} 1 + basecount(s, b_2) & \text{when } b_1 = b_2 \\ basecount(s, b_2) & \text{when } b_1 \neq b_2 \end{cases} \end{array}$$

L with domain $S \times \mathbb{Z}^+$ is defined by, for $s \in S$ and $n \in \mathbb{Z}^+$, BC with domain _____ is defined
 S and $b \in B$ and $n \in \mathbb{N}$,

$$L(s, n) = \begin{cases} T & \text{if } rnalen(s) = n \\ F & \text{otherwise} \end{cases} \qquad BC(s, b, n) = \begin{cases} T & \text{if } basecount(s, b) = n \\ F & \text{otherwise} \end{cases}$$

Element where L evaluates to T : _____

Element where BC evaluates to T : _____

Element where L evaluates to F : _____

Element where BC evaluates to F : _____

Notation: for a predicate P with domain $X_1 \times \dots \times X_n$ and a n -tuple (x_1, \dots, x_n) with each $x_i \in X$, we write $P(x_1, \dots, x_n)$ to mean $P((x_1, \dots, x_n))$.

$\exists t BC(t)$ In English: _____
 Witness that proves this existential quantification is true: _____
 $\forall (s, b, n) (BC(s, b, n))$ In English: _____
 Counterexample that proves this universal quantification is false: _____
New predicates from old $BC(s, b, n)$ means *basecount*(s, b) = n .

Predicate	Domain	Example domain element where predicate is T
$basecount(s, b) = 3$		
$basecount(s, A) = n$		
$\exists n \in \mathbb{N} (basecount(s, b) = n)$		
$\forall b \in B (basecount(s, b) = 1)$		

Alternating quantifiers

$$\forall s \exists n BC(s, A, n)$$

In English: _____

$$\exists n \forall s BC(s, U, n)$$

In English: _____

Evaluate each quantified statement as T or F .

$\forall s \forall b \exists n BC(s, b, n)$	$\forall s \forall n \exists b BC(s, b, n)$	$\forall b \forall n \exists s BC(s, b, n)$
$\exists s \forall b \exists n BC(s, b, n)$	$\forall s \exists n \forall b BC(s, b, n)$	$\exists b \exists n \forall s BC(s, b, n)$

Extra example: Write the negation of each of the statements above, and use De Morgan's law to find a logically equivalent version where the negation is applied only to the BC predicate (not next to a quantifier).

Review quiz questions

1. **Monday** Real-life representations are often prone to corruptions. Biological codes, like RNA, may mutate naturally¹ and during measurement; cosmic radiation and other ambient noise can flip bits in computer storage². One way to recover from corrupted data is to exploit redundancy. Consider the following algorithm to introduce redundancy in a string of 0s and 1s.

Create redundancy by repeating each bit three times

```
1 procedure redun3( $a_{k-1} \dots a_0$ : a binary string)
2 for  $i := 0$  to  $k-1$ 
3    $c_{3i} := a_i$ 
4    $c_{3i+1} := a_i$ 
5    $c_{3i+2} := a_i$ 
6 return  $c_{3k-1} \dots c_0$ 
```

Decode sequence of bits using majority rule on consecutive three bit sequences

```
1 procedure decode3( $c_{3k-1} \dots c_0$ : a binary string whose length is an integer multiple of 3)
2 for  $i := 0$  to  $k-1$ 
3   if exactly two or three of  $c_{3i}, c_{3i+1}, c_{3i+2}$  are set to 1
4      $a_i := 1$ 
5   else
6      $a_i := 0$ 
7 return  $a_{k-1} \dots a_0$ 
```

For each of the following, type in your answers precisely including all notational punctuation.

- (a) Give the output of *redun3*(100).
- (b) If the output of running *redun3* is 000000111000111, what was its input?
- (c) Give the output of *decode3*(100).
- (d) How many distinct possible inputs to *decode3* give the output 01?

2. **Wednesday** Consider the following predicates, each of which has as its domain the set of all bitstrings whose leftmost bit is 1

¹Mutations of specific RNA codons have been linked to many disorders and cancers.

²This RadioLab podcast episode goes into more detail on bit flips: <https://www.wnycstudios.org/story/bit-flip>

$E(x)$ is T exactly when $(x)_2$ is even, and is F otherwise

$L(x)$ is T exactly when $(x)_2 < 3$, and is F otherwise

$M(x)$ is T exactly when $(x)_2 > 256$ and is F otherwise.

(a) What is $E(110)$?

(b) Why is $L(00)$ undefined?

- i. Because the domain of L is infinite
- ii. Because 00 does not have 1 in the leftmost position
- iii. Because 00 has length 2, not length 3
- iv. Because $(00)_{2,3} = 0$ which is less than 3

(c) Is there a bitstring of width (where width is the number of bits) 6 at which $M(x)$ evaluates to T ?

3. **Friday** Recall that S is defined as the set of all RNA strands, strings made of the bases in $B = \{\mathbf{A}, \mathbf{U}, \mathbf{G}, \mathbf{C}\}$. Define the functions *mutation*, *insertion*, and *deletion* as described by the pseudocode below:

```

1  procedure mutation( $b_1 \cdots b_n$ : a RNA strand,  $k$ : a positive integer,  $b$ : an element of  $B$ )
2  for  $i := 1$  to  $n$ 
3      if  $i = k$ 
4           $c_i := b$ 
5      else
6           $c_i := b_i$ 
7  return  $c_1 \cdots c_n$  {The return value is a RNA strand made of the  $c_i$  values}

```

```

1  procedure insertion( $b_1 \cdots b_n$ : a RNA strand,  $k$ : a positive integer,  $b$ : an element of  $B$ )
2  if  $k > n$ 
3      for  $i := 1$  to  $n$ 
4           $c_i := b_i$ 
5       $c_{n+1} := b$ 
6  else
7      for  $i := 1$  to  $k - 1$ 
8           $c_i := b_i$ 
9       $c_k := b$ 
10     for  $i := k + 1$  to  $n + 1$ 
11          $c_i := b_{i-1}$ 
12 return  $c_1 \cdots c_{n+1}$  {The return value is a RNA strand made of the  $c_i$  values}

```

```

1  procedure deletion( $b_1 \cdots b_n$ : a RNA strand,  $k$ : a positive integer)
2  if  $k > n$ 
3       $m := n$ 
4      for  $i := 1$  to  $n$ 
5           $c_i := b_i$ 
6  else
7       $m := n - 1$ 
8      for  $i := 1$  to  $k - 1$ 
9           $c_i := b_i$ 
10     for  $i := k$  to  $n - 1$ 
11          $c_i := b_{i+1}$ 
12 return  $c_1 \cdots c_m$  {The return value is a RNA strand made of the  $c_i$  values}

```

For this question, we will use the following predicates.

$F_{\mathbf{A}}$ with domain S is defined recursively by:

Basis step: $F_{\mathbf{A}}(\mathbf{A}) = T$, $F_{\mathbf{A}}(\mathbf{C}) = F_{\mathbf{A}}(\mathbf{G}) = F_{\mathbf{A}}(\mathbf{U}) = F$

Recursive step: If $s \in S$ and $b \in B$, then $F_{\mathbf{A}}(sb) = F_{\mathbf{A}}(s)$

P_{AUC} with domain S is defined as the predicate whose truth set is the collection of RNA strands where the string **AUC** is a substring (appears inside s , in order and consecutively)

L with domain $S \times \mathbb{Z}^+$ is defined by, for $s \in S$ and $n \in \mathbb{Z}^+$,

$$L(s, n) = \begin{cases} T & \text{if } rnalen(s) = n \\ F & \text{otherwise} \end{cases}$$

Mut with domain $S \times S$ is defined by, for $s_1 \in S$ and $s_2 \in S$,

$$Mut(s_1, s_2) = \exists k \in \mathbb{Z}^+ \exists b \in B(\text{mutation}(s_1, k, b) = s_2)$$

Ins with domain $S \times S$ is defined by, for $s_1 \in S$ and $s_2 \in S$,

$$Ins(s_1, s_2) = \exists k \in \mathbb{Z}^+ \exists b \in B(\text{insertion}(s_1, k, b) = s_2)$$

Del with domain $S \times S$ is defined by, for $s_1 \in S$ and $s_2 \in S$,

$$Del(s_1, s_2) = \exists k \in \mathbb{Z}^+ (\text{deletion}(s_1, k) = s_2)$$

(a) Which of the following is true? (Select all and only that apply.)

- i. $F_A(\mathbf{AA})$
- ii. $F_A(\mathbf{AC})$
- iii. $F_A(\mathbf{AG})$
- iv. $F_A(\mathbf{AU})$
- v. $F_A(\mathbf{CA})$
- vi. $F_A(\mathbf{CC})$
- vii. $F_A(\mathbf{CG})$
- viii. $F_A(\mathbf{CU})$

(b) Which of the following is true? (Select all and only that apply.)

- i. $\exists s \in S \exists n \in \mathbb{Z}^+ (L(s, n))$
- ii. $\exists s \in S \forall n \in \mathbb{Z}^+ (L(s, n))$
- iii. $\forall n \in \mathbb{Z}^+ \exists s \in S (L(s, n))$
- iv. $\forall s \in S \exists n \in \mathbb{Z}^+ (L(s, n))$
- v. $\exists n \in \mathbb{Z}^+ \forall s \in S (L(s, n))$
- vi. $\forall s \in S \forall n \in \mathbb{Z}^+ (L(s, n))$

(c) Which of the following is true? (Select all and only that apply.)

- i. $\exists s \in S Mut(s, s)$

- ii. $\forall s \in S \text{ } Mut(s, s)$
- iii. $\exists s \in S \text{ } Ins(s, \mathbf{A})$
- iv. $\exists s \in S \text{ } Ins(\mathbf{A}, s)$
- v. $\exists s \in S \text{ } Del(s, \mathbf{A})$
- vi. $\forall s \in S \text{ } Del(s, \mathbf{A})$