# Programming Language—Common Lisp

17. Sequences

# 17.1 Sequence Concepts

A sequence is an ordered collection of elements, implemented as either a vector or a list.

Sequences can be created by the function make-sequence, as well as other functions that create objects of types that are subtypes of sequence (e.g., list, make-list, mapcar, and vector).

A **sequence function** is a function defined by this specification or added as an extension by the *implementation* that operates on one or more *sequences*. Whenever a *sequence function* must construct and return a new *vector*, it always returns a *simple vector*. Similarly, any *strings* constructed will be *simple strings*.

	1 /1	
concatenate	length	remove
copy-seq	map	${f remove-duplicates}$
count	map-into	remove-if
count-if	merge	${f remove-if-not}$
count-if-not	${f mismatch}$	replace
delete	notany	reverse
delete-duplicates	$\mathbf{notevery}$	search
delete-if	nreverse	some
delete-if-not	${f nsubstitute}$	$\operatorname{sort}$
elt	${f nsubstitute} ext{-if}$	${f stable} ext{-sort}$
every	${f nsubstitute} ext{-if-not}$	${f subseq}$
fill	position	${f substitute}$
find	position-if	${f substitute-if}$
find-if	position-if-not	${f substitute-if-not}$
find-if-not	reduce	

Figure 17-1. Standardized Sequence Functions

# 17.1.1 General Restrictions on Parameters that must be Sequences

In general, lists (including  $association\ lists$  and  $property\ lists$ ) that are treated as sequences must be  $proper\ lists$ .

## 17.2 Rules about Test Functions

## 17.2.1 Satisfying a Two-Argument Test

When an object O is being considered iteratively against each element  $E_i$  of a sequence S by an operator F listed in Figure 17–2, it is sometimes useful to control the way in which the presence of O is tested in S is tested by F. This control is offered on the basis of a function designated with either a :test-not argument.

adjoin	nset-exclusive-or	search
assoc	nsublis	set-difference
$\operatorname{\mathbf{count}}$	$\mathbf{nsubst}$	${f set} ext{-exclusive-or}$
$\mathbf{delete}$	${f nsubstitute}$	$\operatorname{sublis}$
$\operatorname{find}$	nunion	$\operatorname{\mathbf{subsetp}}$
intersection	position	$\operatorname{subst}$
member	pushnew	substitute
${f mismatch}$	rassoc	tree-equal
nintersection	remove	union
nset-difference	remove-duplicates	

Figure 17-2. Operators that have Two-Argument Tests to be Satisfied

The object O might not be compared directly to  $E_i$ . If a :key argument is provided, it is a designator for a function of one argument to be called with each  $E_i$  as an argument, and yielding an object  $Z_i$  to be used for comparison. (If there is no :key argument,  $Z_i$  is  $E_i$ .)

The function designated by the :key argument is never called on O itself. However, if the function operates on multiple sequences (e.g., as happens in **set-difference**), O will be the result of calling the :key function on an element of the other sequence.

A :test argument, if supplied to F, is a designator for a function of two arguments, O and  $Z_i$ . An  $E_i$  is said (or, sometimes, an O and an  $E_i$  are said) to satisfy the test if this :test function returns a generalized boolean representing true.

A :test-not argument, if supplied to F, is designator for a function of two arguments, O and  $Z_i$ . An  $E_i$  is said (or, sometimes, an O and an  $E_i$  are said) to satisfy the test if this :test-not function returns a generalized boolean representing false.

If neither a :test nor a :test-not argument is supplied, it is as if a :test argument of #'eql was supplied.

The consequences are unspecified if both a :test and a :test-not argument are supplied in the same call to F.

## 17.2.1.1 Examples of Satisfying a Two-Argument Test

```
(remove "F00" '(foo bar "F00" "BAR" "foo" "bar") :test #'equal)
\rightarrow (foo bar "BAR" "foo" "bar")
(remove "F00" '(foo bar "F00" "BAR" "foo" "bar") :test #'equalp)

ightarrow (foo bar "BAR" "bar")
(remove "F00" '(foo bar "F00" "BAR" "foo" "bar") :test #'string-equal)
\rightarrow (bar "BAR" "bar")
(remove "F00" '(foo bar "F00" "BAR" "foo" "bar") :test #'string=)

ightarrow (BAR "BAR" "foo" "bar")
 (remove 1 '(1 1.0 #C(1.0 0.0) 2 2.0 #C(2.0 0.0)) :test-not #'eql)
 (remove 1 '(1 1.0 #C(1.0 0.0) 2 2.0 #C(2.0 0.0)) :test-not #'=)
\rightarrow (1 1.0 #C(1.0 0.0))
 (remove 1 '(1 1.0 #C(1.0 0.0) 2 2.0 #C(2.0 0.0)) :test (complement #'=))
\rightarrow (1 1.0 #C(1.0 0.0))
 (count 1 '((one 1) (uno 1) (two 2) (dos 2)) :key #'cadr) 
ightarrow 2
 (count 2.0 '(1 2 3) :test #'eql :key #'float) 
ightarrow 1
 (count "F00" (list (make-pathname :name "F00" :type "X")
                      (make-pathname :name "FOO" :type "Y"))
        :key #'pathname-name
        :test #'equal)
\rightarrow 2
```

## 17.2.2 Satisfying a One-Argument Test

When using one of the functions in Figure 17–3, the elements E of a sequence S are filtered not on the basis of the presence or absence of an object O under a two argument predicate, as with the functions described in Section 17.2.1 (Satisfying a Two-Argument Test), but rather on the basis of a one argument predicate.

assoc-if	member-if	rassoc-if
assoc-if-not	${f member-if-not}$	rassoc-if-not
count-if	nsubst-if	remove-if
count-if-not	nsubst-if-not	${f remove-if-not}$
delete-if	${f nsubstitute-if}$	$\operatorname{subst-if}$
delete-if-not	${f nsubstitute}$ -if-not	$\mathbf{subst} ext{-}\mathbf{if} ext{-}\mathbf{not}$
find-if	position-if	${f substitute}$ -if
find-if-not	position-if-not	${f substitute}$ -if-not

Figure 17-3. Operators that have One-Argument Tests to be Satisfied

The element  $E_i$  might not be considered directly. If a :key argument is provided, it is a designator for a function of one argument to be called with each  $E_i$  as an argument, and yielding an object  $Z_i$  to be used for comparison. (If there is no :key argument,  $Z_i$  is  $Z_i$ .)

Functions defined in this specification and having a name that ends in "-if" accept a first argument that is a designator for a function of one argument,  $Z_i$ . An  $E_i$  is said to satisfy the test if this :test function returns a generalized boolean representing true.

Functions defined in this specification and having a name that ends in "-if-not" accept a first argument that is a designator for a function of one argument,  $Z_i$ . An  $E_i$  is said to satisfy the test if this :test function returns a generalized boolean representing false.

## 17.2.2.1 Examples of Satisfying a One-Argument Test

```
(count-if #'zerop '(1 #C(0.0 0.0) 0 0.0d0 0.0s0 3)) → 4

(remove-if-not #'symbolp '(0 1 2 3 4 5 6 7 8 9 A B C D E F))

→ (A B C D E F)
(remove-if (complement #'symbolp) '(0 1 2 3 4 5 6 7 8 9 A B C D E F))

→ (A B C D E F)

(count-if #'zerop '("foo" "" "bar" "" "baz" "quux") :key #'length)

→ 3
```

**sequence** System Class

## Class Precedence List:

sequence, t

#### **Description:**

Sequences are ordered collections of objects, called the elements of the sequence.

The types vector and the type list are disjoint subtypes of type sequence, but are not necessarily an exhaustive partition of sequence.

When viewing a vector as a sequence, only the active elements of that vector are considered elements of the sequence; that is, sequence operations respect the fill pointer when given sequences represented as vectors.

copy-seq Function

## **Syntax:**

copy-seq sequence o copied-sequence

## **Arguments and Values:**

 ${\it sequence} {\rm --a} \ proper \ sequence.$ 

copied-sequence—a proper sequence.

#### **Description:**

Creates a copy of *sequence*. The *elements* of the new *sequence* are the *same* as the corresponding *elements* of the given *sequence*.

If sequence is a vector, the result is a fresh simple array of rank one that has the same actual array element type as sequence. If sequence is a list, the result is a fresh list.

#### **Examples:**

```
(setq str "a string") \to "a string" (equalp str (copy-seq str)) \to true (eql str (copy-seq str)) \to false
```

## **Exceptional Situations:**

Should be prepared to signal an error of type type-error if sequence is not a proper sequence.

#### See Also:

copy-list

#### Notes:

From a functional standpoint,  $(copy-seq x) \equiv (subseq x 0)$ 

However, the programmer intent is typically very different in these two cases.

elt

## Syntax:

```
elt sequence index \rightarrow object (setf (elt sequence index) new-object)
```

## **Arguments and Values:**

```
sequence—a proper sequence.
index—a valid sequence index for sequence.
object—an object.
new-object—an object.
```

## **Description:**

Accesses the element of sequence specified by index.

#### **Examples:**

```
(setq str (copy-seq "0123456789")) \to "0123456789" (elt str 6) \to #\6 (setf (elt str 0) #\#) \to #\# str \to "#123456789"
```

## **Exceptional Situations:**

Should be prepared to signal an error of *type* **type-error** if *sequence* is not a *proper sequence*. Should signal an error of *type* **type-error** if *index* is not a *valid sequence index* for *sequence*.

## See Also:

```
aref, nth, Section 3.2.1 (Compiler Terminology)
```

## Notes:

aref may be used to access vector elements that are beyond the vector's fill pointer.

**fill** Function

## Syntax:

fill sequence item &key start end  $\rightarrow$  sequence

## **Arguments and Values:**

```
sequence—a proper sequence.
```

item—a sequence.

start, end—bounding index designators of sequence. The defaults for start and end are 0 and nil, respectively.

## **Description:**

Replaces the *elements* of *sequence bounded* by *start* and *end* with *item*.

#### **Examples:**

```
(fill (list 0 1 2 3 4 5) '(444)) \rightarrow ((444) (444) (444) (444) (444) (444) (444) (fill (copy-seq "01234") #\e :start 3) \rightarrow "012ee" (setq x (vector 'a 'b 'c 'd 'e)) \rightarrow #(A B C D E) (fill x 'z :start 1 :end 3) \rightarrow #(A Z Z D E) \times \rightarrow #(A Z Z D E) (fill x 'p) \rightarrow #(P P P P P) \times \rightarrow #(P P P P P)
```

## Side Effects:

Sequence is destructively modified.

#### **Exceptional Situations:**

Should be prepared to signal an error of *type* **type-error** if *sequence* is not a *proper sequence*. Should signal an error of *type* **type-error** if *start* is not a non-negative *integer*. Should signal an error of *type* **type-error** if *end* is not a non-negative *integer* or **nil**.

#### See Also:

replace, nsubstitute

## Notes:

(fill sequence item) ≡(nsubstitute-if item (constantly t) sequence)

# make-sequence

# make-sequence

Function

## **Syntax:**

make-sequence result-type size &key initial-element  $\rightarrow$  sequence

## **Arguments and Values:**

```
result-type—a sequence type specifier.

size—a non-negative integer.

initial-element—an object. The default is implementation-dependent.

sequence—a proper sequence.
```

## **Description:**

Returns a sequence of the type result-type and of length size, each of the elements of which has been initialized to initial-element.

If the *result-type* is a *subtype* of **list**, the result will be a *list*.

If the *result-type* is a *subtype* of **vector**, then if the implementation can determine the element type specified for the *result-type*, the element type of the resulting array is the result of *upgrading* that element type; or, if the implementation can determine that the element type is unspecified (or \*), the element type of the resulting array is t; otherwise, an error is signaled.

## **Examples:**

## Affected By:

The implementation.

## **Exceptional Situations:**

The consequences are unspecified if *initial-element* is not an *object* which can be stored in the resulting *sequence*.

17–8 Programming Language—Common Lisp

An error of type type-error must be signaled if the result-type is neither a recognizable subtype of list, nor a recognizable subtype of vector.

An error of type type-error should be signaled if result-type specifies the number of elements and size is different from that number.

#### See Also:

make-array, make-list

#### **Notes:**

(make-sequence 'string 5) ≡ (make-string 5)

subseq

Accessor

## Syntax:

```
subseq sequence start &optional end \rightarrow subsequence (setf (subseq sequence start &optional end) new-subsequence)
```

## **Arguments and Values:**

```
sequence—a proper sequence.
```

start, end—bounding index designators of sequence. The default for end is nil.

subsequence—a proper sequence.

new-subsequence—a proper sequence.

#### Description:

**subseq** creates a *sequence* that is a copy of the subsequence of *sequence bounded* by *start* and *end*.

*Start* specifies an offset into the original *sequence* and marks the beginning position of the subsequence. *end* marks the position following the last element of the subsequence.

subseq always allocates a new *sequence* for a result; it never shares storage with an old *sequence*. The result subsequence is always of the same *type* as *sequence*.

If sequence is a vector, the result is a fresh simple array of rank one that has the same actual array element type as sequence. If sequence is a list, the result is a fresh list.

setf may be used with subseq to destructively replace elements of a subsequence with elements taken from a sequence of new values. If the subsequence and the new sequence are not of equal length, the shorter length determines the number of elements that are replaced. The remaining elements at the end of the longer sequence are not modified in the operation.

## **Examples:**

```
(setq str "012345") \rightarrow "012345" (subseq str 2) \rightarrow "2345" (subseq str 3 5) \rightarrow "34" (setf (subseq str 4) "abc") \rightarrow "abc" str \rightarrow "0123ab" (setf (subseq str 0 2) "A") \rightarrow "A" str \rightarrow "A123ab"
```

## **Exceptional Situations:**

Should be prepared to signal an error of type **type-error** if sequence is not a proper sequence. Should be prepared to signal an error of type **type-error** if new-subsequence is not a proper sequence.

#### See Also:

replace

map Function

## Syntax:

map result-type function &rest sequences $^+$  o result

#### Arguments and Values:

```
result-type – a sequence type specifier, or nil.
```

function—a function designator. function must take as many arguments as there are sequences. sequence—a proper sequence.

result—if result-type is a type specifier other than nil, then a sequence of the type it denotes; otherwise (if the result-type is nil), nil.

#### Description:

Applies function to successive sets of arguments in which one argument is obtained from each sequence. The function is called first on all the elements with index 0, then on all those with index 1, and so on. The result-type specifies the type of the resulting sequence.

map returns nil if *result-type* is nil. Otherwise, map returns a *sequence* such that element j is the result of applying *function* to element j of each of the *sequences*. The result *sequence* is as long as the shortest of the *sequences*. The consequences are undefined if the result of applying *function* to the successive elements of the *sequences* cannot be contained in a *sequence* of the *type* given by *result-type*.

If the result-type is a subtype of list, the result will be a list.

If the *result-type* is a *subtype* of **vector**, then if the implementation can determine the element type specified for the *result-type*, the element type of the resulting array is the result of *upgrading* that element type; or, if the implementation can determine that the element type is unspecified (or \*), the element type of the resulting array is t; otherwise, an error is signaled.

## **Examples:**

#### **Exceptional Situations:**

An error of type **type-error** must be signaled if the *result-type* is not a *recognizable subtype* of **list**, not a *recognizable subtype* of **vector**, and not **nil**.

Should be prepared to signal an error of type type-error if any sequence is not a proper sequence.

An error of *type* **type-error** should be signaled if *result-type* specifies the number of elements and the minimum length of the *sequences* is different from that number.

#### See Also:

Section 3.6 (Traversal Rules and Side Effects)

# map-into

map-into Function

#### **Syntax:**

 $ext{map-into}$  result-sequence function &rest sequences o result-sequence

#### **Arguments and Values:**

result-sequence—a proper sequence.

function—a designator for a function of as many arguments as there are sequences.

sequence—a proper sequence.

## **Description:**

Destructively modifies *result-sequence* to contain the results of applying *function* to each element in the argument *sequences* in turn.

result-sequence and each element of sequences can each be either a list or a vector. If result-sequence and each element of sequences are not all the same length, the iteration terminates when the shortest sequence (of any of the sequences or the result-sequence) is exhausted. If result-sequence is a vector with a fill pointer, the fill pointer is ignored when deciding how many iterations to perform, and afterwards the fill pointer is set to the number of times function was applied. If result-sequence is longer than the shortest element of sequences, extra elements at the end of result-sequence are left unchanged. If result-sequence is nil, map-into immediately returns nil, since nil is a sequence of length zero.

If *function* has side effects, it can count on being called first on all of the elements with index 0, then on all of those numbered 1, and so on.

#### **Examples:**

```
(setq a (list 1 2 3 4) b (list 10 10 10 10)) \rightarrow (10 10 10 10) (map-into a #'+ a b) \rightarrow (11 12 13 14) a \rightarrow (11 12 13 14) b \rightarrow (10 10 10 10) (setq k '(one two three)) \rightarrow (ONE TWO THREE) (map-into a #'cons k a) \rightarrow ((ONE . 11) (TWO . 12) (THREE . 13) 14) (map-into a #'gensym) \rightarrow (#:G9090 #:G9091 #:G9092 #:G9093) a \rightarrow (#:G9090 #:G9091 #:G9092 #:G9093)
```

#### **Exceptional Situations:**

Should be prepared to signal an error of type **type-error** if result-sequence is not a proper sequence. Should be prepared to signal an error of type **type-error** if sequence is not a proper sequence.

#### **Notes:**

map-into differs from map in that it modifies an existing sequence rather than creating a new

#### 17–12 Programming Language—Common Lisp

one. In addition,  $\mathbf{map\text{-}into}$  can be called with only two arguments, while  $\mathbf{map}$  requires at least three arguments.

map-into could be defined by:

reduce Function

## Syntax:

 ${f reduce}$  function sequence &key key from-end start end initial-value ightarrow result

## **Arguments and Values:**

function—a designator for a function that might be called with either zero or two arguments.

sequence—a proper sequence.

key—a designator for a function of one argument, or nil.

from-end—a generalized boolean. The default is false.

start, end—bounding index designators of sequence. The defaults for start and end are 0 and nil, respectively.

initial-value—an object.

result—an object.

#### **Description:**

**reduce** uses a binary operation, function, to combine the elements of sequence bounded by start and end.

The *function* must accept as *arguments* two *elements* of *sequence* or the results from combining those *elements*. The *function* must also be able to accept no arguments.

If key is supplied, it is used is used to extract the values to reduce. The key function is applied exactly once to each element of sequence in the order implied by the reduction order but not to

the value of *initial-value*, if supplied. The *key* function typically returns part of the *element* of *sequence*. If *key* is not supplied or is **nil**, the *sequence element* itself is used.

The reduction is left-associative, unless from-end is true in which case it is right-associative.

If initial-value is supplied, it is logically placed before the subsequence (or after it if from-end is true) and included in the reduction operation.

In the normal case, the result of **reduce** is the combined result of *function*'s being applied to successive pairs of *elements* of *sequence*. If the subsequence contains exactly one *element* and no *initial-value* is given, then that *element* is returned and *function* is not called. If the subsequence is empty and an *initial-value* is given, then the *initial-value* is returned and *function* is not called. If the subsequence is empty and no *initial-value* is given, then the *function* is called with zero arguments, and **reduce** returns whatever *function* does. This is the only case where the *function* is called with other than two arguments.

## **Examples:**

```
(reduce #'* '(1 2 3 4 5)) 
ightarrow 120
 (reduce #'append '((1) (2)) :initial-value '(i n i t)) \rightarrow (I N I T 1 2)
 (reduce \#'append '((1) (2)) :from-end t
                                  :initial-value '(i n i t)) 
ightarrow (1 2 I N I T)
 (reduce #'- '(1 2 3 4)) \equiv (- (- (- 1 2) 3) 4) \rightarrow -8
 (reduce #'- '(1 2 3 4) :from-end t)
                                              ;Alternating sum.
\equiv (- 1 (- 2 (- 3 4))) 
ightarrow -2
 (reduce #'+ '()) \rightarrow 0
 (reduce #'+ '(3)) 
ightarrow 3
 (reduce #'+ '(foo)) 
ightarrow FOO
 (reduce #'list '(1 2 3 4)) \rightarrow (((1 2) 3) 4)
 (reduce #'list '(1 2 3 4) :from-end t) 
ightarrow (1 (2 (3 4)))
 (reduce #'list '(1 2 3 4) :initial-value 'foo) \rightarrow (((foo 1) 2) 3) 4)
 (reduce #'list '(1 2 3 4)
         :from-end t :initial-value 'foo) \rightarrow (1 (2 (3 (4 foo))))
```

## **Exceptional Situations:**

Should be prepared to signal an error of type type-error if sequence is not a proper sequence.

#### See Also:

Section 3.6 (Traversal Rules and Side Effects)

# count, count-if, count-if-not

# count, count-if, count-if-not

*Function* 

## Syntax:

```
count item sequence &key from-end start end key test test-not \rightarrow n count-if predicate sequence &key from-end start end key \rightarrow n count-if-not predicate sequence &key from-end start end key \rightarrow n
```

## **Arguments and Values:**

```
item—an object.
```

sequence—a proper sequence.

predicate—a designator for a function of one argument that returns a generalized boolean.

from-end—a generalized boolean. The default is false.

test—a designator for a function of two arguments that returns a generalized boolean.

test-not—a designator for a function of two arguments that returns a generalized boolean.

start, end—bounding index designators of sequence. The defaults for start and end are 0 and nil, respectively.

key—a designator for a function of one argument, or nil.

n—a non-negative integer less than or equal to the length of sequence.

## **Description:**

**count**, **count-if**, and **count-if-not** count and return the number of *elements* in the **sequence** bounded by **start** and **end** that satisfy the test.

The from-end has no direct effect on the result. However, if from-end is true, the elements of sequence will be supplied as arguments to the test, test-not, and key in reverse order, which may change the side-effects, if any, of those functions.

## **Examples:**

```
(count #\a "how many A's are there in here?") \to 2 (count-if-not #'oddp '((1) (2) (3) (4)) :key #'car) \to 2 (count-if #'upper-case-p "The Crying of Lot 49" :start 4) \to 2
```

#### **Exceptional Situations:**

Should be prepared to signal an error of type type-error if sequence is not a proper sequence.

#### See Also:

Section 17.2 (Rules about Test Functions), Section 3.6 (Traversal Rules and Side Effects)

#### Notes:

The :test-not argument is deprecated.

The function **count-if-not** is deprecated.

length

## Syntax:

```
length sequence \rightarrow n
```

## **Arguments and Values:**

```
{\it sequence} {\rm --a} \ proper \ sequence.
```

n—a non-negative integer.

## **Description:**

Returns the number of elements in sequence.

If sequence is a vector with a fill pointer, the active length as specified by the fill pointer is returned.

## **Examples:**

```
(length "abc") \to 3
(setq str (make-array '(3) :element-type 'character :initial-contents "abc" :fill-pointer t)) \to "abc" (length str) \to 3
(setf (fill-pointer str) 2) \to 2
(length str) \to 2
```

#### **Exceptional Situations:**

Should be prepared to signal an error of type type-error if sequence is not a proper sequence.

## See Also:

list-length, sequence

## reverse, nreverse

## reverse, nreverse

Function

## Syntax:

```
reverse sequence \rightarrow reversed-sequence nreverse sequence \rightarrow reversed-sequence
```

## **Arguments and Values:**

```
sequence—a proper sequence.
reversed-sequence—a sequence.
```

## **Description:**

reverse and nreverse return a new sequence of the same kind as sequence, containing the same elements, but in reverse order.

reverse and nreverse differ in that reverse always creates and returns a new sequence, whereas nreverse might modify and return the given sequence. reverse never modifies the given sequence.

For **reverse**, if **sequence** is a **vector**, the result is a **fresh simple array** of **rank** one that has the same **actual array element type** as **sequence**. If **sequence** is a **list**, the result is a **fresh list**.

For **nreverse**, if **sequence** is a **vector**, the result is a **vector** that has the same **actual array element type** as **sequence**. If **sequence** is a **list**, the result is a **list**.

For **nreverse**, **sequence** might be destroyed and re-used to produce the result. The result might or might not be *identical* to **sequence**. Specifically, when **sequence** is a **list**, **nreverse** is permitted to **setf** any part, **car** or **cdr**, of any **cons** that is part of the **list structure** of **sequence**. When **sequence** is a **vector**, **nreverse** is permitted to re-order the elements of **sequence** in order to produce the resulting **vector**.

## **Examples:**

```
(setq str "abc") \rightarrow "abc"

(reverse str) \rightarrow "cba"

str \rightarrow "abc"

(setq str (copy-seq str)) \rightarrow "abc"

(nreverse str) \rightarrow "cba"

str \rightarrow implementation-dependent

(setq 1 (list 1 2 3)) \rightarrow (1 2 3)

(nreverse 1) \rightarrow (3 2 1)

1 \rightarrow implementation-dependent
```

## **Side Effects:**

 ${\bf nreverse}$  might either create a new  ${\it sequence},$  modify the argument  ${\it sequence},$  or both. (reverse does not modify  ${\it sequence}.$ )

## **Exceptional Situations:**

Should be prepared to signal an error of type type-error if sequence is not a proper sequence.

# sort, stable-sort

Function

## Syntax:

sort sequence predicate &key  $key \rightarrow sorted$ -sequence stable-sort sequence predicate &key  $key \rightarrow sorted$ -sequence

## **Arguments and Values:**

sequence—a proper sequence.

predicate—a designator for a function of two arguments that returns a generalized boolean.

key—a designator for a function of one argument, or nil.

*sorted-sequence*—a *sequence*.

## **Description:**

sort and stable-sort destructively sort sequences according to the order determined by the predicate function.

If sequence is a vector, the result is a vector that has the same actual array element type as sequence. If sequence is a list, the result is a list.

sort determines the relationship between two elements by giving keys extracted from the elements to the *predicate*. The first argument to the *predicate* function is the part of one element of *sequence* extracted by the *key* function (if supplied); the second argument is the part of another element of *sequence* extracted by the *key* function (if supplied). *Predicate* should return *true* if and only if the first argument is strictly less than the second (in some appropriate sense). If the first argument is greater than or equal to the second (in the appropriate sense), then the *predicate* should return *false*.

The argument to the *key* function is the *sequence* element. The return value of the *key* function becomes an argument to *predicate*. If *key* is not supplied or **nil**, the *sequence* element itself is used. There is no guarantee on the number of times the *key* will be called.

If the key and predicate always return, then the sorting operation will always terminate, producing a sequence containing the same elements as sequence (that is, the result is a permutation of sequence). This is guaranteed even if the predicate does not really consistently represent a total order (in which case the elements will be scrambled in some unpredictable way, but no element will be lost). If the key consistently returns meaningful keys, and the predicate does reflect some total ordering criterion on those keys, then the elements of the sorted-sequence will be properly sorted according to that ordering.

# sort, stable-sort

The sorting operation performed by **sort** is not guaranteed stable. Elements considered equal by the *predicate* might or might not stay in their original order. The *predicate* is assumed to consider two elements x and y to be equal if (funcall *predicate* x y) and (funcall *predicate* y x) are both false. **stable-sort** guarantees stability.

The sorting operation can be destructive in all cases. In the case of a *vector* argument, this is accomplished by permuting the elements in place. In the case of a *list*, the *list* is destructively reordered in the same manner as for **nreverse**.

## Examples:

```
(setq tester (copy-seq "lkjashd")) \rightarrow "lkjashd"
 (sort tester #'char-lessp) \rightarrow "adhjkls"
 (\text{setq tester (list '(1 2 3) '(4 5 6) '(7 8 9))}) \rightarrow ((1 2 3) (4 5 6) (7 8 9))
 (sort tester \#'> :key \#'car) \rightarrow ((7 8 9) (4 5 6) (1 2 3))
 (setq tester (list 1 2 3 4 5 6 7 8 9 0)) \rightarrow (1 2 3 4 5 6 7 8 9 0)
 (stable-sort tester #'(lambda (x y) (and (oddp x) (evenp y))))
\rightarrow (1 3 5 7 9 2 4 6 8 0)
 (sort (setq committee-data
              (vector (list (list "JonL" "White") "Iteration")
                      (list (list "Dick" "Waters") "Iteration")
                      (list (list "Dick" "Gabriel") "Objects")
                      (list (list "Kent" "Pitman") "Conditions")
                      (list (list "Gregor" "Kiczales") "Objects")
                      (list (list "David" "Moon") "Objects")
                      (list (list "Kathy" "Chapman") "Editorial")
                      (list (list "Larry" "Masinter") "Cleanup")
                      (list (list "Sandra" "Loosemore") "Compiler")))
       #'string-lessp :key #'cadar)

ightarrow #((("Kathy" "Chapman") "Editorial")
     (("Dick" "Gabriel") "Objects")
     (("Gregor" "Kiczales") "Objects")
     (("Sandra" "Loosemore") "Compiler")
     (("Larry" "Masinter") "Cleanup")
     (("David" "Moon") "Objects")
     (("Kent" "Pitman") "Conditions")
     (("Dick" "Waters") "Iteration")
     (("JonL" "White") "Iteration"))
 ;; Note that individual alphabetical order within 'committees'
 ;; is preserved.
 (setq committee-data
       (stable-sort committee-data #'string-lessp :key #'cadr))
\rightarrow #((("Larry" "Masinter") "Cleanup")
     (("Sandra" "Loosemore") "Compiler")
     (("Kent" "Pitman") "Conditions")
     (("Kathy" "Chapman") "Editorial")
```

```
(("Dick" "Waters") "Iteration")
(("JonL" "White") "Iteration")
(("Dick" "Gabriel") "Objects")
(("Gregor" "Kiczales") "Objects")
(("David" "Moon") "Objects"))
```

#### **Exceptional Situations:**

Should be prepared to signal an error of type type-error if sequence is not a proper sequence.

#### See Also:

merge, Section 3.2.1 (Compiler Terminology), Section 3.6 (Traversal Rules and Side Effects), Section 3.7 (Destructive Operations)

#### Notes:

If **sequence** is a **vector**, the result might or might not be simple, and might or might not be **identical** to **sequence**.

# find, find-if, find-if-not

**Function** 

## Syntax:

find item sequence &key from-end test test-not start end key ightarrow element

 $ext{find-if}$  predicate sequence &key from-end start end key o element

 $\mathbf{find\text{-}if\text{-}not} \ \textit{predicate sequence} \ \&\texttt{key} \ \textit{from\text{-}end start} \ \textit{end key} \quad \rightarrow \ \textit{element}$ 

## **Arguments and Values:**

```
item—an object.
```

sequence—a proper sequence.

predicate—a designator for a function of one argument that returns a generalized boolean.

from-end—a generalized boolean. The default is false.

test—a designator for a function of two arguments that returns a generalized boolean.

test-not—a designator for a function of two arguments that returns a generalized boolean.

start, end—bounding index designators of sequence. The defaults for start and end are 0 and nil, respectively.

key—a designator for a function of one argument, or nil.

element—an element of the sequence, or nil.

## **Description:**

find, find-if, and find-if-not each search for an element of the sequence bounded by start and end that satisfies the predicate predicate or that satisfies the test test or test-not, as appropriate.

If from-end is true, then the result is the rightmost element that satisfies the test.

If the sequence contains an element that satisfies the test, then the leftmost or rightmost sequence element, depending on from-end, is returned; otherwise nil is returned.

## **Examples:**

```
(find #\d "here are some letters that can be looked at" :test #'char>) \rightarrow #\Space (find-if #'oddp '(1 2 3 4 5) :end 3 :from-end t) \rightarrow 3 (find-if-not #'complexp '#(3.5 2 #C(1.0 0.0) #C(0.0 1.0)) :start 2) \rightarrow NIL
```

## **Exceptional Situations:**

Should be prepared to signal an error of type type-error if sequence is not a proper sequence.

#### See Also:

**position**, Section 17.2 (Rules about Test Functions), Section 3.6 (Traversal Rules and Side Effects)

#### Notes:

The :test-not argument is deprecated.

The function find-if-not is deprecated.

# position, position-if, position-if-not

Function

#### Syntax:

```
position item sequence &key from-end test test-not start end key \rightarrow position position-if predicate sequence &key from-end start end key \rightarrow position position-if-not predicate sequence &key from-end start end key \rightarrow position
```

## **Arguments and Values:**

```
item—an object.
sequence—a proper sequence.
```

predicate—a designator for a function of one argument that returns a generalized boolean.

from-end—a generalized boolean. The default is false.

test—a designator for a function of two arguments that returns a generalized boolean.

test-not—a designator for a function of two arguments that returns a generalized boolean.

start, end—bounding index designators of sequence. The defaults for start and end are 0 and nil, respectively.

key—a designator for a function of one argument, or nil.

position—a bounding index of sequence, or nil.

#### **Description:**

position, position-if, and position-if-not each search sequence for an element that satisfies the test.

The position returned is the index within sequence of the leftmost (if from-end is true) or of the rightmost (if from-end is false) element that satisfies the test; otherwise nil is returned. The index returned is relative to the left-hand end of the entire sequence, regardless of the value of start, end, or from-end.

## **Examples:**

```
(position #\a "baobab" :from-end t) \rightarrow 4 (position-if #'oddp '((1) (2) (3) (4)) :start 1 :key #'car) \rightarrow 2 (position 595 '()) \rightarrow NIL (position-if-not #'integerp '(1 2 3 4 5.0)) \rightarrow 4
```

#### **Exceptional Situations:**

Should be prepared to signal an error of type type-error if sequence is not a proper sequence.

#### See Also:

find, Section 3.6 (Traversal Rules and Side Effects)

#### Notes:

The :test-not argument is deprecated.

The function position-if-not is deprecated.

## search

search

## Syntax:

search sequence-1 sequence-2 &key from-end test test-not key start1 start2 end1 end2

 $\rightarrow$  position

## **Arguments and Values:**

Sequence-1—a sequence.

Sequence-2—a sequence.

from-end—a generalized boolean. The default is false.

test—a designator for a function of two arguments that returns a generalized boolean.

test-not—a designator for a function of two arguments that returns a generalized boolean.

key—a designator for a function of one argument, or nil.

start1, end1—bounding index designators of sequence-1. The defaults for start1 and end1 are 0 and nil, respectively.

start2, end2—bounding index designators of sequence-2. The defaults for start2 and end2 are 0 and nil, respectively.

position—a bounding index of sequence-2, or nil.

#### **Description:**

Searches sequence-2 for a subsequence that matches sequence-1.

The implementation may choose to search <code>sequence-2</code> in any order; there is no guarantee on the number of times the test is made. For example, when <code>start-end</code> is <code>true</code>, the <code>sequence</code> might actually be searched from left to right instead of from right to left (but in either case would return the rightmost matching subsequence). If the search succeeds, <code>search</code> returns the offset into <code>sequence-2</code> of the first element of the leftmost or rightmost matching subsequence, depending on <code>from-end</code>; otherwise <code>search</code> returns <code>nil</code>.

If from-end is true, the index of the leftmost element of the rightmost matching subsequence is returned.

#### **Examples:**

```
(search "dog" "it's a dog's life") 
ightarrow 7 (search '(0 1) '(2 4 6 1 3 5) :key #'oddp) 
ightarrow 2
```

#### See Also:

Section 3.6 (Traversal Rules and Side Effects)

#### Notes:

The :test-not argument is deprecated.

mismatch Function

## Syntax:

 $\begin{array}{l} \mathbf{mismatch} \ \ \textit{sequence-1} \ \ \textit{sequence-2} \ \& \texttt{key} \ \ \textit{from-end test test-not key start1} \ \ \textit{start2} \ \ \textit{end1} \ \ \textit{end2} \\ \rightarrow \ \ \textit{position} \end{array}$ 

## **Arguments and Values:**

Sequence-1—a sequence.

Sequence-2—a sequence.

from-end—a generalized boolean. The default is false.

test—a designator for a function of two arguments that returns a generalized boolean.

test-not—a designator for a function of two arguments that returns a generalized boolean.

start1, end1—bounding index designators of sequence-1. The defaults for start1 and end1 are 0 and nil, respectively.

start2, end2—bounding index designators of sequence-2. The defaults for start2 and end2 are 0 and nil, respectively.

key—a designator for a function of one argument, or nil.

position—a bounding index of sequence-1, or nil.

#### **Description:**

The specified subsequences of *sequence-1* and *sequence-2* are compared element-wise.

The key argument is used for both the sequence-1 and the sequence-2.

If sequence-1 and sequence-2 are of equal length and match in every element, the result is false. Otherwise, the result is a non-negative integer, the index within sequence-1 of the leftmost or rightmost position, depending on from-end, at which the two subsequences fail to match. If one subsequence is shorter than and a matching prefix of the other, the result is the index relative to sequence-1 beyond the last position tested.

If *from-end* is *true*, then one plus the index of the rightmost position in which the *sequences* differ is returned. In effect, the subsequences are aligned at their right-hand ends; then, the last elements are compared, the penultimate elements, and so on. The index returned is an index relative to *sequence-1*.

## **Examples:**

```
(mismatch "abcd" "ABCDE" :test #'char-equal) \rightarrow 4 (mismatch '(3 2 1 1 2 3) '(1 2 3) :from-end t) \rightarrow 3 (mismatch '(1 2 3) '(2 3 4) :test-not #'eq :key #'oddp) \rightarrow NIL (mismatch '(1 2 3 4 5 6) '(3 4 5 6 7) :start1 2 :end2 4) \rightarrow NIL
```

#### See Also:

Section 3.6 (Traversal Rules and Side Effects)

#### **Notes:**

The :test-not argument is deprecated.

replace

Function

#### **Syntax:**

replace sequence-1 sequence-2 kkey start1 end1 start2 end2  $\rightarrow$  sequence-1

## Arguments and Values:

 ${\it sequence-1} {\it --} a \ {\it sequence}.$ 

sequence-2—a sequence.

start1, end1—bounding index designators of sequence-1. The defaults for start1 and end1 are 0 and nil, respectively.

start2, end2—bounding index designators of sequence-2. The defaults for start2 and end2 are 0 and nil, respectively.

#### **Description:**

Destructively modifies sequence-1 by replacing the elements of subsequence-1 bounded by start1 and end1 with the elements of subsequence-2 bounded by start2 and end2.

Sequence-1 is destructively modified by copying successive elements into it from sequence-2. Elements of the subsequence of sequence-2 bounded by start2 and end2 are copied into the subsequence of sequence-1 bounded by start1 and end1. If these subsequences are not of the same length, then the shorter length determines how many elements are copied; the extra elements near the end of the longer subsequence are not involved in the operation. The number of elements copied can be expressed as:

```
(min (- end1 start1) (- end2 start2))
```

If sequence-1 and sequence-2 are the same object and the region being modified overlaps the region being copied from, then it is as if the entire source region were copied to another place and only then copied back into the target region. However, if sequence-1 and sequence-2 are not the same, but the region being modified overlaps the region being copied from (perhaps because of shared list structure or displaced arrays), then after the replace operation the subsequence of sequence-1 being modified will have unpredictable contents. It is an error if the elements of sequence-2 are not of a type that can be stored into sequence-1.

#### **Examples:**

```
(replace "abcdefghij" "0123456789" :start1 4 :end1 7 :start2 4) \to "abcd456hij" (setq lst "012345678") \to "012345678" (replace lst lst :start1 2 :start2 0) \to "010123456" lst \to "010123456"
```

#### Side Effects:

The *sequence-1* is modified.

#### See Also:

fill

# substitute, substitute-if, substitute-if-not, nsubstitute, nsubstitute-if, nsubstitute-if-not Function

#### Syntax:

```
substitute newitem olditem sequence &key from-end test test-not start end count key
```

ightarrow result-sequence

 $\textbf{substitute-if} \ \textit{newitem predicate sequence} \ \&\texttt{key} \ \textit{from-end start end count key} \\ \rightarrow \textit{result-sequence}$ 

 $\mathbf{substitute\text{-}if\text{-}not}\ \textit{newitem predicate sequence}\ \&\texttt{key}\ \textit{from\text{-}end start end count key} \\ \rightarrow \textit{result\text{-}sequence}$ 

 $\textbf{nsubstitute} \ \textit{newitem olditem sequence} \ \& \texttt{key} \ \textit{from-end test test-not start end count key} \\ \rightarrow \textit{sequence}$ 

# substitute, substitute-if, substitute-if-not, ...

 $\begin{tabular}{ll} {\bf nsubstitute-if} \end{tabular} \begin{tabular}{ll} {\bf nsubstitute-if} \end{tabular} \begin{tabular}{ll} {\bf newitem} \end{tabular} \begin{tabular}{ll} {\bf predicate} \end{tabular} \begin{tabular}{ll} {\bf sequence} \end{tabular} \begin{tabu$ 

 $\begin{array}{l} \textbf{nsubstitute-if-not} \ \textit{newitem predicate sequence \&key} \ \textit{from-end start end count key} \\ \rightarrow \textit{sequence} \end{array}$ 

## **Arguments and Values:**

newitem—an object.

olditem—an object.

sequence—a proper sequence.

predicate—a designator for a function of one argument that returns a generalized boolean.

from-end—a generalized boolean. The default is false.

test—a designator for a function of two arguments that returns a generalized boolean.

test-not—a designator for a function of two arguments that returns a generalized boolean.

start, end—bounding index designators of sequence. The defaults for start and end are 0 and nil, respectively.

count—an integer or nil. The default is nil.

key—a designator for a function of one argument, or nil.

result-sequence—a sequence.

#### Description:

substitute, substitute-if, and substitute-if-not return a copy of sequence in which each element that satisfies the test has been replaced with newitem.

nsubstitute, nsubstitute-if, and nsubstitute-if-not are like substitute, substitute-if, and substitute-if-not respectively, but they may modify sequence.

If sequence is a vector, the result is a vector that has the same actual array element type as sequence. If sequence is a list, the result is a list.

Count, if supplied, limits the number of elements altered; if more than count elements satisfy the test, then of these elements only the leftmost or rightmost, depending on from-end, are replaced, as many as specified by count. If count is supplied and negative, the behavior is as if zero had been supplied instead. If count is nil, all matching items are affected.

Supplying a *from-end* of *true* matters only when the *count* is provided (and *non-nil*); in that case, only the rightmost *count* elements satisfying the test are removed (instead of the leftmost).

predicate, test, and test-not might be called more than once for each sequence element, and their side effects can happen in any order.

# substitute, substitute-if, substitute-if-not, ...

The result of all these functions is a *sequence* of the same *type* as *sequence* that has the same elements except that those in the subsequence *bounded* by *start* and *end* and *satisfying the test* have been replaced by *newitem*.

substitute, substitute-if, and substitute-if-not return a sequence which can share with sequence or may be *identical* to the input sequence if no elements need to be changed.

nsubstitute and nsubstitute-if are required to setf any car (if sequence is a list) or aref (if sequence is a vector) of sequence that is required to be replaced with newitem. If sequence is a list, none of the cdrs of the top-level list can be modified.

## **Examples:**

```
(substitute #\. #\SPACE "0 2 4 6") 
ightarrow "0.2.4.6"
 (substitute 9 4 '(1 2 4 1 3 4 5)) \rightarrow (1 2 9 1 3 9 5)
 (substitute 9 4 '(1 2 4 1 3 4 5) :count 1) \rightarrow (1 2 9 1 3 4 5)
 (substitute 9 4 '(1 2 4 1 3 4 5) :count 1 :from-end t)
\rightarrow (1 2 4 1 3 9 5)
 (substitute 9 3 '(1 2 4 1 3 4 5) :test \#'>) \rightarrow (9 9 4 9 3 4 5)
 (substitute-if 0 #'evenp '((1) (2) (3) (4)) :start 2 :key #'car)
\rightarrow ((1) (2) (3) 0)
 (substitute-if 9 #'oddp '(1 2 4 1 3 4 5)) \rightarrow (9 2 4 9 9 4 9)
 (substitute-if 9 #'evenp '(1 2 4 1 3 4 5) :count 1 :from-end t)
\rightarrow (1 2 4 1 3 9 5)
 (setq some-things (list 'a 'car 'b 'cdr 'c)) 
ightarrow (A CAR B CDR C)
 (nsubstitute-if "function was here" #'fboundp some-things
                   :count 1 :from-end t) 
ightarrow (A CAR B "function was here" C)
 some-things 
ightarrow (A CAR B "function was here" C)
 (setq alpha-tester (copy-seq "ab ")) 
ightarrow "ab "
 (nsubstitute-if-not \ "alpha-char-p alpha-tester) \rightarrow "abz"
 alpha-tester 
ightarrow "abz"
```

#### **Side Effects:**

nsubstitute, nsubstitute-if, and nsubstitute-if-not modify sequence.

## **Exceptional Situations:**

Should be prepared to signal an error of type type-error if sequence is not a proper sequence.

#### See Also:

subst, nsubst, Section 3.2.1 (Compiler Terminology), Section 3.6 (Traversal Rules and Side Effects)

#### **Notes:**

If sequence is a vector, the result might or might not be simple, and might or might not be

#### 17–28 Programming Language—Common Lisp

identical to sequence.

The :test-not argument is deprecated.

The functions substitute-if-not and nsubstitute-if-not are deprecated.

nsubstitute and nsubstitute-if can be used in for-effect-only positions in code.

Because the side-effecting variants (e.g., nsubstitute) potentially change the path that is being traversed, their effects in the presence of shared or circular structure may vary in surprising ways when compared to their non-side-effecting alternatives. To see this, consider the following side-effect behavior, which might be exhibited by some implementations:

concatenate Function

## Syntax:

 $\mathbf{concatenate} \ \mathit{result-type} \ \mathtt{\&rest} \ \mathit{sequences} \ \rightarrow \mathit{result-sequence}$ 

## **Arguments and Values:**

```
result-type—a sequence type specifier.
sequences—a sequence.
result-sequence—a proper sequence of type result-type.
```

#### **Description:**

**concatenate** returns a *sequence* that contains all the individual elements of all the *sequences* in the order that they are supplied. The *sequence* is of type *result-type*, which must be a *subtype* of *type* **sequence**.

All of the *sequences* are copied from; the result does not share any structure with any of the *sequences*. Therefore, if only one *sequence* is provided and it is of type *result-type*, **concatenate** is required to copy *sequence* rather than simply returning it.

It is an error if any element of the sequences cannot be an element of the sequence result.

If the *result-type* is a *subtype* of **list**, the result will be a *list*.

If the *result-type* is a *subtype* of **vector**, then if the implementation can determine the element type specified for the *result-type*, the element type of the resulting array is the result of *upgrading* that element type; or, if the implementation can determine that the element type is unspecified (or \*), the element type of the resulting array is t; otherwise, an error is signaled.

#### **Examples:**

```
(concatenate 'string "all" " "together" " " "now") \rightarrow "all together now" (concatenate 'list "ABC" '(d e f) #(1 2 3) #*1011) \rightarrow (#\A #\B #\C D E F 1 2 3 1 0 1 1) (concatenate 'list) \rightarrow NIL (concatenate '(vector * 2) "a" "bc") should signal an error
```

#### **Exceptional Situations:**

An error is signaled if the *result-type* is neither a *recognizable subtype* of **list**, nor a *recognizable subtype* of **vector**.

An error of *type* **type-error** should be signaled if *result-type* specifies the number of elements and the sum of *sequences* is different from that number.

## See Also:

append

merge

#### **Syntax:**

merge result-type sequence-1 sequence-2 predicate  $kkey \ key \ o \ result-sequence$ 

#### **Arguments and Values:**

```
result-type—a sequence type specifier.

sequence-1—a sequence.

sequence-2—a sequence.

predicate—a designator for a function of two arguments that returns a generalized boolean.

key—a designator for a function of one argument, or nil.

result-sequence—a proper sequence of type result-type.
```

## **Description:**

Destructively merges *sequence-1* with *sequence-2* according to an order determined by the *predicate*. **merge** determines the relationship between two elements by giving keys extracted from the sequence elements to the *predicate*.

The first argument to the *predicate* function is an element of *sequence-1* as returned by the *key* (if supplied); the second argument is an element of *sequence-2* as returned by the *key* (if supplied). *Predicate* should return *true* if and only if its first argument is strictly less than the second (in some appropriate sense). If the first argument is greater than or equal to the second (in the appropriate sense), then *predicate* should return *false*. **merge** considers two elements x and y to be equal if (funcall predicate x y) and (funcall predicate y x) both *yield false*.

The argument to the *key* is the *sequence* element. Typically, the return value of the *key* becomes the argument to *predicate*. If *key* is not supplied or nil, the sequence element itself is used. The *key* may be executed more than once for each *sequence element*, and its side effects may occur in any order.

If key and predicate return, then the merging operation will terminate. The result of merging two  $sequences \ x$  and y is a new sequence of type  $result-type \ z$ , such that the length of z is the sum of the lengths of x and y, and z contains all the elements of x and y. If x1 and x2 are two elements of x, and x1 precedes x2 in x, then x1 precedes x2 in z, and similarly for elements of y. In short, z is an interleaving of x and y.

If x and y were correctly sorted according to the *predicate*, then z will also be correctly sorted. If x or y is not so sorted, then z will not be sorted, but will nevertheless be an interleaving of x and y.

The merging operation is guaranteed stable; if two or more elements are considered equal by the *predicate*, then the elements from *sequence-1* will precede those from *sequence-2* in the result.

sequence-1 and/or sequence-2 may be destroyed.

If the result-type is a subtype of list, the result will be a list.

If the *result-type* is a *subtype* of **vector**, then if the implementation can determine the element type specified for the *result-type*, the element type of the resulting array is the result of *upgrading* that element type; or, if the implementation can determine that the element type is unspecified (or \*), the element type of the resulting array is t; otherwise, an error is signaled.

## **Examples:**

```
(setq test1 (list 1 3 4 6 7))

(setq test2 (list 2 5 8))

(merge 'list test1 test2 #'<) \rightarrow (1 2 3 4 5 6 7 8)

(setq test1 (copy-seq "BOY"))

(setq test2 (copy-seq :nosy"))

(merge 'string test1 test2 #'char-lessp) \rightarrow "BnOosYy"
```

```
(setq test1 (vector ((red . 1) (blue . 4))))
(setq test2 (vector ((yellow . 2) (green . 7))))
(merge 'vector test1 test2 #'< :key #'cdr)
→ #((RED . 1) (YELLOW . 2) (BLUE . 4) (GREEN . 7))

(merge '(vector * 4) '(1 5) '(2 4 6) #'<) should signal an error</pre>
```

#### **Exceptional Situations:**

An error must be signaled if the *result-type* is neither a *recognizable subtype* of **list**, nor a *recognizable subtype* of **vector**.

An error of *type* **type-error** should be signaled if *result-type* specifies the number of elements and the sum of the lengths of *sequence-1* and *sequence-2* is different from that number.

#### See Also:

sort, stable-sort, Section 3.2.1 (Compiler Terminology), Section 3.6 (Traversal Rules and Side Effects)

# remove, remove-if, remove-if-not, delete, delete-if, delete-if-not

## Syntax:

remove item sequence &key from-end test test-not start end count key  $\rightarrow$  result-sequence remove-if test sequence &key from-end start end count key  $\rightarrow$  result-sequence remove-if-not test sequence &key from-end start end count key  $\rightarrow$  result-sequence delete item sequence &key from-end test test-not start end count key  $\rightarrow$  result-sequence delete-if test sequence &key from-end start end count key  $\rightarrow$  result-sequence delete-if-not test sequence &key from-end start end count key  $\rightarrow$  result-sequence

#### **Arguments and Values:**

```
item—an object.
sequence—a proper sequence.
test—a designator for a function of one argument that returns a generalized boolean.
from-end—a generalized boolean. The default is false.
```

# remove, remove-if, remove-if-not, delete, delete-if, ...

test—a designator for a function of two arguments that returns a generalized boolean.

test-not—a designator for a function of two arguments that returns a generalized boolean.

start, end—bounding index designators of sequence. The defaults for start and end are 0 and nil, respectively.

count—an integer or nil. The default is nil.

key—a designator for a function of one argument, or nil.

result-sequence—a sequence.

#### **Description:**

remove, remove-if, and remove-if-not return a sequence from which the elements that satisfy the test have been removed.

delete, delete-if, and delete-if-not are like remove, remove-if, and remove-if-not respectively, but they may modify sequence.

If sequence is a vector, the result is a vector that has the same actual array element type as sequence. If sequence is a list, the result is a list.

Supplying a *from-end* of *true* matters only when the *count* is provided; in that case only the rightmost *count* elements *satisfying the test* are deleted.

Count, if supplied, limits the number of elements removed or deleted; if more than count elements satisfy the test, then of these elements only the leftmost or rightmost, depending on from-end, are deleted or removed, as many as specified by count. If count is supplied and negative, the behavior is as if zero had been supplied instead. If count is nil, all matching items are affected.

For all these functions, elements not removed or deleted occur in the same order in the result as they did in *sequence*.

remove, remove-if, remove-if-not return a sequence of the same type as sequence that has the same elements except that those in the subsequence bounded by start and end and satisfying the test have been removed. This is a non-destructive operation. If any elements need to be removed, the result will be a copy. The result of remove may share with sequence; the result may be identical to the input sequence if no elements need to be removed.

delete, delete-if, and delete-if-not return a sequence of the same type as sequence that has the same elements except that those in the subsequence bounded by start and end and satisfying the test have been deleted. Sequence may be destroyed and used to construct the result; however, the result might or might not be identical to sequence.

**delete**, when *sequence* is a *list*, is permitted to **setf** any part, **car** or **cdr**, of the top-level list structure in that *sequence*. When *sequence* is a *vector*, **delete** is permitted to change the dimensions of the *vector* and to slide its elements into new positions without permuting them to produce the resulting *vector*.

# remove, remove-if, remove-if-not, delete, delete-if, ...

delete-if is constrained to behave exactly as follows:

```
(delete nil sequence
          :test #'(lambda (ignore item) (funcall test item))
          ...)
```

## **Examples:**

```
(remove 4 '(1 3 4 5 9)) \rightarrow (1 3 5 9)
 (remove 4 , (1 2 4 1 3 4 5)) \rightarrow (1 2 1 3 5)
 (remove 4 '(1 2 4 1 3 4 5) :count 1) \rightarrow (1 2 1 3 4 5)
 (remove 4 '(1 2 4 1 3 4 5) :count 1 :from-end t) \rightarrow (1 2 4 1 3 5)
 (remove 3 '(1 2 4 1 3 4 5) :test #'>) \rightarrow (4 3 4 5)
 (setq lst '(list of four elements)) 
ightarrow (LIST OF FOUR ELEMENTS)
 (setq 1st2 (copy-seq 1st)) 
ightarrow (LIST OF FOUR ELEMENTS)
 (\texttt{setq 1st3 (delete 'four 1st)}) \ \rightarrow \ (\texttt{LIST OF ELEMENTS})
 (equal 1st 1st2) \rightarrow false
 (remove-if #'oddp '(1 2 4 1 3 4 5)) 
ightarrow (2 4 4)
 (remove-if #'evenp '(1 2 4 1 3 4 5) :count 1 :from-end t)
\rightarrow (1 2 4 1 3 5)
 (remove-if-not #'evenp '(1 2 3 4 5 6 7 8 9) :count 2 :from-end t)
\rightarrow (1 2 3 4 5 6 8)
 (setq tester (list 1 2 4 1 3 4 5)) \rightarrow (1 2 4 1 3 4 5)
 (delete 4 tester) \rightarrow (1 2 1 3 5)
 (setq tester (list 1 2 4 1 3 4 5)) \rightarrow (1 2 4 1 3 4 5)
 (delete 4 tester :count 1) \rightarrow (1 2 1 3 4 5)
 (setq tester (list 1 2 4 1 3 4 5)) \rightarrow (1 2 4 1 3 4 5)
 (delete 4 tester :count 1 :from-end t) 
ightarrow (1 2 4 1 3 5)
 (setq tester (list 1 2 4 1 3 4 5)) 
ightarrow (1 2 4 1 3 4 5)
 (delete 3 tester :test #'>) 
ightarrow (4 3 4 5)
 (setq tester (list 1 2 4 1 3 4 5)) \rightarrow (1 2 4 1 3 4 5)
 (delete-if #'oddp tester) \rightarrow (2 4 4)
 (setg tester (list 1 2 4 1 3 4 5)) \rightarrow (1 2 4 1 3 4 5)
 (delete-if #'evenp tester :count 1 :from-end t) 
ightarrow (1 2 4 1 3 5)
 (setq tester (list 1 2 3 4 5 6)) \rightarrow (1 2 3 4 5 6)
 (delete-if #'evenp tester) 
ightarrow (1 3 5)
 tester \rightarrow implementation-dependent
 (setq foo (list 'a 'b 'c)) 
ightarrow (A B C)
 (\texttt{setq bar (cdr foo)}) \, \rightarrow \, (\texttt{B C})
 (setq foo (delete 'b foo)) 
ightarrow (A C)
 bar \rightarrow ((C)) or ...
 (eq (cdr foo) (car bar)) \rightarrow T or ...
```

#### Side Effects:

For delete, delete-if, and delete-if-not, sequence may be destroyed and used to construct the result.

#### **Exceptional Situations:**

Should be prepared to signal an error of type type-error if sequence is not a proper sequence.

#### See Also:

Section 3.2.1 (Compiler Terminology), Section 3.6 (Traversal Rules and Side Effects)

## Notes:

If *sequence* is a *vector*, the result might or might not be simple, and might or might not be *identical* to *sequence*.

The :test-not argument is deprecated.

The functions delete-if-not and remove-if-not are deprecated.

# remove-duplicates, delete-duplicates

Function

## Syntax:

remove-duplicates sequence &key from-end test test-not start end key

ightarrow result-sequence

delete-duplicates sequence &key from-end test test-not start end key

ightarrow result-sequence

#### **Arguments and Values:**

sequence—a proper sequence.

from-end—a generalized boolean. The default is false.

test—a designator for a function of two arguments that returns a generalized boolean.

test-not—a designator for a function of two arguments that returns a generalized boolean.

start, end—bounding index designators of sequence. The defaults for start and end are 0 and nil, respectively.

key—a designator for a function of one argument, or nil.

# remove-duplicates, delete-duplicates

result-sequence—a sequence.

## **Description:**

**remove-duplicates** returns a modified copy of *sequence* from which any element that matches another element occurring in *sequence* has been removed.

If sequence is a vector, the result is a vector that has the same actual array element type as sequence. If sequence is a list, the result is a list.

delete-duplicates is like remove-duplicates, but delete-duplicates may modify sequence.

The elements of *sequence* are compared *pairwise*, and if any two match, then the one occurring earlier in *sequence* is discarded, unless *from-end* is *true*, in which case the one later in *sequence* is discarded.

**remove-duplicates** and **delete-duplicates** return a *sequence* of the same *type* as *sequence* with enough elements removed so that no two of the remaining elements match. The order of the elements remaining in the result is the same as the order in which they appear in *sequence*.

**remove-duplicates** returns a *sequence* that may share with *sequence* or may be *identical* to *sequence* if no elements need to be removed.

**delete-duplicates**, when *sequence* is a *list*, is permitted to **setf** any part, **car** or **cdr**, of the top-level list structure in that *sequence*. When *sequence* is a *vector*, **delete-duplicates** is permitted to change the dimensions of the *vector* and to slide its elements into new positions without permuting them to produce the resulting *vector*.

## Examples:

```
(remove-duplicates "aBcDAbCd" :test #'char-equal :from-end t) \rightarrow "aBcD" (remove-duplicates '(a b c b d d e)) \rightarrow (A C B D E) (remove-duplicates '(a b c b d d e) :from-end t) \rightarrow (A B C D E) (remove-duplicates '((foo #\a) (bar #\%) (baz #\A)) :test #'char-equal :key #'cadr) \rightarrow ((BAR #\%) (BAZ #\A)) (remove-duplicates '((foo #\a) (bar #\%) (baz #\A)) :test #'char-equal :key #'cadr :from-end t) \rightarrow ((F00 #\a) (BAR #\%)) (setq tester (list 0 1 2 3 4 5 6)) (delete-duplicates tester :key #'oddp :start 1 :end 6) \rightarrow (0 4 5 6)
```

#### **Side Effects:**

delete-duplicates might destructively modify sequence.

#### **Exceptional Situations:**

Should signal an error of type type-error if sequence is not a proper sequence.

#### See Also:

Section 3.2.1 (Compiler Terminology), Section 3.6 (Traversal Rules and Side Effects)

# remove-duplicates, delete-duplicates

#### Notes:

If **sequence** is a **vector**, the result might or might not be simple, and might or might not be **identical** to **sequence**.

The :test-not argument is deprecated.

These functions are useful for converting sequence into a canonical form suitable for representing a set.