- ${\it Copyright: https://github.com/bringhurst/tlaplus/blob/master/org.lamport.tla.toolbox.uitest/farsite/AdditionalSequenceOperators and the control of the$
- EXTENDS FiniteSets, Sequences, TLC, AdditionalSetOperators, AdditionalFunctionOperators
- 8 LOCAL INSTANCE Naturals

The TLA+ Sequences module defines the operators Head and Tail for retrieving the first element of a sequence and all-but-the-first elements of a sequence, respectively. This module provides four operators that slightly generalize the notions of Head and Tail:

First returns the first element of a sequence, equivalently to Head. Last returns the last element of a sequence. AllButFirst returns all-but-the-first elements of a sequence, equivalently to Tail.

AllButLast returns all-but-the-last elements of a sequence.

This module also provides several additional operators on sequences: IsElementInSeq is a predicate that is true when the specified value is an element of the specified sequence. IsSequenceOfSetElements is a predicate that is true when the specified sequence contains all and only elements of the specified set. IsSortedSequenceOfSetElements is a predicate that is true when the IsSequenceOfSetElements is true and the sequence is also sorted in increasing order. DeleteElement produces a sequence by deleting an indicated element from another sequence.

```
Prepend(s, e) \stackrel{\Delta}{=} \langle e \rangle \circ s
    First(seq) \stackrel{\triangle}{=} seq[1]
    Last(seq) \stackrel{\Delta}{=} seq[Len(seq)]
     AllButFirst(seq) \stackrel{\Delta}{=} [i \in 1 .. (Len(seq) - 1) \mapsto seq[(i + 1)]]
     AllButLast(seq) \triangleq [i \in 1 ... (Len(seq) - 1) \mapsto seq[i]]
     DoesSeqPrefixSeq(seq1, seq2) \triangleq
42
        \land Len(seq1) \le Len(seq2)
43
        \land (\forall \, i \in 1 \dots Len(seq1) : seq1[i] = seq2[i])
44
     DoesSeqProperlyPrefixSeq(seq1, seq2) \stackrel{\Delta}{=}
        \land Len(seq1) < Len(seq2)
47
        \land (\forall i \in 1 ... Len(seq1) : seq1[i] = seq2[i])
48
     IsElementInSeq(el, seq) \triangleq \exists i \in DOMAIN seq : seq[i] = el
     IsSequenceOfSetElements(seq, set) \triangleq
52
        \wedge Len(seq) = Cardinality(set)
53
        \land (\forall el \in set : IsElementInSeq(el, seq))
54
     IsSortedSequenceOfSetElements(seq, set) \triangleq
        \land IsSequenceOfSetElements(seq, set)
57
        \land (\forall i \in \text{DOMAIN } seq, j \in \text{DOMAIN } seq: i < j \Rightarrow seq[i] < seq[j])
58
    DeleteElement(seq, index) \triangleq
```

```
[i \in 1..(Len(seq) - 1) \mapsto \text{if } i < index \text{ then } seq[i] \text{ else } seq[(i+1)]]
     Retain only the elements in R in their original order in seq.
     RECURSIVE Retain(_, _)
 66
 67
     Retain(seq, R) \triangleq
          IF seq = \langle \rangle
 68
           THEN ()
 69
           ELSE LET h \stackrel{\triangle}{=} Head(seq)
 70
                    IN IF h \in R
 71
                           THEN h \circ Retain(Tail(seq), R)
 72
                           ELSE Retain(Tail(seq), R)
 73
     It requires that index \geq 1.
     If index > Len(seq) + 1, then it appends the element to seq.
     (ADDED by hengxin; July 04, 2018)
     InsertElement(seq, elem, index) \triangleq
 81
       [i \in 1 ... (Len(seq) + 1) \mapsto IF \ i < index
 82
                                           THEN IF i = (Len(seq) + 1)
 83
                                                    THEN elem
 84
                                                    ELSE seq[i]
 85
                                           ELSE IF i = index
 86
                                                    THEN elem
 87
                                                    ELSE seq[(i-1)] i > index
 88
     IsSorted2Partition(n, seq1, seq2) \stackrel{\Delta}{=}
        \land seq1 \in Seq(1 \dots n)
 91
 92
        \land seq2 \in Seq(1 \dots n)
        \wedge n = Len(seq1) + Len(seq2)
 93
        \land (\forall i \in \text{DOMAIN } seq1, j \in \text{DOMAIN } seq1: i < j \Rightarrow seq1[i] < seq1[j])
 94
        \land (\forall i \in \text{DOMAIN } seq2, j \in \text{DOMAIN } seq2: i < j \Rightarrow seq2[i] < seq2[j])
 95
        \land (\forall i \in \text{DOMAIN } seq1, j \in \text{DOMAIN } seq2 : seq1[i] \neq seq2[j])
 96
      IsSequenceInterleaving(seq, subSeq1, subSeq2, indSeq1, indSeq2) \stackrel{\Delta}{=}
 98
        \land indSeq1 \in Seq(Nat)
 99
        \land indSeq2 \in Seq(Nat)
100
        \land IsSorted2Partition(Len(seq), indSeq1, indSeq2)
101
        \wedge Len(indSeq1) = Len(subSeq1)
102
103
        \wedge Len(indSeq2) = Len(subSeq2)
        \land (\forall i \in DOMAIN \ indSeq1 : seq[(indSeq1[i])] = subSeq1[i])
104
        \land (\forall i \in DOMAIN \ indSeq2 : seq[(indSeq2[i])] = subSeq2[i])
105
     Sequences up to length n, including the empty sequence \langle \rangle.
     Copyright: https://www.learntla.com/libraries/sequences/
112 SegMaxLen(S, n) \triangleq UNION \{[1 ... m \rightarrow S] : m \in 0 ... n\}
```

```
Map on a sequence.
     Copyright: https://www.learntla.com/libraries/sequences/
119 SeqMap(Op(\_), seq) \stackrel{\triangle}{=} [x \in DOMAIN \ seq \mapsto Op(seq[x])]
121 PermsWithin(S) \triangleq \{s \in \text{UNION } \{[1 ... m \to S] : m \in 0 ... Cardinality(S)\} : Cardinality(Range(s)) = Cardinality(S)\} \}
     All possible permutations generated based on sequence T.
     Copyright: https://learntla.com/tla/functions/
128 PermutationKey(n) \triangleq \{key \in [1 ... n \rightarrow 1 ... n] : Range(key) = 1 ... n\}
Permutations Of(T) \triangleq \{[x \in 1 ... Len(T) \mapsto T[P[x]]] : P \in Permutation Key(Len(T))\}
     Get the index of the first occurrence of elem in seq.
     Precondition: elem \in SeqImage(seq).
     ADDED by hengxin; Aug. 12, 2018
   RECURSIVE FirstIndexOfElement(_, _)
137
     FirstIndexOfElement(seq, elem) \stackrel{\Delta}{=}
138
         IF Head(seq) = elem
139
140
          THEN 1
          ELSE 1 + FirstIndexOfElement(Tail(seq), elem)
141
     Check if two sequences are compatible.
     Precondition: No duplication in each individual sequence.
     Two sequences are compatible if and only if for any two common elements in both sequences, the
     relative order of them in the two sequences are the same.
     ADDED by hengxin; Aug. 12, 2018
     Compatible(seq1, seq2) \stackrel{\Delta}{=}
154
          \lor seq1 = seq2
155
          \vee LET commonElements \stackrel{\Delta}{=} Range(seq1) \cap Range(seq2)
156
            IN \forall e1, e2 \in commonElements:
157
                    \vee e1 = e2
158
                    \vee FirstIndexOfElement(seq1, e1) < FirstIndexOfElement(seq1, e2)
159
                       \equiv FirstIndexOfElement(seq2, e1) < FirstIndexOfElement(seq2, e2)
160
     The length of the longest common subsequence of two sequences seq1 and seq2.
     ADDED by hengxin; Aug. 12, 2018
    RECURSIVE LCS(\_,\_)
167
     LCS(seq1, seq2) \triangleq
168
           IF seq1 = \langle \rangle \lor seq2 = \langle \rangle
169
            THEN 0
170
            ELSE IF Last(seq1) = Last(seq2)
171
                    THEN 1 + LCS(AllButLast(seq1), AllButLast(seq2))
172
                    ELSE MaxOfSet(\{LCS(AllButLast(seq1), seq2), LCS(seq1, AllButLast(seq2))\})
173
```

 $LCSCompatible(seq1, seq2) \stackrel{\Delta}{=}$

```
Compatible(seq1, seq2) \equiv LCS(seq1, seq2) = Cardinality(Range(seq1) \cap Range(seq2))
178 \quad LCSCompatibleTest(S) \triangleq \\ \forall seq1, seq2 \in PermsWithin(S) : LCSCompatible(seq1, seq2)
180 \quad \\ \land \text{Modification History} \\ \land \text{Last modified Tue Sep 25 19:09:57 CST 2018 by hengxin} \\ \land \text{Created Tue Jul 03 15:21:02 CST 2018 by hengxin}
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