ARR v1.0

```
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The ARR library provides basic functions for working with arrays in AWK.

The library includes 25 functions for array operations:

- 5 functions for combining array elements into a string:
 - ret::a return array elements
 - ret::as return array elements with a separator
 - ret::ad return array elements
 - ret::ab return elements of two arrays
 - ret::abs return elements of two arrays with a separator

These functions were created using elements of linear programming technologies, which is why they offer high performance and are designed for large data volumes and high numbers of merged indices.

See the "ARRAY JOINING" section.

See the "PERFORMANCE" section.

- 3 functions for declaring subarrays within arrays and removing indices from arrays:

def::ia declare an empty subarray in array[index]

let::ia declare a subarray in array[index] if it has not been

declared as an array yet

del::ia delete array[index]

- 2 special functions

arr::same two DIFFERENT arrays?

arr::name defining the name of a global array

- 2 functions for generating visualization of array contents

dump::a array dump

dump::ab dump of two arrays

- 15 functions for copying data between arrays and their elements:

let::aa

def::aa array1 < array2

mov::aa

let::iav

def::iav array[index] < array/value

mov::iav

let::iai

def::iai array[index1] < array[index2]

mov::iai

let::aia

def::aia array1[index] < array2[index]

mov::aia

let::iaia

def::iaia array1[index1] < array2[index2]

mov::iaia

The **let**, **def**, and **mov** functions differ in behavior when copying arrays into arrays:

mov-functions simply copy data from the source to the recipient: when copying an array into an already existing array, the data in it is preserved except for those that will be overwritten by the source indices during the copying process.

def-functions always clear the target: array or array[index]: exact copying.

let-functions copy only what is not present in the recipient: that is, only those source indices that are not in the recipient are copied.

```
USE
```

To use it, execute:

```
@include
                        "arr.lib"
      Библиотека ..
FUNCSET
del::ia( i, A )
      Delete A[i] (if exist)
      Returns i.
let::ia( i, A )
def::ia(i, A)
      Define A[i] as the subarray.
      Returns i.
      Specifuc:
             def::ia
                          always clear array A[i]
      example:
             ... patsplit( ..., A[ def::a( "A", A ) ], ..., A[ def::a( "B", A ) ] ) ...
             patsplit will be targeted to arrays: A[ "A" ] and A[ "B" ]
```

let::aa(D, S) def::aa(D, S) mov::aa(D, S)

Copy all indexes from the source array S to destination array D.

if S and D is the same array then do nothing.

Returns null

Specifuc:

def::aa pre-clear array D

let::aa indexes of S are will be copied in case if it's not exist in D.

example:

let::iav(i, A, V) def::iav(i, A, V) mov::iav(i, A, V)

Copy V to A[i]

Returns i

V may be any of type (array or other)

Specific:

def::iav A[i] will be always deleted before V copied

def::iav if V is untyped then A[i] will be deleted

example:

let::iai(d, A, s) def::iai(d, A, s) mov::iai(d, A, s)

ZZ

example:

let::aia(D, i, S)
def::aia(D, i, S)
mov::aia(D, i, S)

ZZ

example:

let::iaia(d, D, s, S) def::iaia(d, D, s, S) mov::iaia(d, D, s, S)

ZZ

example:

arr::**same**(*A*, *B*)

Returns true (1) in case if the arrays A and B are the same array. Otherwise returns 0.

arr::name(A)

Returns the name of the given global array A.

If A is not global array then returns null.

```
dump::a(A, name )
```

Returns dump of the given array A

example:

ZZ

dump::ab(A, B)

Returns dump of the two given arrays A and B

example:

ARRAY JOINING

This library includes the implementation of four operations related to combining array elements into a string:

RETA	объединение элементов массива			
	A[a] $A[a+=x]$ $A[a+=x]$			
RETAS	объединение элементов массива через сепаратор			
	A[a] sep A[a+=x] sep A[a+=x]			
RETAB	объединение пар элементов двух массивов			
	A[a]B[b] $A[a+=x]B[b+=y]$ $A[a+=x]B[b+=y]$			
RETABS	объединение пар элементов двух массивов через сепаратор			
	A[a]B[b] sep A[a+=x]B[b+=y] sep A[a+=x]B[b+=y]			

In the process of combining elements into a string, the data from the combined elements is read and copied into the resulting string. This process is called **data pass**.

Data pass refers to when some data is read and copied somewhere else. This process is what consumes performance during the joining of array elements.

When creating this library, the primary goal was to minimize the number of data passes for the same data. The number of required data passes directly depends on the number of indices being combined.

Below is a table informing about the number of data passes for various amounts of combined indices:

indexes	data pass
1 64	1
65 64K	2
64K 4M	3
4M 256M	4
256M 16G	5

- 1 64 indices are returned in a single data pass—i.e., directly from the source arrays.
- 65 64K combined indices are returned in two data passes: the data is copied from the source arrays to a temporary array and then returned from it.
- 64K 4M indices are returned in 3 data passes: the data is copied from the source arrays to a temporary array, the temporary array is compressed 64 times (AIR), and then the data is returned from it.

More than 4M and up to 256M combined indices are returned in 4 data passes: data is copied from the source arrays to a temporary array, the temporary array is compressed 64 times twice (AIR), and then the data is returned from it.

More than 256M and up to 16G (billions) of indices will require 5 data passes: data is copied from the source arrays to a temporary array, the temporary array is compressed 64 times three times (AIR), and then the data is returned from it.

The RETA operation has a double number of indices for each data pass—i.e., it returns up to 128 indices in one data pass and up to 128K in two data passes, and so on.

```
ret::a( A, a, q, sa, sep )
ret::ad( A, sa, a, q, sep )
ret::as( A, sep, a, q, sa )
```

All three functions are doing the same things and differs from each other only by parameters order.

Returns specified number of indexes of an array with separator string between each index:

```
return A[a]sep A[a+=sa]sep ... A[a+=sa]
```

A is the source array

the first index in array is calculated by the following

```
first = ! (0 in A)
```

first index is equals to 0 in case if index 0 is exist in array A otherwise first index equals to 1

last index in an array is calculated by the following:

```
last = first + length(A) - 1
```

a is the start index in array A

if a is omitted (== "") then it's will be calculated depending from sa parameter (see below):

if index modifier (sa) is equals to zero or positive number (≥ 0) then start index a will be first index in array A

otherwise if index modifier (sa) is less than zero (< 0) then start index a will be last index in array A

sa is the index modifier for an array A

if sa is omitted (== "") then default value 1 will be used

q is the number of indexes to return

if q is omitted (== "") then it's will be calculated for covering rest of the elements of the array A starting from determinated start index a and using determinated index modifier sa

sep is the separator string that will be placed between the content of the each index joined

Below is the examples of usage functions:

```
example 1:
     t = "ABCDE"
     split( t, T, "")
                                  # splitting t for each character
     t2 = ret::a( T )
                                   # joining all elements of the array T
     # t == t2 == "ABCDE"
example 2:
     t = "ABCDE"
     split( t, T, "")
                                  # splitting t for each character
     t2 = ret::ad( T, -1 )
                                   # joining all elements of the array T
                                   # in reverse order
     # t == "ABCDE"
     # t2 == "EDCBA"
example 3:
     split(t, L, /\n/)
                                  # splitting multilined t for each line
     t2 = ret::as( L, "\n")
                                   # joining all elements (lines of t) of
                                   # the array L with separator: "\n" (eol)
     \# t == t2
example 4:
     t = "ABCDE"
     split( t, T, "")
                                  # splitting t for each character
     t2 = ret::a( T, 3 )
                                   # joining rest of the elements of the
                                   # array T starting from index 3 using
                                   # default index modifier (+1) with no
                                   # separator
     # t2 == "CDE"
example 5:
     t = "ABCDE"
     split( t, T, "")
                                  # splitting t for each character
     t2 = ret::ad(T, -1, 3)
                                   # joining rest of the elements of the
                                   # array T starting from index 3 and using
                                   # index modifier -1 with no separator
     # t2 == "CBA"
```

```
ret::ab( A, B, a, q, b, sa, sb, sep )
ret::abs( A, B, sep, a, q, b, sa, sb )
```

Both functions are doing the same things and differs from each other only by parameters order.

Returns specified number of index-pairs of the two arrays with separator string between each pair:

```
return A[a]B[b]sep A[a+=sa]B[b+=sb]sep ... A[a+=sa]B[b+=sb]
```

A, B are the source arrays

the first index in array is calculated by the following

```
first = ! ( 0 in A/B )
```

first index is equals to 0 in case if index 0 is exist in array A/B otherwise first index equals to 1

last index in an array is calculated by the following:

```
last = first + length (A/B) - 1
```

a, b are the start indexes of the array A, B respectively

if a/b is omitted (== "") then it's will be calculated depending from sa/sb parameter (see below):

if index modifier is equals to zero or positive number (≥ 0) then start index a/b will be first index in an array A/B

otherwise if index modifier is less than zero (< 0) then start index a/b will be last index in an array A/B

sa, sb are the index modifiers for array A, B respectively

if sa/sb is omitted (== "") then default value 1 will be used

q is the number of index-pairs to return

if q is omitted (== "") then it's will be calculated for covering rest of the elements of the array A starting from determinated start index a and using determinated index modifier sa

sep is the separator string that will be placed between the content of the each index-pair joined Below is the example of usage functions:

```
example:
```

```
f = "somefunc"

if ( q = patsplit( t, T, /.../, D ) ) {
    for ( t = 1; t <= q; t++ )
        T[ t ] = @f( T[ t ] )

    t = ret::ab( T, D ) }</pre>
```

The provided example shows a fragment of typical pointer processing in a string: we split the source string into two arrays: some pointers and the data between them. Then we call a certain function for each of the found pointers, and each time the result of this function is saved in the same array instead of the body of the found pointer. When we have "gone through" all the found pointers, we assemble the string back together by combining the two arrays into a string using the ret::ab function.

ARRAYS IN AWK

to define names (A,B,C,...) as the arrays use:

simpliest data accummulation in an array A is possible using

$$A[length(A)] = new added data$$

please note that A MUST to be defined as an array before

to pass undefined subarray A[i] use the def::ia/let::ia functions:

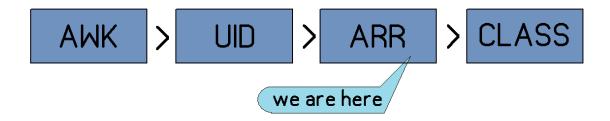
ret::ab() function is designed as the opposite to built-in split/patsplit functions called with the two target arrays specified

PERFORMANCE

PROSPECTS

The next release will be the CLASS – the OOP implementation for AWK: the fundamentals of basic techniques and a description of OOP practices in AWK.

A hypothetical publication plan for AWK libraries for the years 2024-2025:



AUTHOR

RESPECT DUE