

Repairing solutions

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1 Introduction

There are several approaches for including constraints into heuristics, see Chapter~12 of Gilli et al. [2011]. The notes in this vignette give some examples for simple repair mechanisms. These can be called in DEopt, GAopt and PSopt through the repair function; in LSopt/TAopt, they could be included in the neighbourhood function.

```
> set.seed(112233)
> options(digits = 3)
```

2 Upper and lower limits

Suppose the solution x is to satisfy $\text{all}(x \geq lo)$ and $\text{all}(x \leq up)$, with lo and up being vectors of $\text{length}(x)$.

2.1 Setting values to the boundaries

One strategy is to replace elements of x that violate a constraint with the boundary value. Such a repair function can be implemented very concisely. An example:

```
> up <- rep(1, 4L)
> lo <- rep(0, 4L)
> x <- rnorm(4L)
> x
```



```
[1] 2.127 -0.380 0.167 1.600
```

Three of the elements of x actually violate the constraints.

```
> repair1a <- function(x, up, lo) pmin(up, pmax(lo, x))
> x
```



```
[1] 2.127 -0.380 0.167 1.600
```



```
> repair1a(x, up, lo)
```



```
[1] 1.000 0.000 0.167 1.000
```

We see that indeed all values greater than 1 are replaced with 1, and those smaller than 0 become 0. Two other possibilities that achieve the same result:

```
> repair1b <- function(x, up, lo) {
+   ii <- x > up
+   x[ii] <- up[ii]
+   ii <- x < lo
+   x[ii] <- lo[ii]
+   x
```

```

+ }
> repair1c <- function(x, up, lo) {
+   xadjU <- x - up
+   xadjU <- xadjU + abs(xadjU)
+   xadjL <- lo - x
+   xadjL <- xadjL + abs(xadjL)
+   x <- (xadjU - xadjL)/2
+ }
```

The function `repair1b` uses comparisons to replace only the relevant elements in `x`. The function `repair1c` uses the ‘trick’ that

$$\text{pmax}(x, y) = \frac{x+y}{2} + \left| \frac{x-y}{2} \right|,$$

$$\text{pmin}(x, y) = \frac{x+y}{2} - \left| \frac{x-y}{2} \right|.$$

```

> repair1a(x, up, lo)
[1] 1.000 0.000 0.167 1.000

> repair1b(x, up, lo)
[1] 1.000 0.000 0.167 1.000

> repair1c(x, up, lo)
[1] 1.000 0.000 0.167 1.000

> trials <- 10000L
> strials <- seq_len(trials)
> system.time(for(i in strials) y1 <- repair1a(x, up, lo))

  user  system elapsed
0.416   0.000   0.416

> system.time(for(i in strials) y2 <- repair1b(x, up, lo))

  user  system elapsed
0.152   0.000   0.151

> system.time(for(i in strials) y3 <- repair1c(x, up, lo))

  user  system elapsed
0.084   0.004   0.089
```

The third of these functions would also work on matrices if `up` or `lo` were scalars.

```

> X <- array(rnorm(25L), dim = c(5L, 5L))
> X

 [,1]  [,2]  [,3]  [,4]  [,5]
[1,]  0.1962 0.434 -2.155 -1.5881 -1.029
[2,]  0.2284 1.231  0.975  0.0682  1.818
[3,] -1.1492 0.580 -0.711 -0.4457 -1.315
[4,] -0.0712 0.246  0.628  1.4662  0.511
[5,] -0.5619 0.388 -0.136 -0.8412  1.337
```

```

> repair1c(X, up = 0.5, lo = -0.5)

      [,1]   [,2]   [,3]   [,4]   [,5]
[1,]  0.1962  0.434 -0.500 -0.5000 -0.5
[2,]  0.2284  0.500  0.500  0.0682  0.5
[3,] -0.5000  0.500 -0.500 -0.4457 -0.5
[4,] -0.0712  0.246  0.500  0.5000  0.5
[5,] -0.5000  0.388 -0.136 -0.5000  0.5

```

The considerable speedup comes at a price, of course, since there is no checking (eg, for NA values) in `repair1b` and `repair1c`. We could also define new functions `pmin2` and `pmax2`.

```

> pmax2 <- function(x1, x2) ( (x1 + x2) + abs(x1 - x2) ) / 2
> pmin2 <- function(x1, x2) ( (x1 + x2) - abs(x1 - x2) ) / 2

```

A test follows.

```

> x1 <- rnorm(100L)
> x2 <- rnorm(100L)
> t1 <- system.time(for (i in trials) z1 <- pmax(x1,x2) )
> t2 <- system.time(for (i in trials) z2 <- pmax2(x1,x2))
> t1[[3L]]/t2[[3L]] ## speedup

[1] 3.12

> all.equal(z1, z2)

[1] TRUE

> t1 <- system.time(for (i in trials) z1 <- pmin(x1,x2) )
> t2 <- system.time(for (i in trials) z2 <- pmin2(x1,x2))
> t1[[3L]]/t2[[3L]] ## speedup

[1] 3.49

> all.equal(z1, z2)

[1] TRUE

```

One downside of this repair mechanism is that a solution may quickly become stuck at the boundaries (but of course, in some cases this is exactly what we want).

2.2 Reflecting values into the feasible range

The function `repair2` reflects a value that is too large or too small around the boundary. It restricts the change in a variable `x[i]` to the range `up[i] - lo[i]`.

```

> repair2 <- function(x,up,lo) {
+   done <- TRUE
+   e <- sum(x - up + abs(x - up) + lo - x + abs(lo - x))
+   if (e > 1e-12) done <- FALSE
+   r <- up - lo
+   while (!done) {
+     adjU <- x - up
+     adjU <- adjU + abs(adjU)

```

```

+     adjU <- adjU + r - abs(adjU - r)
+
+     adjL <- lo - x
+     adjL <- adjL + abs(adjL)
+     adjL <- adjL + r - abs(adjL - r)
+
+     x <- x - (adjU - adjL)/2
+     e <- sum(x - up + abs(x - up) + lo - x + abs(lo - x))
+     if (e < 1e-12) done <- TRUE
+
+   }
+
+ x
+ }
> x

[1] 2.127 -0.380 0.167 1.600

> repair2(x, up, lo)

[1] 0.873 0.380 0.167 0.600

> system.time(for (i in trials) y4 <- repair2(x, up, lo))

  user  system elapsed
0.624    0.000    0.623

```

2.3 Adjusting a cardinality limit

Let x be a logical vector.

```

> T <- 20L
> x <- logical(T)
> x[runif(T) < 0.4] <- TRUE
> x

[1] FALSE TRUE TRUE FALSE TRUE TRUE FALSE FALSE FALSE FALSE FALSE
[13] TRUE TRUE FALSE FALSE TRUE FALSE FALSE TRUE

```

Suppose we want to impose a minimum and maximum cardinality, k_{\min} and k_{\max} .

```

> kmax <- 5L
> kmin <- 3L

```

We could use an approach like the following (for the definition of `resample`, see `?sample`):

```

> resample <- function(x, ...) x[sample.int(length(x), ...)]
> repairK <- function(x, kmax, kmin) {
+   sx <- sum(x)
+   if (sx > kmax) {
+     i <- resample(which(x), sx - kmax)
+     x[i] <- FALSE
+   } else if (sx < kmin) {
+     i <- resample(which(!x), kmin - sx)
+     x[i] <- TRUE
+   }
+ x

```

```

+ }
> printK <- function(x)
+   cat(paste(ifelse(x, "o", "."), collapse = ""),
+        "-- cardinality", sum(x), "\n")

```

For kmax:

```

> for (i in 1:10) {
+   if (i==1L) printK(x)
+   x1 <- repairK(x, kmax, kmin)
+   printK(x1)
+ }

.oo.oo.....oo...o -- cardinality 8
.oo.o.....oo..... -- cardinality 5
.o...o.....oo.....o -- cardinality 5
.o..o.....o...o..o -- cardinality 5
..o.oo.....o.....o -- cardinality 5
..o...o.....oo...o... -- cardinality 5
....oo.....oo...o... -- cardinality 5
..o..oo.....o.....o -- cardinality 5
..oo...o.....oo..... -- cardinality 5
..oo..o.....o...o..o -- cardinality 5
..oo...o.....o...o... -- cardinality 5

```

For kmin:

```

> x <- logical(T); x[10L] <- TRUE
> for (i in 1:10) {
+   if (i==1L) printK(x)
+   x1 <- repairK(x, kmax, kmin)
+   printK(x1)
+ }

.....o..... -- cardinality 1
.....oo..o..... -- cardinality 3
o.....o.....o.... -- cardinality 3
...o.....o..o..... -- cardinality 3
.....o..o....o... -- cardinality 3
....o.....oo..... -- cardinality 3
....o...oo..... -- cardinality 3
.....o.o.....o...o -- cardinality 3
.....o..o..o..... -- cardinality 3
....o...o...o..... -- cardinality 3
....o....o...o..... -- cardinality 3

```

References

Manfred Gilli, Dietmar Maringer, and Enrico Schumann. *Numerical Methods and Optimization in Finance*. Elsevier, 2011.