

# RobASt: Implementation of optimally robust estimators

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# Outline

- 1 Introduction
- 2 Important S4 classes
- 3 Implementation of Algorithms
- 4 To-Do list

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## A first simple example

```
R > ## 24 determinations of copper in wholemeal flour
```

```
R > library(MASS)
```

```
R > data(chem)
```

```
R > sort(chem)
```

```
[1] 2.20 2.20 2.40 2.40 2.50 2.70 2.80 2.90 3.03  
[10] 3.03 3.10 3.37 3.40 3.40 3.40 3.50 3.60 3.70  
[19] 3.70 3.70 3.70 3.77 5.28 28.95
```

```
R > library(ROptEst)
```

```
R > robEst <- roptest(x = chem,  
+                   L2Fam = NormLocationScaleFamily(),  
+                   neighbor = ContNeighborhood(), # default  
+                   eps.lower = 0.05, eps.upper = 0.20, # rmx-estimator  
+                   risk = asMSE(), # default  
+                   distance = CvMDist, # default  
+                   steps = 3L, useLast = TRUE)
```

## How to obtain information about robust estimates

```
R > distrModOptions("show.details" = "minimal") # to reduce output  
R > robEst
```

Evaluations of 3-step estimate:

```
-----  
      mean      sd  
3.1649732  0.6668010  
(0.1513134) (0.1127157)
```

```
R > ## confidence intervals  
R > confint(robEst, method = symmetricBias())
```

```
A[n] asymptotic (LAN-based), uniform (bias-aware)  
confidence interval:  
for symmetric Bias  
      2.5 %    97.5 %  
mean 2.7676325 3.5623140  
sd   0.3708158 0.9627862
```

## Some diagnostics

```
R > ## check precision of computations  
R > checkIC(pIC(robEst), out = FALSE)
```

```
maximum deviation  
1.508162e-05
```

### Some diagnostic plots:

**plot:** plot the IC.

**infoPlot:** plot relative and absolute information; cf. [1].

**outlyingPlotIC:** outlyingness plot based on distances applied to ICs.

**comparePlot:** compare influence curves of AL estimators.

## Move IC to another parameter value (or ideal model)

```
R > head(modifyIC(pIC(robEst)), n = 19)
```

```
1 function (L2Fam, IC)
2 {
3   ICL2Fam <- eval(CallL2Fam(IC))
4   if (is(L2Fam, "L2LocationScaleFamily") && is(distribution(L2Fam),
5       class(distribution(ICL2Fam)))) {
6     sdneu <- main(L2Fam)[2]
7     sdalt <- main(ICL2Fam)[2]
8     r <- neighborRadius(IC)
9     w <- weight(IC)
10    clip(w) <- sdneu * clip(w)/sdalt
11    cent(w) <- sdalt * cent(w)/sdneu
12    stand(w) <- sdneu^2 * stand(w)/sdalt^2
13    weight(w) <- getweight(w, neighbor = ContNeighborhood(radius = r),
14        biastype = biastype(IC), normW = normtype(IC))
15    A <- sdneu^2 * stand(IC)/sdalt^2
16    b <- sdneu * clip(IC)/sdalt
17    a <- sdneu * cent(IC)/sdalt
18    mse <- sum(diag(A))
19    Cov <- sdneu^2 * Risks(IC)$asCov/sdalt^2
```

```
R > tail(modifyIC(pIC(robEst)), n = 4)
```

```
31   else {
32     makeIC(IC, L2Fam)
33   }
34 }
```

## Finite-sample correction

```
R > Infos(pIC(robEst))[2,2]

[1] "least favorable radius: 0.526"

R > ## finite-sample correction for normal location and scale
R > library(RobLox)
R > (r.fi <- finiteSampleCorrection(r = 0.526, n = length(chem)))

[1] 1.74

R > ## use argument fsCor or directly
R > robEst.fi <- roptest(x = chem, L2Fam = NormLocationScaleFamily(),
+                       eps = r.fi/sqrt(length(chem)),
+                       steps = 3L, useLast = TRUE)

R > estimate(robEst)

      mean      sd
3.164973 0.666801

R > estimate(robEst.fi)

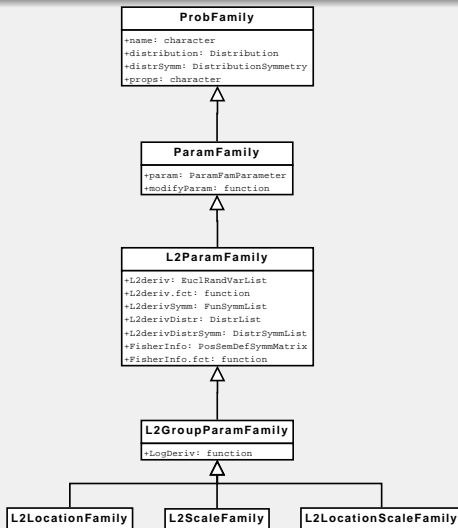
      mean      sd
3.2840688 0.6779314
```



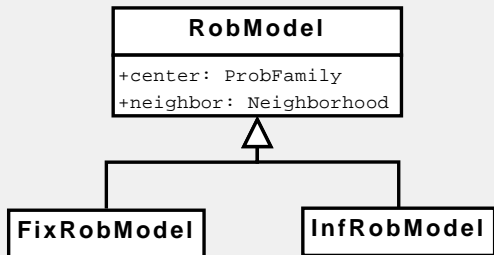
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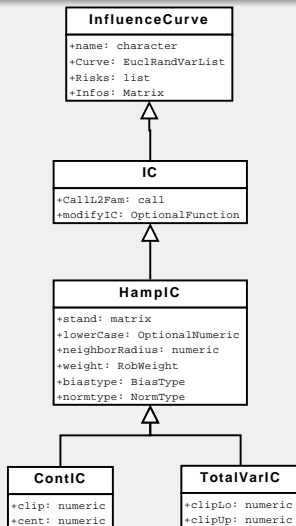
# Ideal family



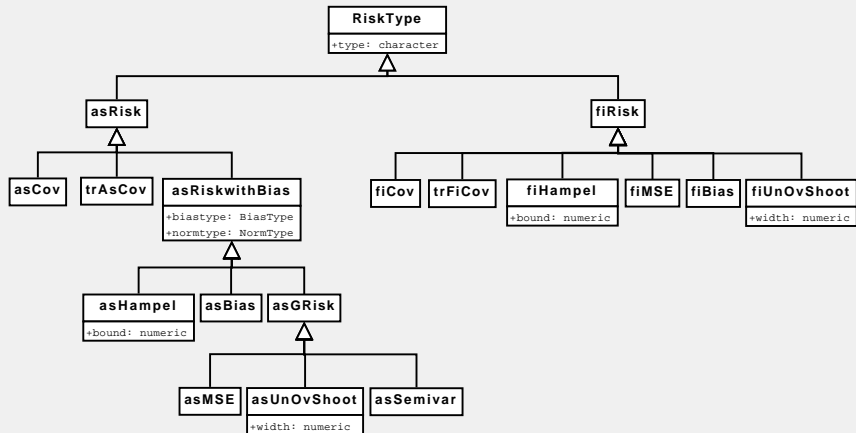
## Robust model



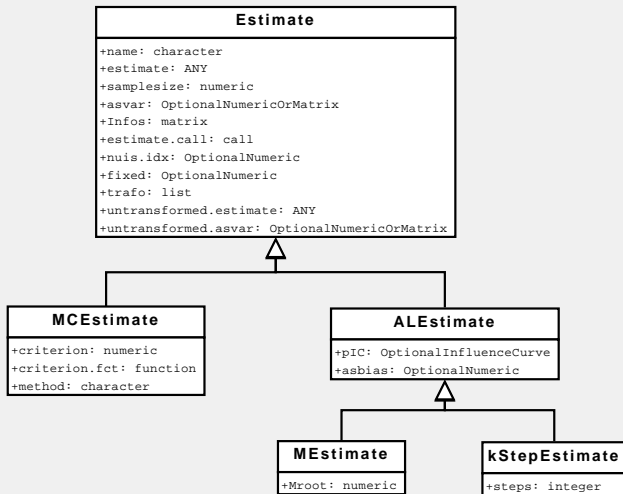
# Influence curve



## Risk type



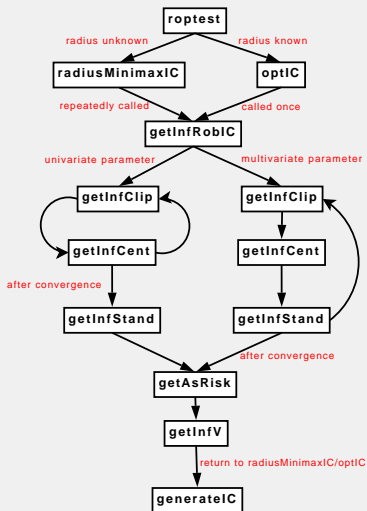
# Estimate



# Outline

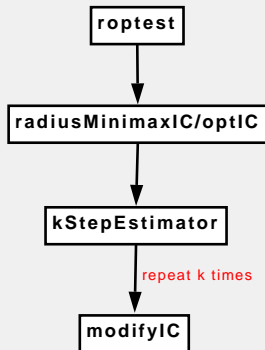
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# Computation of optimally robust influence curves





## Computation of k-step estimator



**Note:** modifyIC is slot of class IC!

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## To-Do list

- Can implementation of L2-derivative be improved?
- Move symmetry information to RandVar and Distribution classes?

# Bibliography I



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