SEP 2016: histplot() and histc() upgrade

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A bit of history

The discussion in the bug report ● #6306 and the ● SEP #110 aimed

- to split the computation of an histogram on one hand, and its display on the other hand.
- to become able to get the histogram's data that -- up to then -- were not available from histplot()
- to introduce a "normalized" way to compute the histogram heights.

It introduced in Scilab 5.4 a new *histc()* macro having the same syntaxes as *histplot()* for input data, and returning histogram's heights and memberships of input data in defined bins. See • https://codereview.scilab.org/#/c/13155/: new histc() + histplot() output added. However,

- The *normalization* option has been badly designed, since several ways to normalize data can be defined, whereas a boolean can take only one active value.
- By the way, these ways to compute the histogram do not *normalize* heights or area, in such a way that the name of this option is misleading. It shall be changed.
- In *histplot()*,
 - *normalization* has been appended to the very long list of input parameters, instead of being inserted after **data** -- as for *histc()* -- and before graphical options.
 - $\circ\,$ the default *normalization* value has been set to %T and then breaks back-compatibility for nothing.

By chance, neither the *normalization* nor the *polygon* options implemented for histplot() were yet documented for it in Scilab 5.5.2.

Moreover, other features are still missing:

- Only one method to compute bins is available. Other methods could be implemented and called through their name as a string.
- A default method to compute bins is missing.
- Neither *histc()* nor *histplot()* return bin's edges when these ones are computed.
- With *histc()*, it is now possible to compute an histogram out of *histplot()*, but there is still no way to make *histplot()* just plotting it without recomputing it.
- As for the *normalization* option, the *histplot() polygon* option has been appended to the list of input parameters instead of being inserted before the *style* option.
- Only vertical-up histograms can be displayed. vertical-down and Horizontal histograms shall become available.
- Polar histograms shall be supported.
- There is no option to display heights values on the bars.
- Processing of *-%inf*, *%inf*, and *%nan* values is undefined. There should be ways to make then counted and become able to ignore them or to take them into account when required.
- *histc()* and *histplot()* do not yet support text data, despite *dsearch()* on witch *histc()* is based was extended to text since Scilab 5.5.0.

New syntaxes proposed

histc()

- Existing syntaxes
 - \circ [heights] = ..
 - [heights, memberships] = ..
 - $\circ .. = histc(nBins, data)$
 - $\circ .. = histc(edges, data)$

• Removed syntaxes

• histc(..., data, normalization)

If backward-compatibility is of concern despite the short history of the *normalization* option, this one may be warn-obsoleted (for some while or forever) and automatically translated using the *histScale* option.

• New syntaxes

- histc(data)
- *histc*(*binsMethod*, *data*) with *binsMethod* = "*sqrt*"(*default*) | "*sturges*" | "*freediac*":
 - *sqrt*: *nbins* = *sqrt*(*size*(*data*, "*")
 - sturges: Sturges criteria: nbins = ceil(1 + log2(size(data, "*")))
 - *freediac*: Freedman-Diaconis criteria: binWidth = 2*iqr(data)*size(data,"*")^(-1/3). This method can't be applied to text data.
- histc(.., data, histScale) with histScale = "counts"/"countsNorm"/"density"/"densityNorm":
 - "counts": the height is the bin's number of members (default)
 - "density": the bin's area is the bin's number of members
 - "countsNorm": as "counts", divided by the total number of data. Discussion: should -%inf, %inf, %nan values, or/and data out of defined bins be taken into account for the "normalization"?
 - "densityNorm": as "density", divided...
- [heights, memberships, **binsDef**, **outside**] = histc(..) with
 - binsDef edges (continuous) or values (discrete)
 - outside = [Nminf, Npinf, Nnan] counts occurrences of -%inf, %inf and %nan values. In the membership array, -%inf will have the index -%inf, %inf will have the index %inf, and %nan will have the index -1

histplot()

- Existing syntaxes
 - [heights, memberships] = histplot(Nbins/edges, data [, <graphical options>..])
- New syntaxes
 - [heights, memberships, binsDef, outside] = histplot(..) : as for histc(..) (see here-above)
 - o histplot(data)
 - histplot(binsMethod, data)
 - histplot(.. data, histScale ..)
 - *histplot(.. data, histScale, dispOptions ..)* with *dispOptions* being a vector of one to five strings *provided in any order*, among the following, specifying options to display the histogram:
 - positions: "bottom" or "" (default) | "up" | "left" | "right" : draw the histogram with its base at the given position wrt its bars. Are these values clear enough?
 - "cumulate": draw the cumulated "staired" histogram instead of the simple one.
 - "*polygon*" : draw as well the polygon of frequencies or densities etc.

- "polar": draw the histogram in polar mode, bins being rescaled over the full [0, 180°] fan. Values of bins edges are displayed on an external graduated half circle.
- "values" : display values of heights on the bars
- *histplot(binsType, bins, heights, dispOptions* ...) to display as is an histogram already computed,
 - with *binsType* = "*binsEdges*" / "*binsValues*" : must be explicit (no default value)
 - Since allowed values of the *binsType* string are all distinct from values of the *binsMethod* string, parsing input arguments can easily detect this specific syntax.

Questions an discussion

- Since now, since Scilab 5.4, *histc()* can compute and return histogram results, and that histplot() can be fed by them, **should** *histplot()* **still return the results?**
- In output, do we shift the *memberships* array (introduced recently, 5.5.0) in 3rd position to put computed *edges* in argout#2?
- How do we manage *%nan* and *%inf*?

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