align: : CHEAT SHEET

Basics

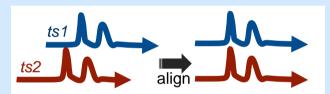
align is an R package for time-series data alignment.

There are multiple reasons why data may not be optimally aligned, e.g.:

- Data is from sensors at different points in a process-line.
- · Data is from sensors with different response characteristics.
- Data logged with incorrect time stamp or insufficient buffering.

Sometimes smaller alignment issues can be worked around by reducing the timeseries resolution, but often extra insights can be gained if you can work at the highest resolution available...

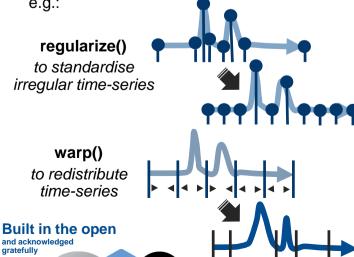
For two unaligned time-series, *ts1* and *ts2*:



There are multiple alignment methods, some better suited to particular applications, and align (hopefully) provides a simple coding structure for implementing, studying and comparing different alignment strategies.

Miscellanea

Although the main align functions expect at least two time-series, some use sub-routines to reshape data that can applied directly. e.g.:



Coding Structure

The main align functions are named [type] align() and intended for use with vectors and elements/columns in data.frames, or object classes converted to either.

General alignment calls: vectors called directly

_align(ts1, ts2, ...)

with data.frames (e.g. **df1** and **df2**)

_align(df1, by=c("ts1", "ts2"), ...) align(df1, df2, by=c("ts1"), ...) align(df1, df2, by=c("ts1", "ts2"), ...)

> by argument identifies columns to be aligned

Catching outputs:

optional argument, output

return <- align(..., output)

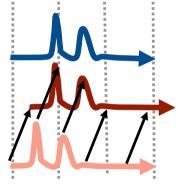
_align()s typically plot fit analysis, report summary to console and return aligned data as a data.frame

NOTE: _align()s are applied to both ts2 and its source data.frame if not also the ts1 source, so they can align data.frames with common (or similar) time-series...

nonto/oblanno in datamamos, or		
Output term	returns	
"plot"	plot(alignment)	
"summary"	summary(alignment)	
"ans"	alignment product, data.frame	
"alignment"	Generic _align() function output; alignment class object	
c(,)	Multiple outputs; any of above; all run but ONLY last returned to catch	

Linear

Linear alignment - applying a fixed offset.



The simplest and least aggressive of the alignments, moves ts2 (and df2) relative to ts1 (and df1), without warping, and pads with NAs.

n_align()

Optional argument n Example (default n=0)

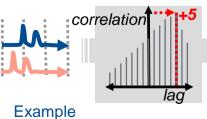
n = 3

like cbind() BUT does not need same or divisible number of columns

moves ts2/df2 forward 5 rows relative to ts1/df1

cor_align()

Like n align() but it uses correlation between ts1 and ts2 to automatically assign n



cor align(..., min.overlap=20)

.optional argument, min.overlap sets the smallest ts1/ts2 overlap, here 20 elements of vector or rows of data.frame

Non-linear

Non-linear alignment - applying a variable offset based on optimising the ts1 and ts2 agreement.

> Arguably the most aggressive and least conservative of the alignment strategies

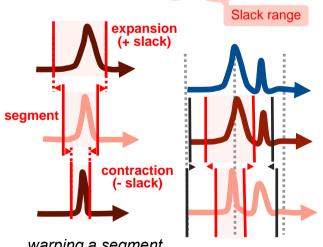
Example:

Correlation optimized warping (COW)

... divides ts2 into a series of subsets (called segments) and concertinas them using an expansion/contraction range (slack) to maximise correlation with ts1

segment width

cow_align(..., seg, slack)



... warping a segment

... and as part of segment series to align ts2 with ts1

NOTE: COW is not the only warping option, other examples include dynamic time warping

Constrained

Constrained alignment - applying an offset, either fixed or variable, based on an assumed relationship between ts1 and ts2 timings.

> Often the preferred option if the nature of offsets are well understood and effects can be mapped from ts1 onto ts2

Example:

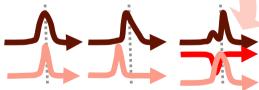
Fitted alignment

The time offset function for ts2/df2

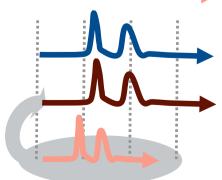
fit align(..., fun, upper, lower)

the upper and lower limits for any constants in fun that require fitting

> transform ts2 using function based on ts2 (or other time-series) and/or diff(ts2), etc



... function applied consistently across ts2 to improve agreement with ts1



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Basics

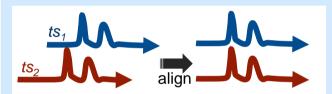
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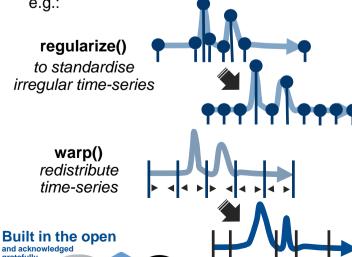
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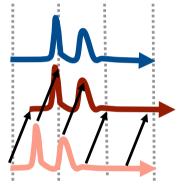
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into/columns in data.names , or		
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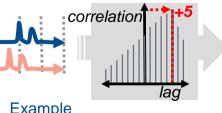
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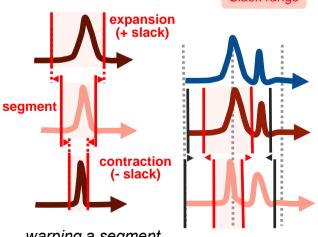
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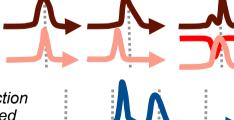
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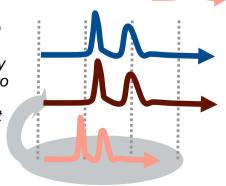
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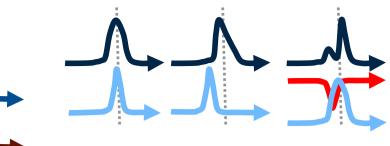
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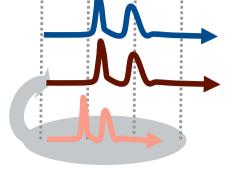


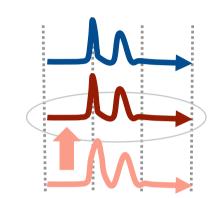
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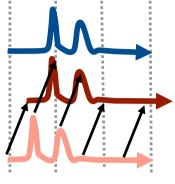


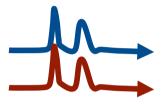


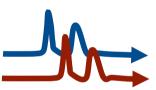


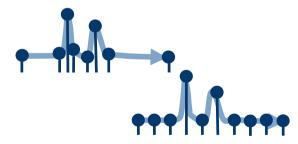


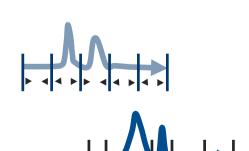


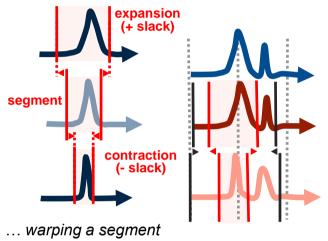




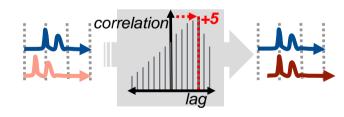


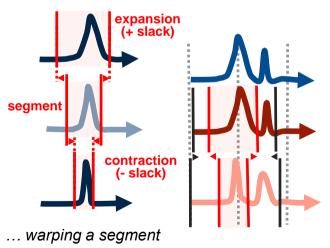






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