

ModulateFlow

Modulating diastolic flow or velocity waveforms

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Abstract.

This report presents **ModulateFlow**, a script for modulating diastolic flow or velocity waveforms.

1. Summary

Clinical blood flow or velocity waveform pre-processing can be difficult due to a low sampling rate (SR) and/or noise. This report describes how **ModulateFlow** deals with these two common problems by (a) using a cubic spline interpolation for low SRs; and (b) modulating the flow or velocity waveform during diastole to eliminate noise after valve closure.

2. Methods

ModulateFlow consists of three stages: (i) waveform interpolation; (ii) noise thresholding; and (iii) downslope projection. The methods used at each stage are now described in turn.

2.1. Waveform interpolation

If the number of waveform datapoints is smaller than 100, a cubic spline interpolation (**spline**) is applied to increase the number of datapoints to 100.

2.2. Noise thresholding

ModulateFlow identifies the maximum flow or velocity datapoint and excludes analyses the waveform from there.

Firstly, the following points are identified on the second derivative (SDPW):

- **a**: The point of maximum value on the SDPW prior to the last local minimum on the SDPW.

2.3. Downslope projection

3. Using ModulateFlow

ModulateFlow can be used to calculate CV indices from either a pulsatile signal containing several pulses, or a single pulse wave. The input data, *S*, should be prepared as a structure with two fields: *S.v*, a vector of signal amplitudes, and *S.fs*, the sampling frequency of the signal. At its simplest, can be called using

```
cv_inds = PulseAnalyse(S);
```

where **cv_inds** is a structure containing individual fields for each of the calculated CV indices (named according to the abbreviations listed in Table ??). Each index's field is itself a structure, containing the calculated value (*e.g.* **cv_inds.SI.v**, which is a median in the case of multiple pulses), and raw values for each pulse (**cv_inds.SI.raw**) if the input signal contains multiple pulses.

Additional functionality can be exploited by specifying additional inputs. Firstly, if the subject's height is provided in *S.ht* then those CV indices which require height will be calculated. Secondly, options can be specified as a second input using

```
cv_inds = PulseAnalyse(S, options);
```

where **options** is a structure containing the following logical fields:

- **exclude_low_quality_data**: whether or not to exclude pulses with a low signal quality in the calculation of a median value for each CV index (only applicable when using an input signal with multiple pulses).
- **do_plot**: whether or not to plot an example pulse with fiducial points annotated (similar to those in Figure ??).

Additional outputs can also be obtained using

```
[cv_inds, fid_pts, pulses, S_filt] = PulseAnalyse(S);
```

where **fid_pts** is a structure containing the indices of each fiducial point; **pulses** is a structure containing the indices of the onsets and peaks of each pulse, and the signal quality of each pulse (a logical where 1 indicates high signal quality); and **S_filt** is the filtered pulsatile signal to which these indices correspond.

4. Further Work

Ideas for improvement.

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