



### Centred vs Uncentred r

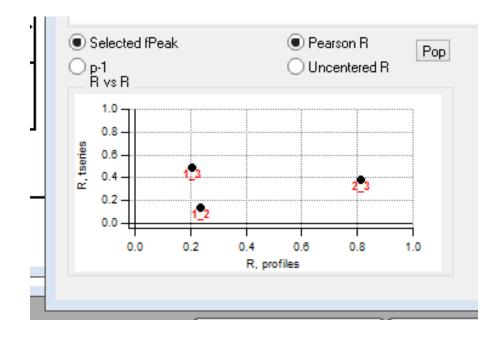
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### Motivation

- Frequently need to compare mass spectra,
  e.g. PMF output with library spectra
- Need a metric that is a good discriminator







## Generic unweighted closeness metric

- Data vectors A and B of length n
- Generate closeness metric S

$$S(A,B) = \frac{\sum_{i=1}^{i=n} (A_i - A')(B_i - B')}{\sqrt{\sum_{i=1}^{i=n} (A_i - A')^2 \sum_{i=1}^{i=n} (B_i - B')^2}}$$

• For *r*:

$$A' = \bar{A} = n^{-1}$$
 (if normalised)

• For uncentred *r*:

$$A'=0$$

– AKA normalised dot product:

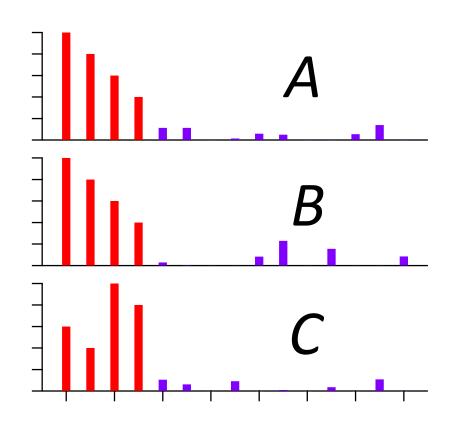
$$S(A,B) = \frac{A \cdot B}{\sqrt{(A \cdot A)(B \cdot B)}}$$

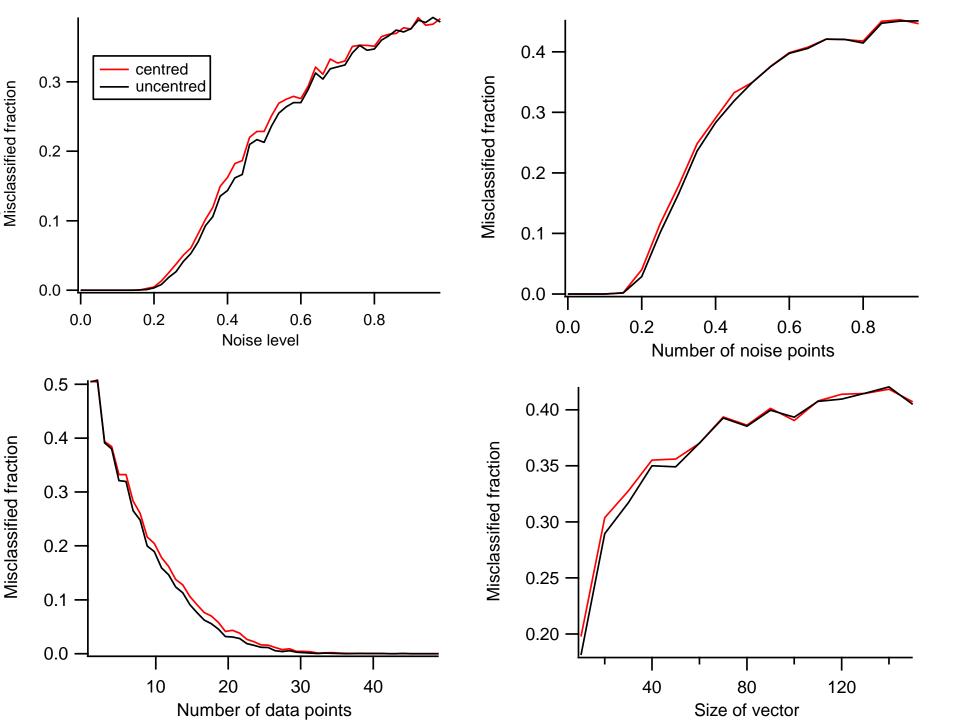




# **Testing Metrics**

- Generate three vectors of synthetic data
  - Part 'real', part 'noise'
- S(A,B) should always
  be greater than S(A,C)
- Repeat multiple times with different random data, record fraction of misclassified results











### Conclusions

- Uncentred r is more mathematically sound and performs better than centred r for comparing mass spectra
  - But not by much

 Centred r remains better for time series as the mean of a time series is meaningful

