

SUPPLEMENTAL MATERIAL

An adaptive and fully-automated baseline correction method for Raman spectroscopy based on morphological operations and mollification

Hao Chen^{1,2}, Weiliang Xu^{1,2,*}, and Neil G. R. Broderick^{3,2}

*Corresponding Author: Weiliang Xu, email: p.xu@auckland.ac.nz

¹Department of Mechanical Engineering, the University of Auckland, Auckland 1010, New Zealand

²The Dodd-Walls Centre for Photonic and Quantum Technologies, PO Box 56, Dunedin 9054, New Zealand

³Department of Physics, the University of Auckland, Auckland 1010, New Zealand

The four types of baselines described in the paper are generated using the following equations:

1) linear baseline:

$$baseline_l = 0.5 \times r, \quad r \in [0, 2000]$$

2) sine baseline:

$$baseline_{sin} = 1000 \times \sin\left(\frac{(r+1000)\pi}{1000}\right), \quad r \in [0, 2000]$$

3) sigmoidal baseline:

$$baseline_{sig} = \frac{1000}{1 + e^{-0.03(r-1000)}}, \quad r \in [0, 2000]$$

4) 4th order polynomial baseline:

$$baseline_p = 80.5 + 0.001519 \times r + 1.6625 \times 10^{-5} \times r^2 + 6.39 \times 10^{-9} \times r^3 - 4.6105 \times 10^{-12} \times r^4, \quad r \in [0, 2000]$$

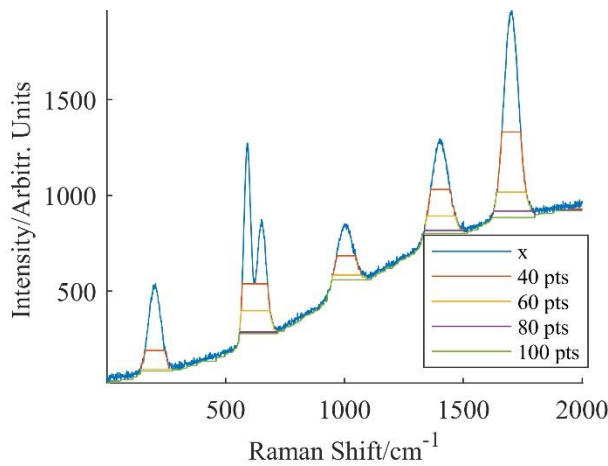


Figure S1 The influence of selected structuring element size on the opening operation

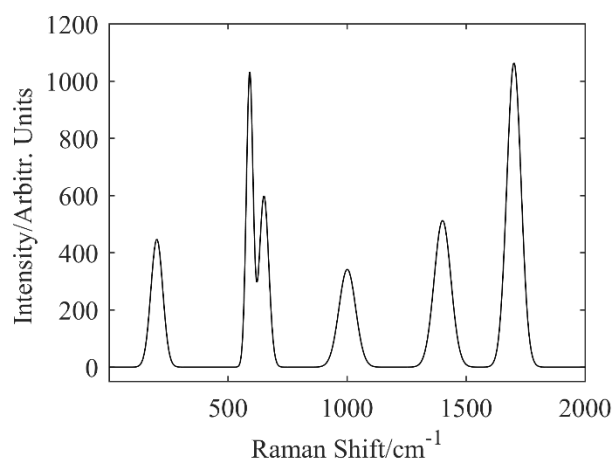


Figure S2 The pure Raman signal generated with Gaussian peaks.