

AFModulus_Flex

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1/11/2021

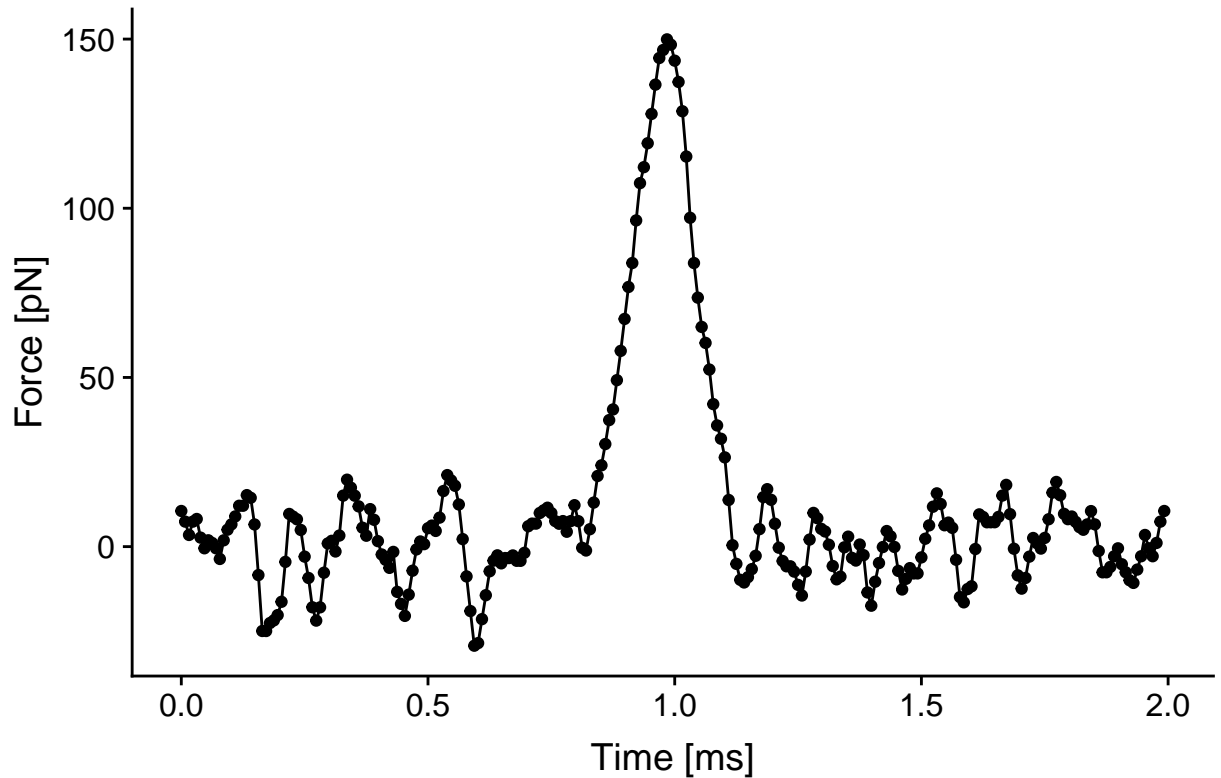
Atomic Force Microscopy - From Topology to Stiffness (Modulus)

The git page of this project can be found here https://github.com/audreyyeoCH/AFModulus_Flex.

Import of AFM curves (force vs. separation distance) for each pixel

Plot 1 curve

is/20200619_.005.pfc-4069_ForceCurveIndex_45647.spm – NanoScope



We will use the force signal between time 0.0 - 0.5 ms as well as 1.5 - 2.0 ms as **baseline**.

We will use a sliding window approach to approximate the **gradient** of the linear slope within the time 0.5 - 1.2 ms.

Extract maximal force (F max) from each graph

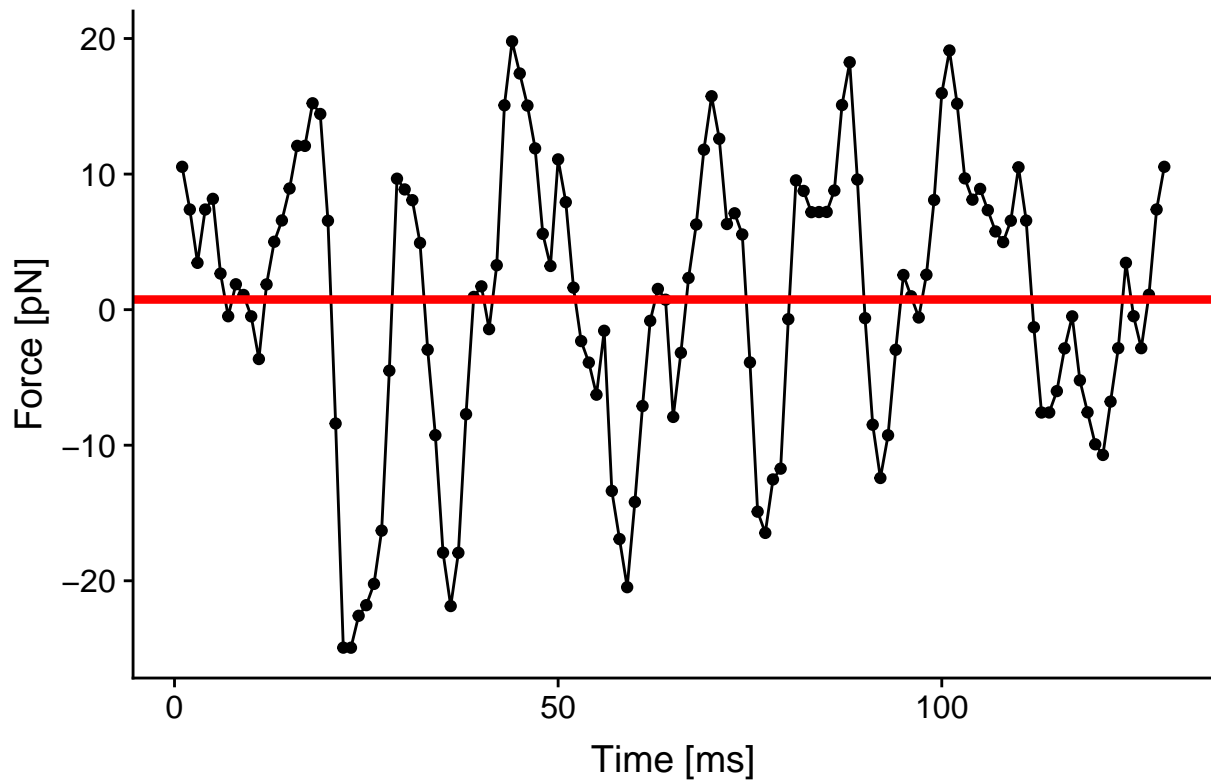
F max = 149.9152, the time of F max = 1.99219.

Compute touching point

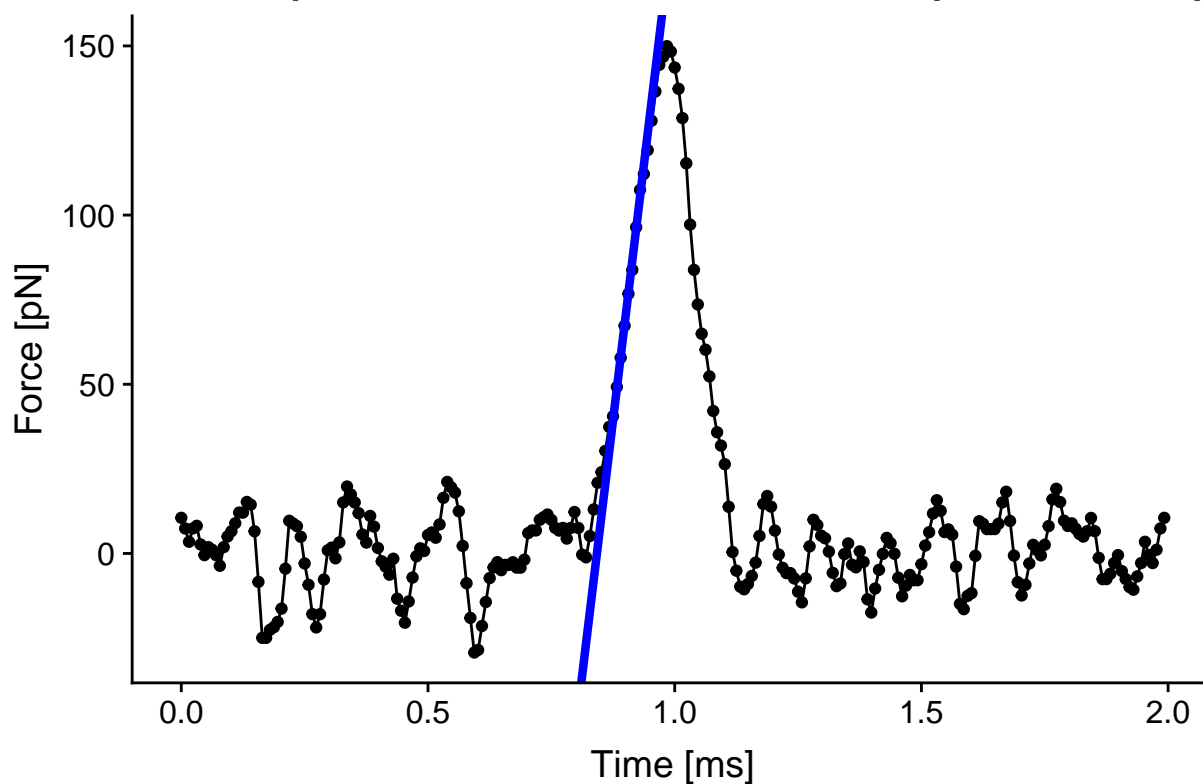
Intersect between baseline and linear gradient

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- We will use a sliding window approach to approximate the **gradient** of the linear slope within the time 0.5 - 1.2 ms.

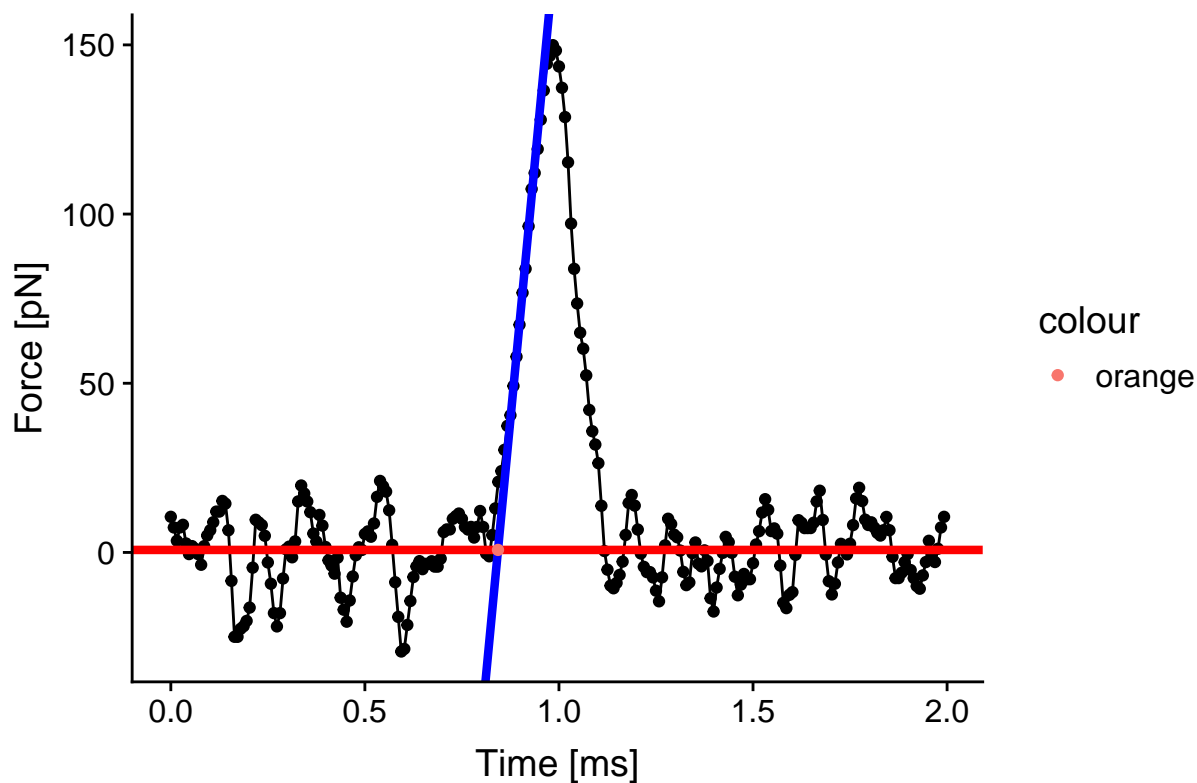
is/20200619_.005.pfc-4069_ForceCurveIndex_45647.spm – NanoScope



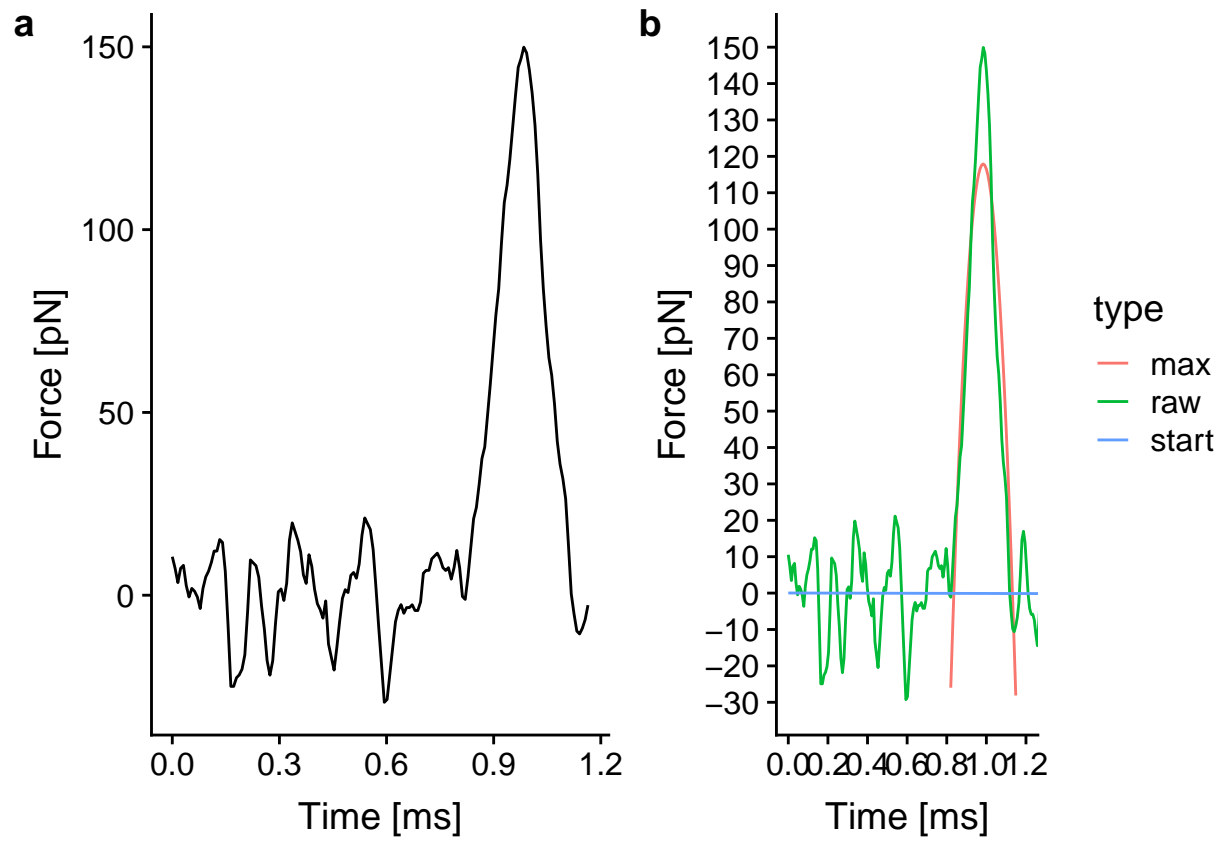
is/20200619_.005.pfc-4069_ForceCurveIndex_45647.spm – NanoScope

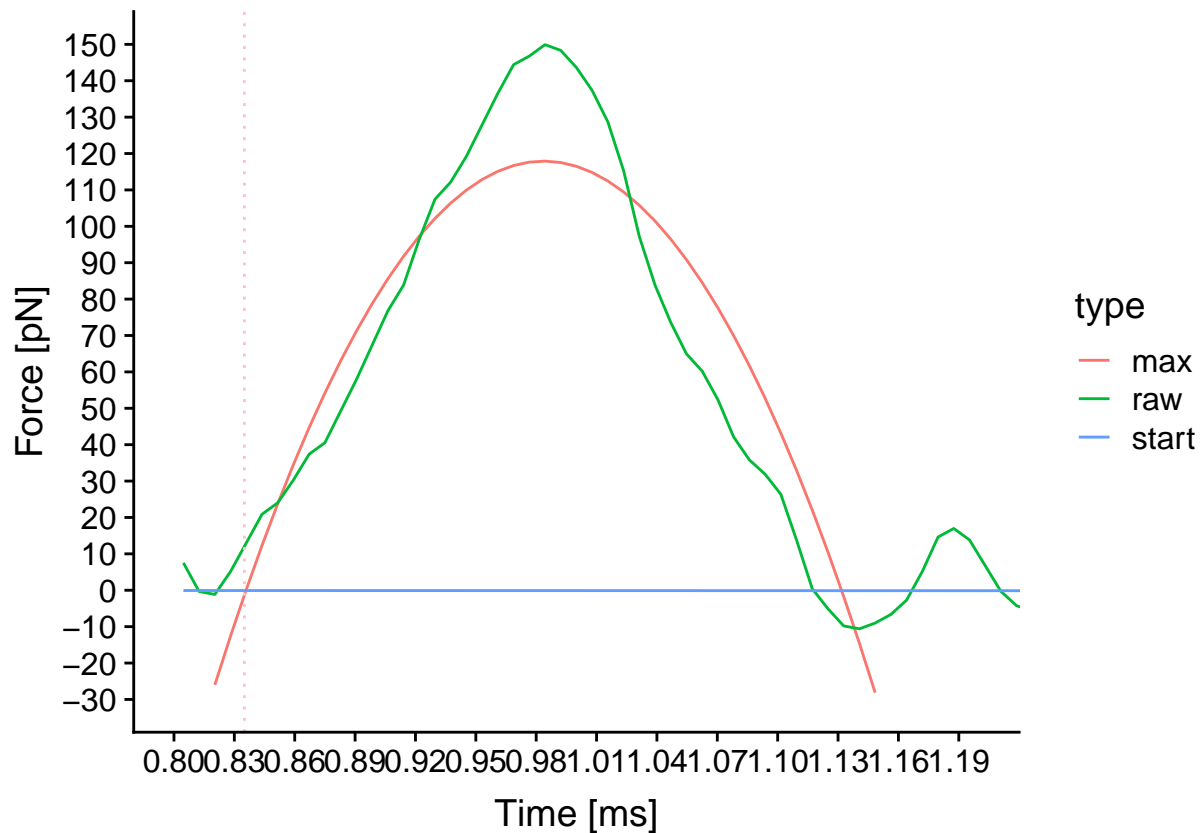


0619_.005.pfc-4069_ForceCurveIndex_45647.spm – NanoScope Analy



Intersect between baseline and parabolic fit





Mean of error increase from baseline (= start of adhesion dent) and error from linear gradient (= end of adhesion dent)

Compute indentation depth (d) from touching point for each pixel

The indentation depth $d = 1.1492608$

Compute modulus (= stiffness, E) for each pixel from F-max and d

Print picture of stiffness (next to topology picture)

```
{python pic, echo=FALSE} #B = r.A + 1 # #
```

Error propagation/ sensitivity analysis of the modulus

```
{r error, echo=FALSE} #C <- py$B + 1 # #
```

Plot topology against modulus? Can this detect 'antibiotics affected areas'?