LocText
HELLO!

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A learning algorithm that extracts the semantic relationship between proteins and their subcellular localization from scientific literature.
What is the status? (LocText)

- Docker: Runs STRING Tagger
  - Input: Scientific literature
  - Output: Proteins and compartment
- Extract localization relationships between proteins and compartments.

TP53 R72P polymorphism modulates DNA methylation in hepatocellular carcinoma.
**LocText**: Ultimate goal

- Helps in the prediction of as many annotations as possible.
- Publicly available

[GitHub](https://github.com/Rostlab/nalaf)

- Open source
THANKS!

Any questions?
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