### Multi-omics abundance matrices as input

```
Omics1
Feature1 ...
Sample1 0.9 ...
```

#### <u>Omics2</u>

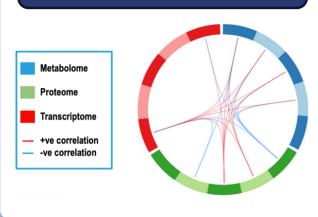
Feature1 .. Sample1 21.3 ..

...

# Discover regulatory features with genomeNLP



## Identifying correlations across multi-omics data



## Overlay functional and regulatory omics signatures

