# In-silico identification of Tyrosine nitration sites in protein peptide sequences

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Thesis motivation
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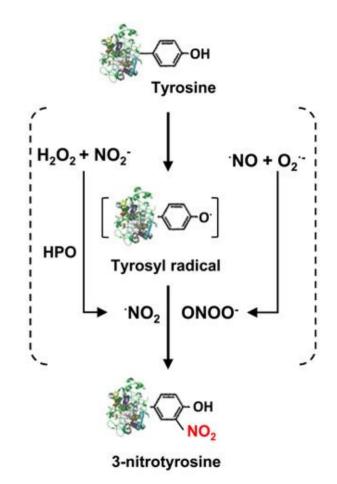
#### **Post-translational modification in Proteins**

- Proteins are biomolecules/macromolecules containing long chains of amino acid residues
  - Perform critical functions within organisms, such as catalyzing metabolic reactions, DNA replication, responding to stimuli, providing structure to cells and organisms, and transporting molecules.
- Protein biosynthesis is a core biological process occurring inside cells
  - Balances the loss of cellular proteins in cells by producing new proteins
  - Occurs in two stages:
    - **Transcription:** A section of DNA encoding a protein (gene), is converted into a messenger RNA (mRNA).
    - Translation: The nucleotide sequence of mRNA is read by ribosomes to determine the sequence of amino acids in the resulting Protein
- Post-translational modification (PTM) is the modification of proteins after protein biosynthesis.
  - Influences normal cell biology and pathogenesis, and increase the functional diversity of the proteome
  - Modifications: covalent addition, proteolytic cleavage of regulatory subunits, or degradation of entire proteins.
  - ~460 different types of PTMs have been identified
- Identifying and understanding PTMs is critical for studying cell biology, and disease treatment or prevention.



## **Tyrosine nitration**

- Nitrotyrosine is the product of Tyrosine nitration
  - It is a covalent Post-translational modification (PTM) in proteins
  - An irreversible nitrative modification of the tyrosine residue of a protein and permanently alters the structure of the protein
  - Result of the substitution of a hydrogen by a nitro group (-NO<sub>2</sub>) at the ortho position of the phenolic ring of tyrosine
    - Two nitrating agents account for the nitration, i.e., **peroxynitrite** (ONOO $\bar{}$ ), and **hemoperoxidases** (HPO) in the presence of hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) and nitrite (NO<sub>2</sub> $\bar{}$ )
  - Tyrosine nitration is selective, and not a random process
    - Depending on the accessibility of the tyrosine residues to the nitrating agents (e.g., tyrosine residues exposed on the surface of proteins can become target)
    - Mostly, the nitrated tyrosine residues are in the vicinity of a site which is generating nitrating agents



## **Cause and effect of Tyrosine nitration**

#### Causes:

- Determined by several factors, such as reactive **species**, reaction kinetics, presence of **antioxidants** and **radical scavengers**, and **compartmentalization**
- Two pathways for nitration:
  - When Nitric Oxide (NO<sup>-</sup>) is generated in high concentration, in the presence of Superoxide anion (O<sub>2</sub><sup>-</sup>) it will lead to the rapid formation of peroxynitrite (ONOO<sup>-</sup>), which causes nitration
  - Peroxynitrite-independent nitration depends on the presence of transition metals such as iron and copper, through either peroxidase-dependent or Fenton-dependent pathways in the presence of hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) and nitrite (NO<sub>2</sub><sup>-</sup>)

#### Effects:

- Substantial changes in the biological function of proteins
- The structural alteration leads to loss or augmentation of protein function
  - Can render a protein inactive
- Inhibits protein phosphorylation that impedes the normal transduction pathways in cell signaling
- Functional loss of proteins in inflammatory diseases
- High correlation with pathogenesis of diseases
- Marker of NO-dependent, reactive nitrogen species (RNS) induced nitrative stress



## **Benefits of identifying Tyrosine nitration sites**

- Causal nitration pathway inducing agents identified in various disease conditions:
  - Nitric Oxide (NO) is produced at a high rate in inflammatory, stimuli-induced conditions
  - Nitrite ion (NO<sub>2</sub><sup>-</sup>) is greatly increased in systemic inflammatory disorders (sepsis, gastroenteritis, & hemolytic diseases)
  - Abnormal elevation of copper ion (Cu<sup>2+</sup>) and free heme in the pathogenesis of type 2 diabetes mellitus, neurological disorders, and severe hemolytic diseases
- Tyrosine nitration is identified in large number of pathological conditions
  - Neurodegenerative diseases Parkinson's and Alzheimer's, degeneration of dopamine neurons, cerebral ischemia and edema
  - Cardiovascular diseases
  - Autoimmune diseases Rheumatoid Arthritis, Systemic Lupus Erythematosus
  - Carcinogenesis Breast, Esophageal and Gastric cancer; Colorectal, Squamous cell, Adeno- and Cholangial carcinoma
- Therefore, **effective interception of protein nitration** can represent novel and critical points of **therapeutic intervention in diseases** associated with protein nitration.
  - Identifying onset and progression of the associated disease
  - Surrogate markers for the design of clinical interventions (therapeutic strategies and drugs)



### Motivation for computational methods

- PTMs of proteins experimentally detected and recorded by a variety of techniques
  - **Experiments**: Immunohistochemical analysis, Chromatography, Mass spectrometry with prior immunoprecipitation, Eastern and Western blotting
  - Datastores: PhosphoSitePlus, ProteomeScout, Human Protein Reference Database, PROSITE, Protein Information Resource (PIR), dbPTM, Uniprot, O-GlcNAc Database
- Huge amounts of data available for knowledge discovery
  - Resulted in increasing use of Machine Learning and Deep Learning approaches
    - Wet lab experimental approaches:
      - Technically challenging with Theoretical limitations
      - Labor intensive
      - Requires skilled laboratory experience
      - Time consuming
      - Expensive
      - Biases in proteome wide identification
      - Generates ground truth data

#### Computational approaches:

- Easier development
- Comparatively, much less labor intensive
- Requires coding skills
- Reusability saves time
- Economical
- Ability to balance biases in imbalanced scenarios
- Requires wet-lab experiments to generate data



# Thank you for your time! Questions?



### References

1. Radi R. (2013). Protein tyrosine nitration: biochemical mechanisms and structural basis of functional effects. *Accounts of chemical research*, *46*(2), 550–559. https://doi.org/10.1021/ar300234c