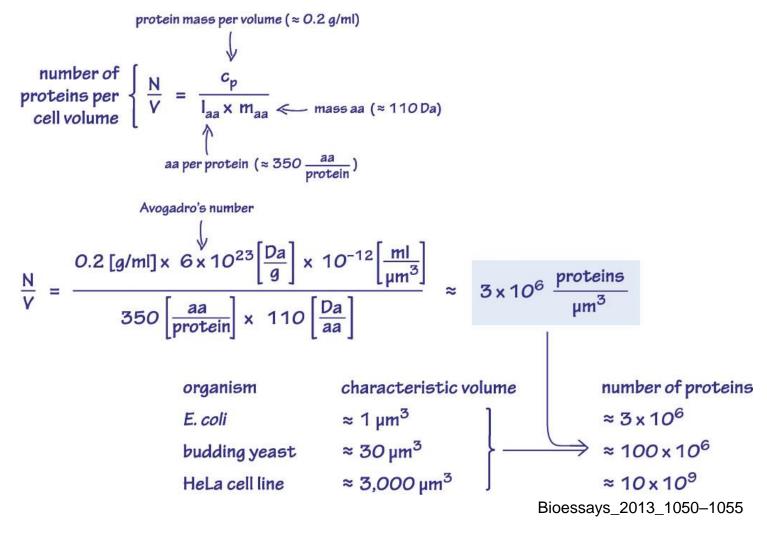
# Proximity Labeling: A Powerful Tool for Protein Complex Purification and Proteomic Mapping

Liu Wenjuan 2021.03.04

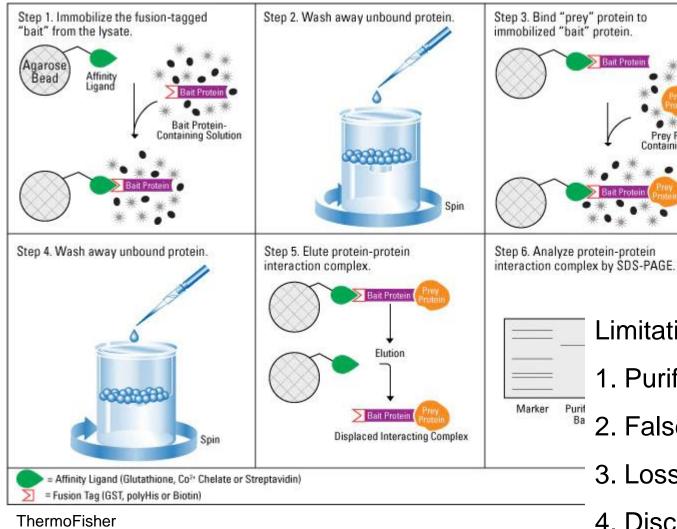
#### estimating the number of proteins per cell volume



- 1. Subcellular regions located
- 2. Interactional
- → Identification of interacting proteins within spatially defined cellular domains is key

#### Traditional:

#### Antibody based affinity purification and mass spectrometry approaches



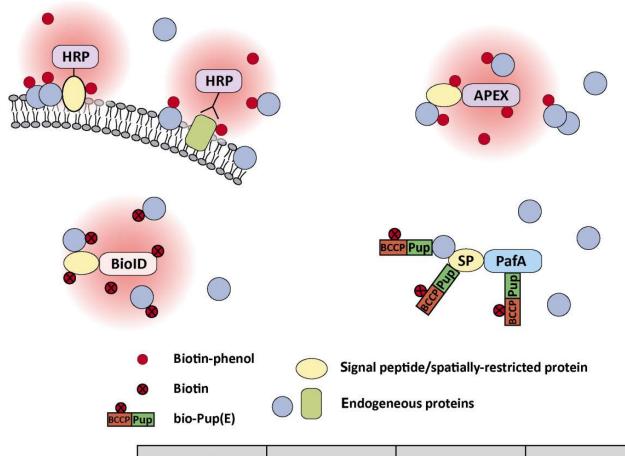
#### Limitation:

- 1. Purification methods
- 2. False positives introduced by cellular disruption
- 3. Loss of components caused by disruption
- 4. Discrete cellular regions cannot be purified by centrifugation

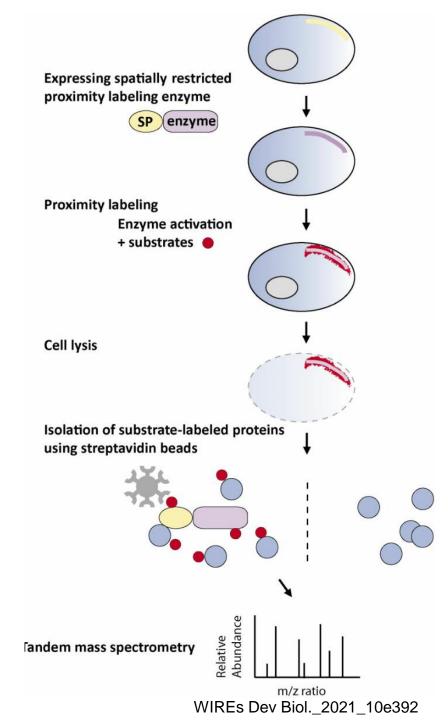
Now:

**Proximity labeling** 

Enzymes catalyze the conversion of a substrate into a reactive radical that covalently tags neighboring proteins

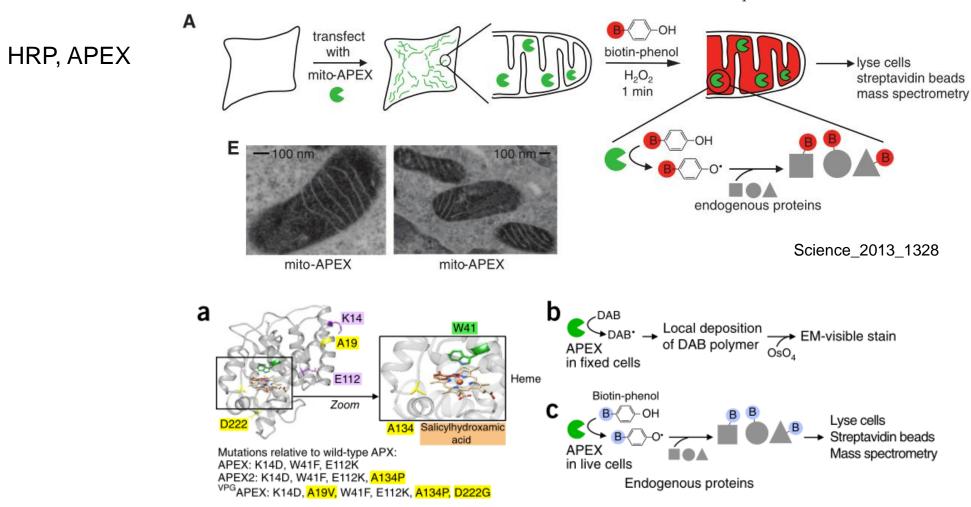


	HRP	APEX	BioID	Pup-IT
Enzymatic Activity	Peroxidase	Peroxidase	Biotin ligase	Pup ligase



### Key Role: Ascorbate peroxidase

基于过氧化物酶催化二氨基联苯胺(Diaminobenzidine, DAB)染色标记方法是在过氧化氢( $H_2O_2$ )存在的条件下,过氧化物酶可以催化DAB形成DAB 聚合物,进而被 $OsO_4$ 染色实现标记。



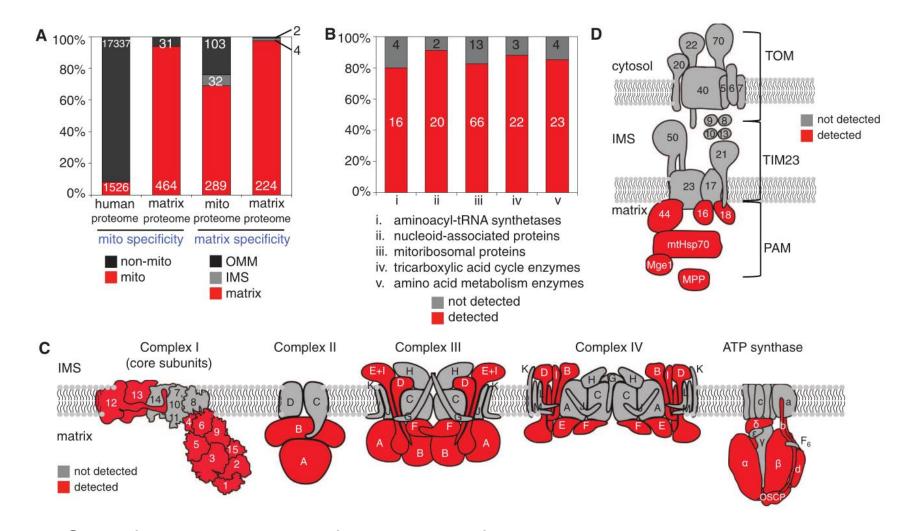
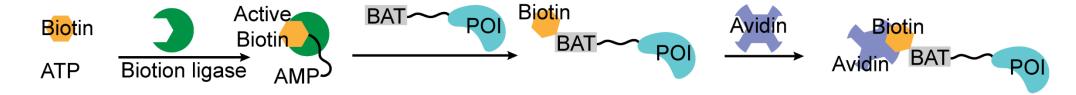


Fig. Specificity and depth of coverage of the mitochondrial matrix proteome.

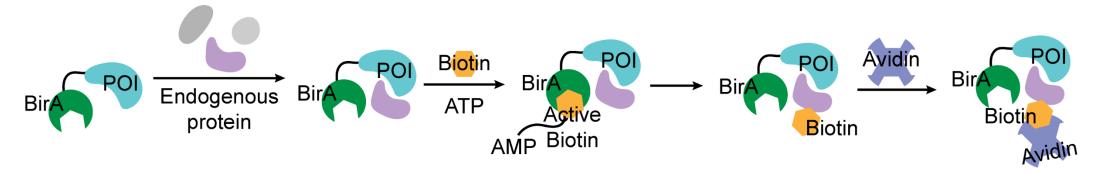
#### Limitation:

- 1. Low sensitivity
- 2. Toxicity of Biotin-phenol reagent

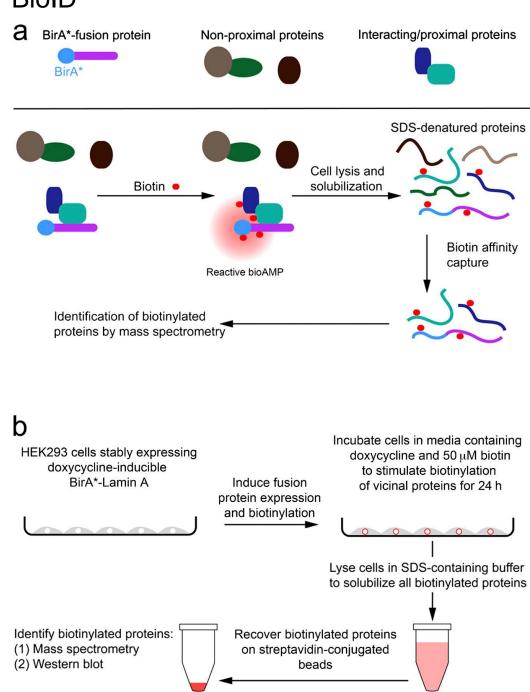
# Key Role: Biotin ligase



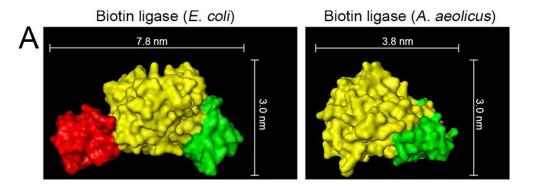
# BioID



#### **BioID**

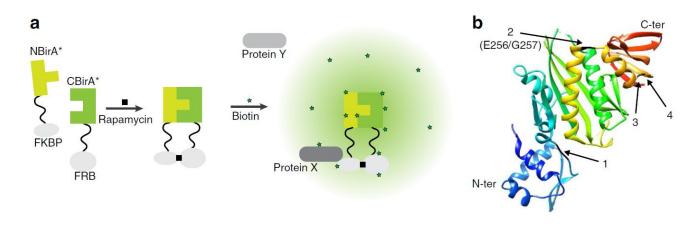


#### BioID2



Molecular Biology of the Cell\_2016\_1188

### Split-BioID

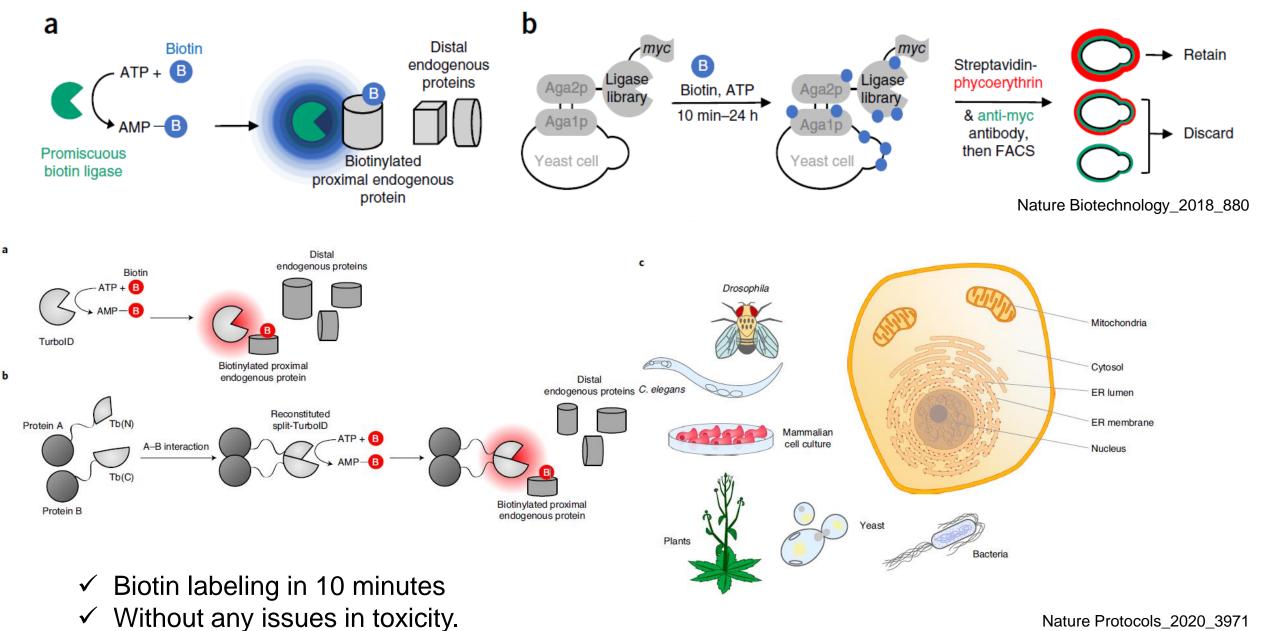


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# Limitation: Long time needed

J. Cell Biol.\_2012\_801

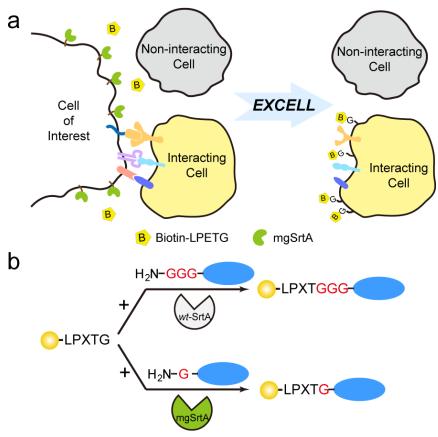
### TurboID and split-TurboID



# RaPID

#### a Streptavidin (SA) capture SA beads BoxB BoxB RNA motif BoxB RNA motif BoxB b \*\*\*\* C RaPID LC-MS/MS 60 Western blot LC-MS/MS CELF1 50 -1.0 -Cutoff 0.9 Hold enrichment 20 20 10 SAINT score 30 0 10 20 40 Fold change bio-RNA RaPID

# **EXCELL**

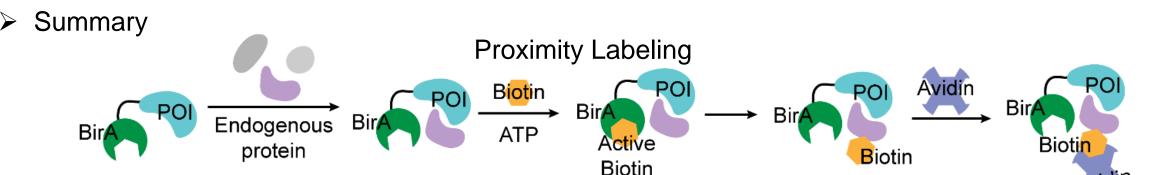


NATURE METHODS\_2018\_207

J. Am. Chem. Soc.\_2019\_1833

# > Summary

	HRP	APEX	BioID
Enzymatic Activity Peroxidase		Peroxidase	Biotin ligase
Labeling Target Electron-rich amino acids		Tryrosine and potentially other electron-rich amino acids	Lysine
Size	44 kD	27 kD	BioID, TurboID: 35kD BASU: 29kD miniTurbo: 28kD BioID2: 27kD
Labeling time	5-10min	1min	BioID:15-24hr TurboID: 10min
Incubation time with substrate 5-10min		30-60min	BioID:15-24hr TurboID: 10min
Activation by H <sub>2</sub> O <sub>2</sub>	Activation by H <sub>2</sub> O <sub>2</sub> Yes		No
Substrates for protein labeling biotin-phenol (and biotin- or fluorescein-acylazide)		biotin-phenol	biotin
Half-life of generated radicals	<1me		mins
Active region	extracellular, secretory pathway (inactive in cytosol)	intracellular	intracellular
Notes	Can be used as EM tag; HRP-conjugated antibodies available	Can be used as EM tag	BioID: reduced activity below 37°C TurboID: evolved in yeast at 30°C



Research Project: Interacting proteins, protein and RNA, cells

Key Factor: Enzyme catalysis

Suitable object: Live cells

Labeling Time: 5-10 min

Combination Techniques: LC/MS/MS, SDS-PAGE, Western Blot, Fluorescence image, EM Affinity labeling

Cell Phenylsulfonate linkage LDT reagent Lgand Protein

Endogenous protein

Tosyl chemistry on protein surface Protein

Functionalized endogenous protein

Research Project: Interested proteins, organelles

Key Factor: Substrate

Suitable object: Live cells

Labeling Time: Several minutes to hours

Combination Techniques: LC/MS/MS, SDS-PAGE, Western

Blot, Fluorescence image