



Stockholm  
University

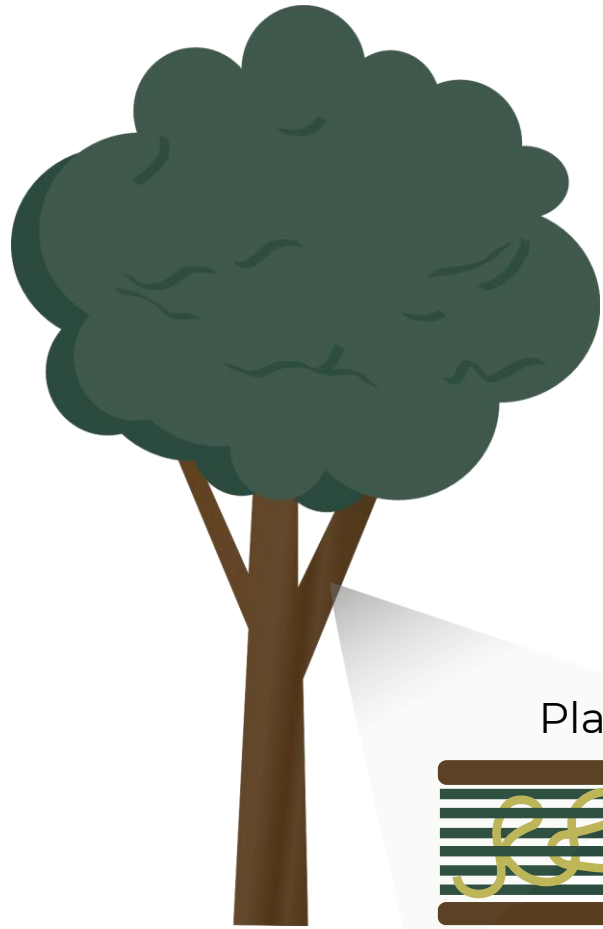
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# Unravelling lignin structure with mass spectrometry and generative modelling

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*The authors declare no competing financial interest.*

# LIGNIN



Plant cell wall



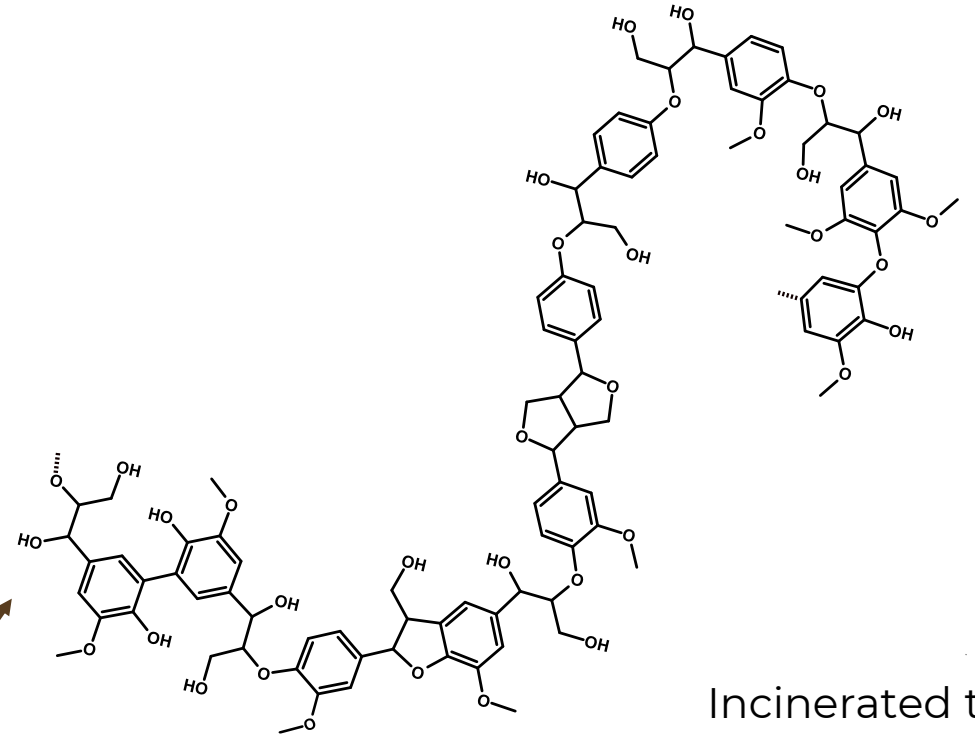
Lignin

Hemicellulose

Cellulose

Paper, textile,  
food, and  
pharmaceutical  
industries

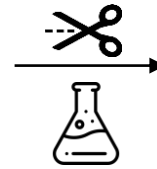
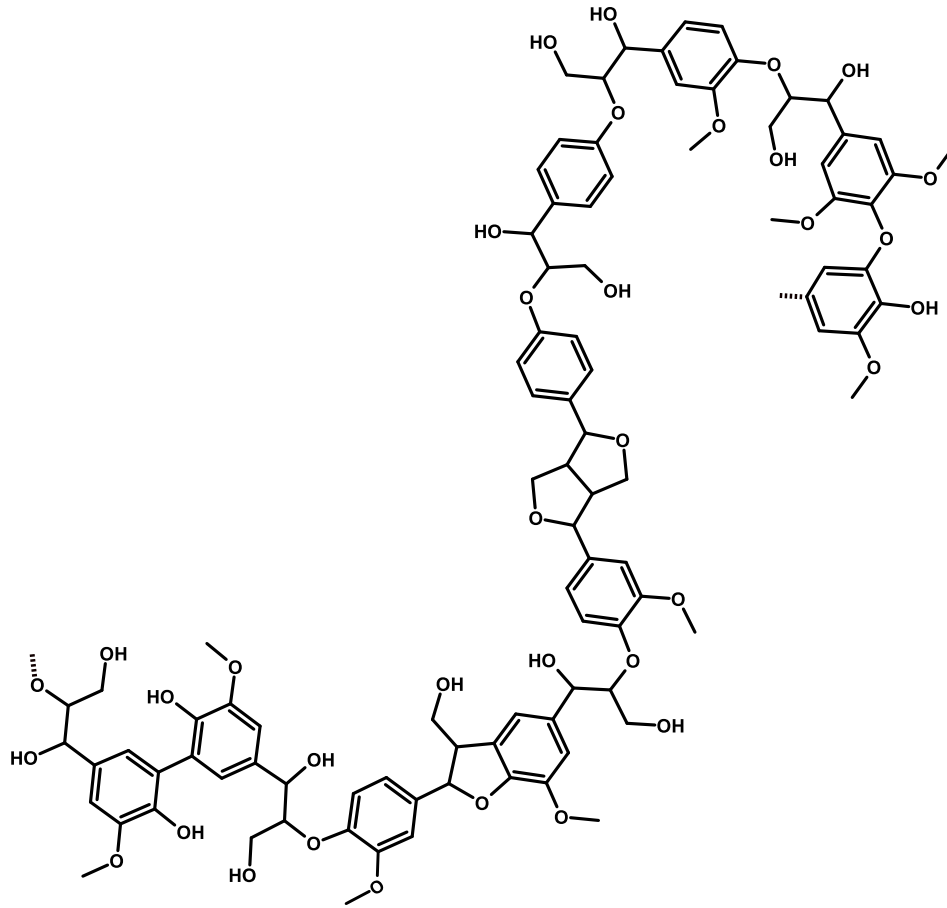
...



- Incinerated to recover chemicals
- produce energy



# LIGNIN



Extraction of compounds



Biofuel

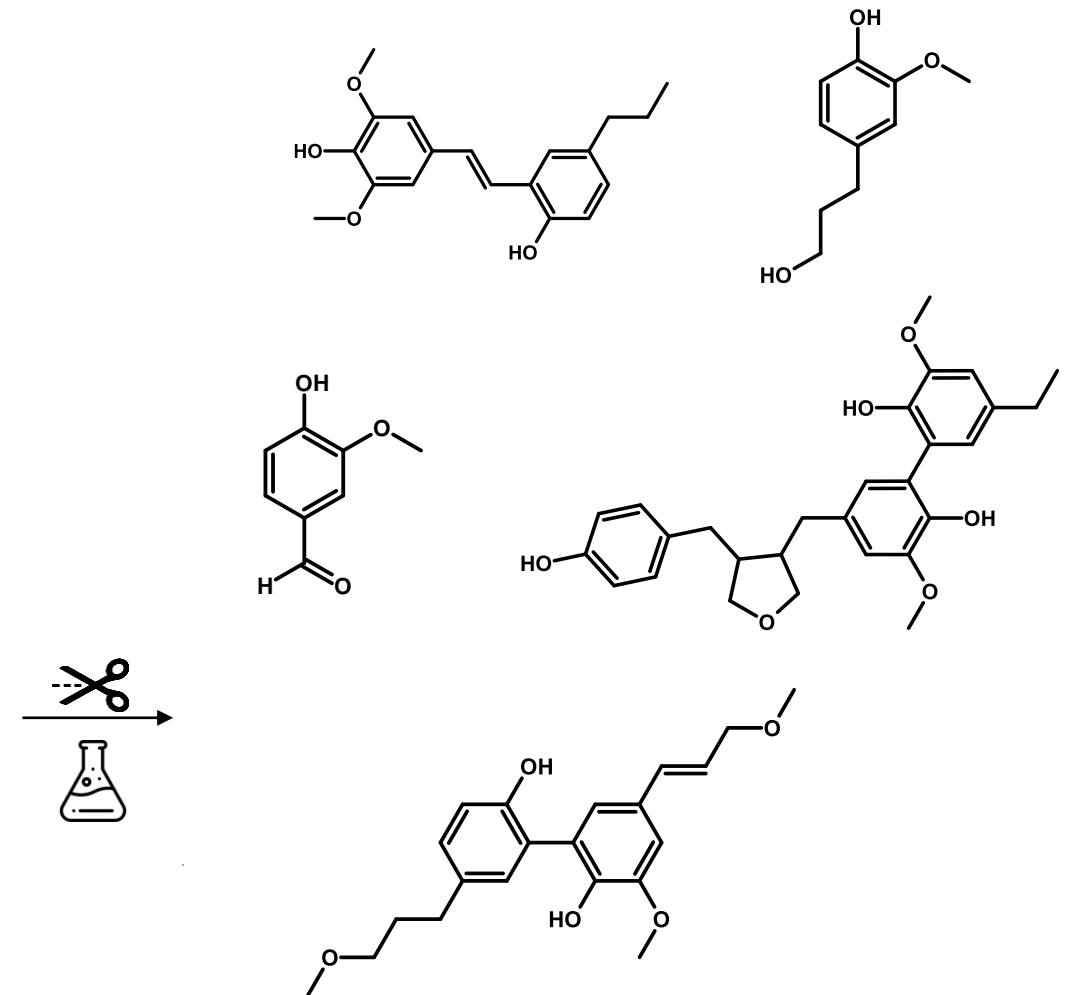
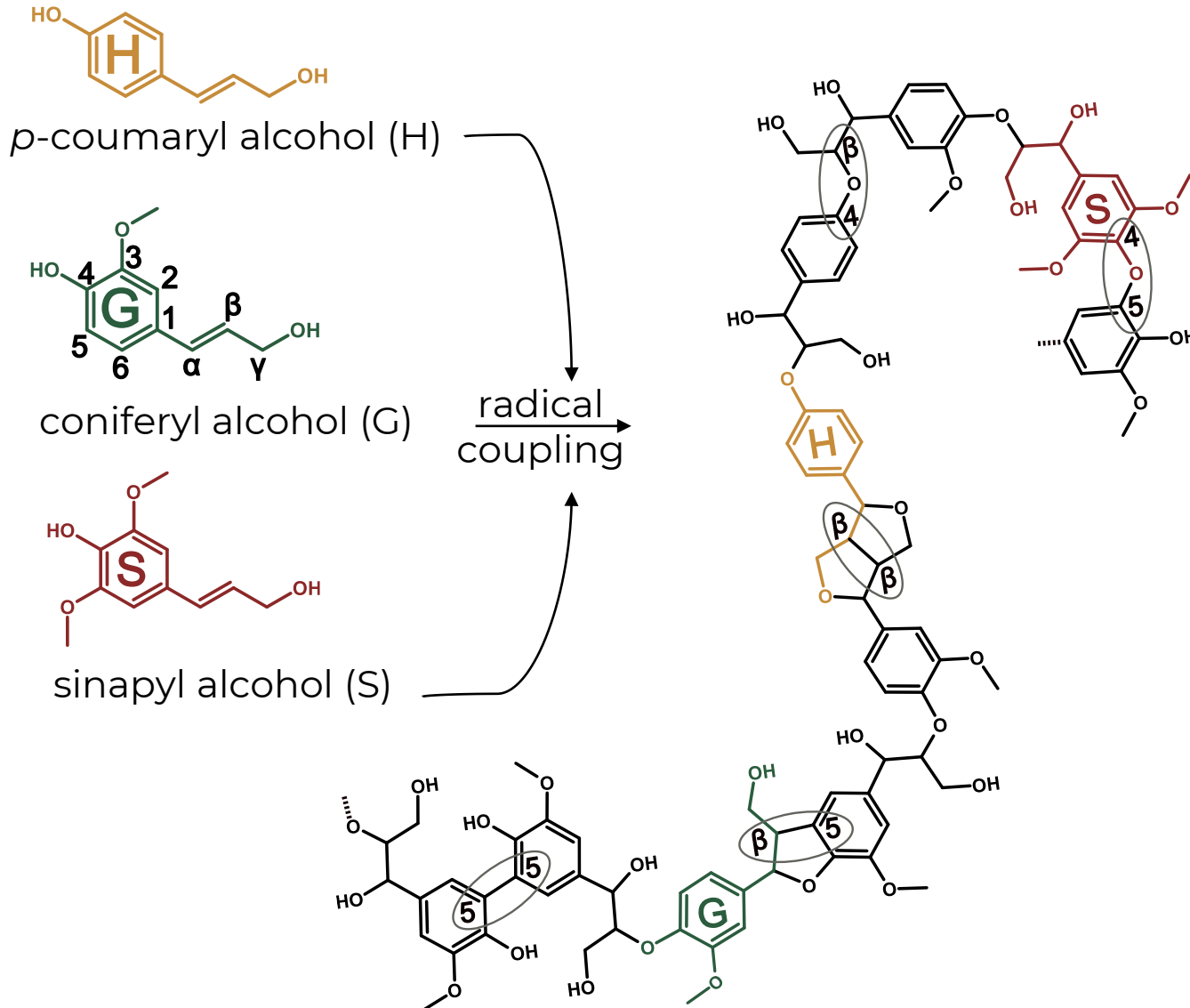


Biomaterials

- The most abundant natural aromatic material

- Around ~2% is commercialised
- Develop new valorisation ways

# LIGNIN



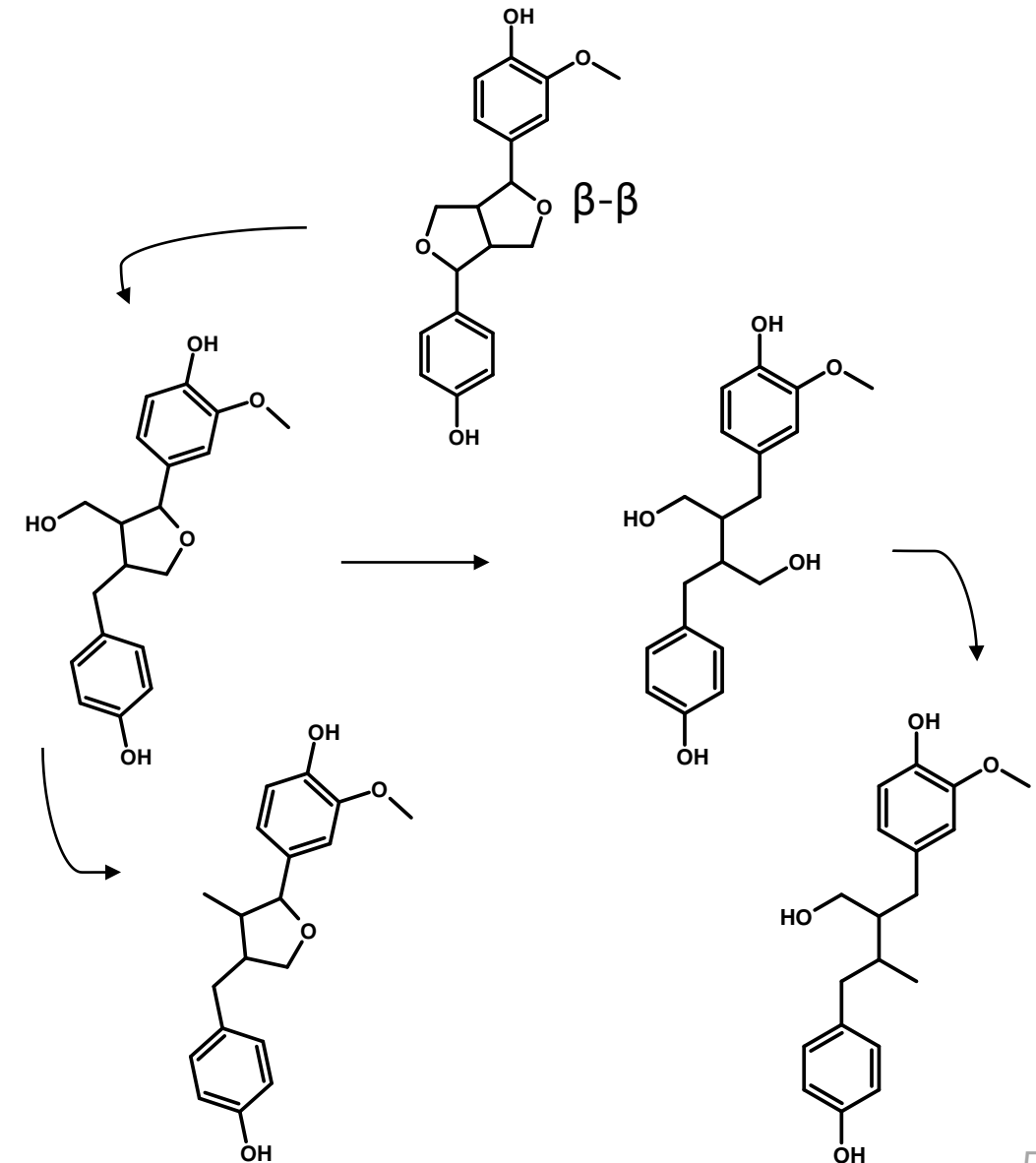
- Mixture containing hundreds of compounds
- Structurally highly similar compounds (including numerous isomers)

# REDUCTIVE CATALYTIC FRACTIONATION (RCF)

- A “lignin-first” extraction and depolymerisation method
- The occurring reactions are quite well known

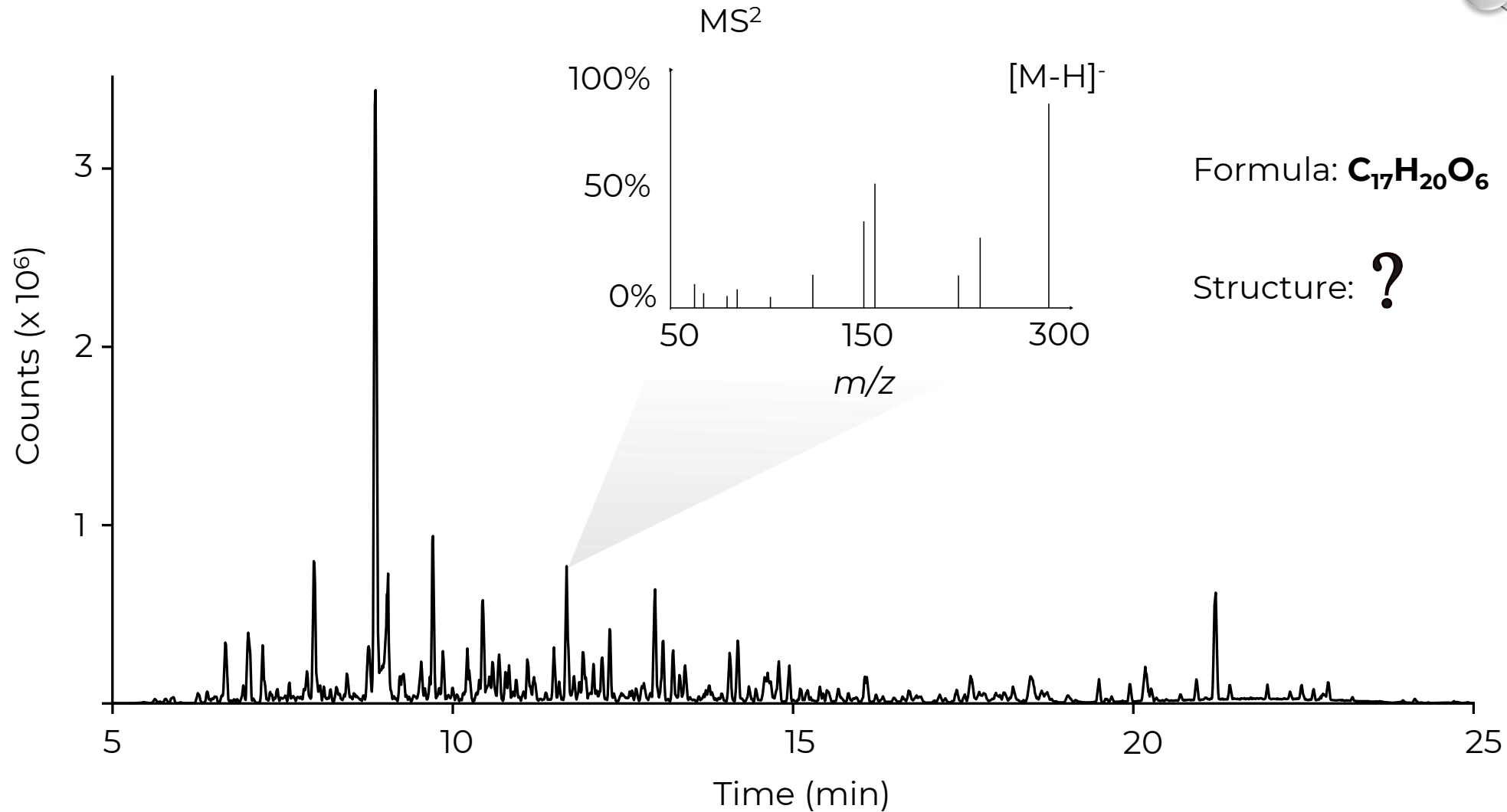
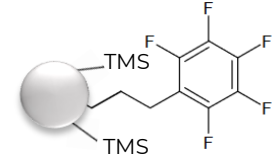
Reactions occurring under reductive conditions

Reaction	Scheme
hydrogenation	$R_1-\text{CH}=\text{CH}_2 \xrightarrow{\text{H}_2} R_1-\text{CH}_2\text{CH}_3$
hydrogenolysis	$R_1-\text{O}-\text{CH}_2-\text{R}_2 \xrightarrow{\text{H}_2} R_1-\text{OH} + \text{CH}_3-\text{R}_2$
decarboxylation	$R_1-\text{COOH} \xrightarrow{\text{H}_2} R_1-\text{H} + \text{CO}_2$
demethoxylation	$R_1-\text{OCH}_3 \xrightarrow{\text{H}_2} R_1-\text{H} + \text{CH}_3-\text{OH}$
demethylation	$R_1-\text{OCH}_3 \xrightarrow{\text{H}_2} R_1-\text{OH} + \text{CH}_4$
hydrodeoxygenation	$R_1-\text{OH} \xrightarrow{\text{H}_2} R_1-\text{H} + \text{H}_2\text{O}$

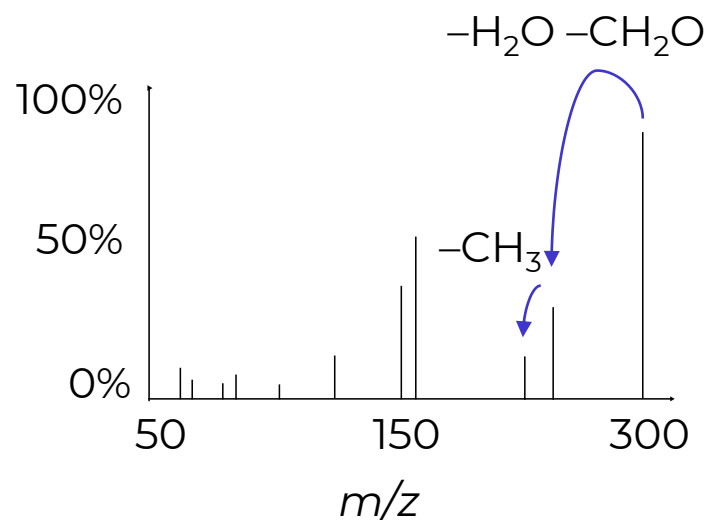


# RCF LIGNIN ANALYSIS WITH LC/HRMS

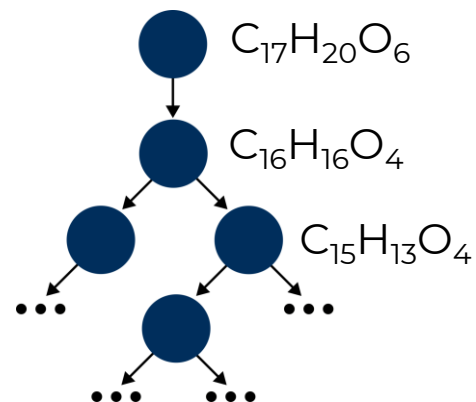
Column: Kinetex **F5** Core-shell



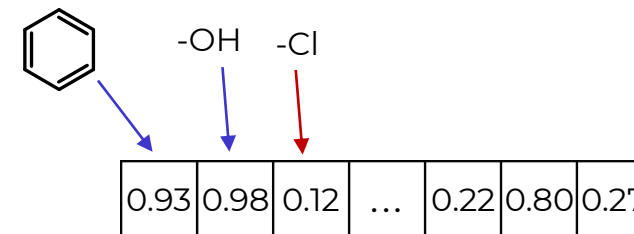
# SIRIUS+CSI:FINGERID



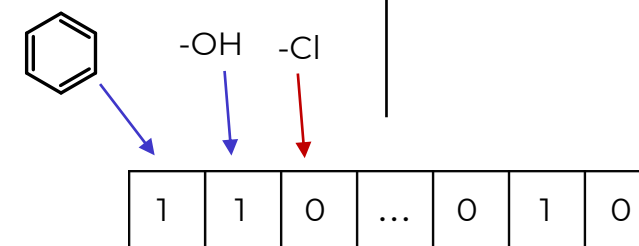
Fragmentation tree



Probabilistic fingerprint

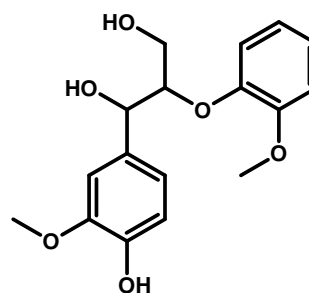
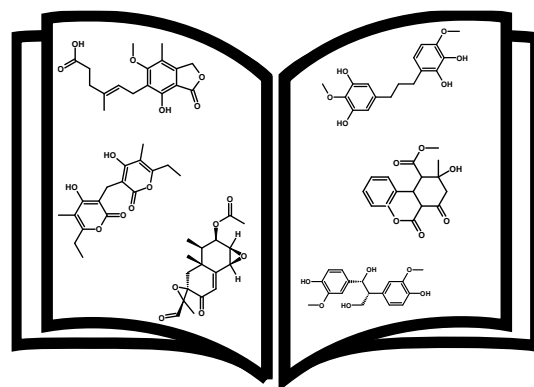


★ Top 1 match ★



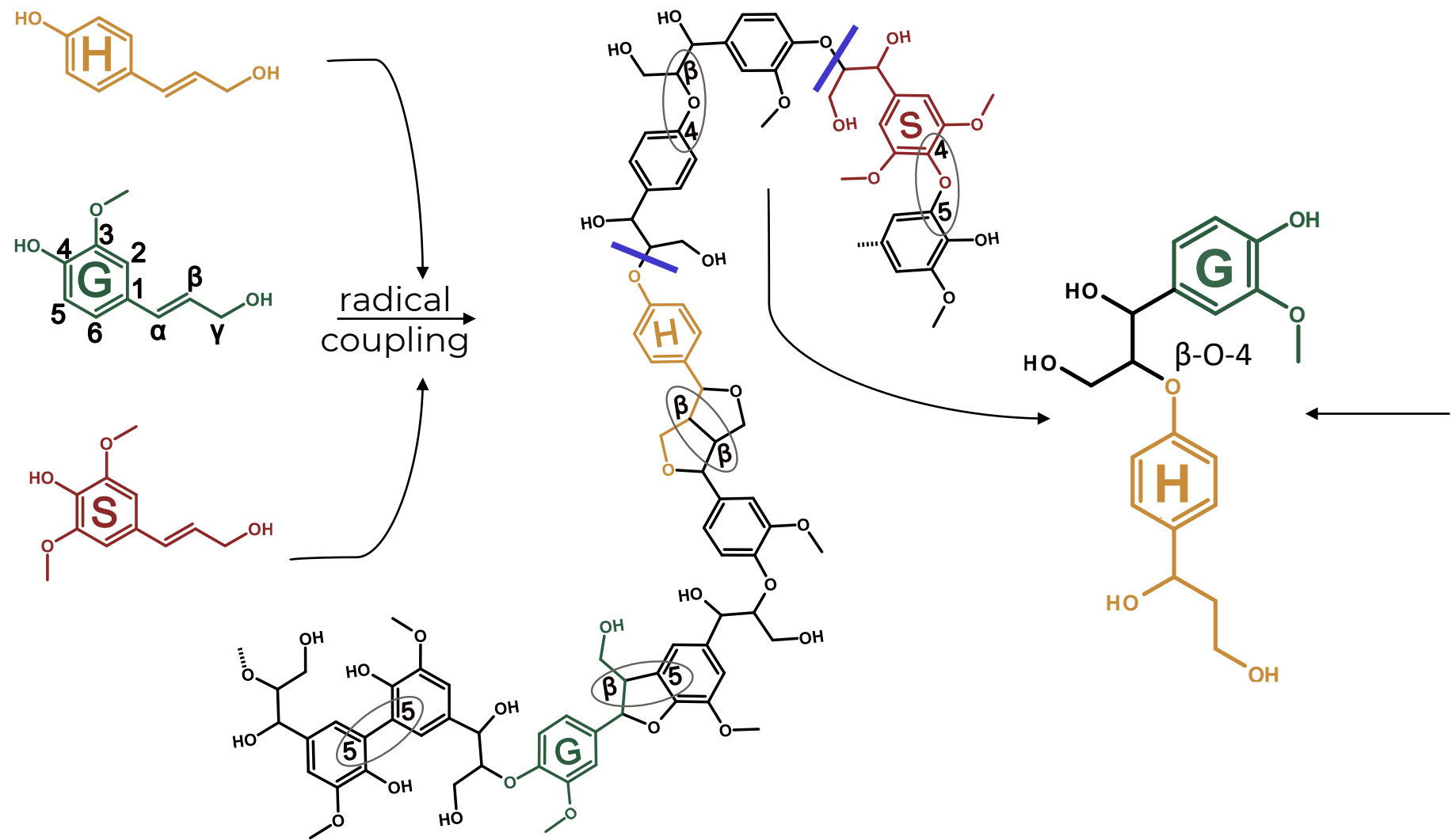
Binary fingerprint

PubChem, COCONUT, ...

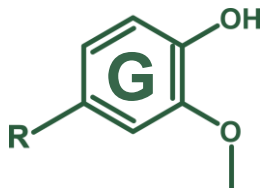


OC(C(OC1CCCC1OC)CO)c1cc(OC)c(O)cc1

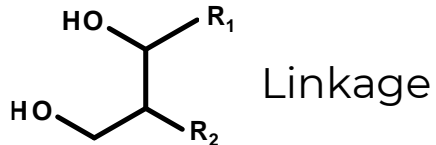
# LIGNIN DIMER STRUCTURES



Monomeric unit 1

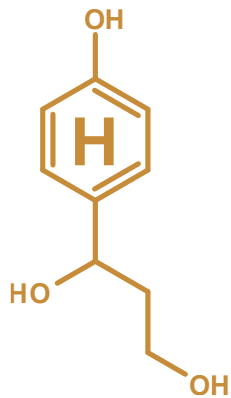


+



+

Monomeric unit 2

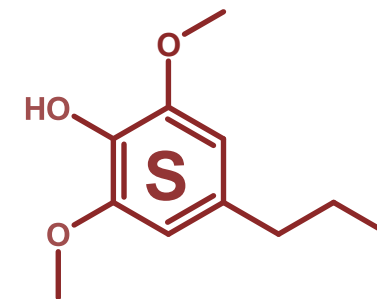
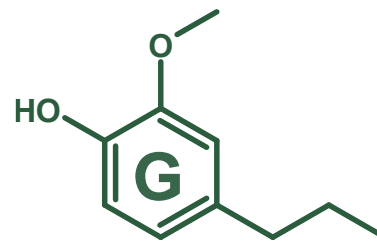
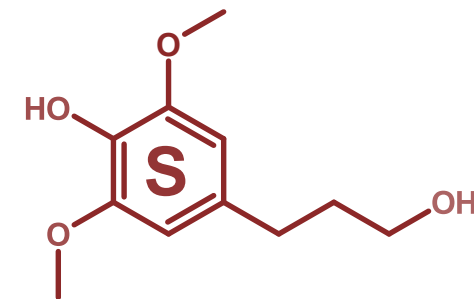
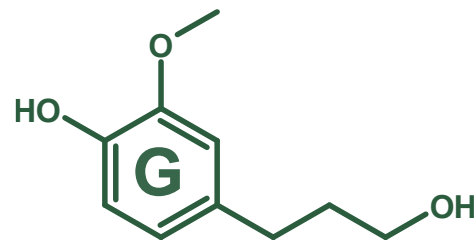
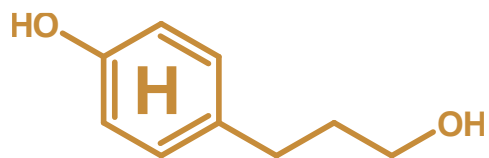
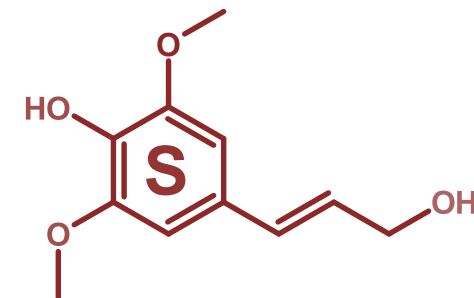
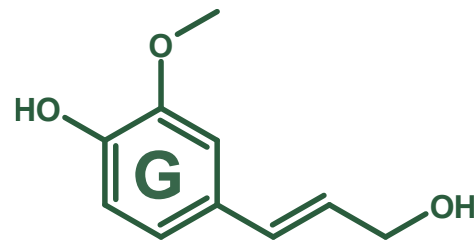
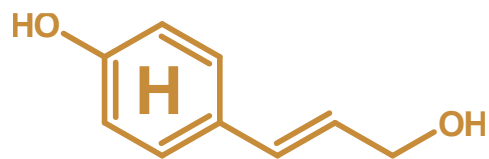




# GENERATING LIGNIN DIMER STRUCTURES

- Monomeric units in the potential RCF lignin dimers

In total: **26 monomeric units**



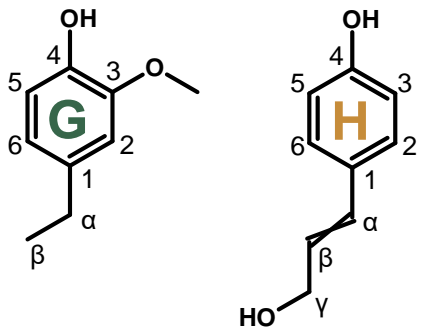
...

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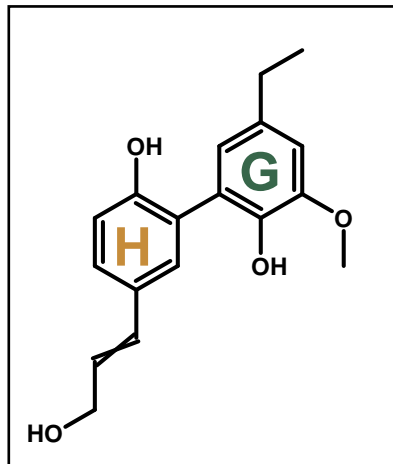
# GENERATING LIGNIN DIMER STRUCTURES

Exemplary monomeric units to show the linkages

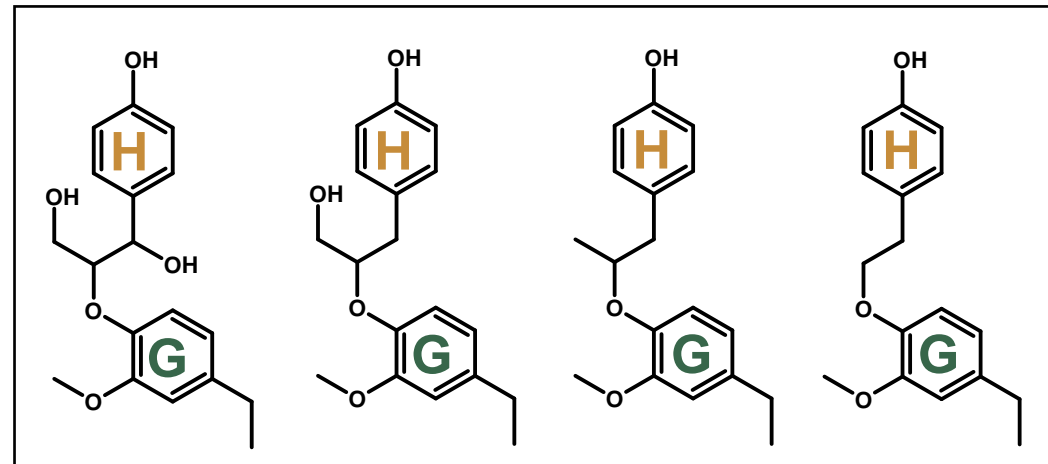


In total: **24 linkages**

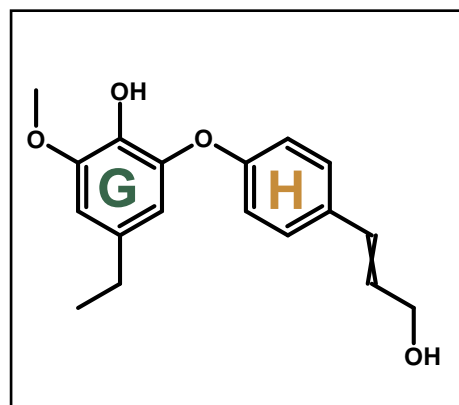
**5-5: 1 linkage**



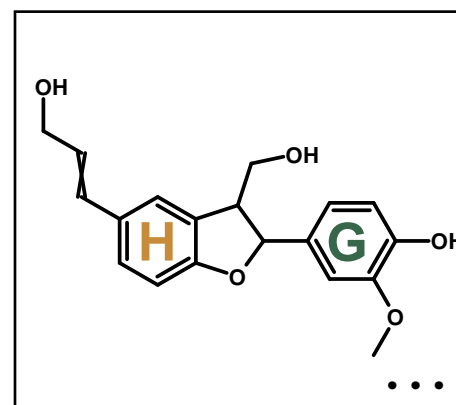
**$\beta$ -O-4: 4 linkages**



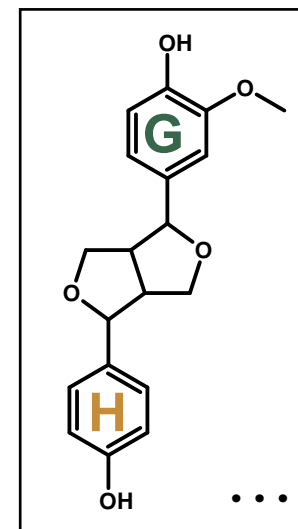
**4-O-5: 1 linkage**



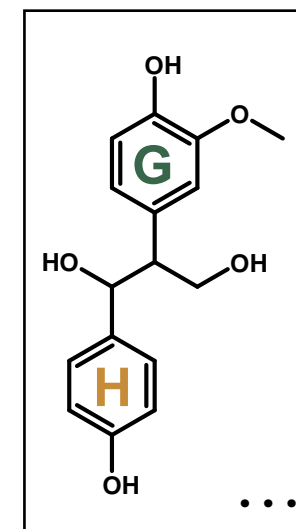
**$\beta$ -5: 6 linkages**



**$\beta$ - $\beta$ : 8 linkages**



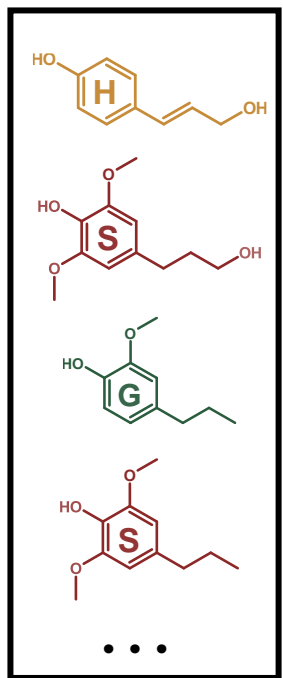
**$\beta$ -1: 4 linkages**



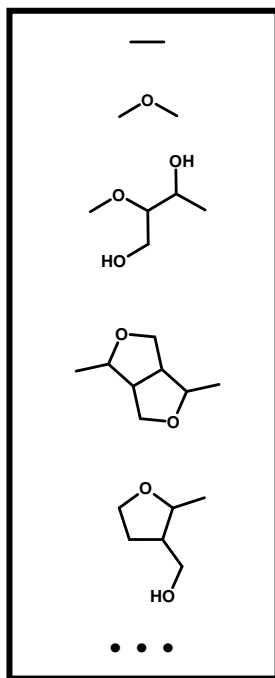
# GENERATING LIGNIN DIMER STRUCTURES

- Python script using **rule-based modelling**
- Implementing RDKit package

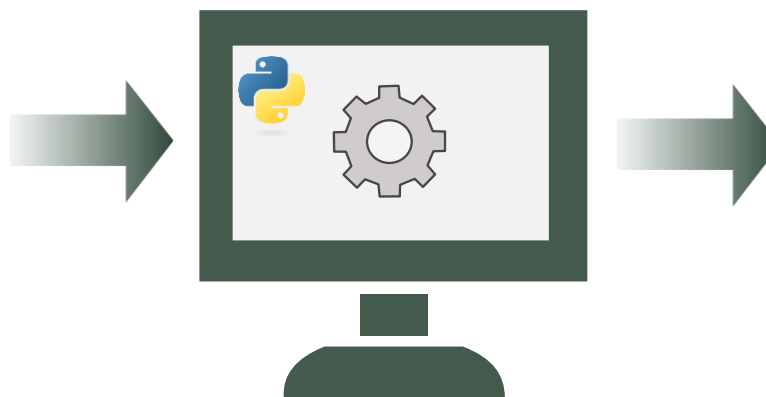
26  
monomers



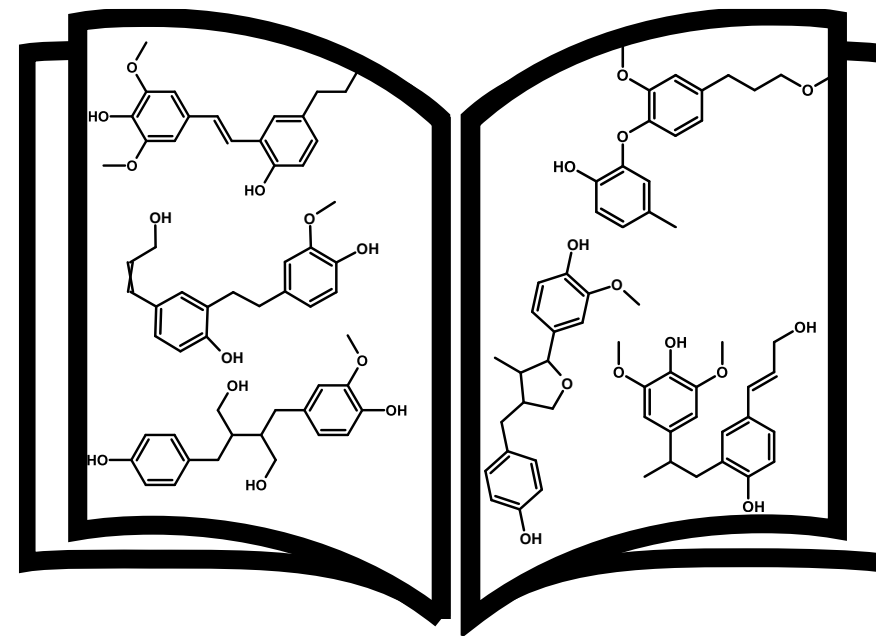
24  
linkages



Generating all possible  
combinations



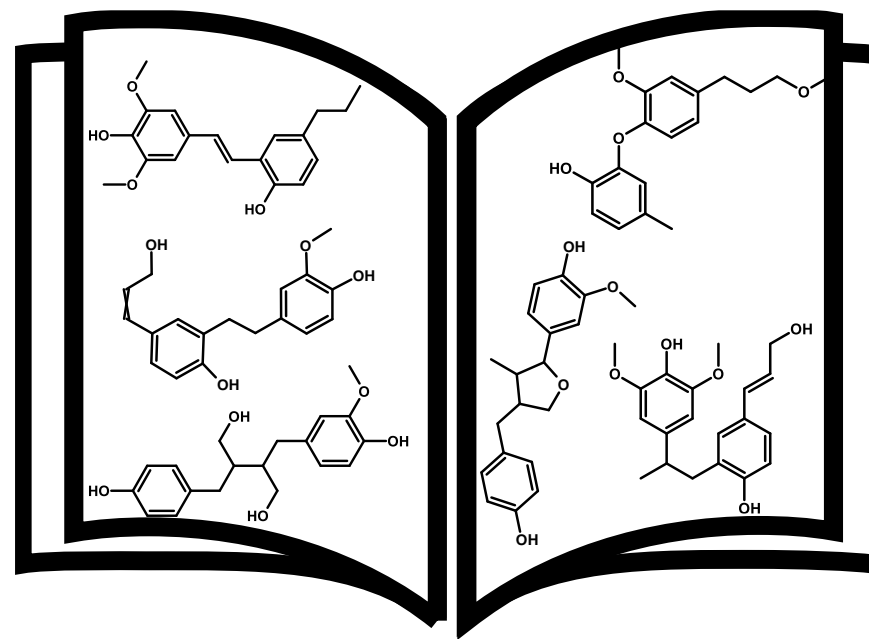
**Library** of generated RCF  
lignin dimer structures



# GENERATING LIGNIN DIMER STRUCTURES

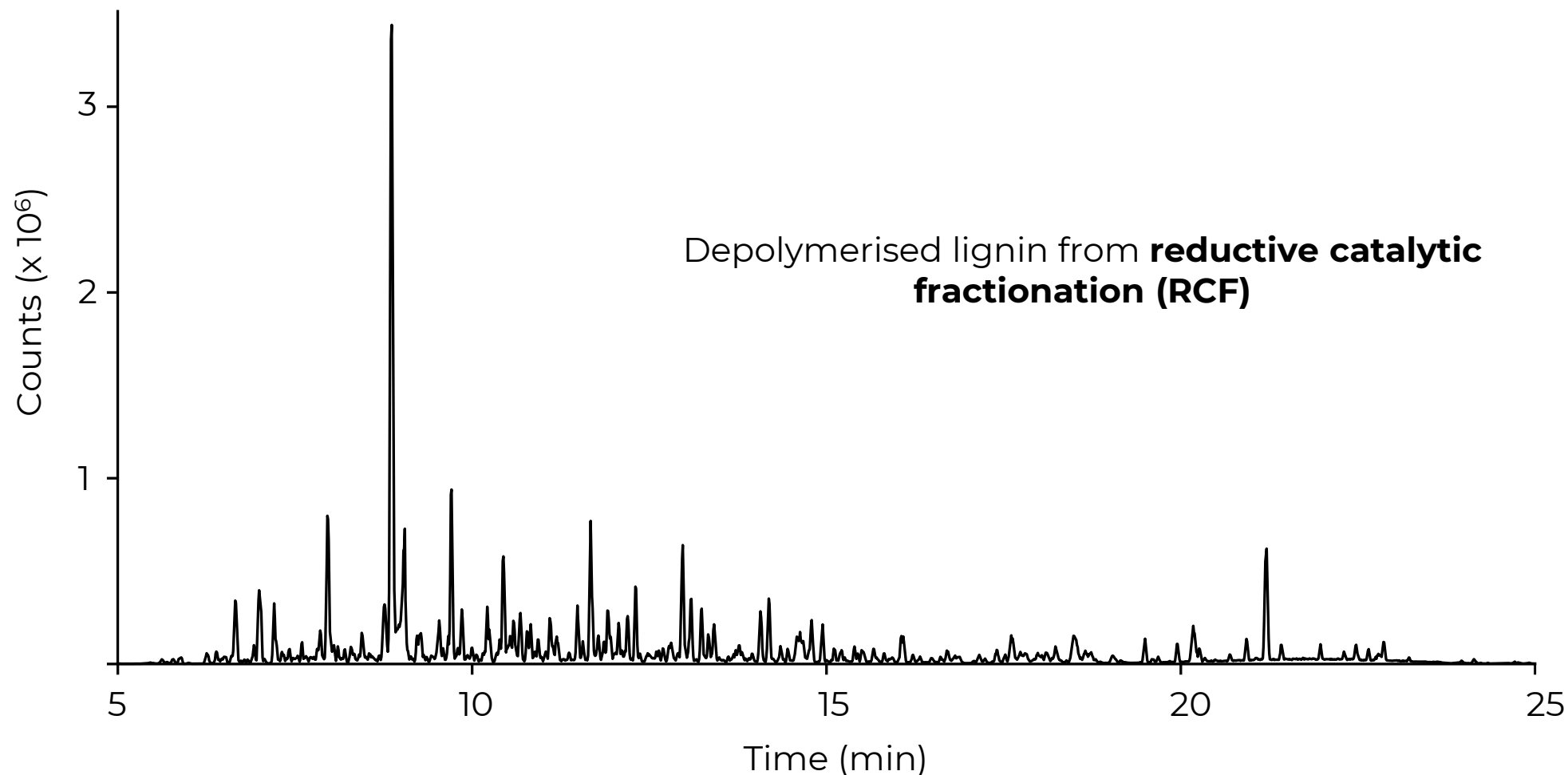
- Contains **1720 dimer** structures
- **SMILES** for each structure
- Only **117 molecular formulas**
- Only **195 (12%)** of them are in **PubChem**
- Structures of **potential** lignin RCF dimers

**Library** of generated RCF  
lignin dimer structures



# APPLYING THE LIGNIN LIBRARY: SUSPECT

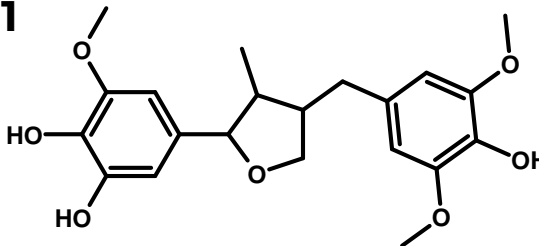
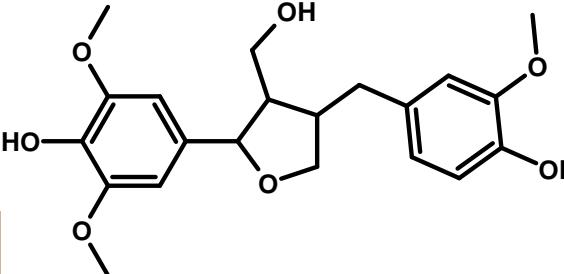
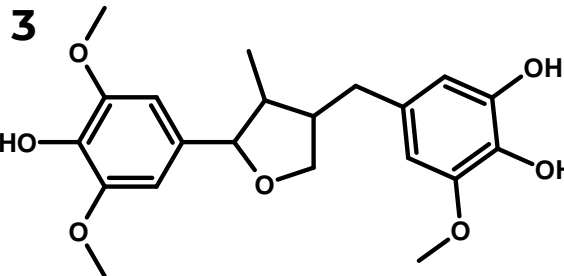
- **DDA (Top 5)** using the **suspect list** of the lignin dimer molecular formulas
- **mzMine** (4.3.0) was used for data processing
- The lignin library was imported into SIRIUS (6.0.4) as a **custom database**



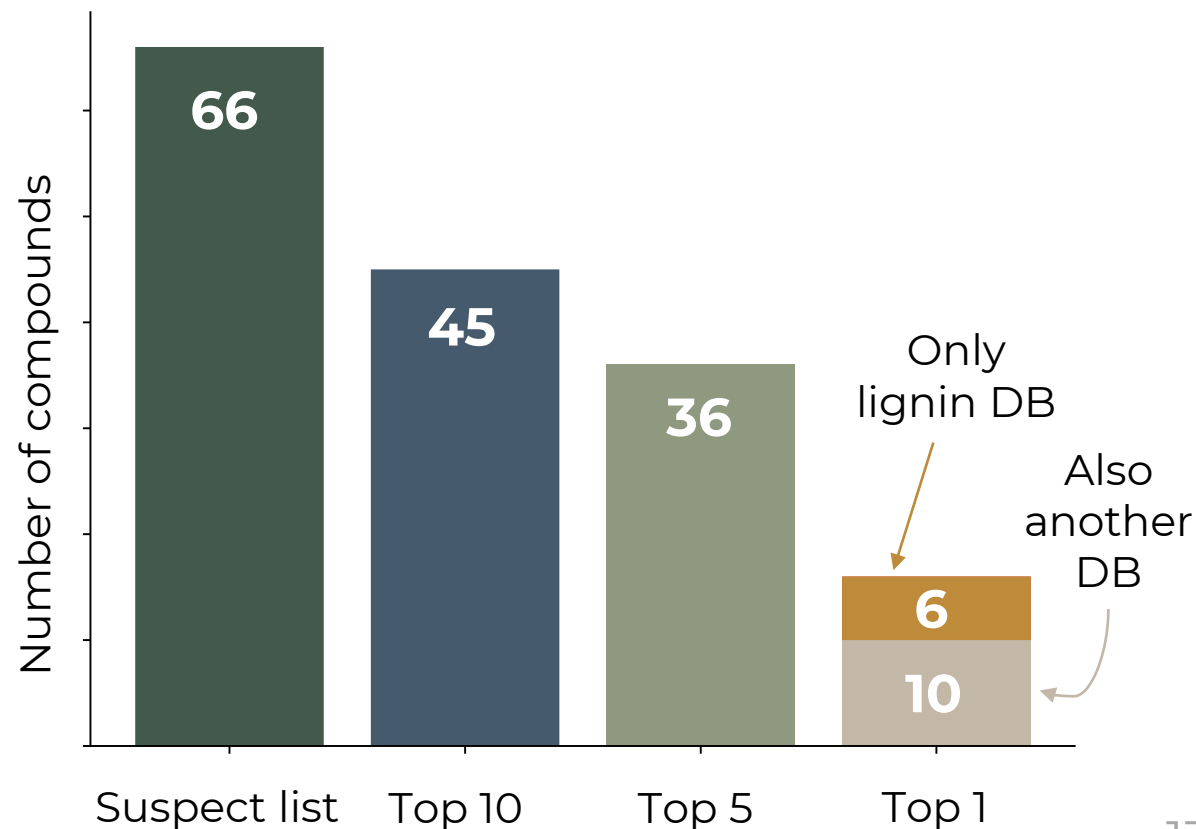
# APPLYING THE LIGNIN LIBRARY: SUSPECT

- **DDA (Top 5)** using the **suspect list** of the lignin dimer molecular formulas
- **mzMine** (4.3.0) was used for data processing
- The lignin library was imported into SIRIUS (6.0.4) as a **custom database**

SIRIUS results for one MS<sup>2</sup>

<p>Sources:</p> <p>Lignin DB</p>	<p><b>Top 1</b></p> 
<p>Sources:</p> <p>Lignin DB</p> <p>SuperNatural</p> <p>PubChem</p> <p>LOTUS</p> <p>COCONUT</p>	<p><b>Top 2</b></p> 
<p>Sources:</p> <p>Lignin DB</p>	<p><b>Top 3</b></p> 

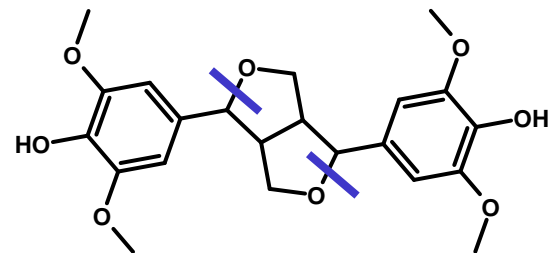
Matches with the lignin library 





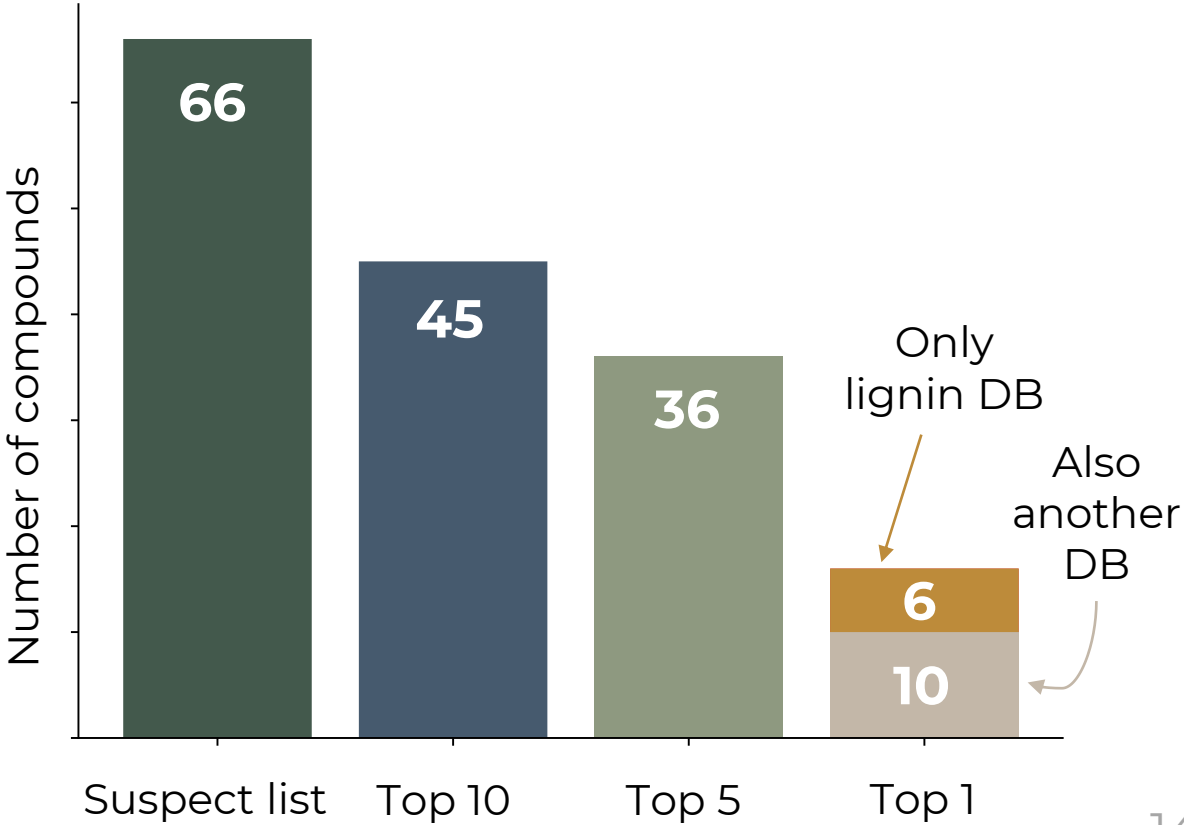
# APPLYING THE LIGNIN LIBRARY: SUSPECT

Original linkage:




Matches with the lignin library

<b>Top 1</b>	
Sources: PubChem	
<b>Top 2</b>	
Sources: Lignin DB PubChem DSSTox Blood Exposome HMDB	
<b>Top 3</b>	
Sources: Lignin DB	



# APPLYING THE LIGNIN LIBRARY: NON-TARGET


- Two HRMS instruments were used for analysis:
- Thermo **Orbitrap MS** (Exploris 480)
  - 0.1% HCOOH in both H<sub>2</sub>O and ACN
  - 5 mM NH<sub>4</sub>HCO<sub>3</sub> in H<sub>2</sub>O
- Waters **Cyclic IMS-TOF**
  - 0.1% HCOOH in both H<sub>2</sub>O and ACN

	Suspect	Non-target
Method	Orbitrap NH <sub>4</sub> HCO <sub>3</sub>	Orbitrap NH <sub>4</sub> HCO <sub>3</sub>
Acquisition	DDA Top 5	DDA Top 5
Features	449	428
Features with MS <sup>2</sup>	66	415
Top 10 match 	45 (68%)	45 (11%)
Top 1 match	16 (24%)	16 (4%)

\*Percentage (%) is calculated taking features with MS<sup>2</sup> as 100%

# APPLYING THE LIGNIN LIBRARY: NON-TARGET

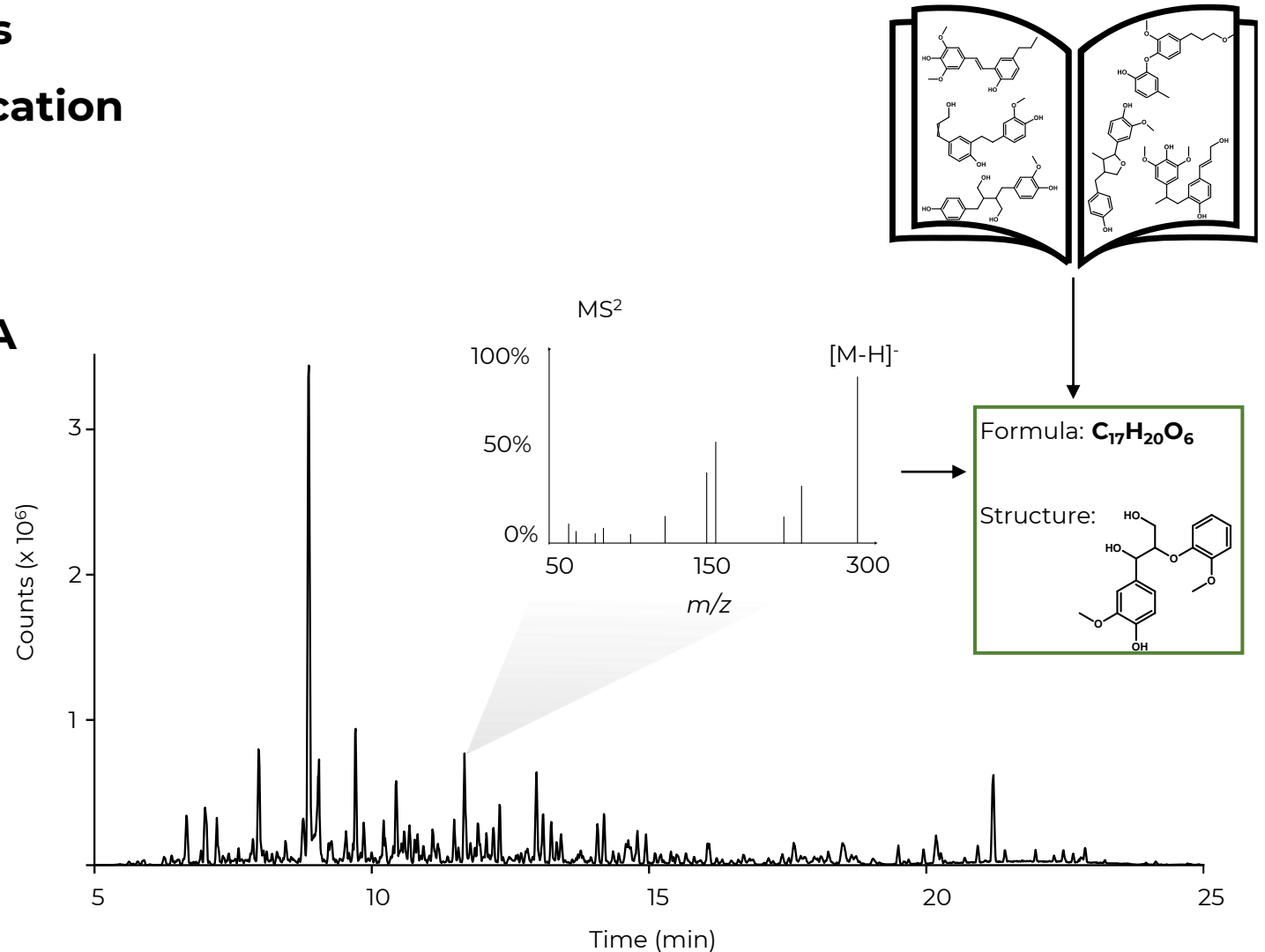
- Two HRMS instruments were used for analysis:
- Thermo **Orbitrap MS** (Exploris 480)
  - 0.1% HCOOH in both H<sub>2</sub>O and ACN
  - 5 mM NH<sub>4</sub>HCO<sub>3</sub> in H<sub>2</sub>O
- Waters **Cyclic IMS-TOF**
  - 0.1% HCOOH in both H<sub>2</sub>O and ACN

	Non-target		
Method	Orbitrap NH <sub>4</sub> HCO <sub>3</sub>	Orbitrap HCOOH	IMS HCOOH
Acquisition	DDA Top 5	DDA Top 5	DIA
Features	428	903	448
Features with MS <sup>2</sup>	415	822	442
Top 10 match 	45 (11%)	72 (9%)	58 (13%)
Top 1 match	16 (4%)	37 (5%)	20 (5%)

\*Percentage (%) is calculated taking features with MS<sup>2</sup> as 100%

# CONCLUSIONS

- **Lignin structures** can be **generated computationally**
- Lignin library contains **1720 dimers**
- Can be used for **tentative identification**
  - **new** structures in databases
  - **confirm** existing structures
- Can be used for both **DIA and DDA**







# THANK YOU!

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