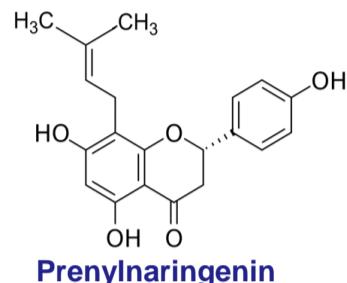


# Construction of an optimized *Escherichia coli* strain able to produce prenylnaringenins

## Motivation



### Potent estrogenic activity

#### Other Biological activities:

- Anticancer
- Antioxidant
- Anti-viral
- Immunomodulatory
- Anti-inflammatory
- Anti-bacterial

Growing interest in its industrial production

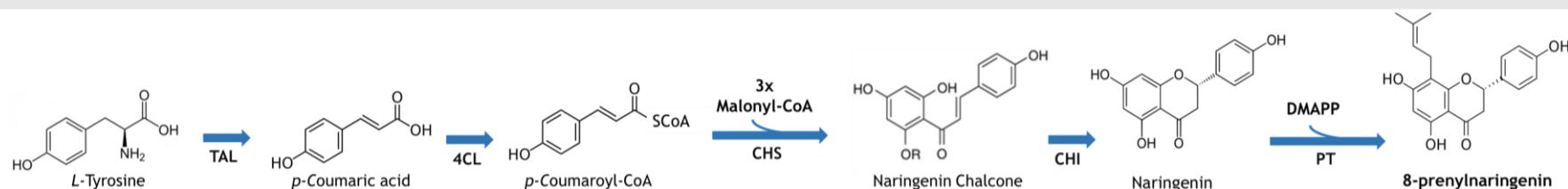
**Problem:** Low accumulation in plants

Find a solution!

**Construction of microorganisms able to produce these compounds**



Herein, we aimed to design, construct, and validate a biosynthetic pathway to produce prenylnaringenins in *Escherichia coli* for the first time



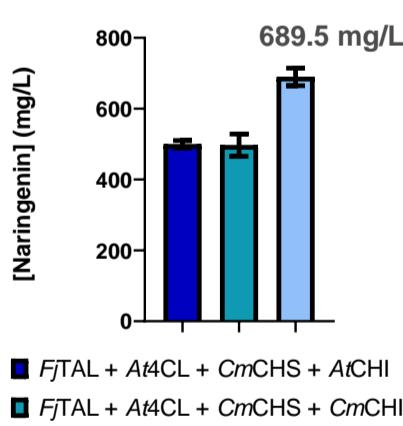
## Methods & Results

### Step 1. Construction of an optimized *E. coli* strain to produce naringenin

*FjTAL + At4CL + CmCHS*  
+  
*AtCHI / CmCHI / MsCHI*

*E. coli* M-PAR-121 – Tyrosine-overproducing strain

Evaluate naringenin production from glucose – shake flask experiments



Platform strain for prenylnaringenin production

### Step 2. Construction of DMAPP-boosted strains and validation of PT step to produce prenylnaringenin

Identification of rate limiting steps on the MEP pathway

*DXS and IDI genes*

Integration of heterologous and native DXS and IDI genes into *E. coli* M-PAR-121 genome

*CRISPR-Cas12a*

1. M-PAR-121:BsDXS
2. M-PAR-121:ScIDI
3. M-PAR-121:B/IDI
4. M-PAR-121:BsDXS:ScIDI
5. M-PAR-121:BsDXS:B/IDI
6. M-PAR-121:EcDXS
7. M-PAR-121:EcIDI
8. M-PAR-121:EcDXS:EcIDI

Transformation with naringenin pathway + PT holding plasmid

- Test a library of PTs:
1. *HlPT*
  2. *SfPT*
  3. *CdpC3PT*
  4. *AnaPT*
  5. *CsPT3*
  6. *coAnaPT*
  7. *CloQ*
  8. *EcPT*
  9. *NphB*
  10. *StrepPT*
  11. *UbiA*

99 combinations of strains/PTs

Evaluate prenylnaringenin production from glucose - 1 mL scale experiments in deep well blocks

#### Results:

*E. coli* M-PAR-121 (WT)

→ No prenylnaringenin detected

DMAPP-boosted M-PAR-121 strains

→ Prenylnaringenin detected in 12 different combinations of strains/PTs

Strain	PT	Prenylnaringenin
	<i>CdpC3PT</i>	21.6 $\mu$ M
	<i>coAnaPT</i>	19.6 $\mu$ M
<i>M-PAR-121:BsDXS</i>	<i>CloQ</i>	3.3 $\mu$ M
	<i>NphB</i>	24.7 $\mu$ M
	<i>StrepPT</i>	37.0 $\mu$ M
	<i>UbiA</i>	20.6 $\mu$ M
<i>M-PAR-121: EcDXS</i>	<i>CoAnaPT</i>	15.3 $\mu$ M
	<i>CloQ</i>	20.6 $\mu$ M
	<i>NphB</i>	27.5 $\mu$ M
	<i>StrepPT</i>	19.6 $\mu$ M
<i>M-PAR-121: BsDXS: ScIDI</i>	<i>CloQ</i>	6.2 $\mu$ M
<i>M-PAR-121: EcDXS: EcIDI</i>	<i>EcPT</i>	49.6 $\mu$ M

## Conclusions

### 1. Construction of an optimized *E. coli* strain to produce naringenin

→ Highest *de novo* production reported so far in *E. coli*

### 2. Construction of DMAPP-boosted strains and validation of PT step to produce prenylnaringenin

→ DMAPP modifications in the M-PAR-121 resulted in prenylnaringenin production  
Next step: Validation at flask scale

Significant step towards *de novo* production of prenylflavonoids in *E. coli*

This study was supported by the Portuguese Foundation for Science and Technology (FCT) under the scope of the strategic funding of UIDB/04469/2020 unit, with DOI 10.54499/UIDB/04469/2020 and by LABBELS – Associate Laboratory in Biotechnology, Bioengineering and Microelectromechanical Systems, LA/P/0029/2020. Daniela Gomes also acknowledge FCT for the fellowship SFRH/BD/04433/2020.

**Acronyms:** 4CL: 4-coumarate-CoA ligase; AnaPT: C3-prenyltransferase from *Neosartorya fischeri*; At: *Arabidopsis thaliana*; CdpC3PT: 7-O-prenyltransferase from *Neosartorya fischeri*; CHS: chalcone synthase; CHI: chalcone isomerase; CloQ: prenyltransferase from *Streptomyces roseochromogenes*; Cm: *Curculbita maxima*; co: codon optimized; Cs: *Candida sativa*; DXS: deoxyxylulose 5-phosphate synthase; Ec: *Escherichia coli*; Fj: *Flavobacterium johnsoniae*; Hl: *Humulus lupulus*; IDI: isopentenyl diphosphate isomerase; MEP: methyl erythritol phosphate; Ms: *Medicago sativa*; NphB: hydroxynaphthalene Ptase; PT: prenyltransferase; Strep: *Streptomyces*; TAL: tyrosine ammonia lyase; UbiA: 4-hydroxybenzoate octaprenyltransferase.