

An Overview of Biosynthesis Pathways – Inspiration for Pharmaceutical and Agrochemical Discovery

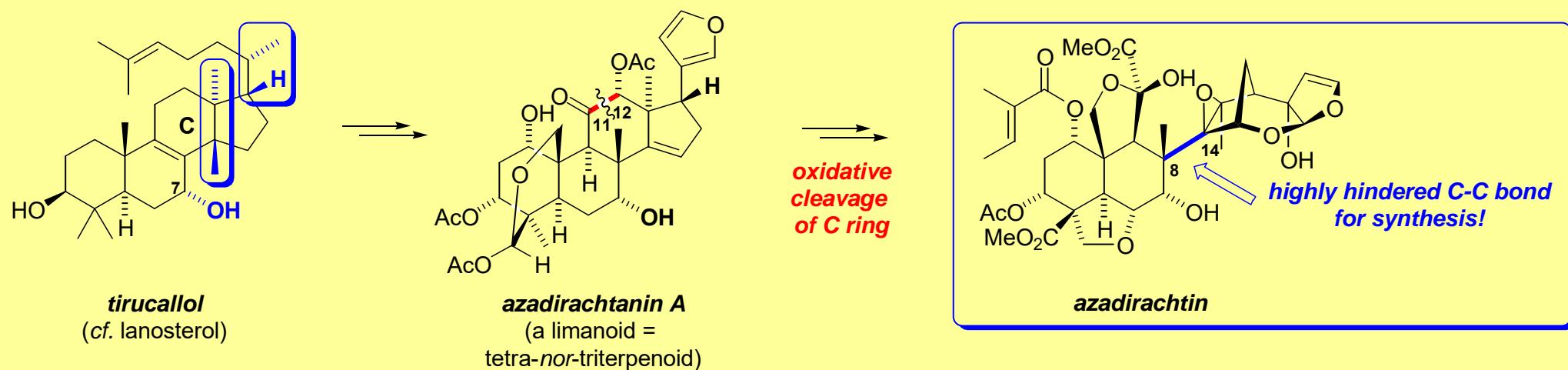
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19th Oct 2019

Lessons in Synthesis - Azadirachtin

- **Azadirachtin** is a potent *insect anti-feedant* from the Indian *neem tree*:
 - exact biogenesis unknown but certainly via steroid modification:



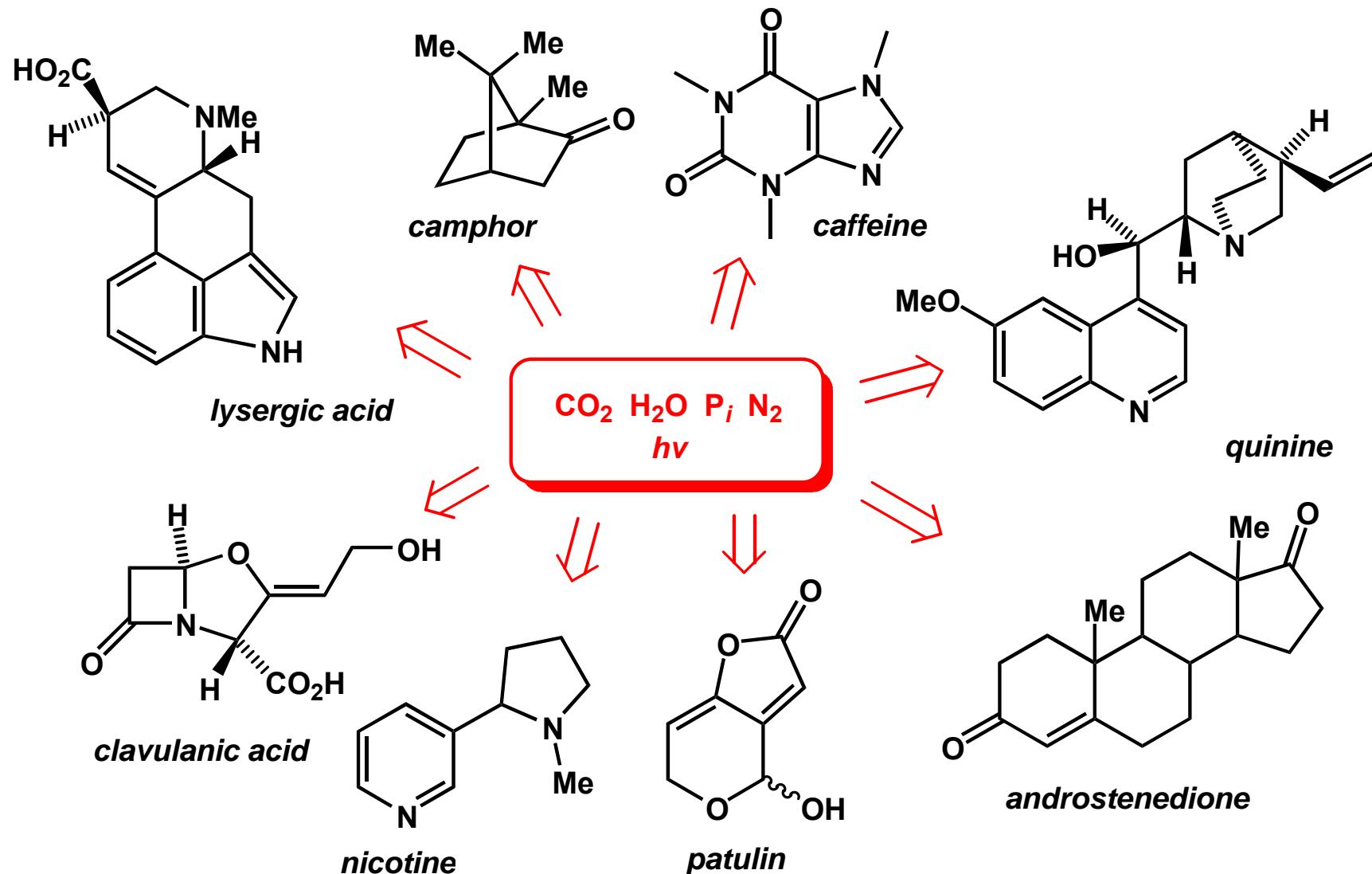
- Intense synthetic efforts by the groups of Nicolaou, Watanabe, Ley and others since structural elucidation in 1987.
- 1st total synthesis achieved in 2007 by Ley following 22 yrs of effort
- **~40 researchers and over 100 person-years of research! – 64-step synthesis**
- Veitch *Angew. Chem. Int. Ed.* **2007**, *46*, 7629 ([DOI](#)) & Veitch *Angew. Chem. Int. Ed.* **2007**, *46*, 7633 ([DOI](#))
- **Review 'The azadirachtin story' see:** Veitch *Angew. Chem. Int. Ed.* **2008**, *47*, 9402 ([DOI](#))

Format & Scope of Presentation

- ***Metabolism & Biosynthesis***
 - some definitions, 1° & 2° metabolites
- ***Shikimate Metabolites***
 - photosynthesis & glycolysis → shikimate formation → shikimate metabolites
 - Glyphosate – a non-selective herbicide
- ***Alkaloids***
 - acetylCoA & the citric acid cycle → α-amino acids → alkaloids
 - Opioids – powerful pain killers
- ***Fatty Acids and Polyketides***
 - acetylCoA → malonylCoA → fatty acids, prostaglandins, polyketides, macrolide antibiotics
 - NSAIDs – anti-inflammatory's
- ***Isoprenoids/terpenes***
 - acetylCoA → mevalonate → isoprenoids, terpenoids, steroids, carotenoids
 - Statins – cholesterol-lowering agents

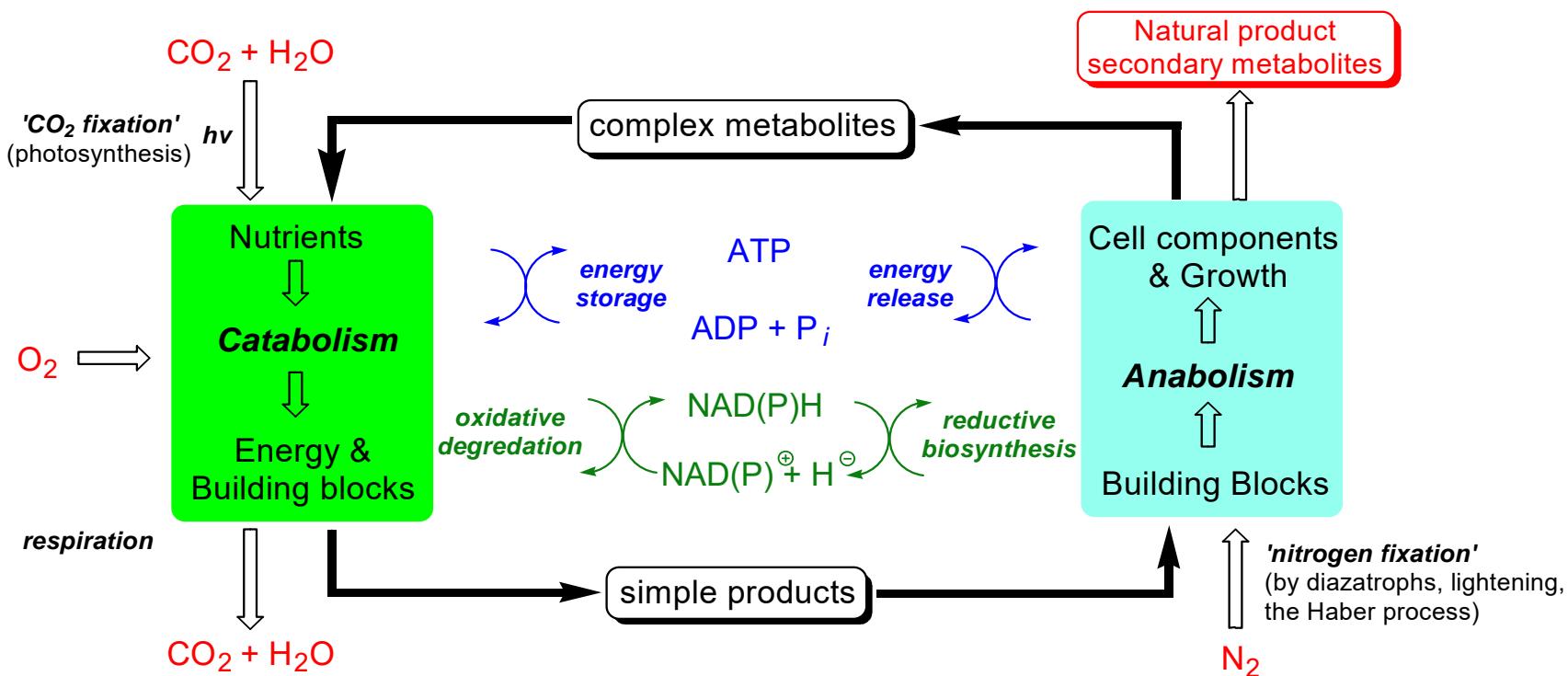
Metabolism and Biosynthesis

Metabolism & Natural Product Diversity

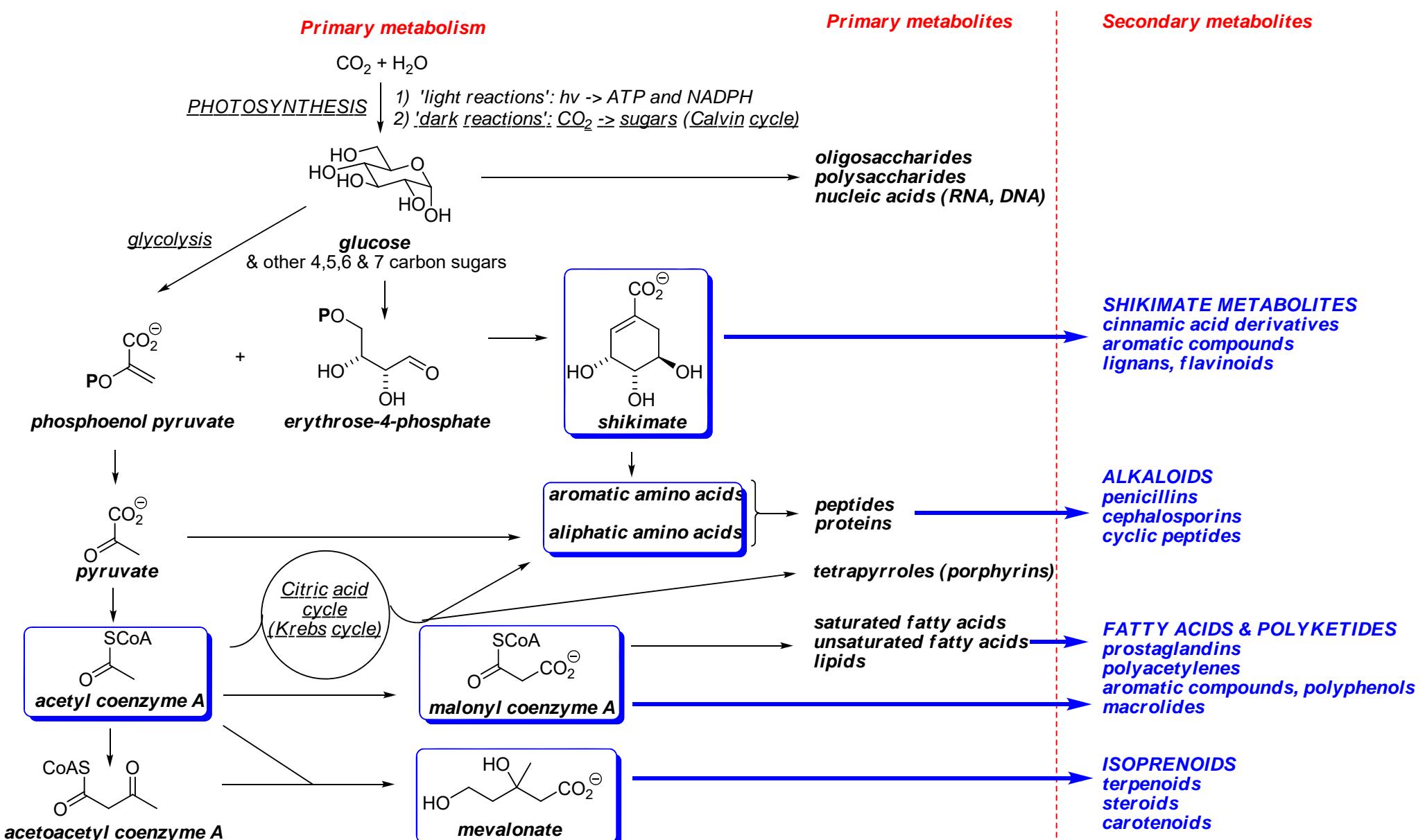


Metabolism

- **Metabolism** is the term used for *in vivo* processes by which compounds are degraded, interconverted and synthesised:
 - **Catabolic** or **degradative**: primarily to release energy and provide building blocks
 - generally **oxidative** processes/sequences (glycolysis, Krebs cycle)
 - **Anabolic** or **biosynthetic**: primarily to create new cellular materials (1° & 2° metabolites)
 - generally **reductive** processes/sequences
- These two types of process are coupled – one provides the driving force for the other:



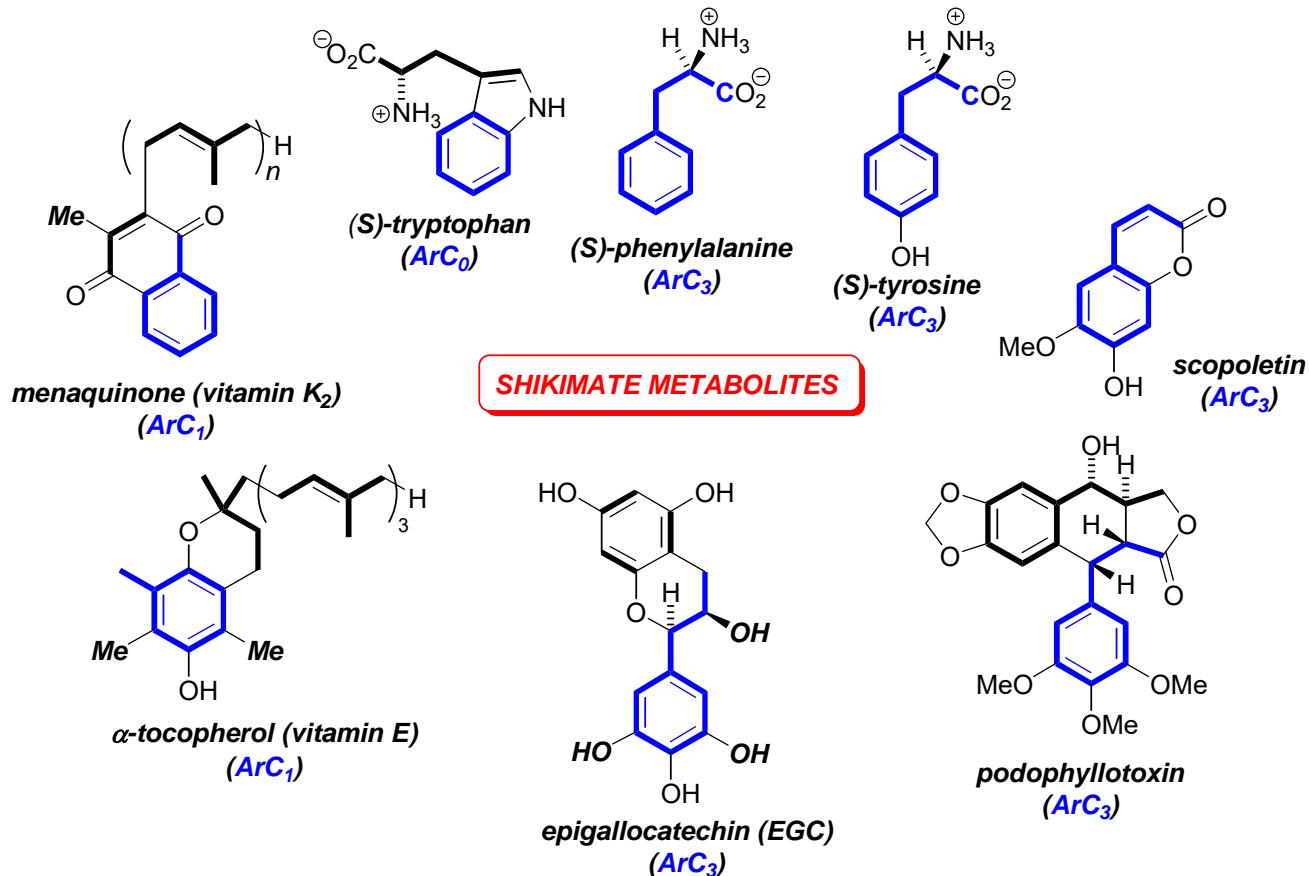
Primary Metabolism - Overview



For interesting animations' of e.g. photosynthesis see: <http://www.johnkyrk.com/index.html>

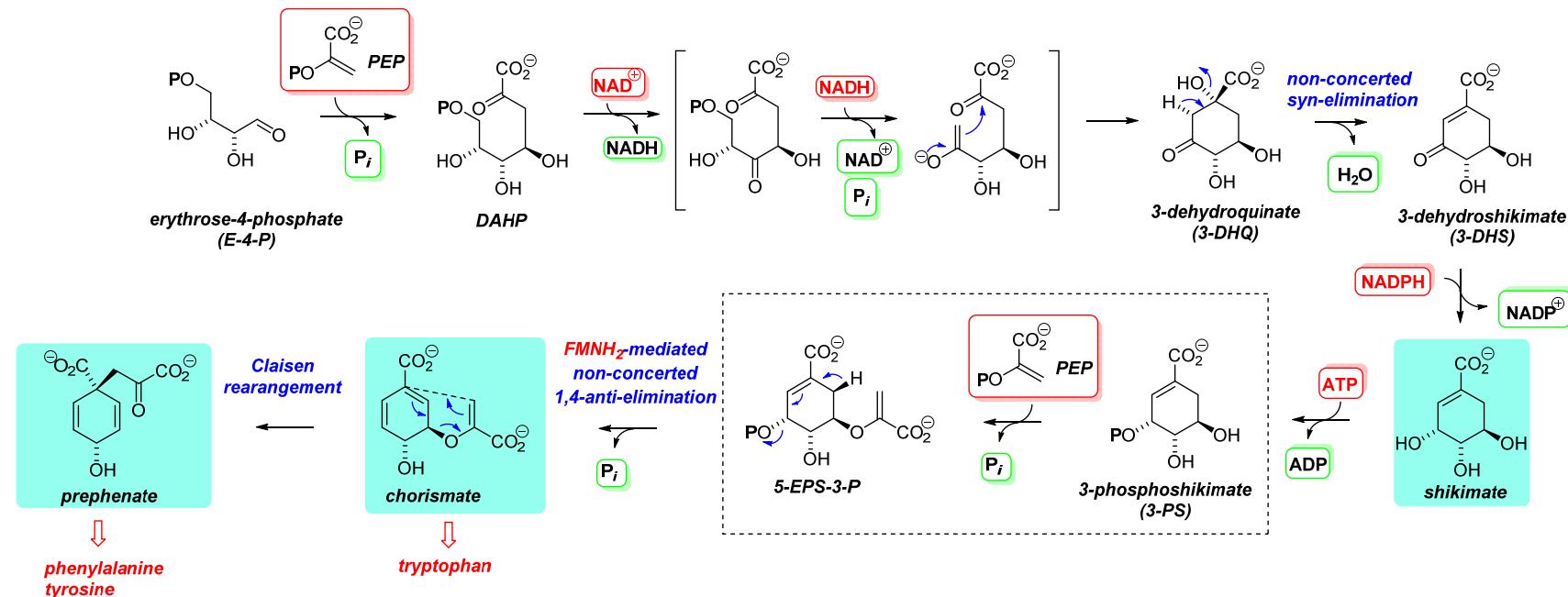
Shikimate Metabolites

Shikimate Metabolites



The Shikimate Biosynthetic Pathway - Overview

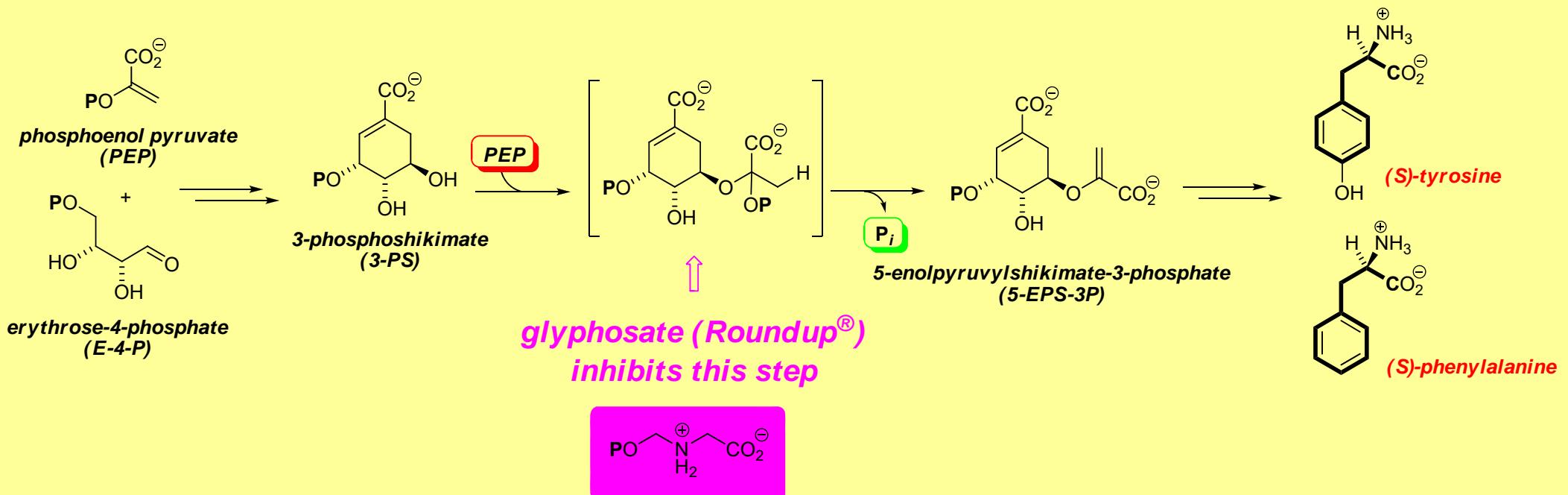
- **Phosphoenol pyruvate & erythrose-4-phosphate → shikimate → chorismate → prephenate:**



- The detailed mechanisms of these steps have been studied intensively. Most are chemically complex and interesting. For additional details see:
 - Mann *Chemical Aspects of Biosynthesis* Oxford Chemistry Primer No. 20, **1994** (*key details*)
 - Haslam *Shikimic Acid – Metabolism and Metabolites* Wiley, **1993** (*full details and primary Lit. citations*)
 - <http://www.chem.qmul.ac.uk/iubmb/enzyme/reaction/misc/shikim.html> (*interesting web-site with many biosynthetic pathways*)

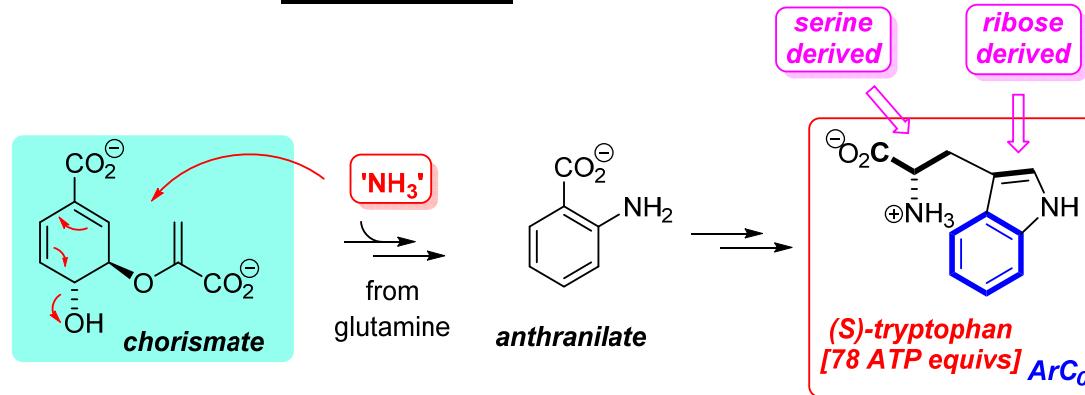
Rational Agrochemical Development – *Shikimate Pathway Intervention*

- **The shikimate biosynthetic pathway is not found in animals/humans – only in plants**
 - selective intervention in these pathways allows development of agrochemicals with minimal human toxicity
- **Glyphosate ('Roundup') – a Monsanto agrochemical is a potent inhibitor of the conversion of 3-phosphoshikimate (3-PS) → 5-enolpyruvylshikimate-3-phosphate (5-EPS-3P)**
 - a non-selective herbicide



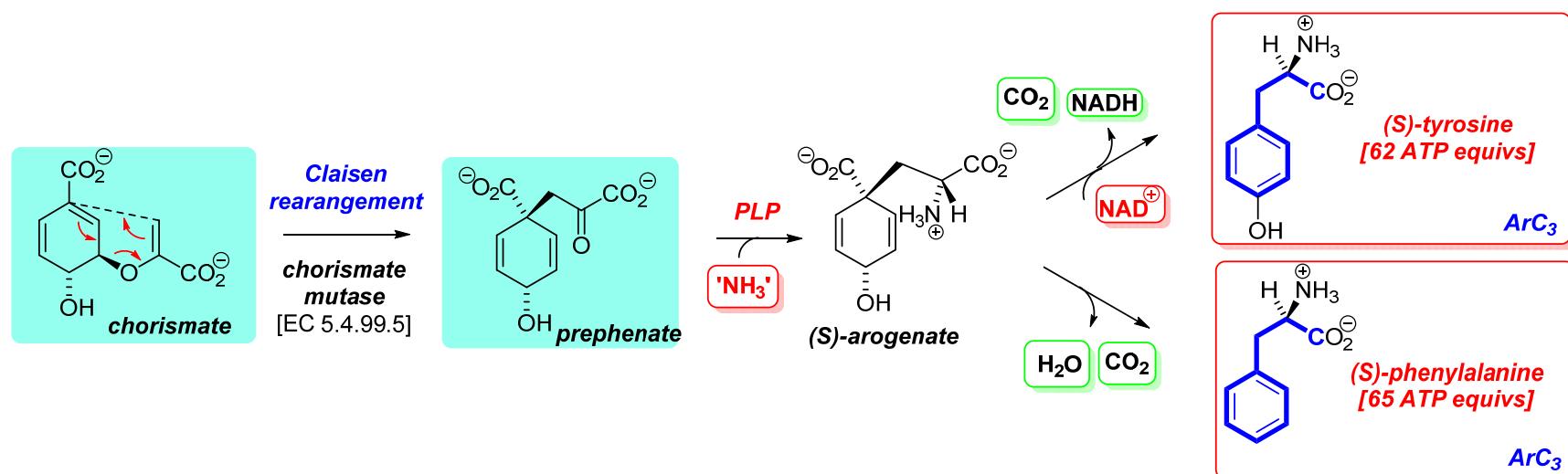
Chorismate → Tryptophan, Tyrosine & Phenylalanine

- **Chorismate → anthranilate → tryptophan**



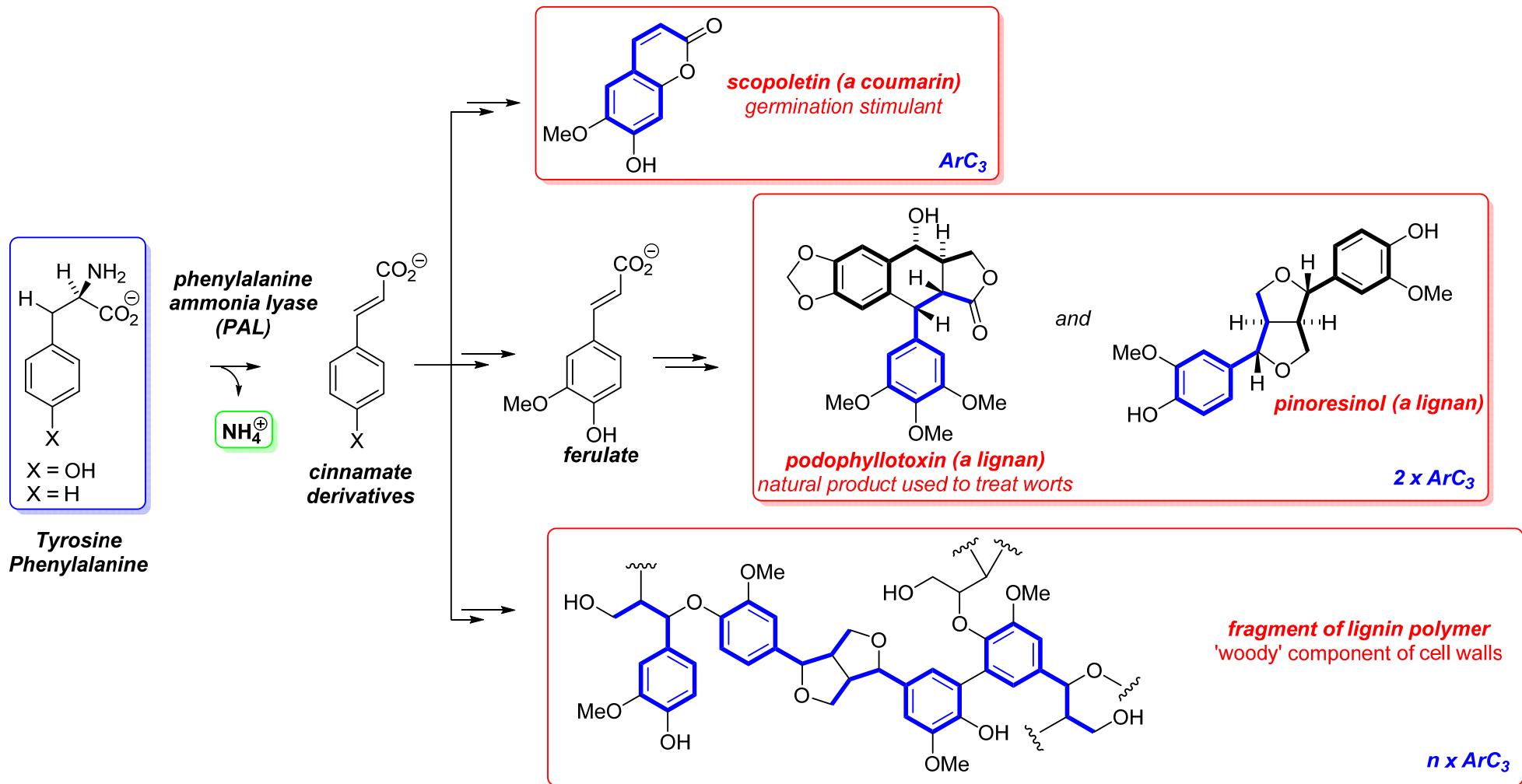
- **Chorismate → prephenate → tyrosine & phenylalanine**

– NB. The enzyme *chorismate mutase* [EC 5.4.99.5] which mediates the conversion of chorismate to prephenate is the only known ‘Claisen rearrangementase’

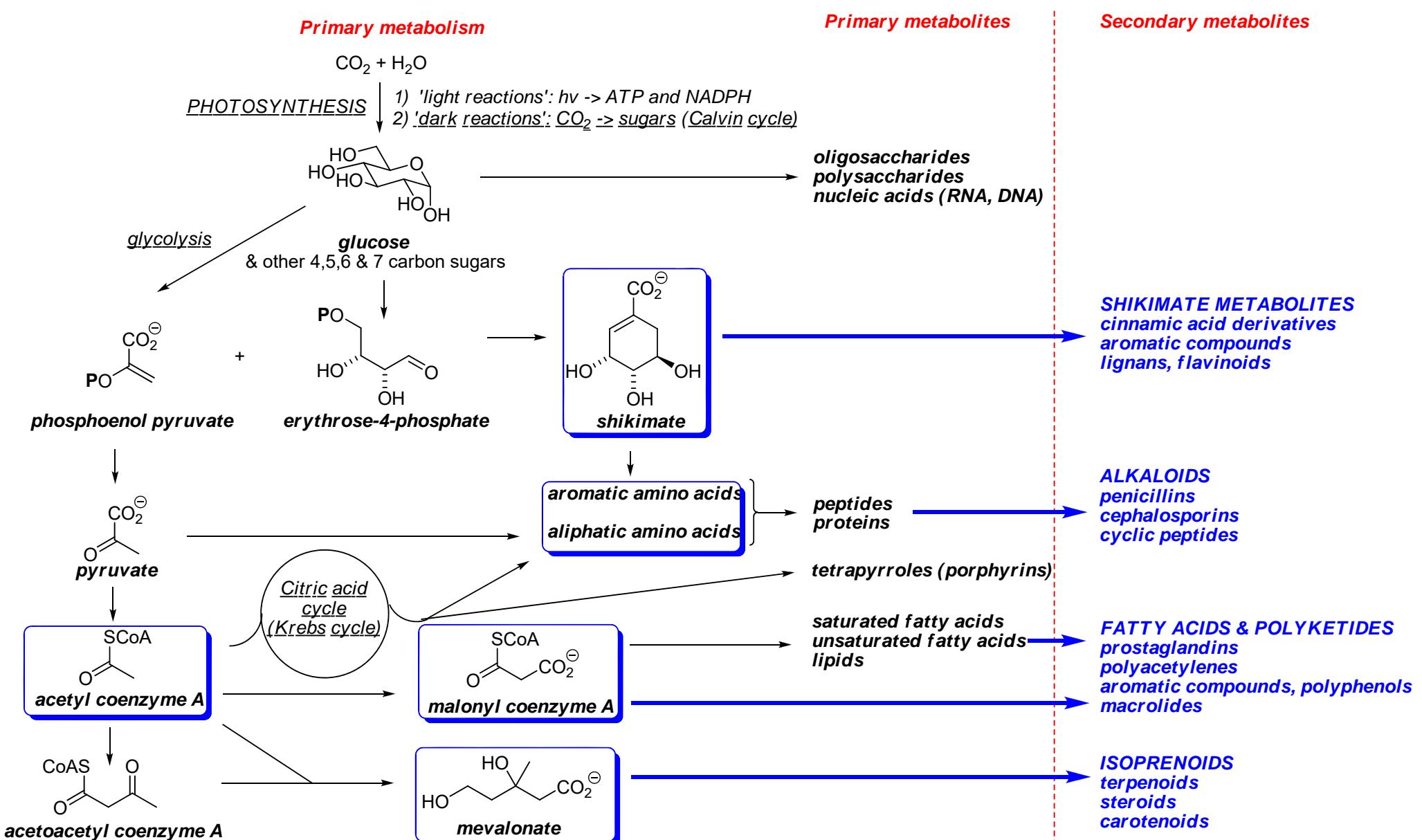


Tyrosine/Phenylalanine → ArC₃ Metabolites

- Tyrosine & phenylalanine → cinnamate derivatives → ArC₃ metabolites
 - coumarins, lignans (stereoselective enzymatic dimerisation) & lignins (stereorandom radical polymerisation)



Primary Metabolism - Overview



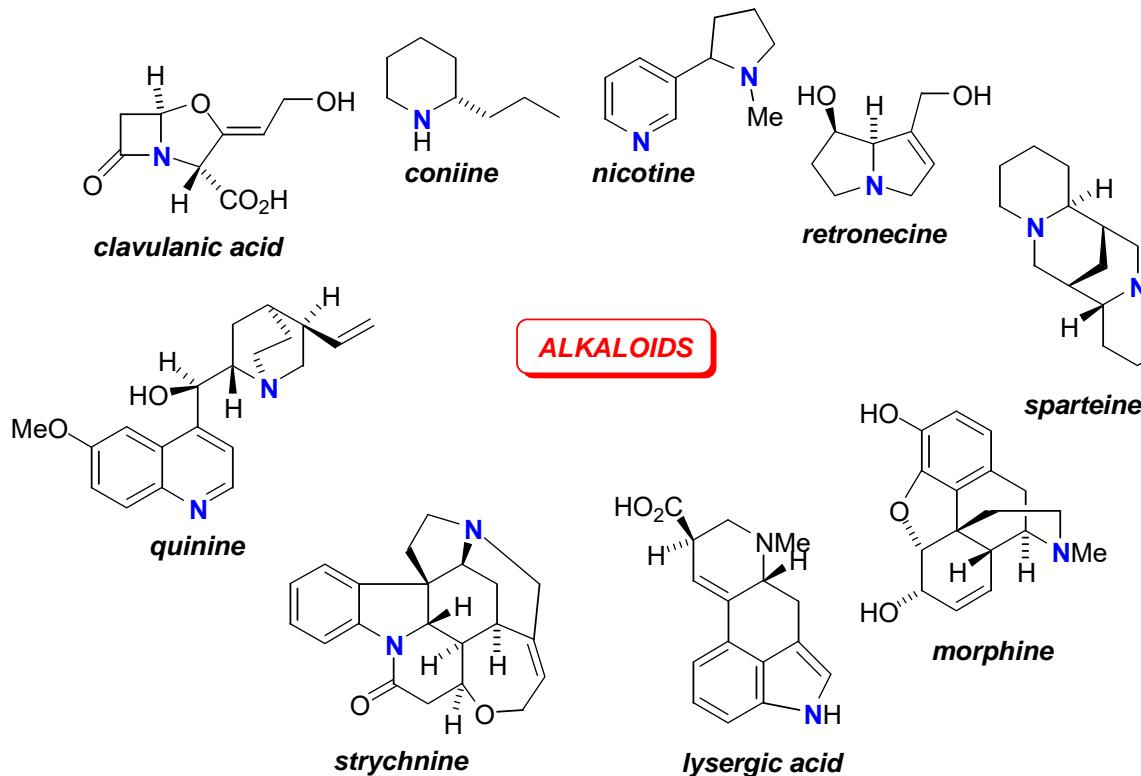
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Alkaloids

Alkaloids

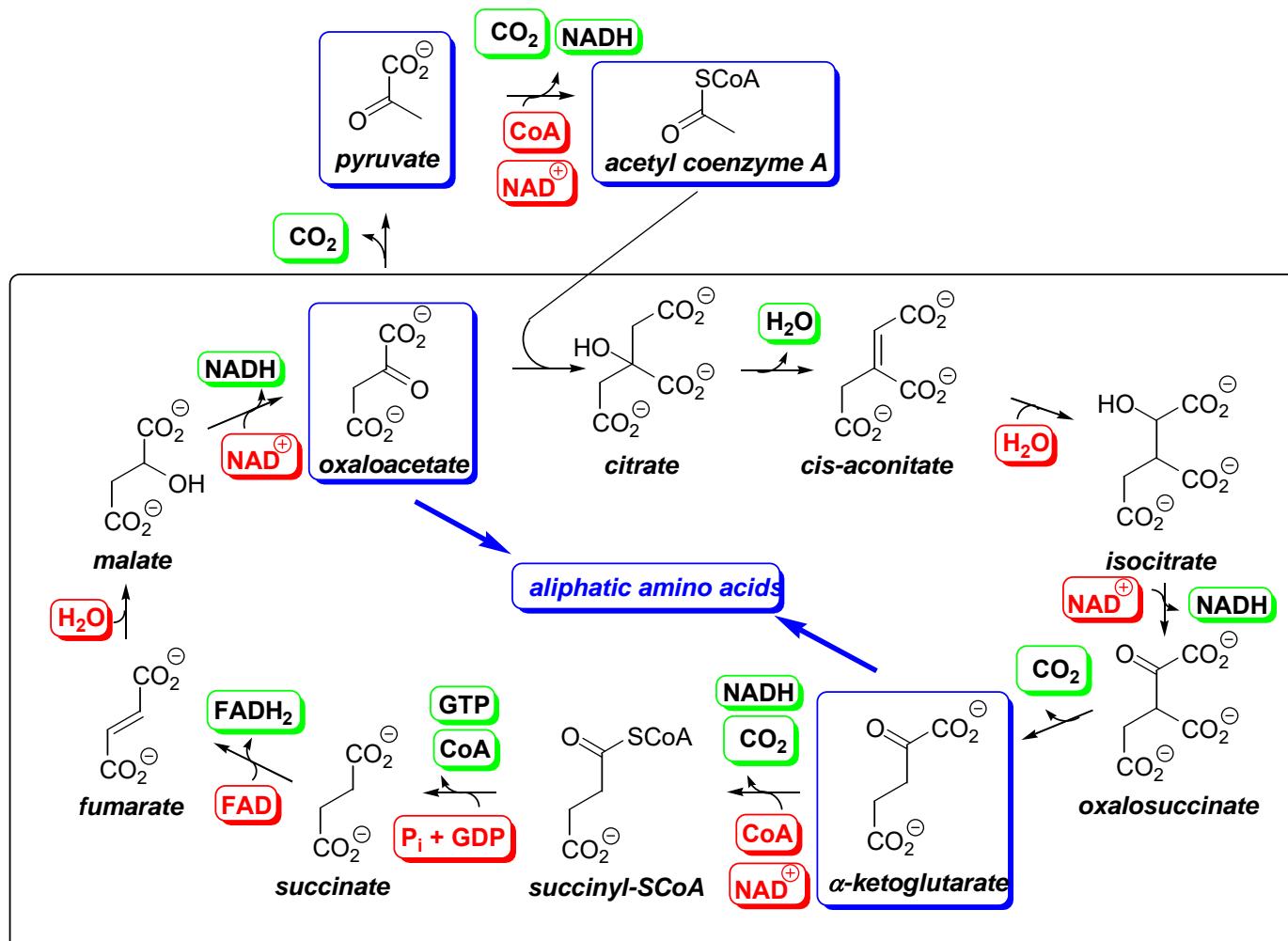
- **Definitions:**

- **originally** – ‘a natural product that could be extracted out of alkaline but not acidic water’ (i.e. containing a basic amine function that protonated in acid)
- **more generally** - ‘any non-peptidic & non-nucleotide nitrogenous secondary metabolite’



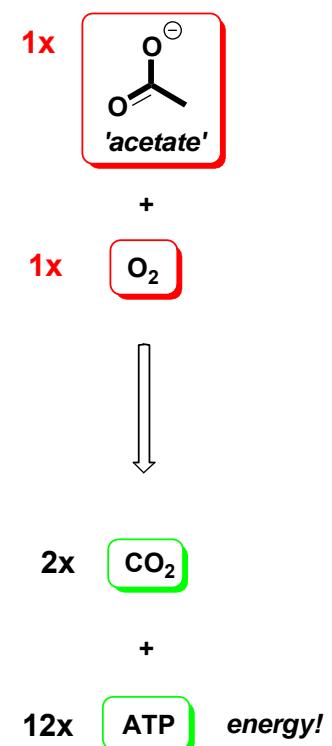
The Citric Acid Cycle

- The citric acid (Krebs) cycle** is a major catabolic pathway of 1° metabolism that provides two key building blocks for aliphatic amino acid biosynthesis - **oxaloacetate** & **α -ketoglutarate**:



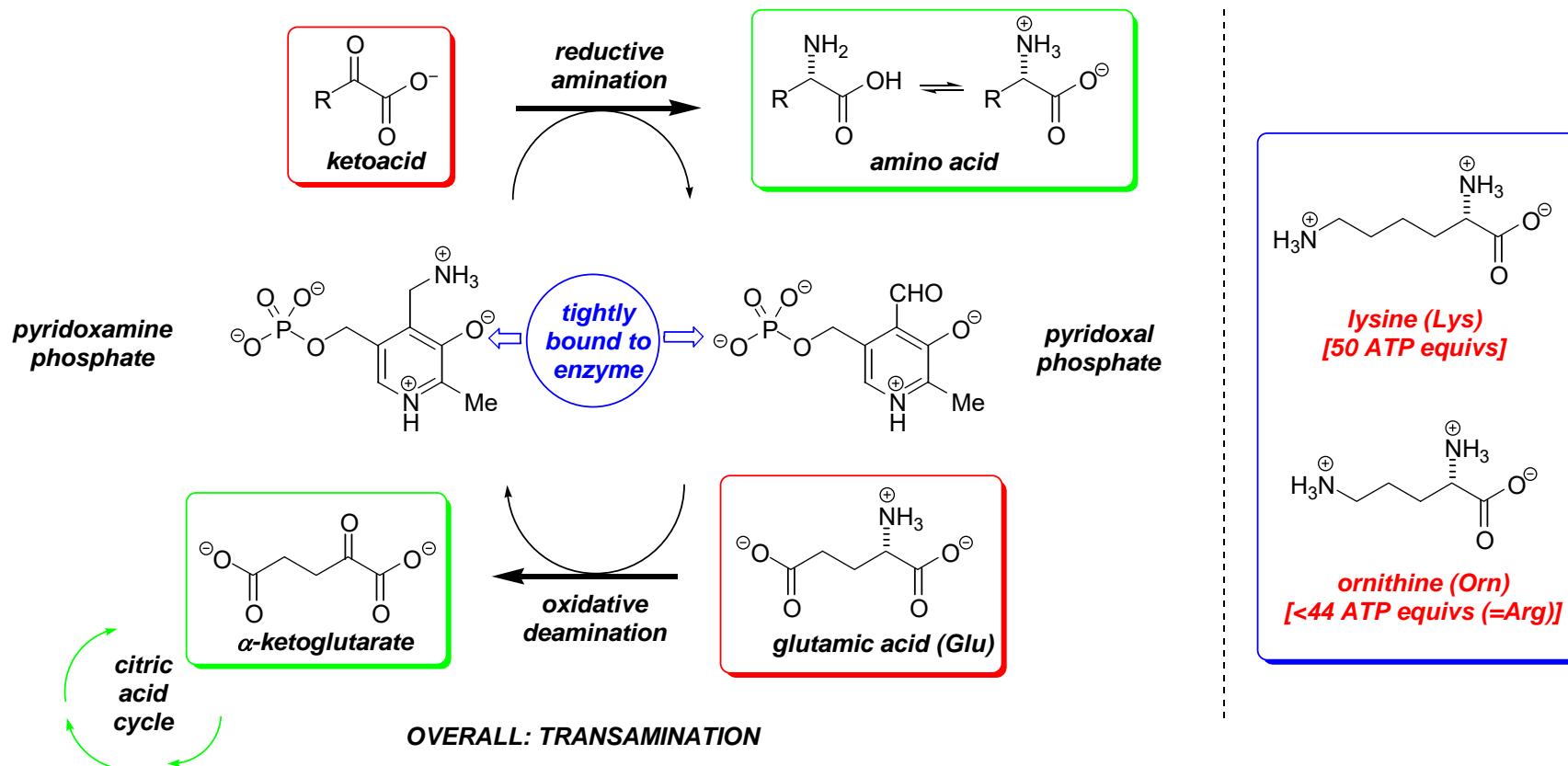
THE CITRIC ACID CYCLE

OVERALL STOICHIOMETRY



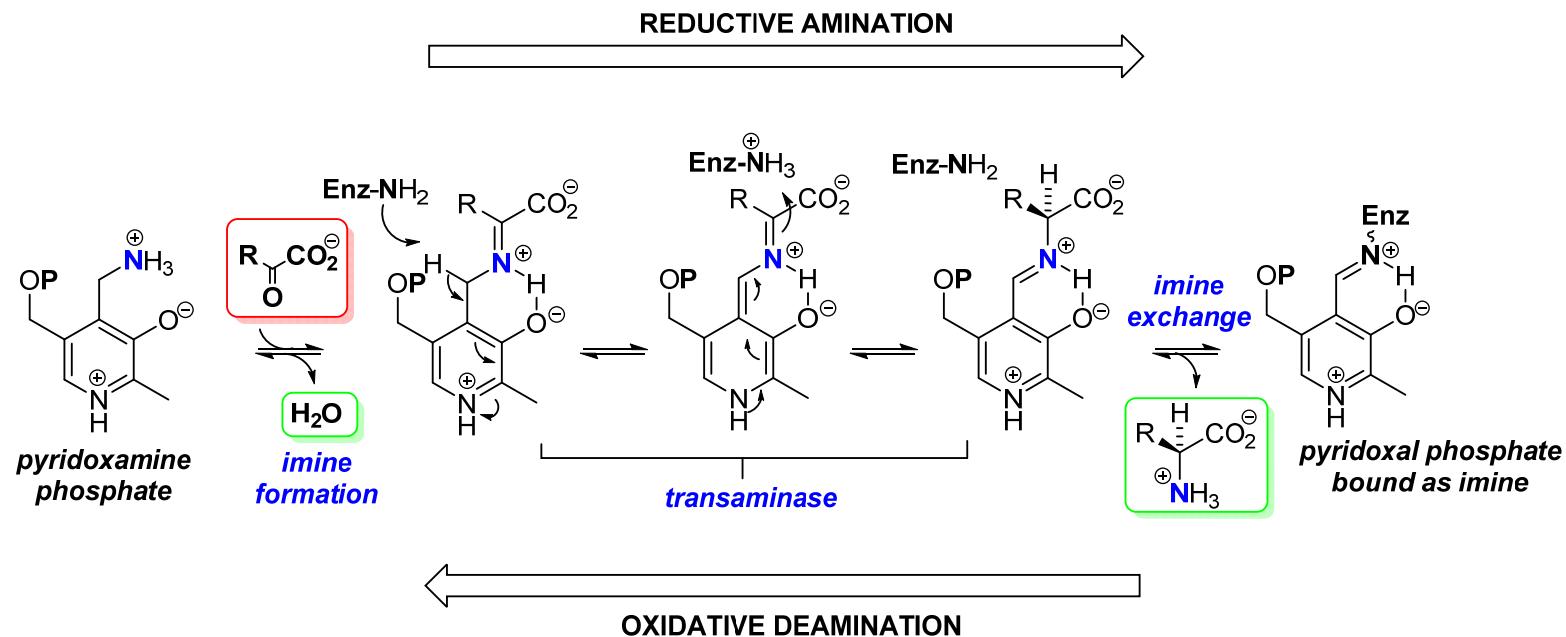
The Biosynthesis of Lysine & Ornithine

- **Lysine & ornithine** - the two most significant, *non-aromatic α-amino acid precursors to alkaloids*:
 - NB. lysine (Lys) is proteinogenic whereas ornithine (Orn) is not
 - phenylalanine (Phe), tyrosine (Tyr) & tryptophan (Trp) from **shikimate** are the other important precursors
 - biosynthesis is via reductive amination of the appropriate α-ketoacid mediated by **pyridoxal-5'-phosphate (PLP)**



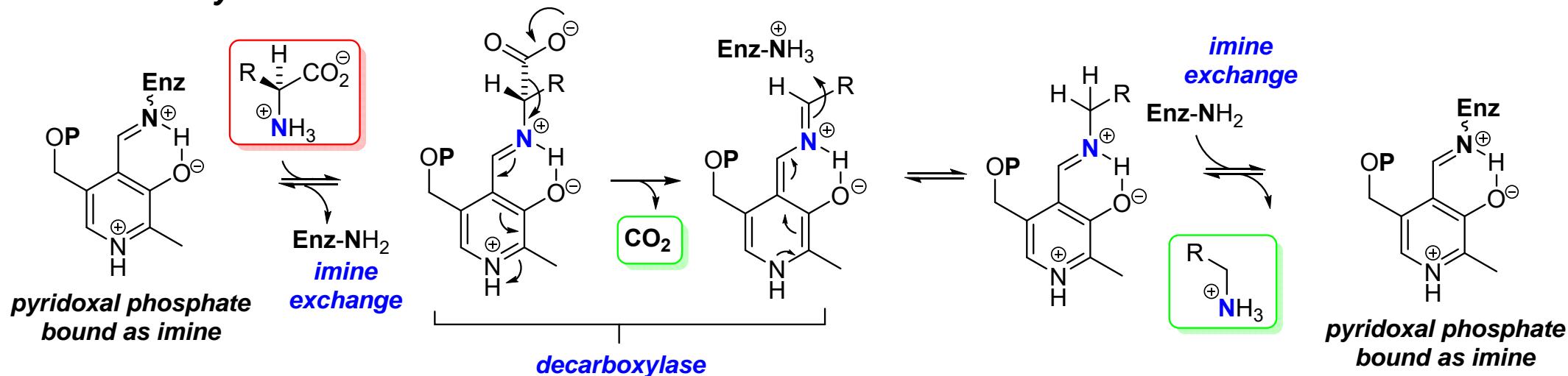
PLP Chemistry – *Transamination & Racemisation*

- **Transamination:**

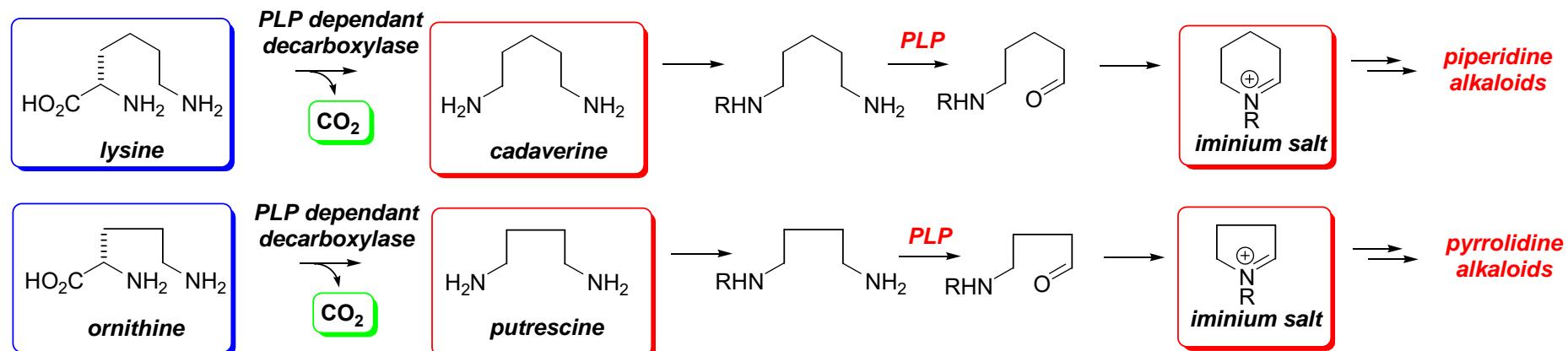


PLP Chemistry – Decarboxylation

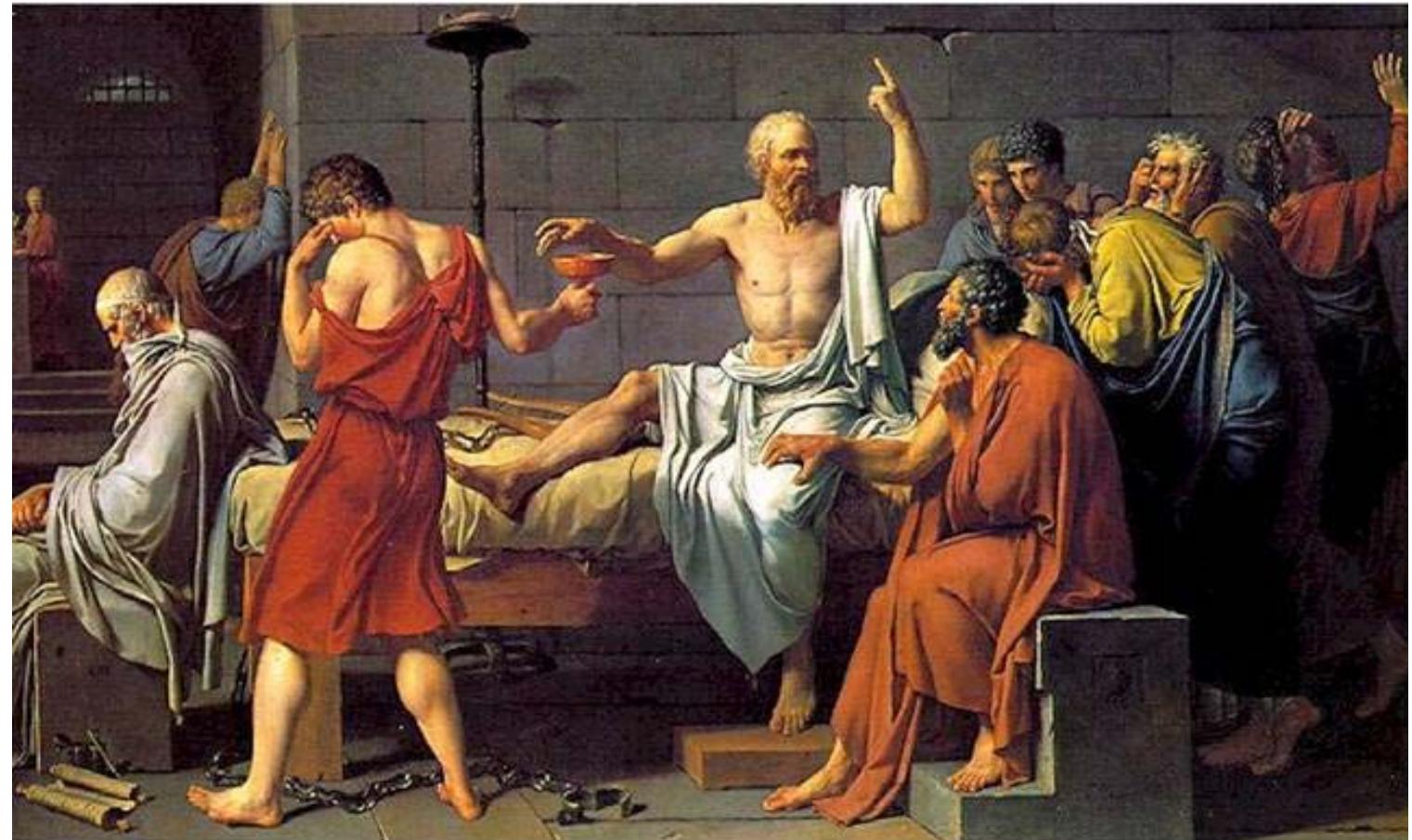
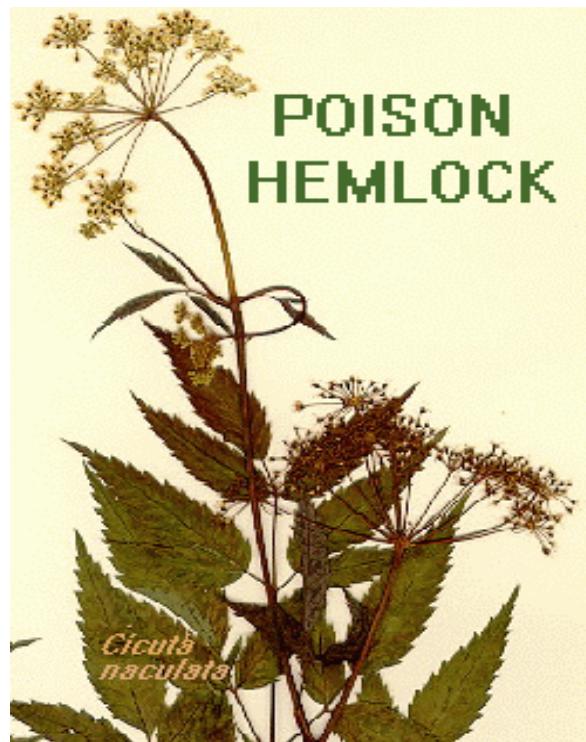
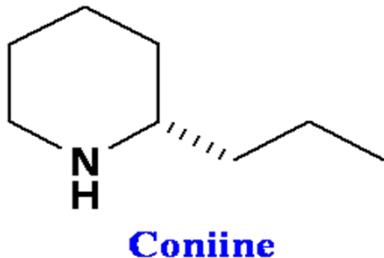
- Decarboxylation:**



- Decarboxylation of **lysine** & **ornithine**:



Lysine-derived Piperidine Alkaloids – *Hemlock!*

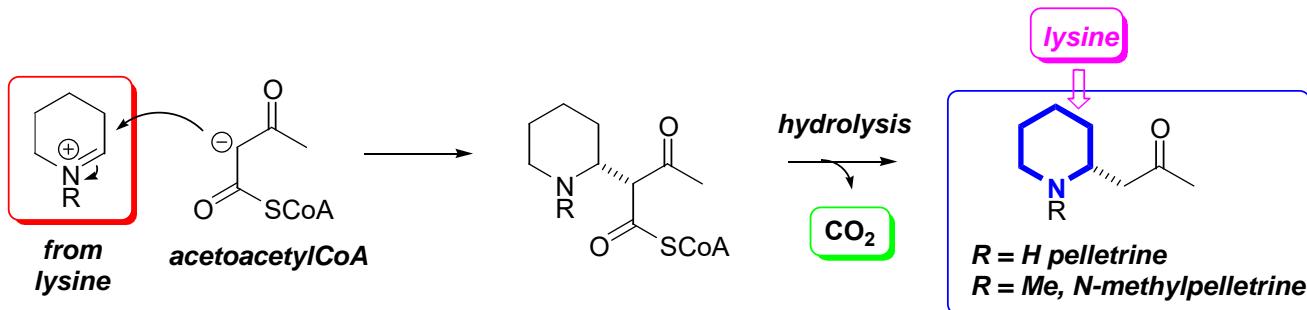


Socrates drinking poison hemlock, 399 B.C.

"The Death of Socrates" by Jacques-Louis David (1787)

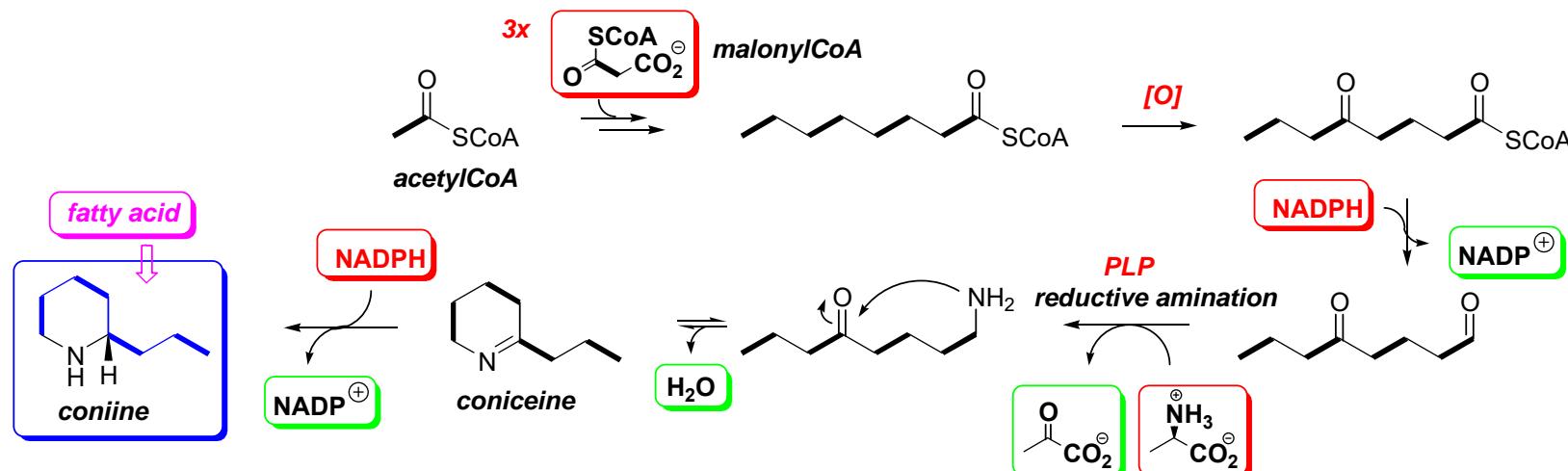
Piperidine Alkaloids – *Pelletierine* & *Coniine*

- ***Pelletierine*:**



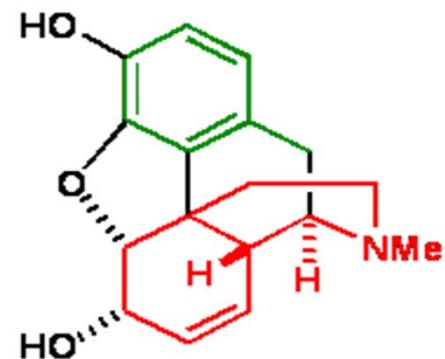
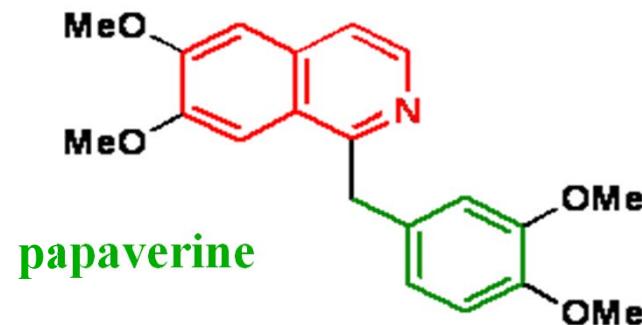
- ***Coniine*:**

- in 399 BC Socrates was sentenced to death for impiety and executed by being forced to drink a potion made from poison hemlock. The toxic component in hemlock is coniine. Although by analogy with the above pathway, biosynthesis from lysine might be suspected, it is in fact of **fatty acid** origin



Tyrosine-derived Alkaloids - Opium Alkaloids

Benzylisoquinoline Alkaloids

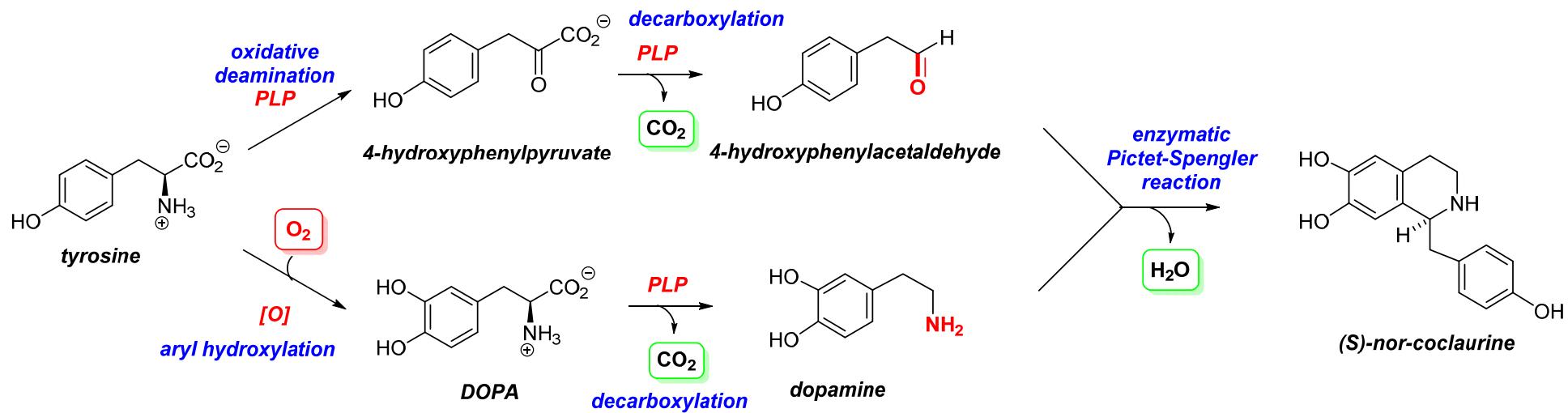


morphine

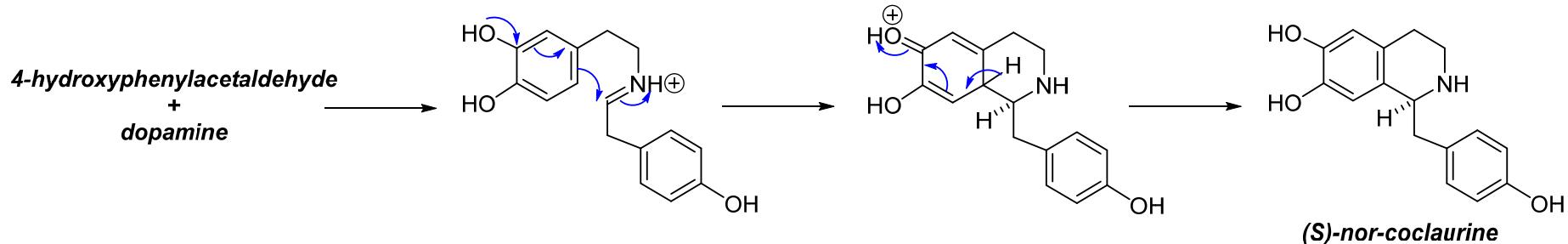


Benzylisoquinoline Alkaloids – Ring Formation

- **Benzylisoquinoline alkaloids** constitute an extremely large and varied group of alkaloids
 - many, particularly the *opium alkaloids* (e.g. papaverine, morphine) are **biosynthesised** from two molecules of tyrosine via ***nor-coclaurine*** (and then ***nor-laudanosoline***).

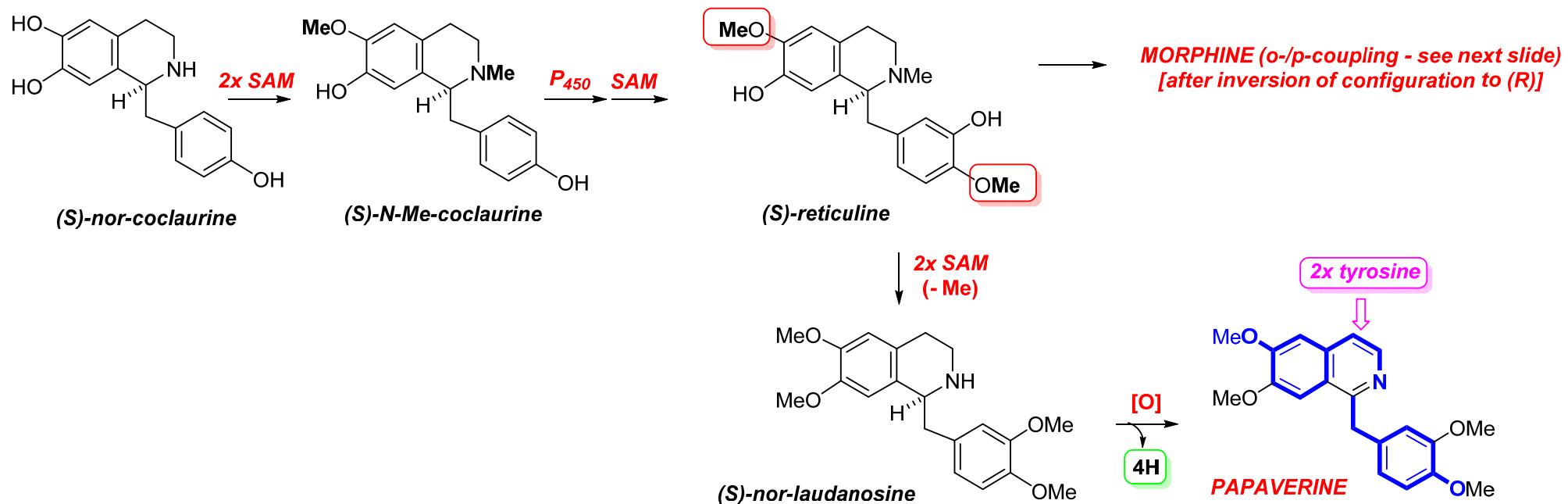


- **Mechanism of Pictet Spengler reaction:**



Benzylisoquinoline Alkaloids - Papaverine

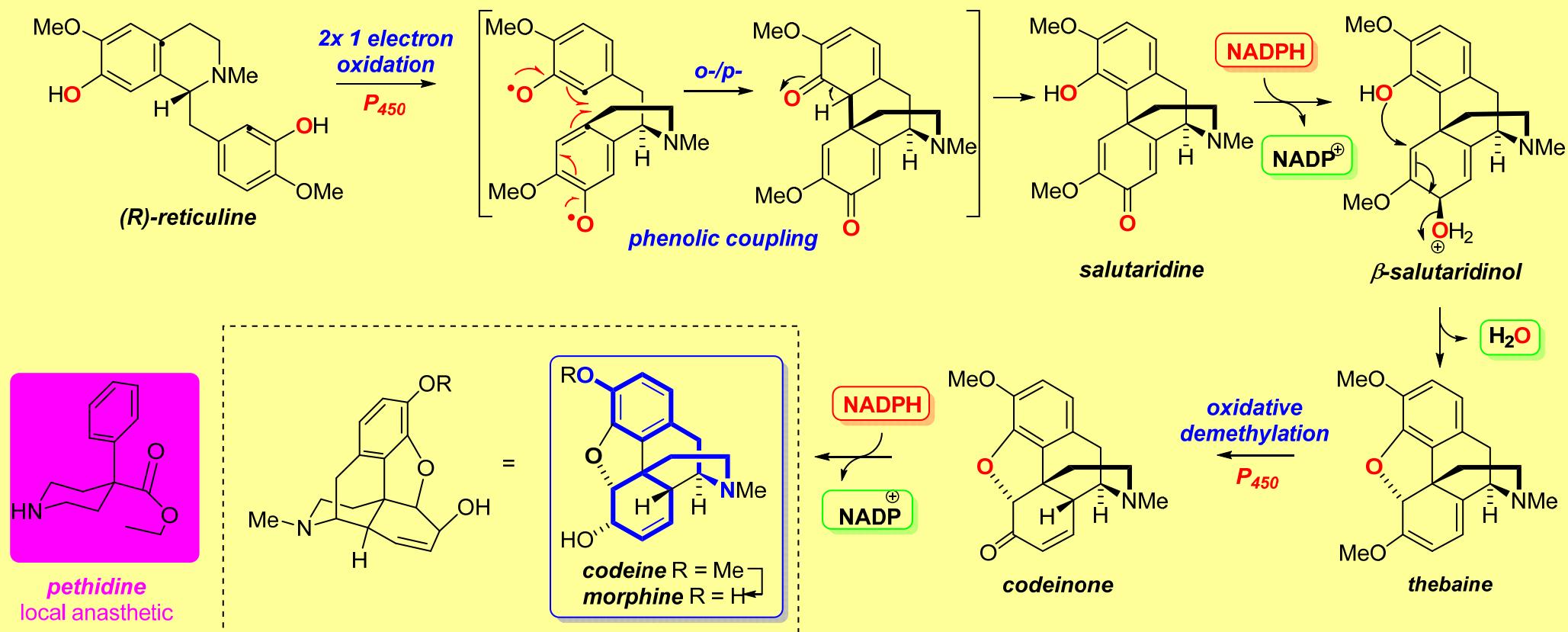
- **Papaverine:** analgesic constituent of the **opium poppy** (*Papaver somniferum*):
 - **biosynthesis:**



- NB. The prefix **nor** means **without a methyl group**. Coclaurine, reticuline and laudanosine are the *N*-methyl compounds

Oxidative Phenolic Coupling – Morphine & Synthetic Opioids

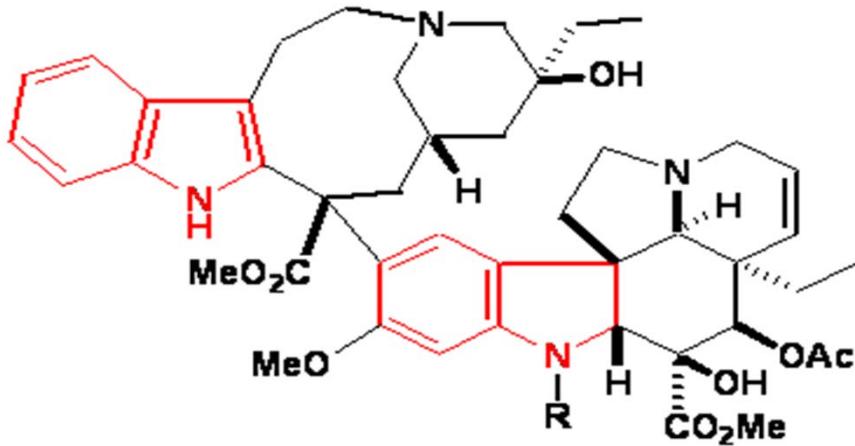
- **Morphine:** analgesic & sedative constituent of the **opium poppy** (*Papaver somniferum*):
 - **biosynthesis:** o-/p- oxidative phenolic coupling of **reticuline**:



- Morphine acts by activating the **opiate receptors** in the brain (IC_{50} 3 nM)
- The natural ligands for these receptors are peptides: e.g. Leu-enkephalin (Tyr–Gly–Gly–Phe–Leu) (IC_{50} 12 nM)

Dimeric Indole Alkaloids – *Vinca* extracts

Dimeric Indole Alkaloids



vinblastine ($\text{R} = \text{Me}$)

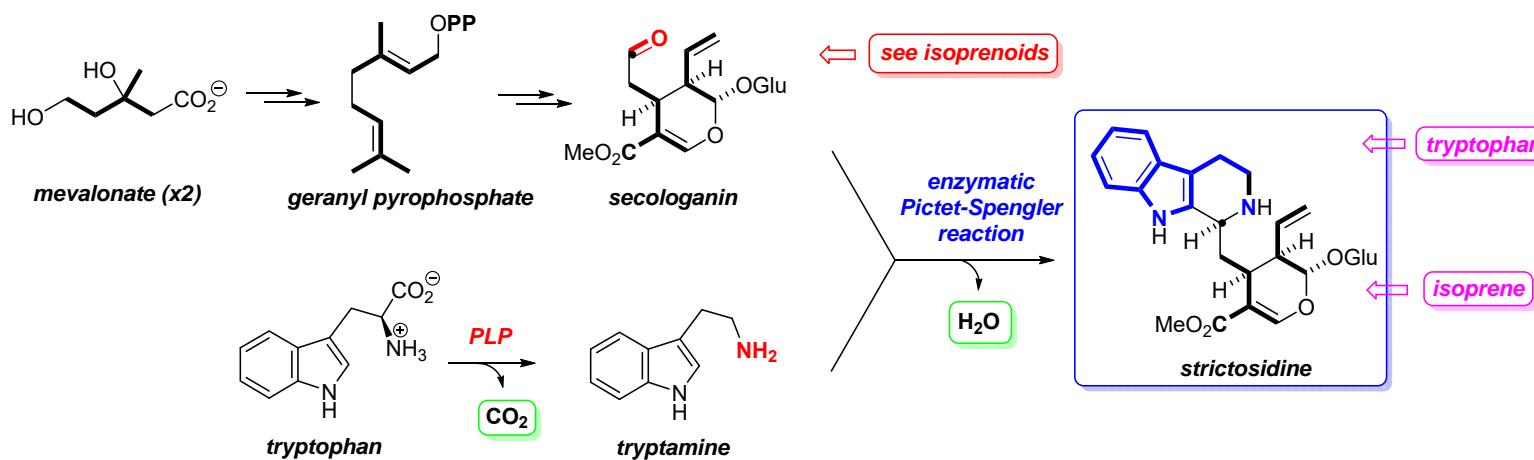
vincristine ($\text{R} = \text{CHO}$)



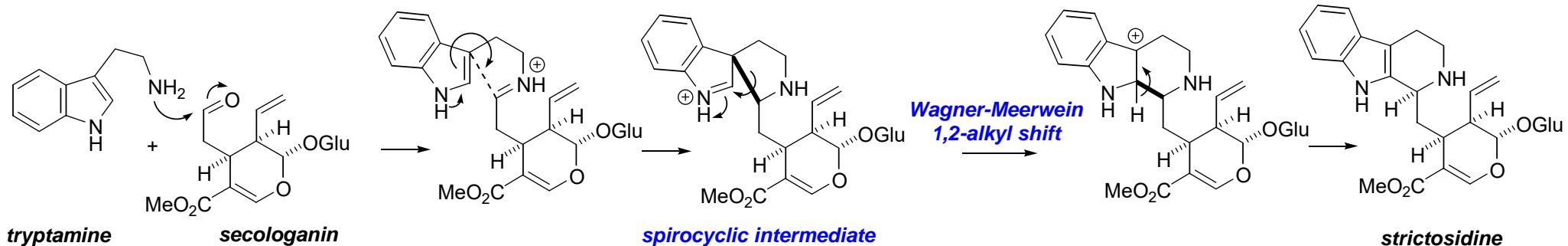
Potent anti tumour alkaloids used in cancer chemotherapy

Tryptamine + Secologanin → Strictosidine

- Most alkaloids of ***mixed Tryptophan/mevalonate biogenesis*** (>1200) are derived from ***strictosidine***:
 - Strictosidine*** is derived from the condensation of ***tryptamine*** with the iridoid C₁₀ monoterpenoid ***secologanin***:

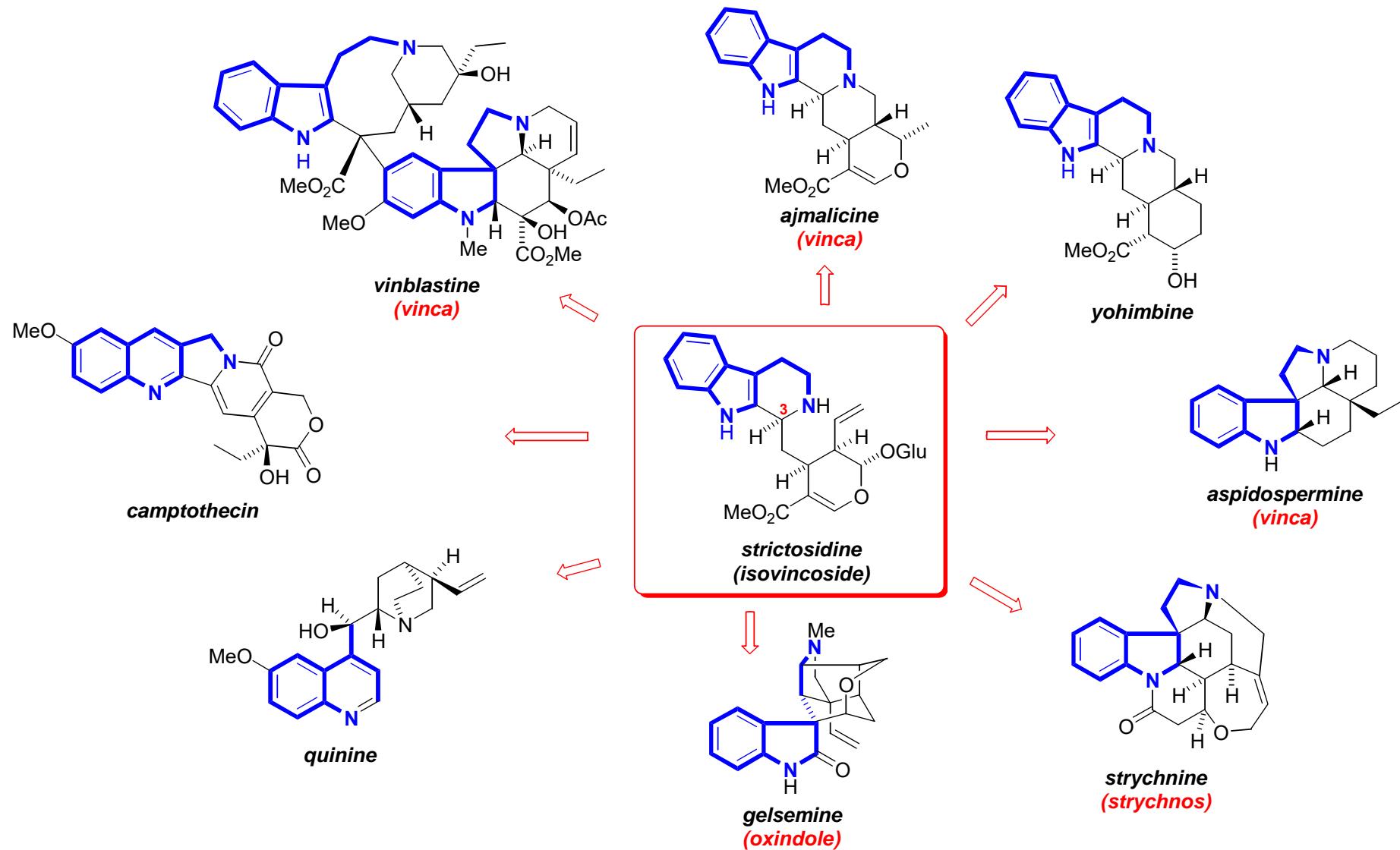


- Mechanism of Pictet-Spengler reaction:
 - via spirocyclic intermediate then Wagner-Meerwein 1,2-alkyl shift:

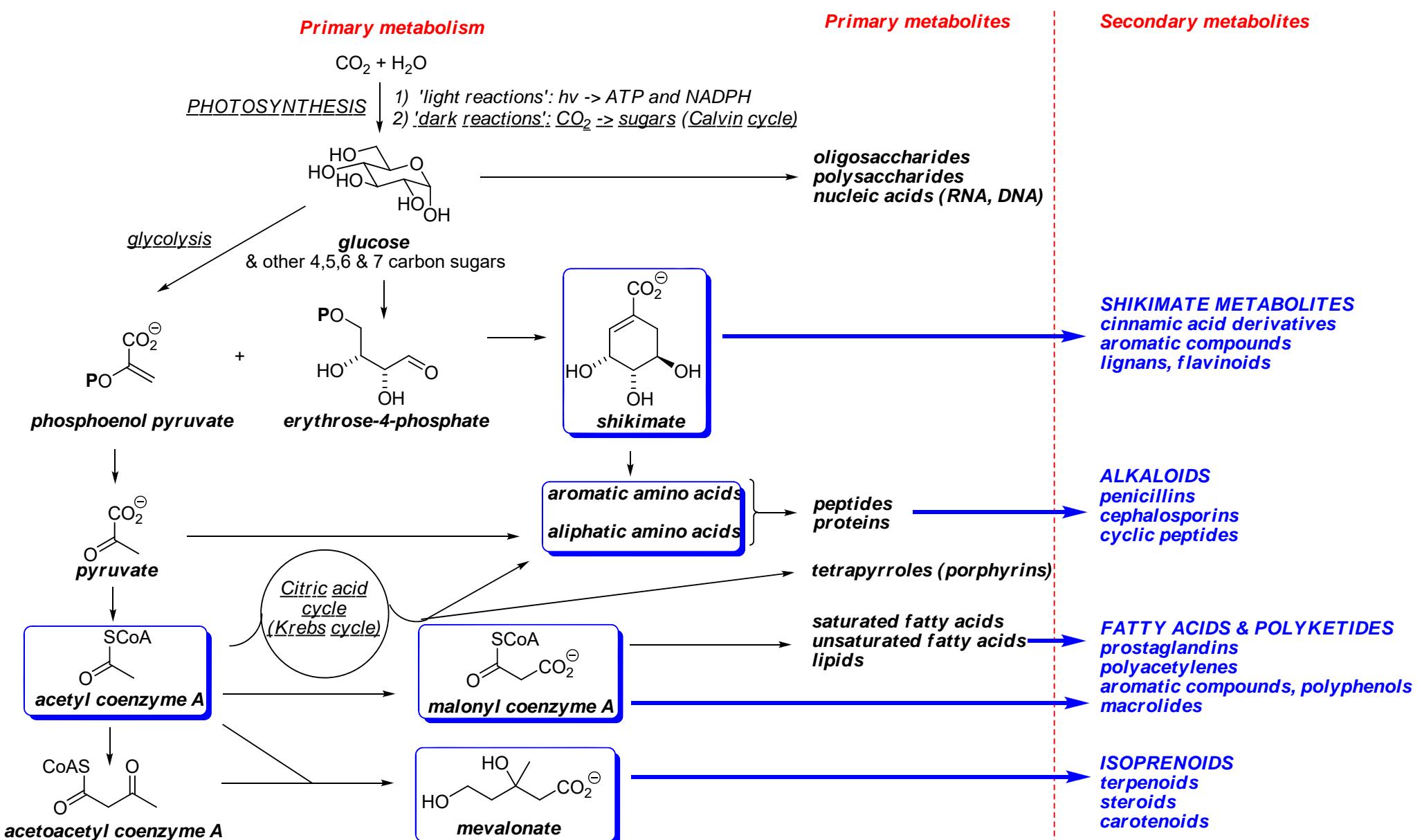


Strictosidine → Vinca, Strychnos, Quinine etc.

- The diversity of alkaloids derived from **strictosidine** is stunning and many pathways remain to be fully elucidated:



Primary Metabolism - Overview

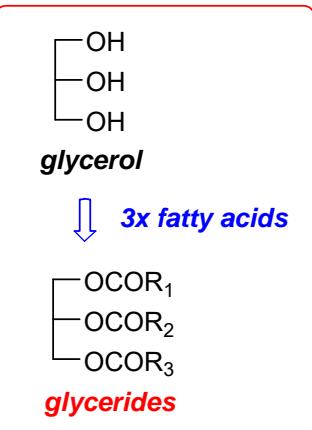


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Fatty Acids

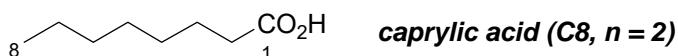
Fatty Acid Primary Metabolites

- **Primary metabolites:**
 - **fully saturated, linear carboxylic acids** & derived **(poly)unsaturated derivatives**:
 - constituents of essential natural waxes, seed oils, **glycerides** (fats) & phospholipids
 - **structural role** – **glycerides** & phospholipids are essential constituents of cell membranes
 - **energy storage** – **glycerides** can also be catabolised into acetate → citric acid cycle
 - **biosynthetic precursors** – for elaboration to secondary metabolites

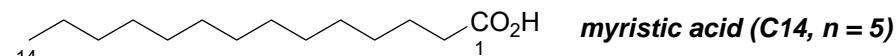


SATURATED ACIDS [$\text{MeCH}_2(\text{CH}_2\text{CH}_2)_n\text{CH}_2\text{CO}_2\text{H}$ ($n = 2-8$)]

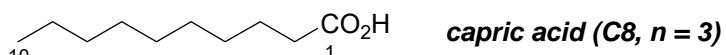
e.g.



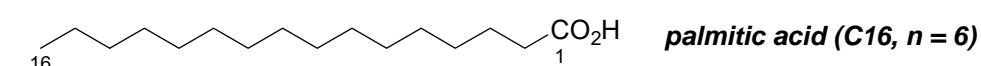
caprylic acid (C8, n = 2)



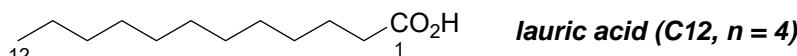
myristic acid (C14, n = 5)



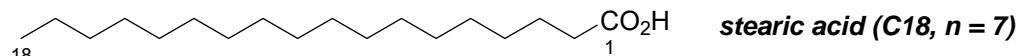
capric acid (C8, n = 3)



palmitic acid (C16, n = 6)

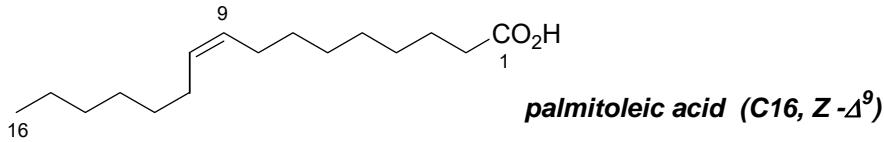


lauric acid (C12, n = 4)

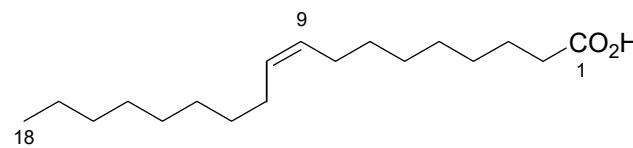


stearic acid (C18, n = 7)

MONO-UNSATURATED ACID DERIVATIVES (MUFAs) e.g.



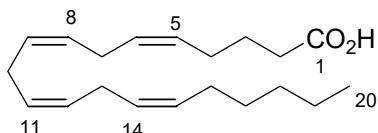
palmitoleic acid (C16, Z -Δ⁹)



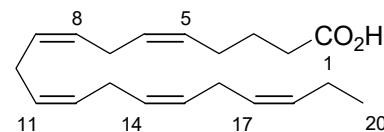
oleic acid (C18, Z-Δ⁹)
(>80% of fat in olive oil)

POLY-UNSATURATED ACID DERIVATIVES (PUFAs)

e.g.



arachidonic acid (AA)
 $(C20, Z\text{-}\Delta^5, Z\text{-}\Delta^8, Z\text{-}\Delta^{11}, Z\text{-}\Delta^{14})$



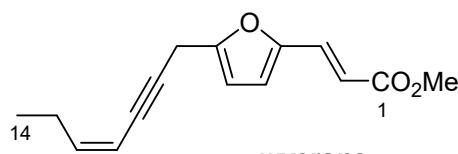
eicosapentaenoic acid (EPA)
 $(C20, Z\text{-}\Delta^5, Z\text{-}\Delta^8, Z\text{-}\Delta^{11}, Z\text{-}\Delta^{14}, Z\text{-}\Delta^{17})$
(in cod liver oil)

Fatty Acids Derivatives – Secondary Metabolites

- **Secondary metabolites**

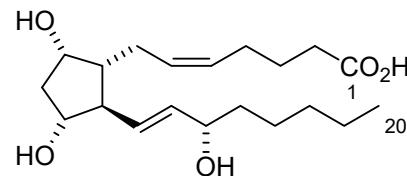
- further **elaborated** derivatives of **polyunsaturated fatty acids (PUFAs)**
 - e.g. polyacetylenes & ‘eicosanoids’ (prostaglandins, thromboxanes & leukotrienes)

POLYACETYLENES
e.g.



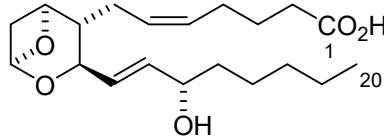
wyerone
anti-fungal

PROSTAGLANDINS
e.g.



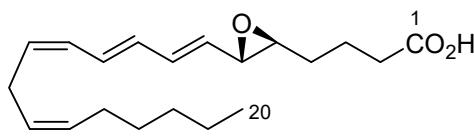
prostaglandin $F_{2\alpha}$ ($\text{PGF}_{2\alpha}$)

THROMBOXANES
e.g.

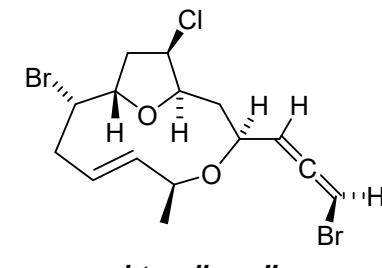


thromboxane A_2 (TXA_2)

LEUKOTRIENES
e.g.



leukotriene A_4 (LTA_4)

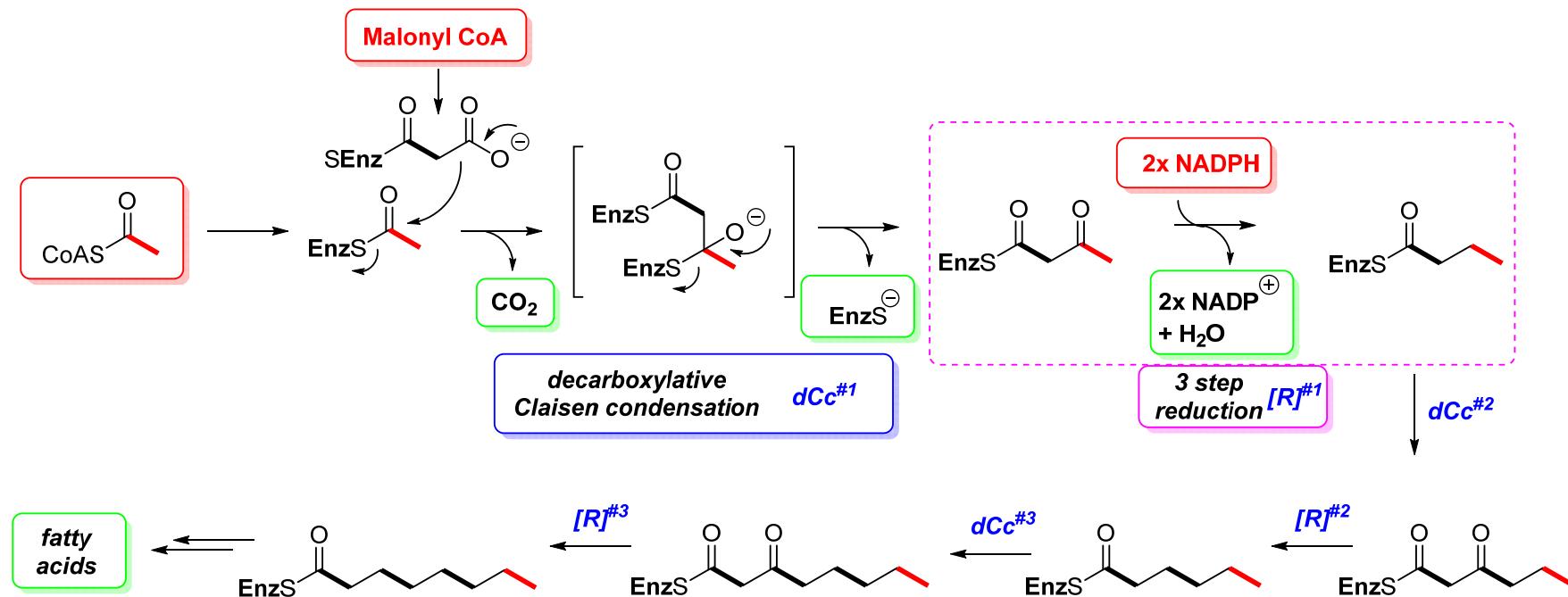


obtusallene II
marine natural product

EICOSANOIDS

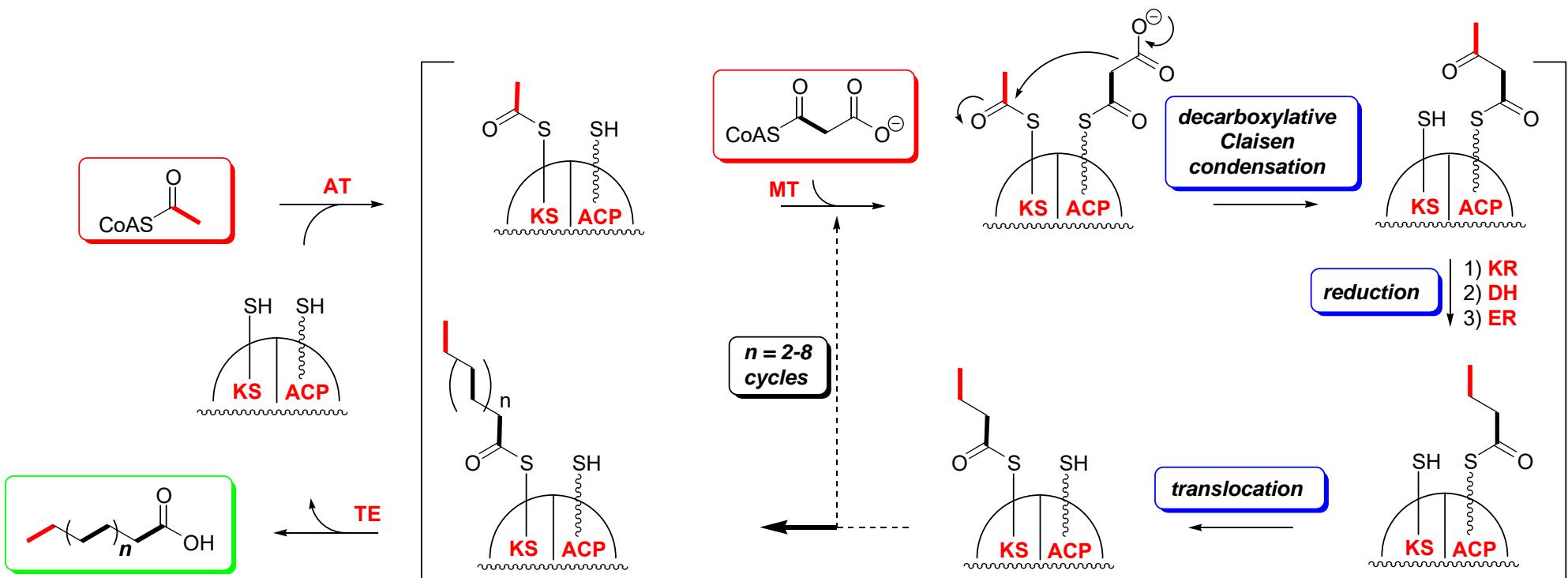
Biosynthesis of Fatty Acids – Iterative Oligomerisation

- **fatty acids** are biosynthesised from **acetyl CoA** as a **starter unit** by **iterative** ‘head-to-tail’ **oligomerisation** involving:
 - condensation with **malonyl CoA** as an **extender unit** (with loss of **CO_2**) – a **decarboxylative Claisen condensation**
 - 3-step **reduction** of the resulting **ketone** → **methylene**
- after **n = 2-8 iterations** the **C8-20 saturated fatty acid** is released from the enzyme(s):



Biosynthesis of Fatty Acids – Overview of FAS

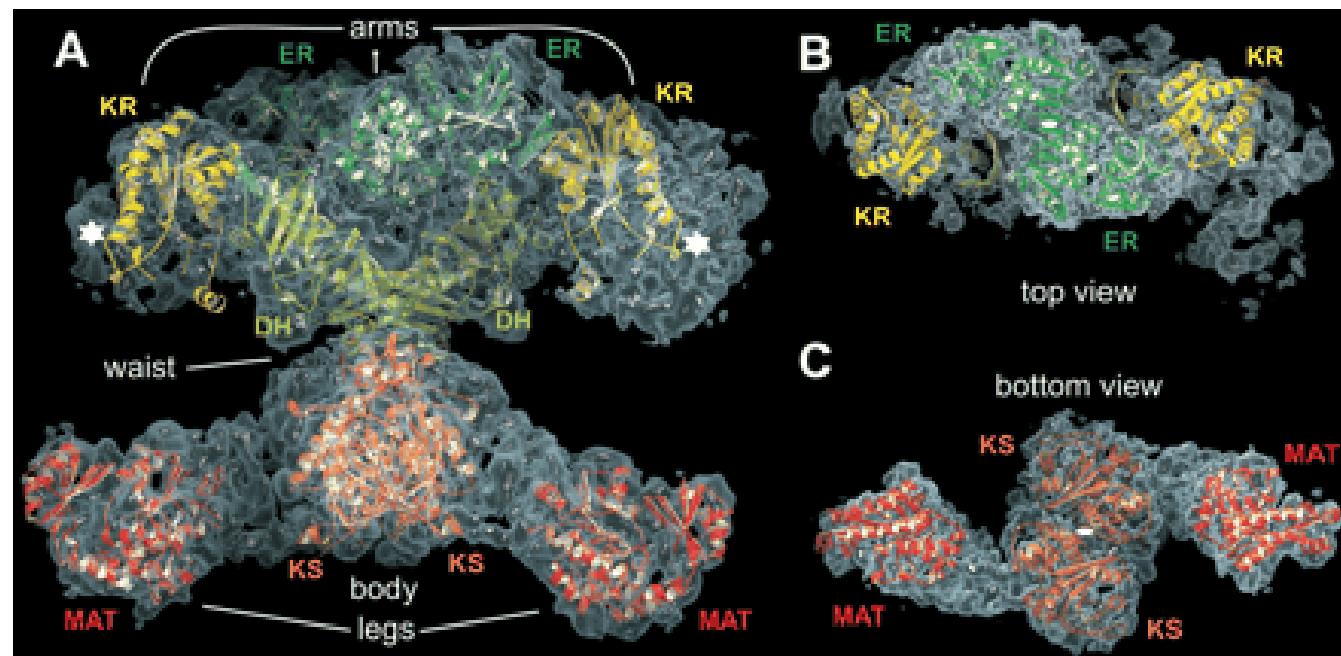
- The *in vivo* process by which all this takes place involves a ‘molecular machine’ - **Fatty Acid Synthase (FAS)**
 - Type I FAS: single multifunctional protein complex** (e.g. in mammals incl. humans)
 - Type II FAS: set of discrete, dissociable single-function proteins** (e.g. in bacteria)
 - All FASs** comprise **8 components** (ACP & 7× catalytic activities): **ACP, KS, AT, MT, KR, DH, ER & [TE]** :



KS = keto synthase (also known as **CE** = condensing enzyme); **AT** = acetyl transferase; **MT** = malonyl transferase;
KR = keto reductase; **DH** = dehydratase; **ER** = enoyl reductase; **TE** = thioesterase; **ACP** = acyl carrier protein

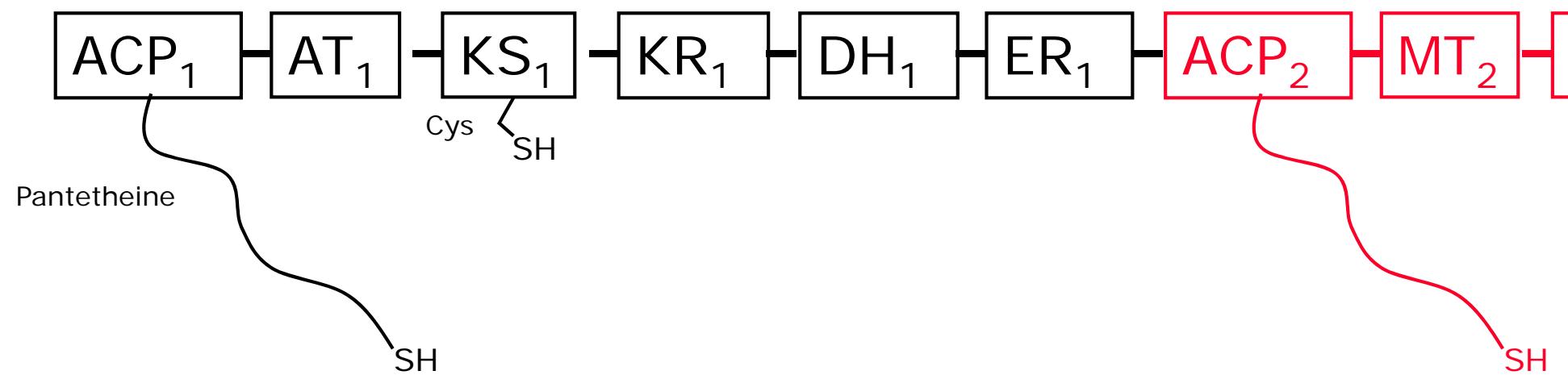
Human Fatty Acid Synthase (FAS)

- the first three-dimensional structure of human fatty acid synthase (272 kDa) at 4.5 Å resolution by X-ray crystallography:
 - Maier, Jenni & Ban *Science* **2006**, 311, 1258 ([DOI](#)) ; also Fungal FAS @ 3.1 Å resolution see: Jenni *et al.* *Science* **2007**, 316, 254 & 288



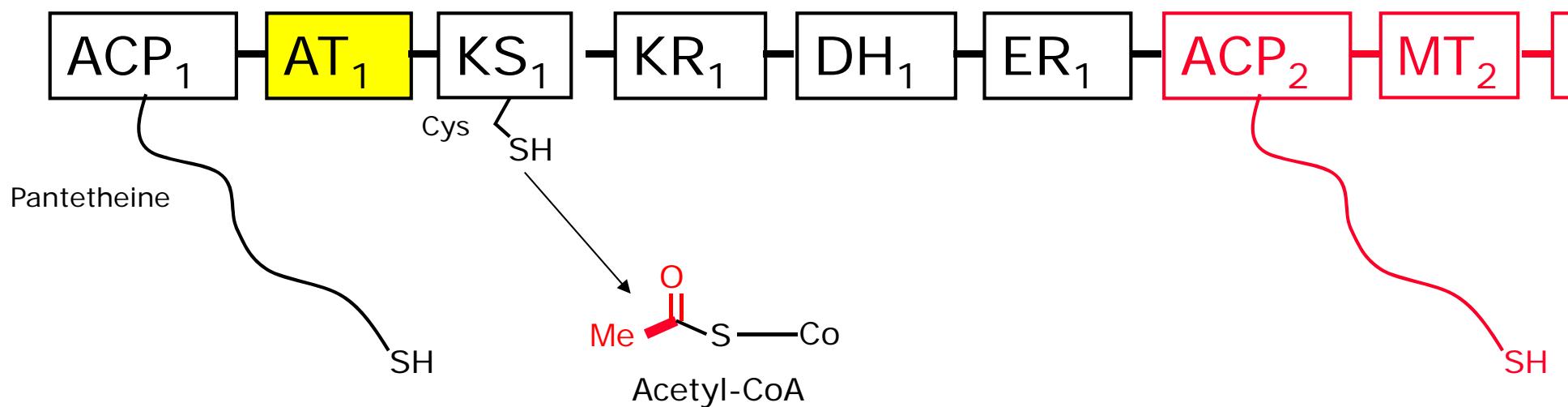
Structural overview. **(A)** Front view: FAS consists of a lower part comprising the KS (lower body) and MAT domains (legs) connected at the waist with an upper part formed by the DH, ER (upper body), and KR domains (arms). **(B)** Top view of FAS with the ER and KR domains resting on the DH domains. **(C)** Bottom view showing the arrangement of the KS and MAT domains and the continuous electron density between the KS and MAT domains

FATTY ACID BIOSYNTHESIS (type II FAS)



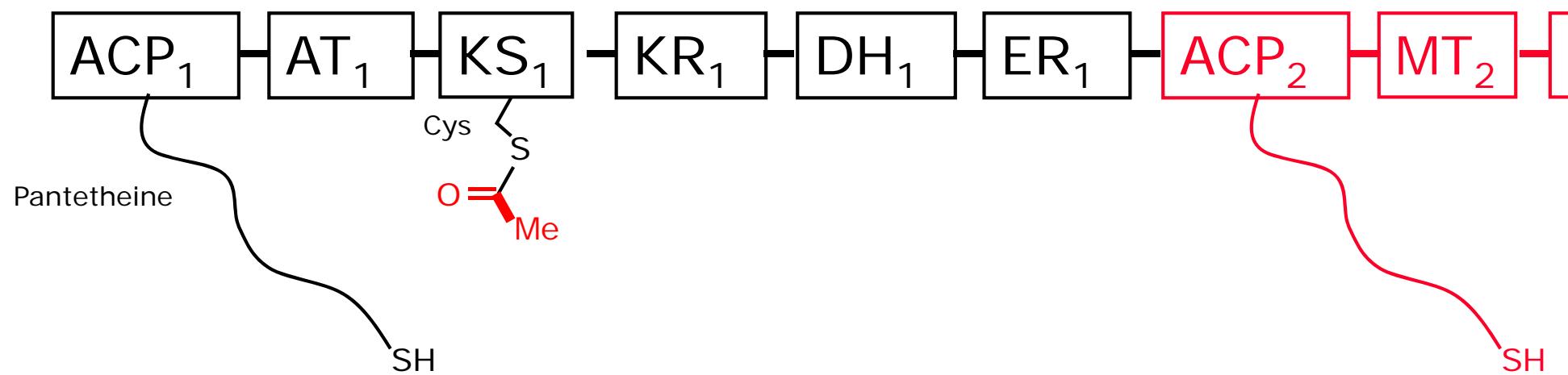
NB. the following sequence of slides have been adapted from: <http://www.courses.fas.harvard.edu/%7echem27/>

FATTY ACID BIOSYNTHESIS

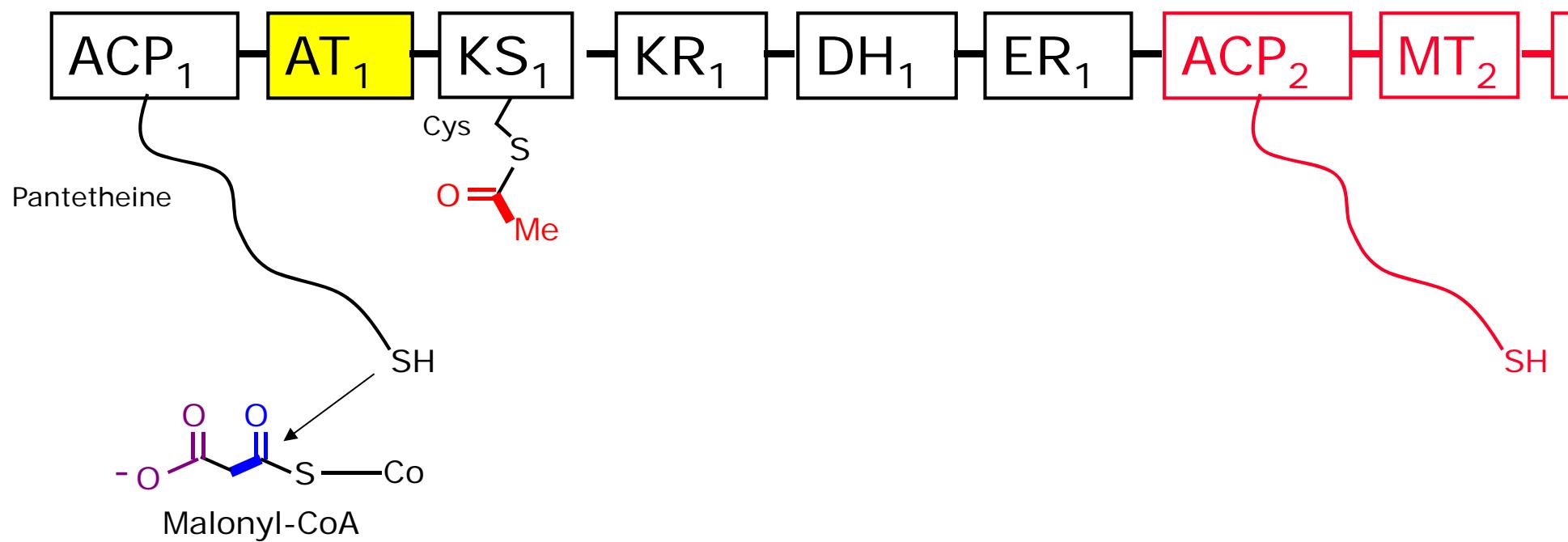


- AT_1 loads acetyl group onto KS_1

FATTY ACID BIOSYNTHESIS

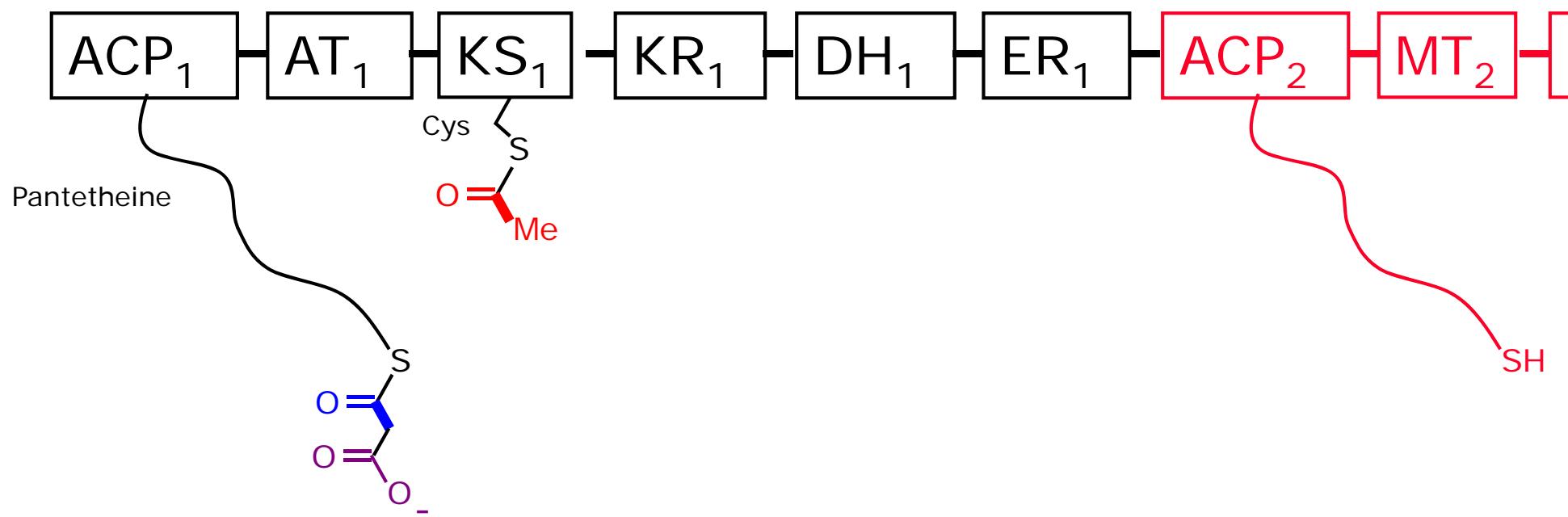


FATTY ACID BIOSYNTHESIS

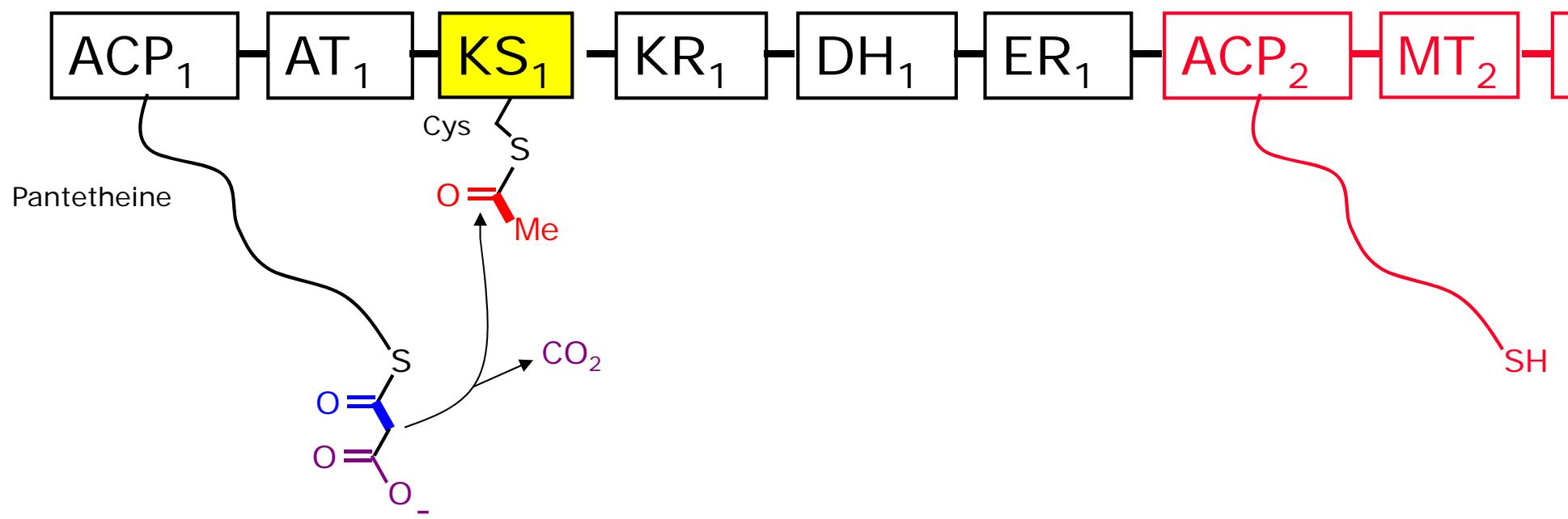


- AT_1 loads malonyl group onto ACP_1

FATTY ACID BIOSYNTHESIS

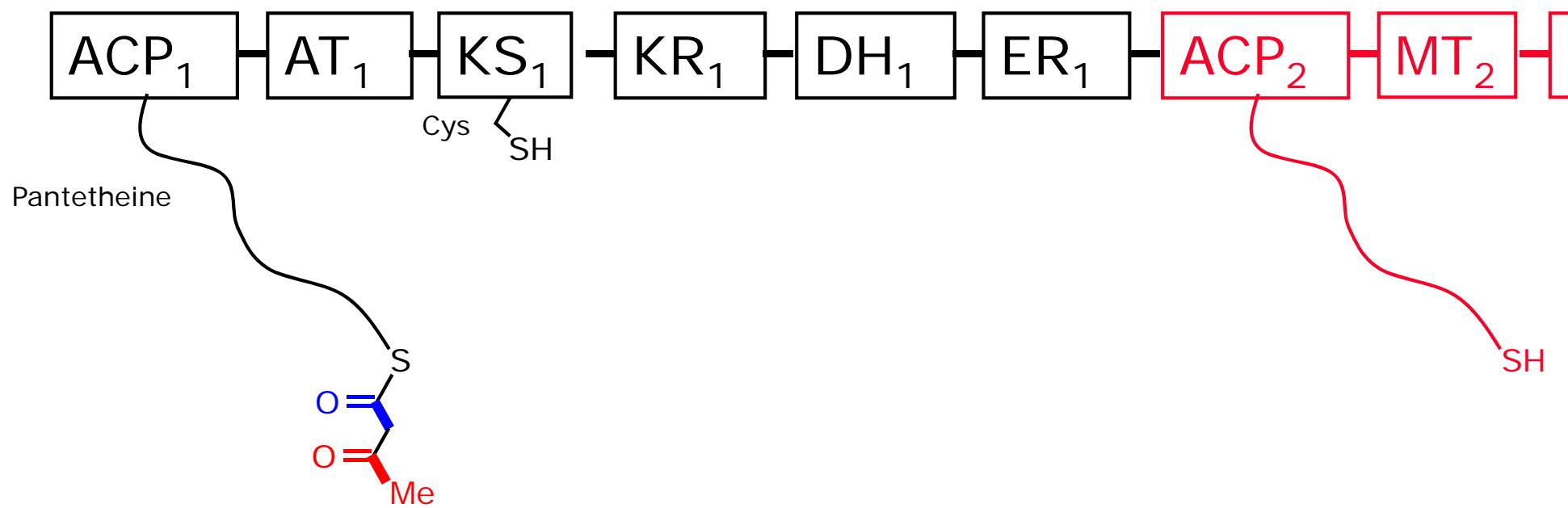


FATTY ACID BIOSYNTHESIS

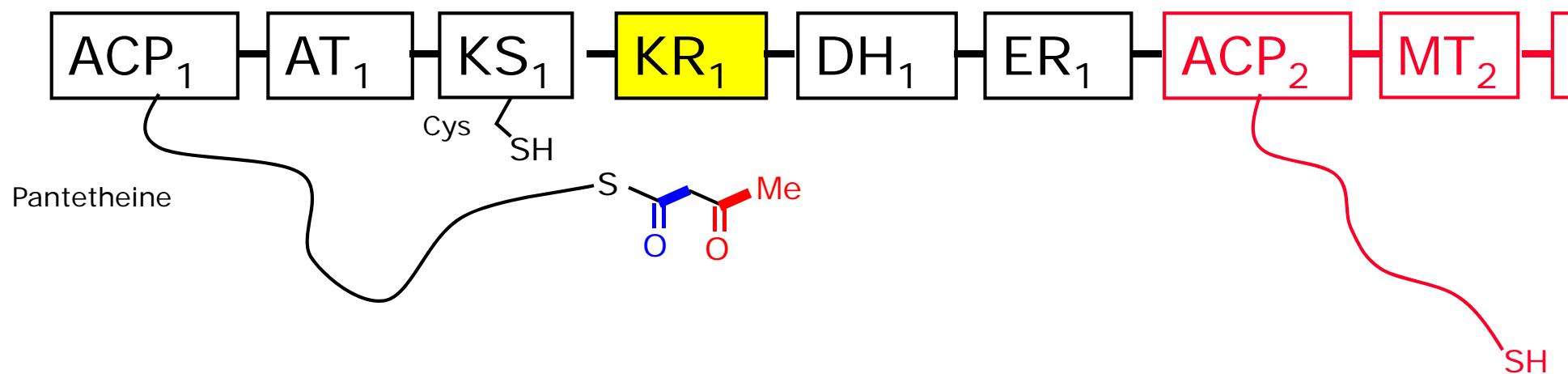


- KS_1 catalyzes Claisen condensation

FATTY ACID BIOSYNTHESIS

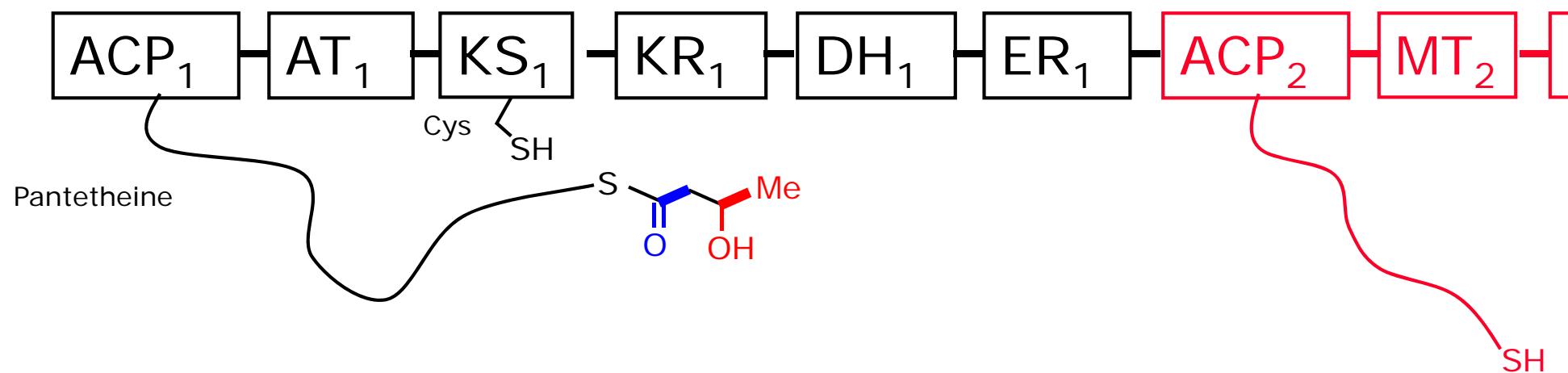


FATTY ACID BIOSYNTHESIS

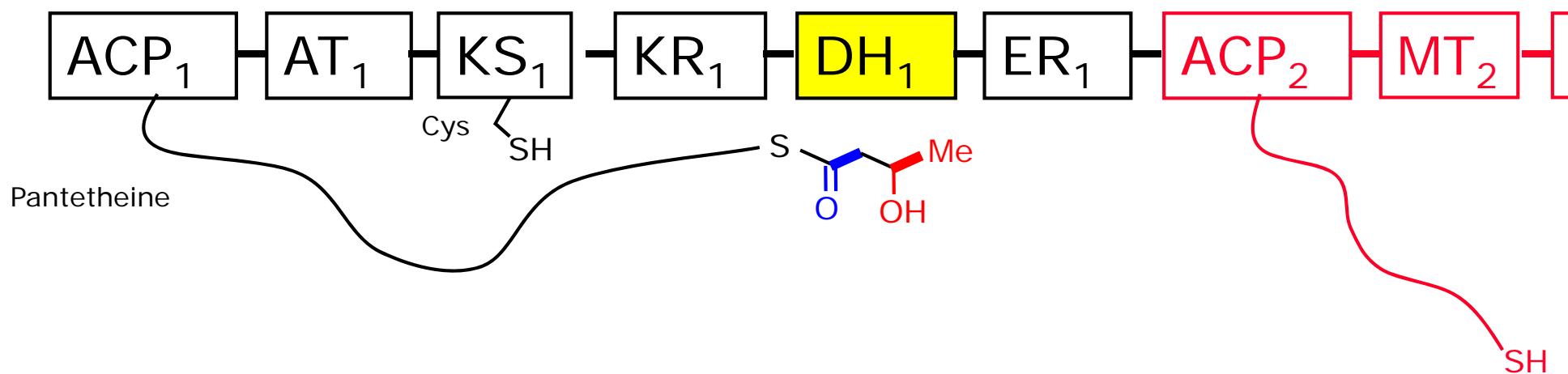


- KR₁ catalyzes reduction of ketone

FATTY ACID BIOSYNTHESIS

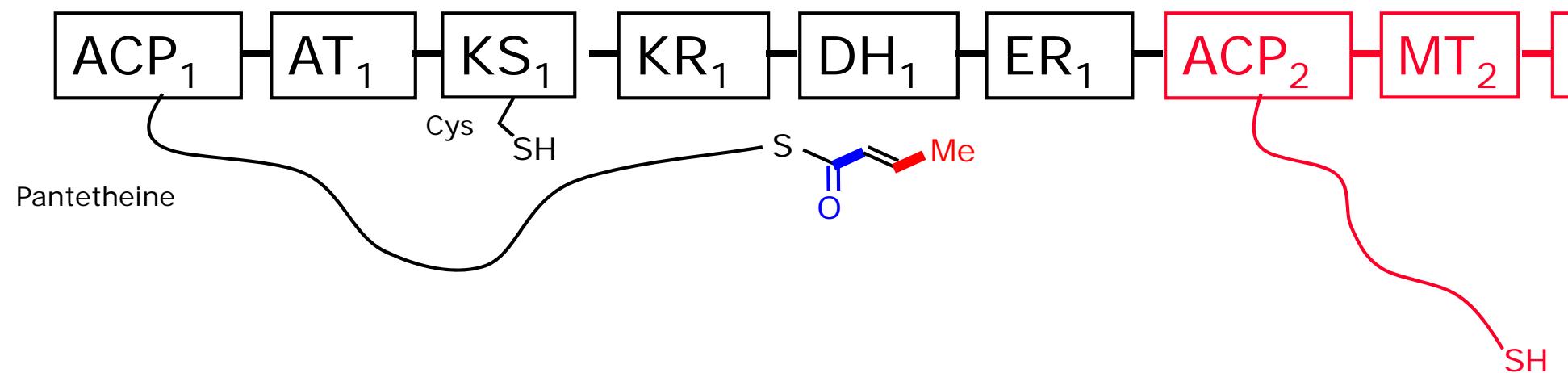


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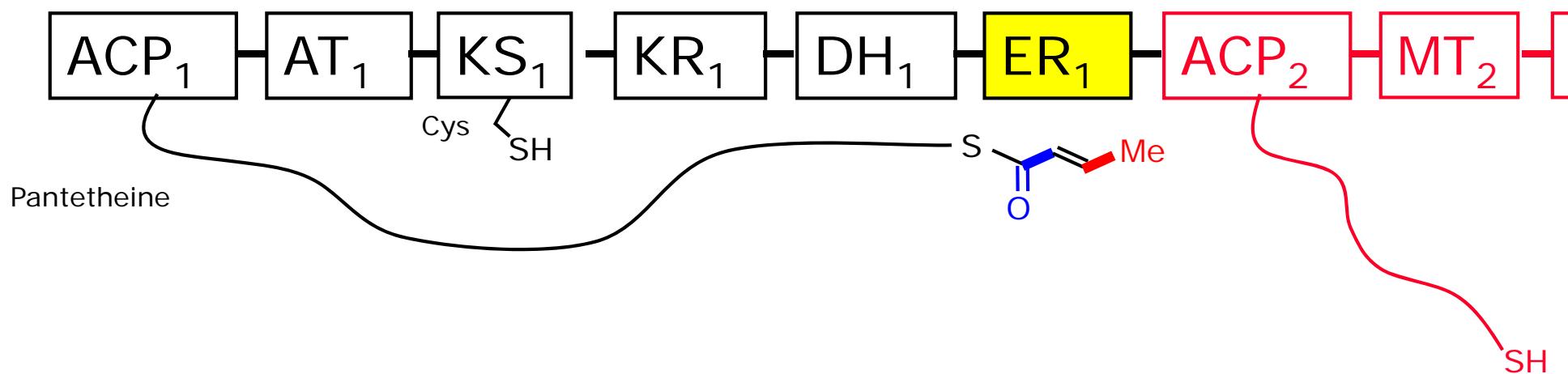


- DH_1 catalyzes dehydration of alcohol

FATTY ACID BIOSYNTHESIS

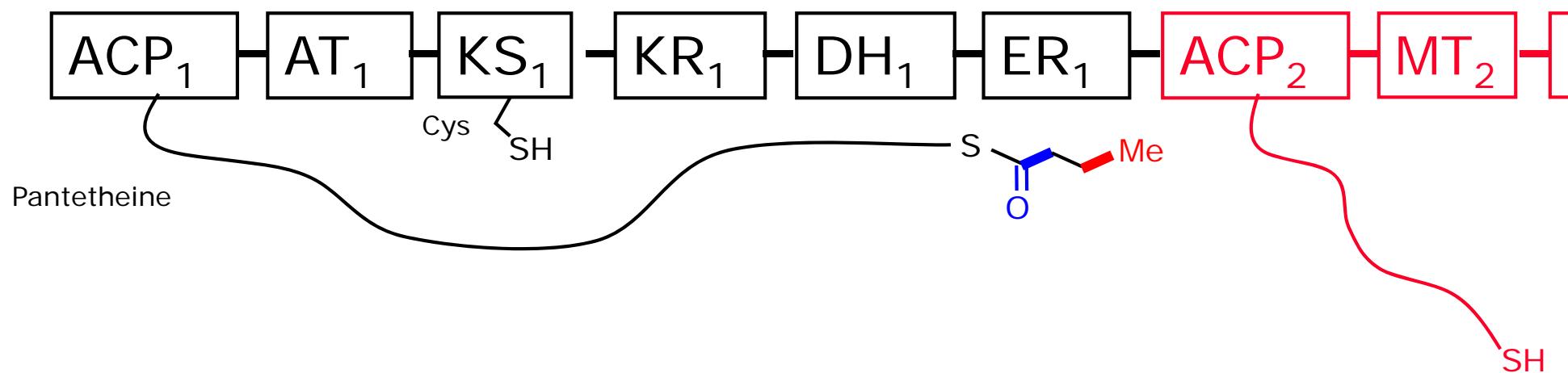


FATTY ACID BIOSYNTHESIS

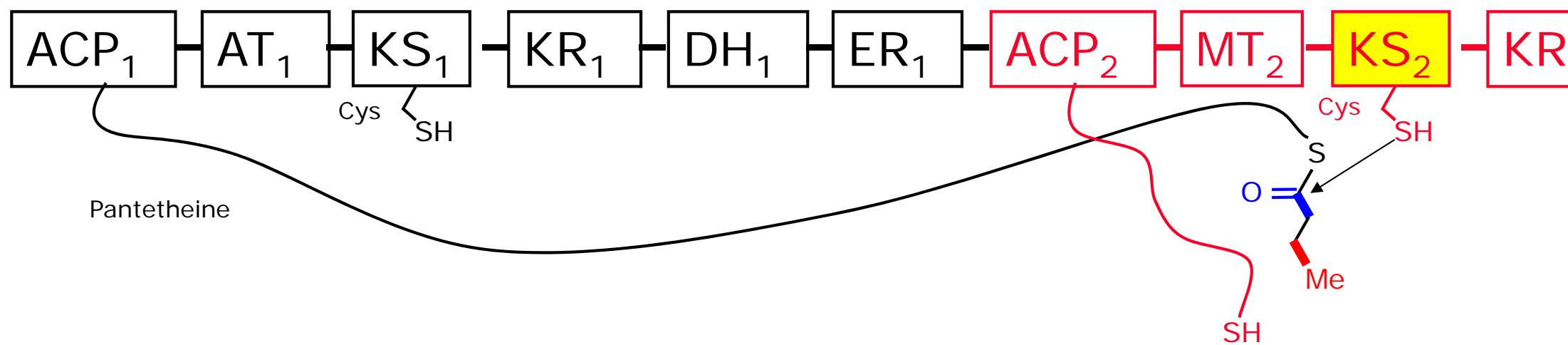


- ER_1 catalyzes reduction of alkene

FATTY ACID BIOSYNTHESIS

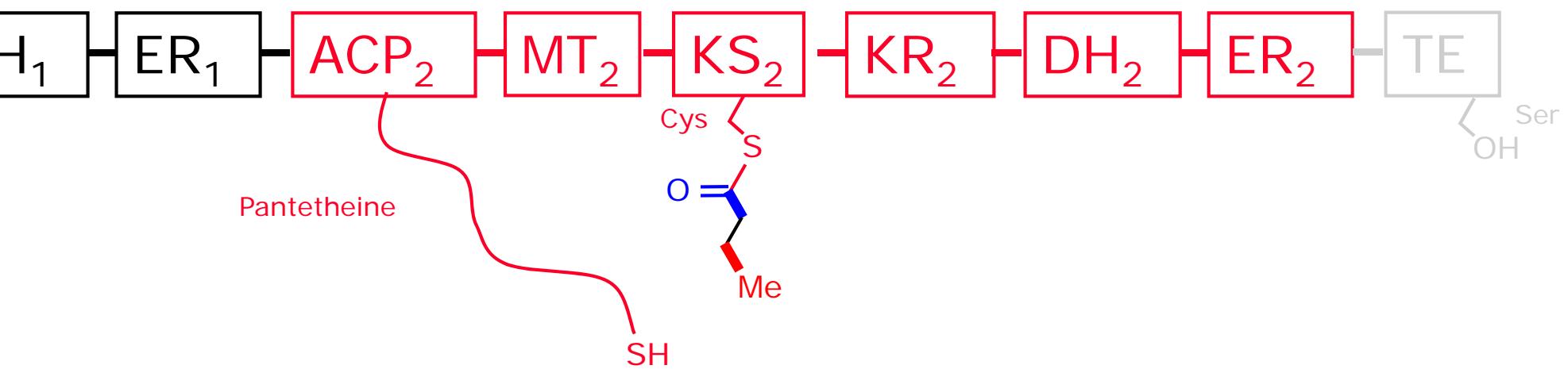


FATTY ACID BIOSYNTHESIS

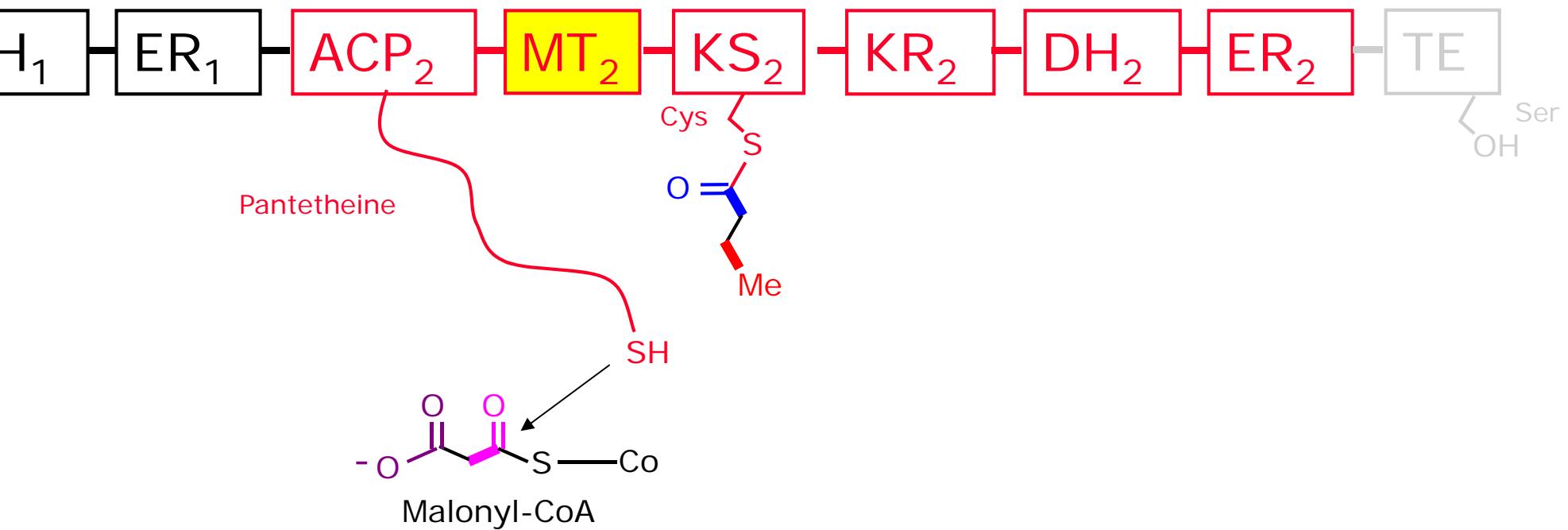


- KS₂ catalyzes translocation to module 2

FATTY ACID BIOSYNTHESIS

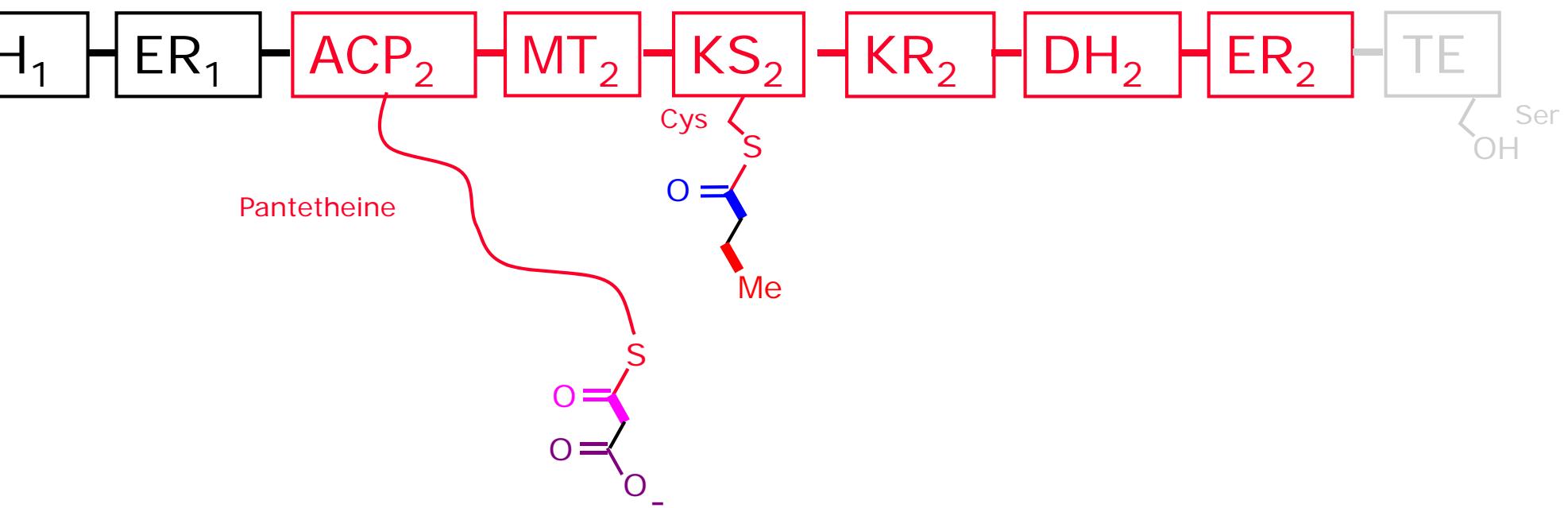


FATTY ACID BIOSYNTHESIS

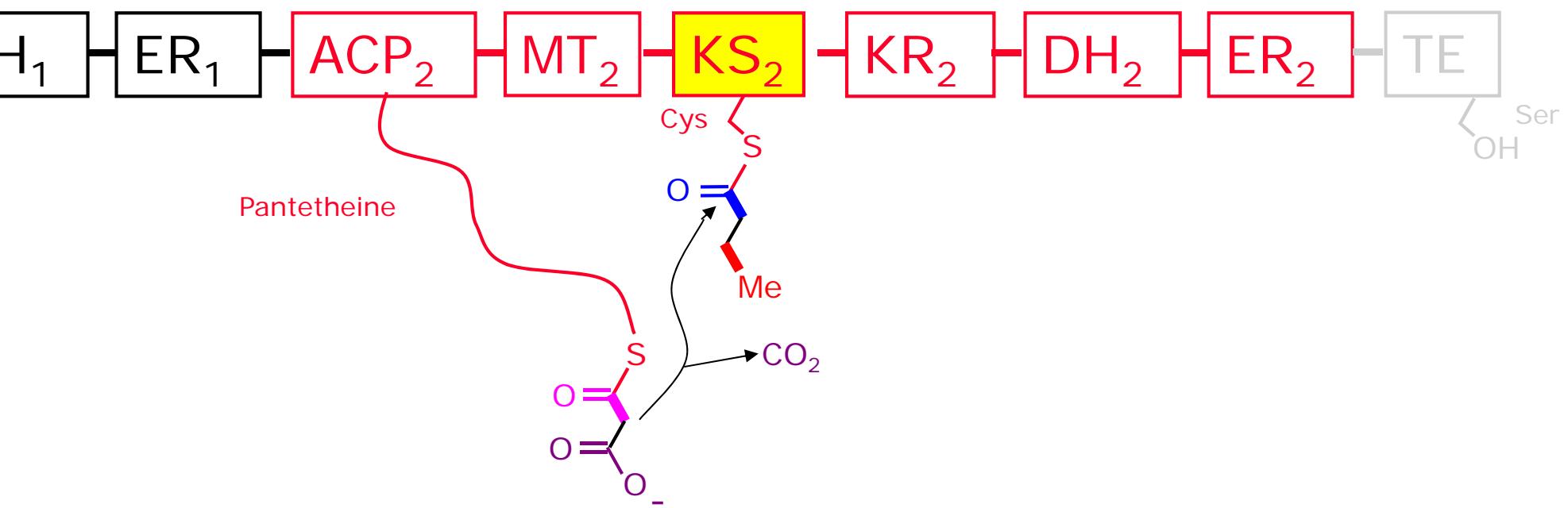


- MT₂ loads malonyl group onto ACP₂

FATTY ACID BIOSYNTHESIS

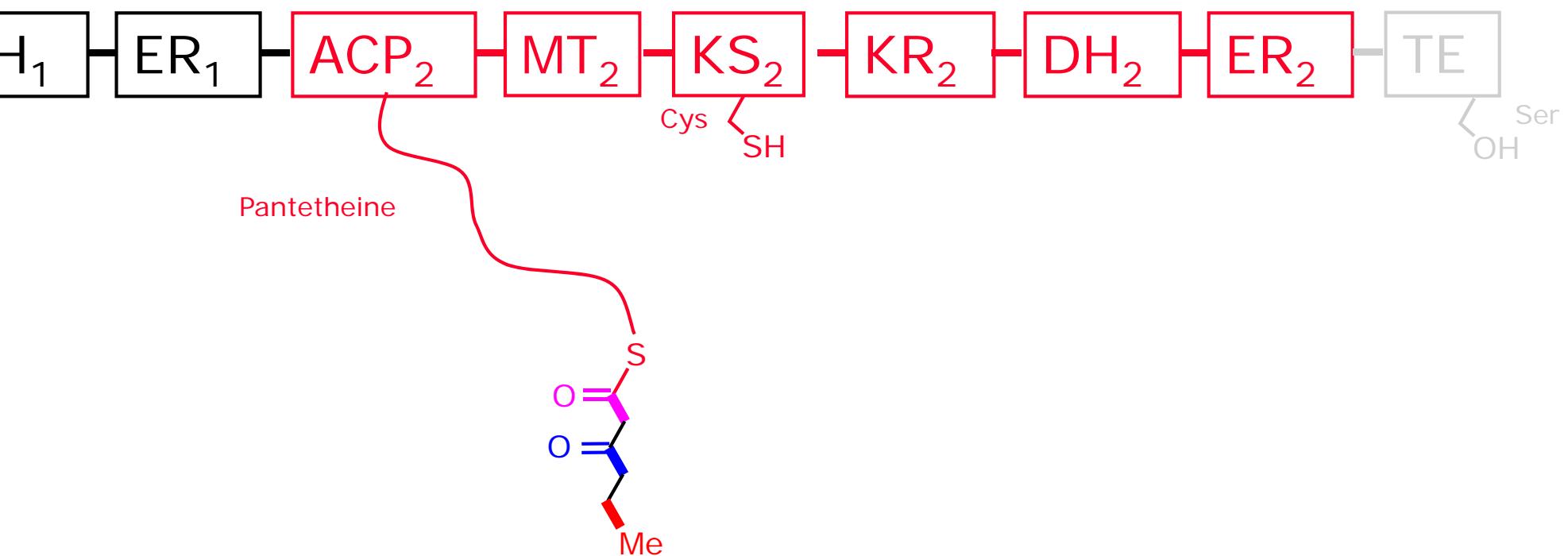


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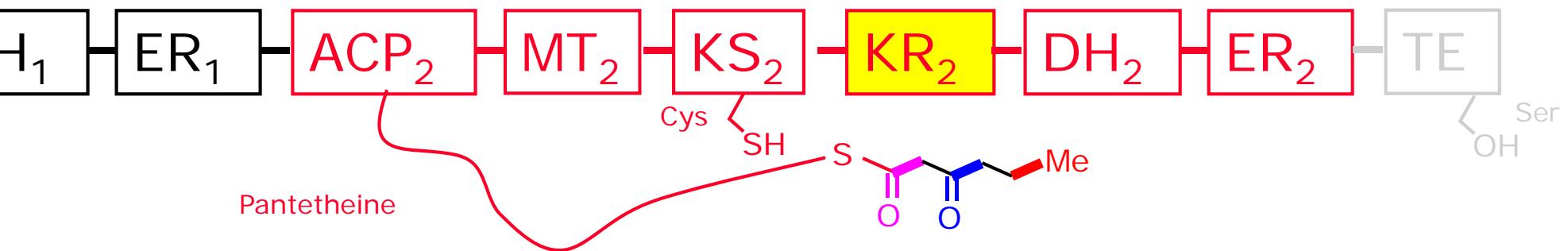


- KS₂ catalyzes Claisen condensation

FATTY ACID BIOSYNTHESIS

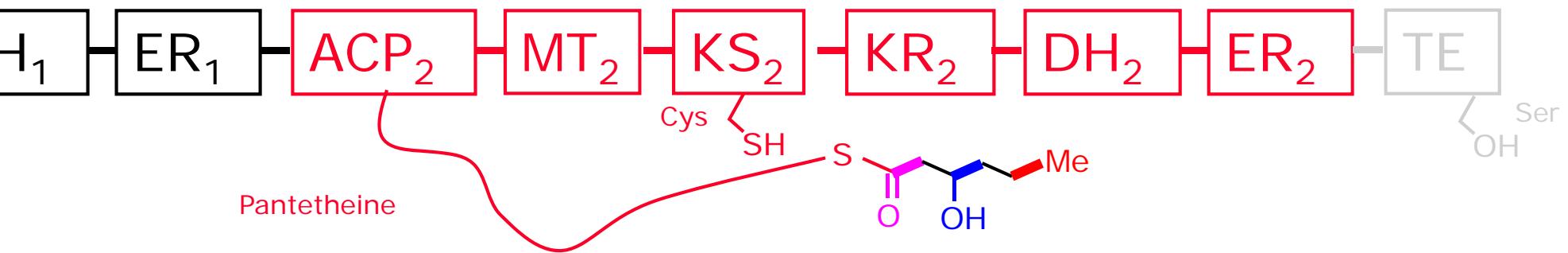


FATTY ACID BIOSYNTHESIS

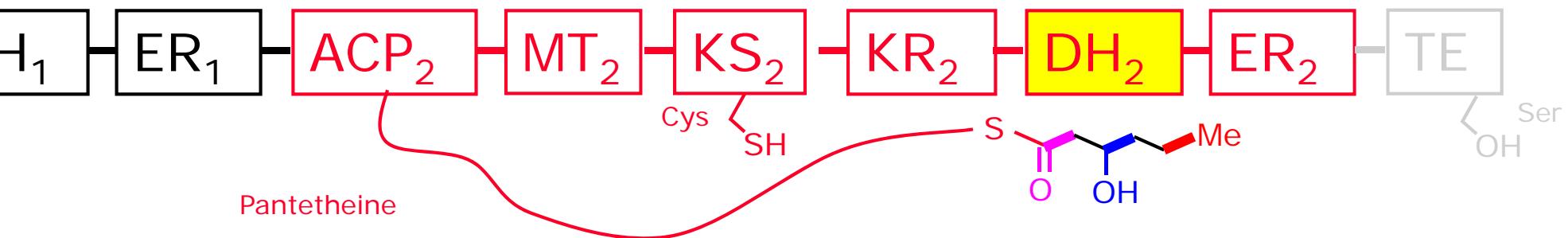


- KR₂ catalyzes reduction of ketone

FATTY ACID BIOSYNTHESIS

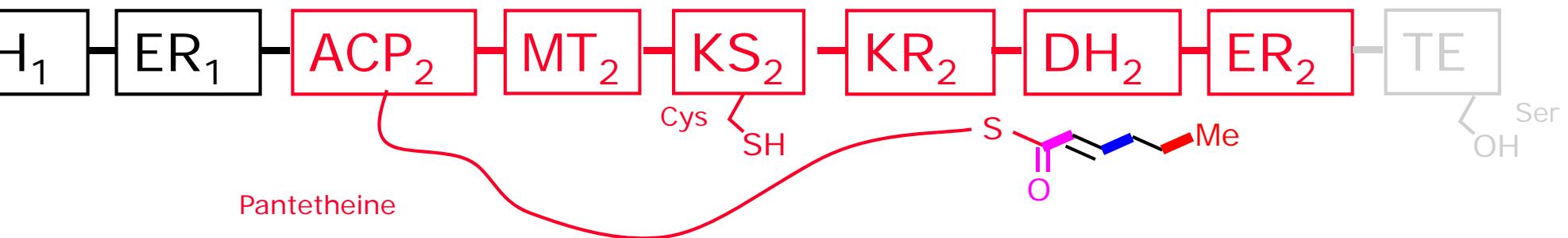


FATTY ACID BIOSYNTHESIS

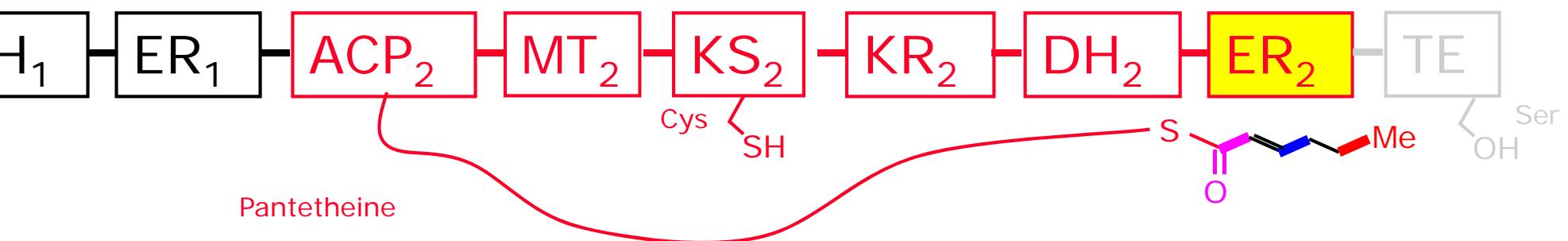


- DH₂ catalyzes dehydration of alcohol

FATTY ACID BIOSYNTHESIS

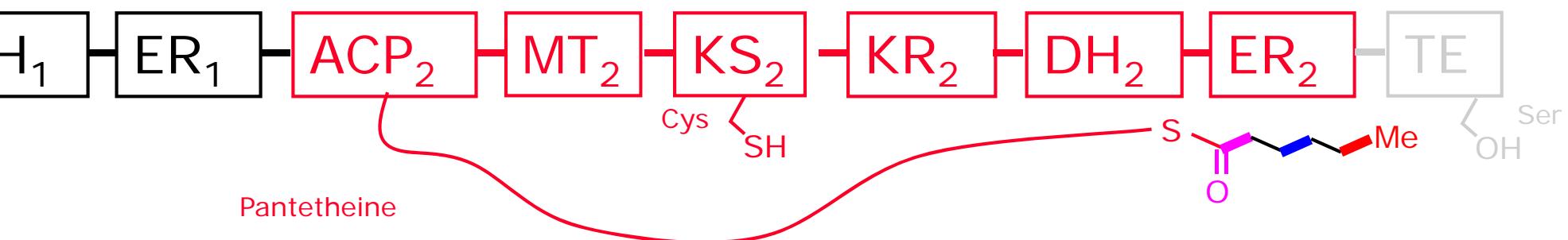


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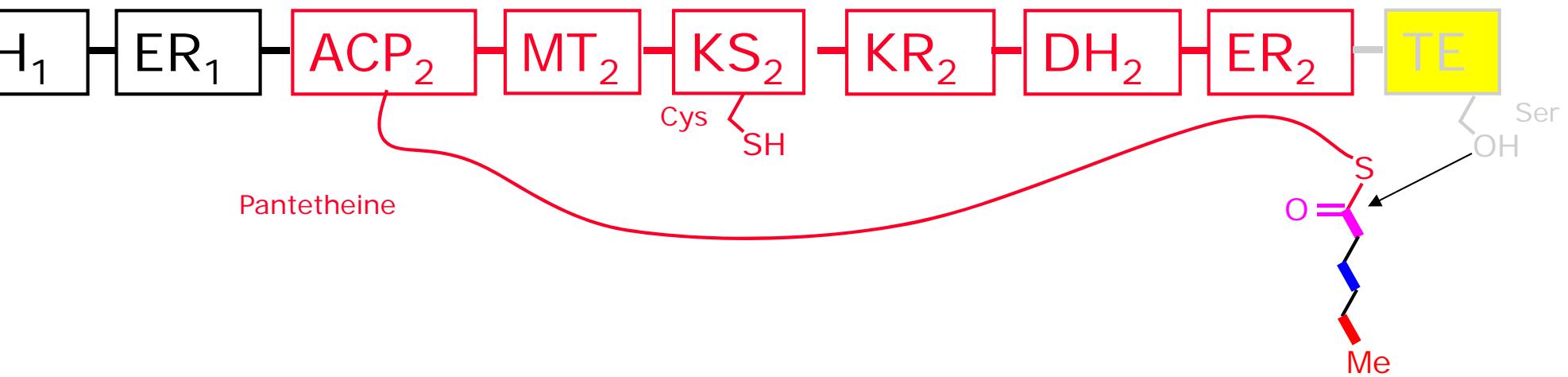


- ER₂ catalyzes reduction of alkene

FATTY ACID BIOSYNTHESIS

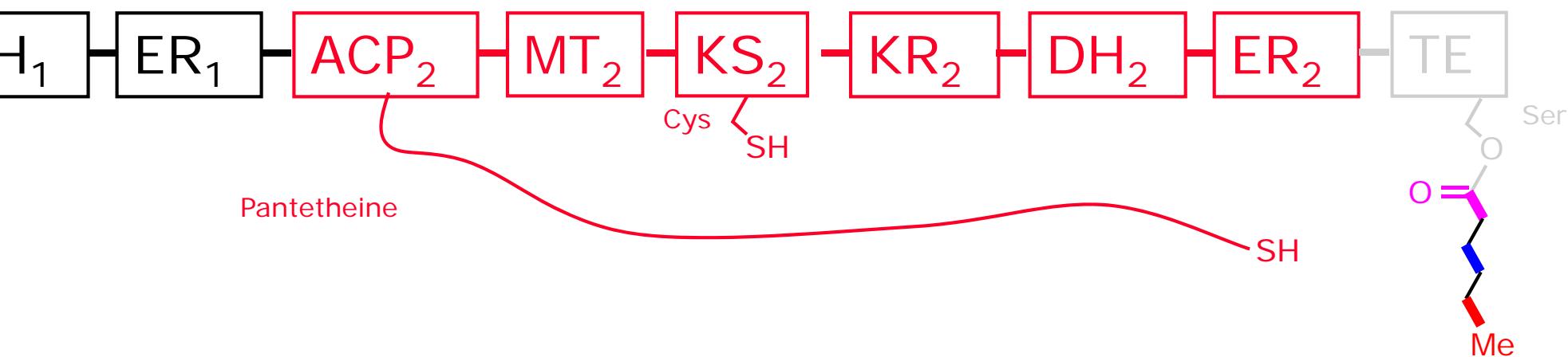


FATTY ACID BIOSYNTHESIS

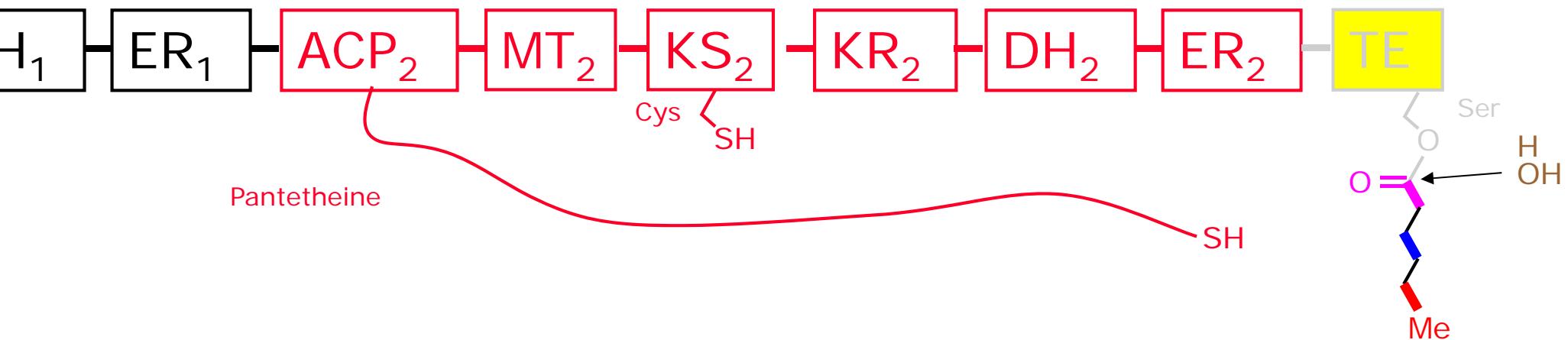


- TE catalyzes transesterification

FATTY ACID BIOSYNTHESIS

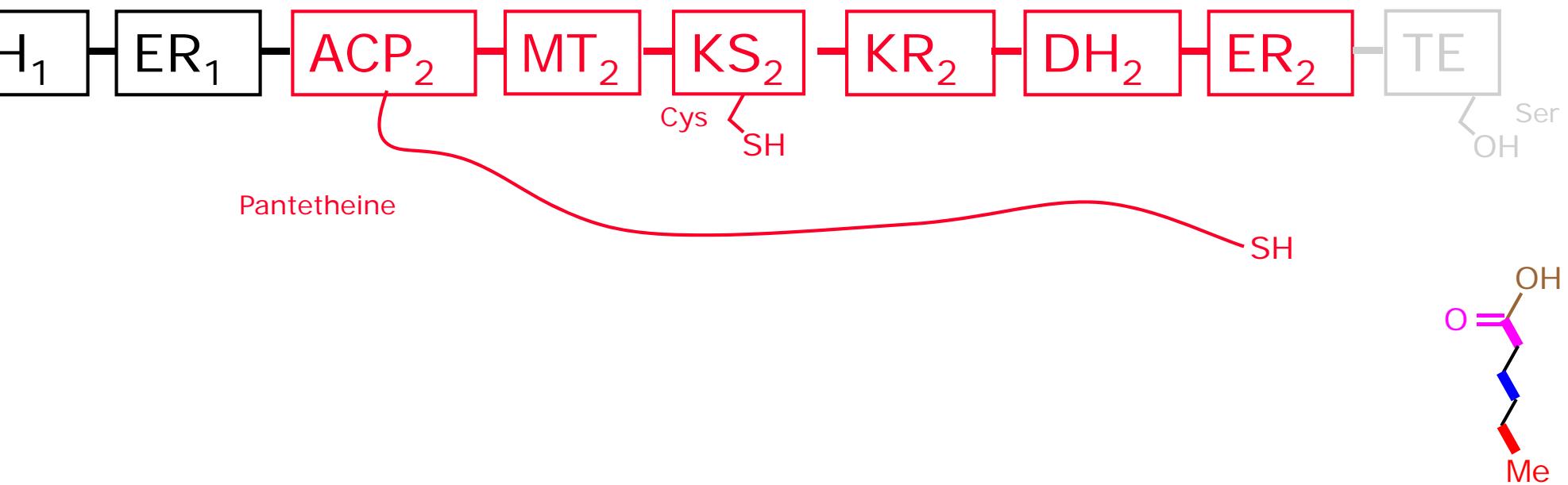


FATTY ACID BIOSYNTHESIS



- TE catalyzes hydrolysis

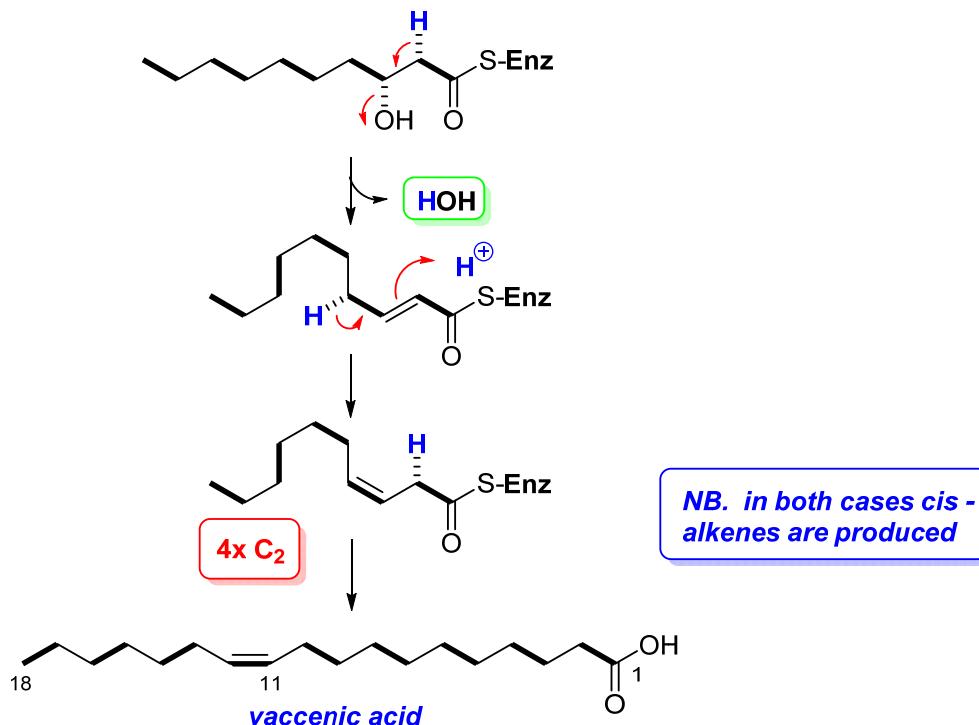
FATTY ACID BIOSYNTHESIS



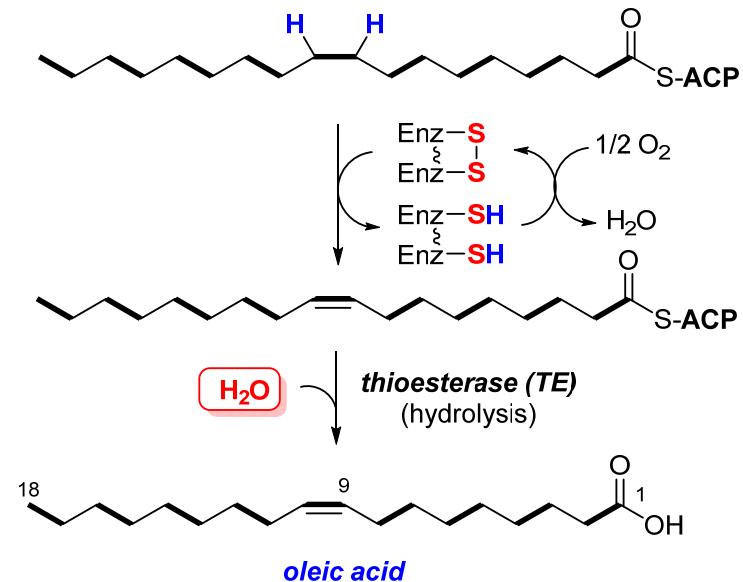
Biosynthesis of Unsaturated Fatty Acids

- **two mechanisms** are known for the introduction of double bonds into fatty acids:
 - in **BACTERIA**: **anaerobic [O]** → monounsaturated FAs (**MUFAs**)
 - in **MAMMALS, INSECTS & PLANTS**: **aerobic [O]** → **MUFAs** & polyunsaturated FAs (**PUFAs**)

ANAEROBIC ROUTE (bacteria)
 (dehydrogenation occurs during chain elongation)
 mainly MUFAs but some PUFAs



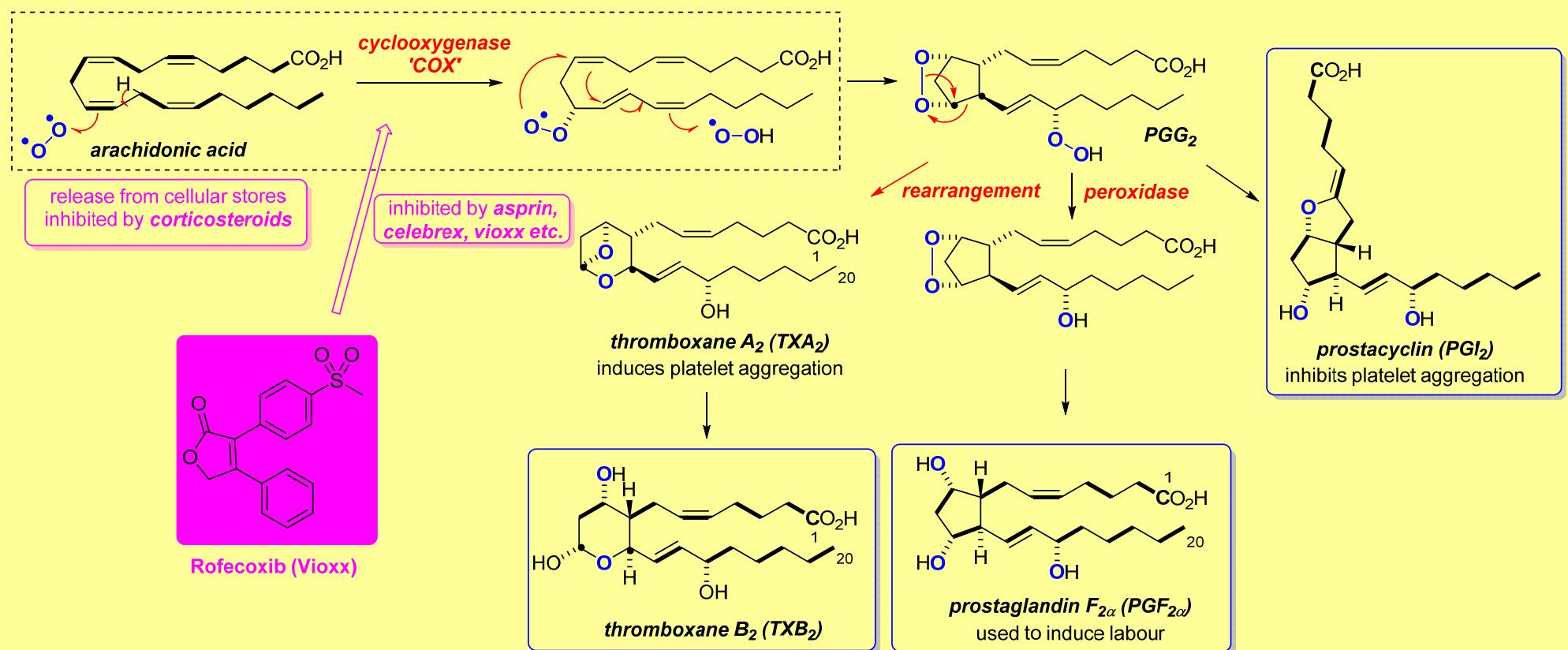
AEROBIC ROUTE (mammals, insects & plants)
 (dehydrogenation occurs after chain elongation)
 MUFAs & PUFAs



Position of alkenes in PUFAs	
animals	1st alkene C ₁₁ -C ₁₂
plants	1st alkene C ₅ -C ₆

Rational Anti-inflammatory Development – Prostaglandin & Thromboxane Pathway Intervention

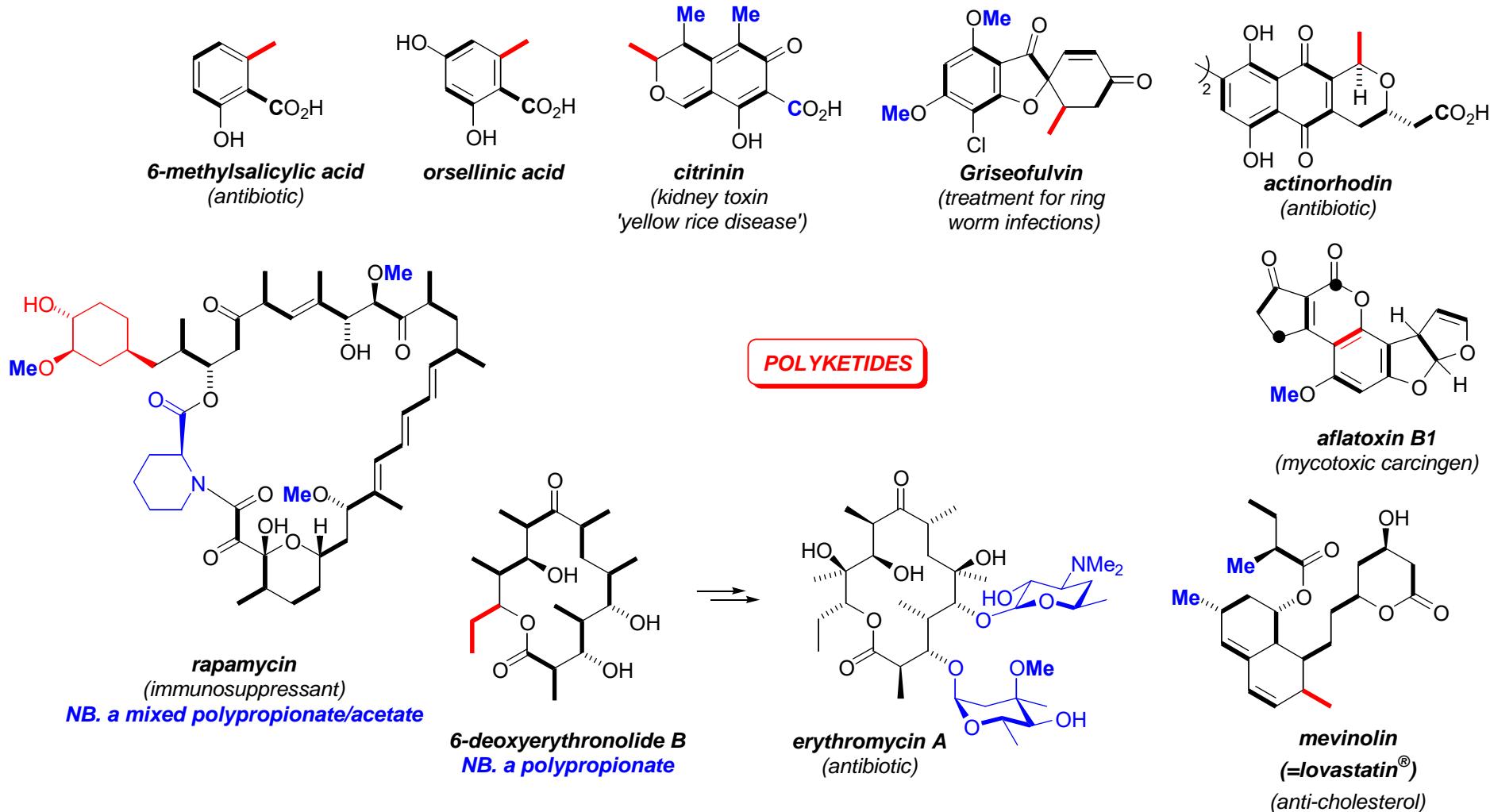
- **prostaglandins & thromboxanes** are derived from further oxidative processing of arachidonic acid
- both are important **hormones** which control e.g. smooth **muscle contractility** (blood pressure), **gastric secretion**, **platelet aggregation** & **inflammation** (<nM activity)
 - various pharmaceuticals including **corticosteroids** & **aspirin** inhibit biosynthetic steps in these pathways



Polyketides

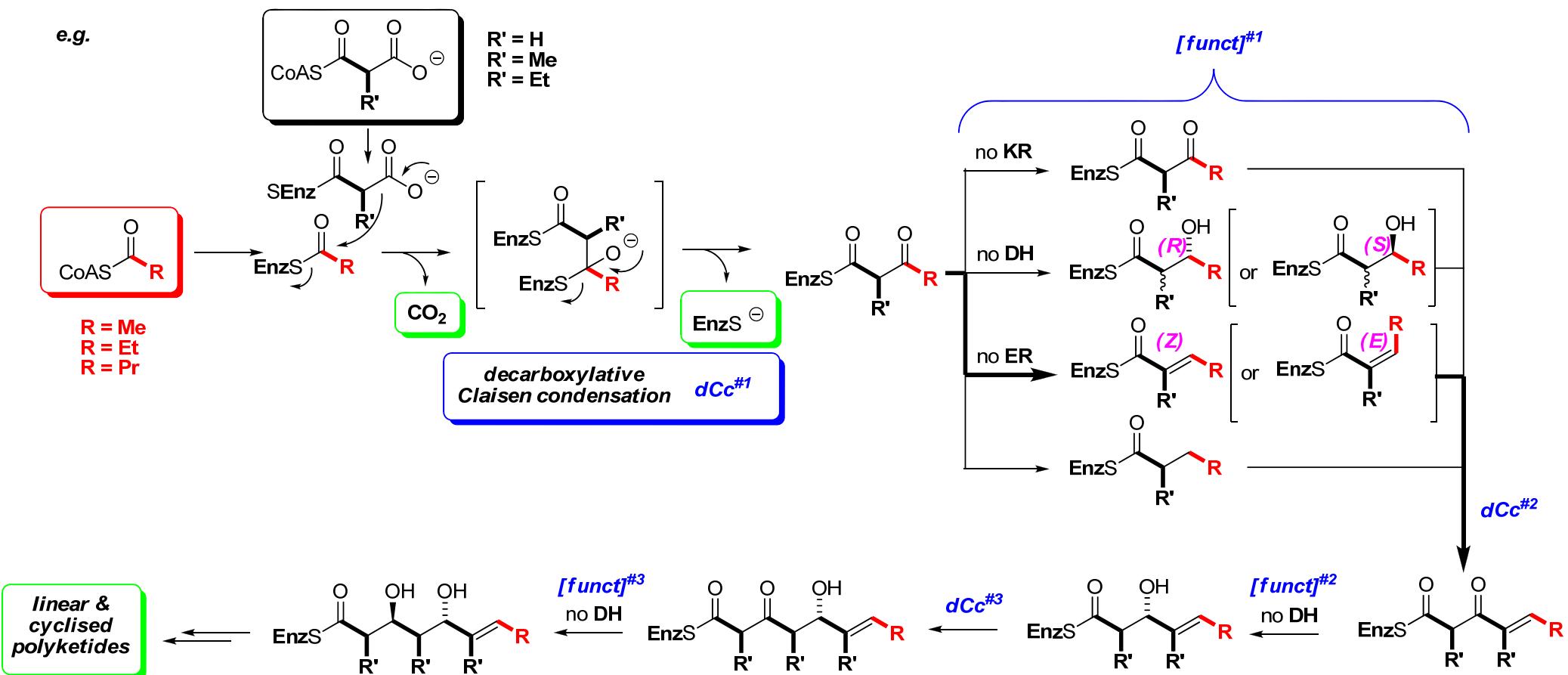
Polyketides

- the structural variety of ***polyketide secondary metabolites*** is very wide:
 - NB. starter units marked in red; extender units in bold black; post oligomerisation appended groups in blue



Biosynthesis of Polyketides – Oligomerisation Steps

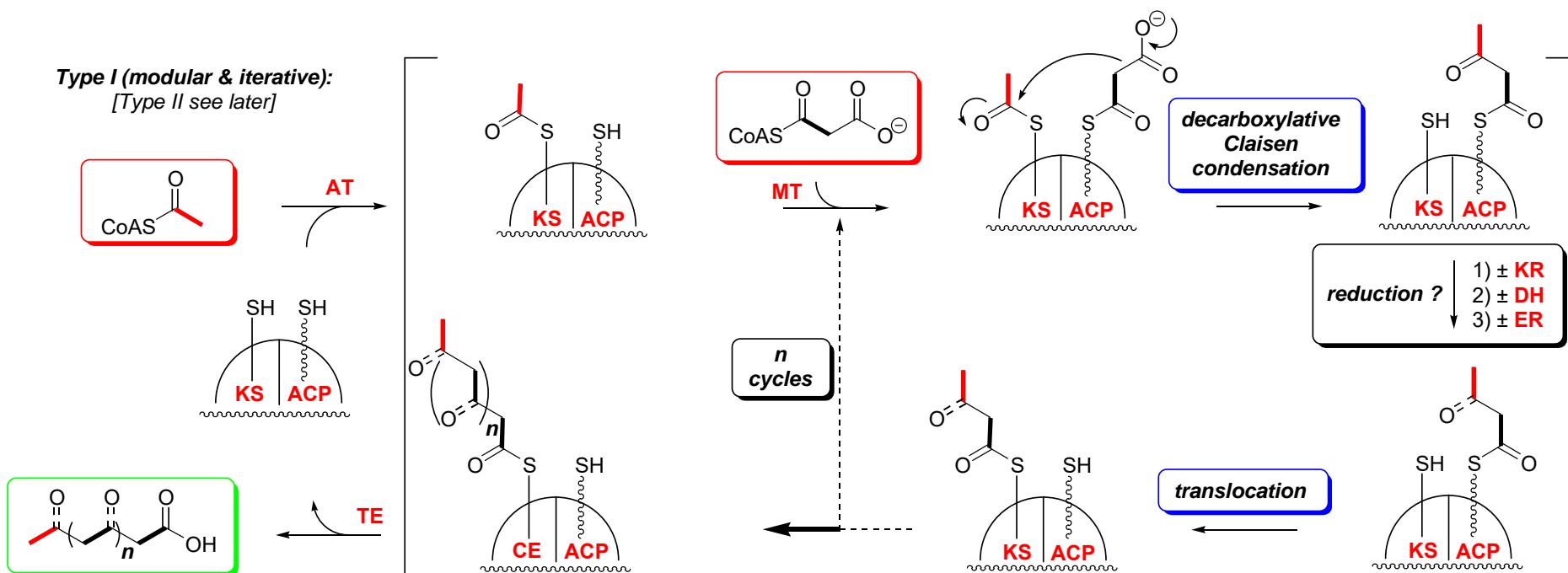
- **polyketides** are biosynthesised by a process very similar to that for **fatty acids**
 - the key **differences** are:
 - greater variety of **starter units**, **extender units** & **termination processes**
 - absent or incomplete reduction of the iteratively introduced β -carbonyl groups: ie. each cycle may differ in terms of **KR**, **DH** & **ER** modules & stereochemistry



- this leads to **enormous diversity...**

Biosynthesis of Polyketides – Overview of PKS

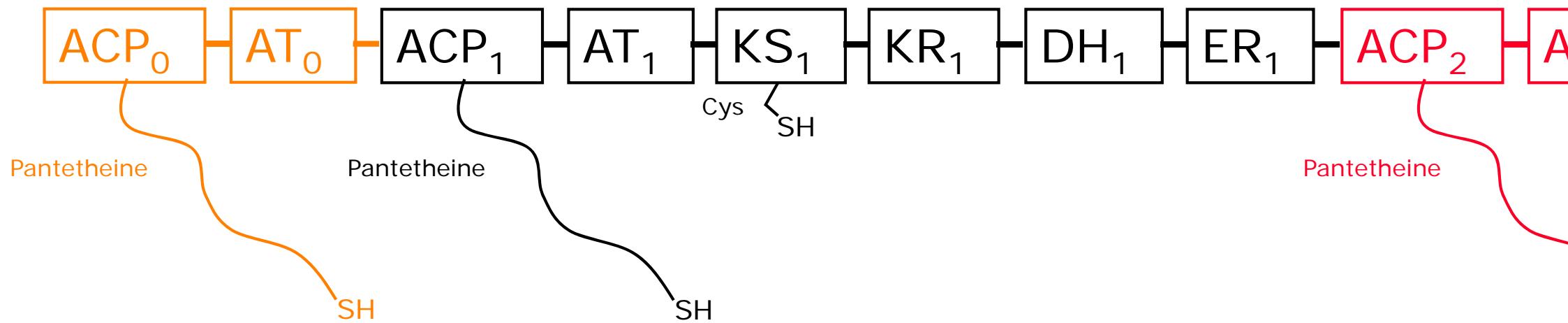
- the *in vivo* process of polyketide synthesis involves **PolyKetide Synthases (PKSs)**:
 - PKSs** (except Type II, see later) comprise the same **8 components** as FASs. i.e. (ACP & 7× catalytic activities): **ACP, KS, AT, MT, [KR, DH, ER & TE]**
 - Type I PKSs: single (or small set of) multifunctional protein complex(es)**
 - modular (microbial)** - each ‘step’ has a dedicated catalytic site (\rightarrow **macrolides**)
 - iterative (fungal)** – single set of catalytic sites, each of which *may* operate in each iteration (cf. FASs) (\rightarrow **aromatics/polyphenols** - generally)
 - Type II PKSs: single set of discrete, dissociable single-function proteins**
 - iterative (microbial)** - each catalytic module *may* operate in each iteration (cf. FASs) (\rightarrow **aromatics/polyphenols**)



KS = keto synthase; **AT** = acetyl transferase; **MT** = malonyl transferase;

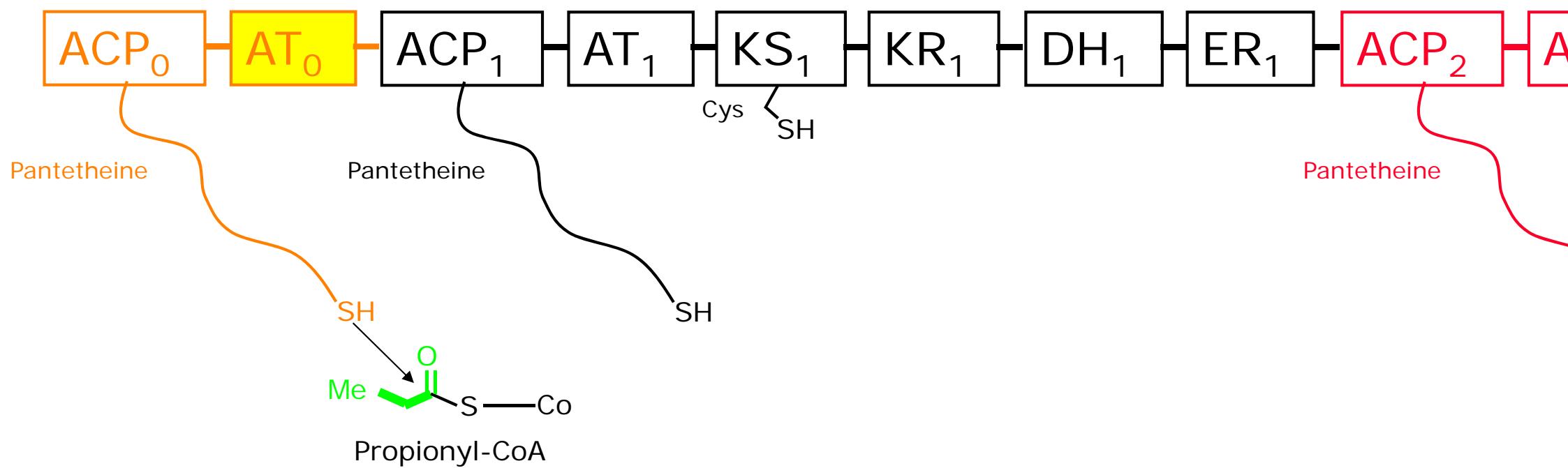
KR = keto reductase; **DH** = dehydratase; **ER** = enoyl reductase; **TE** = thioesterase; **ACP** = acyl carrier protein

POLYKETIDE BIOSYNTHESIS [Type I – (modular)]



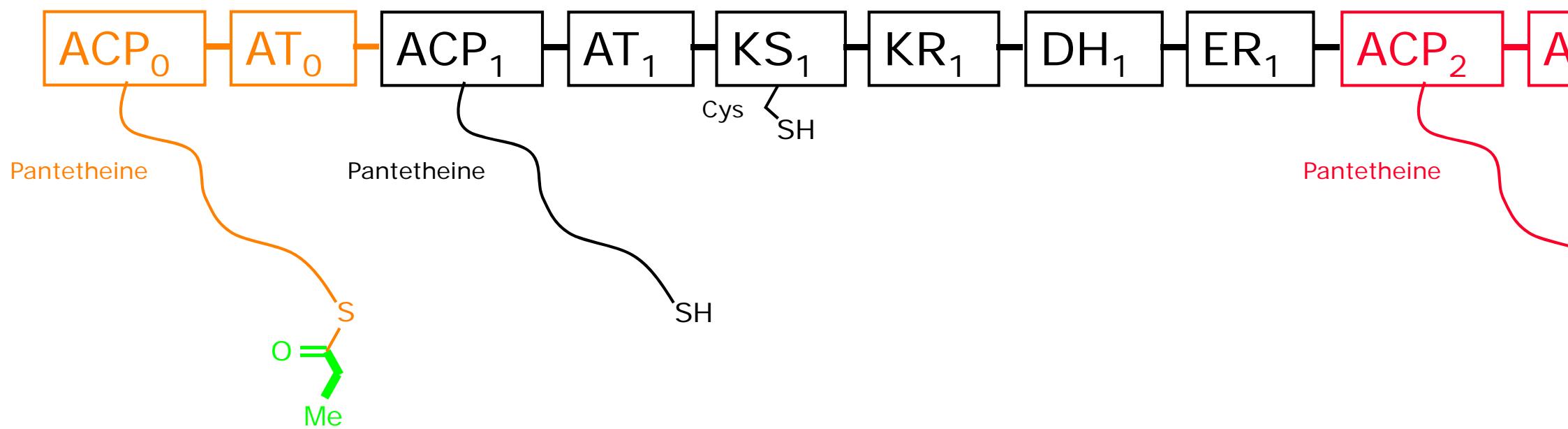
NB. the following sequence of slides has also been adapted from: <http://www.courses.fas.harvard.edu/%7echem27/>

POLYKETIDE BIOSYNTHESIS

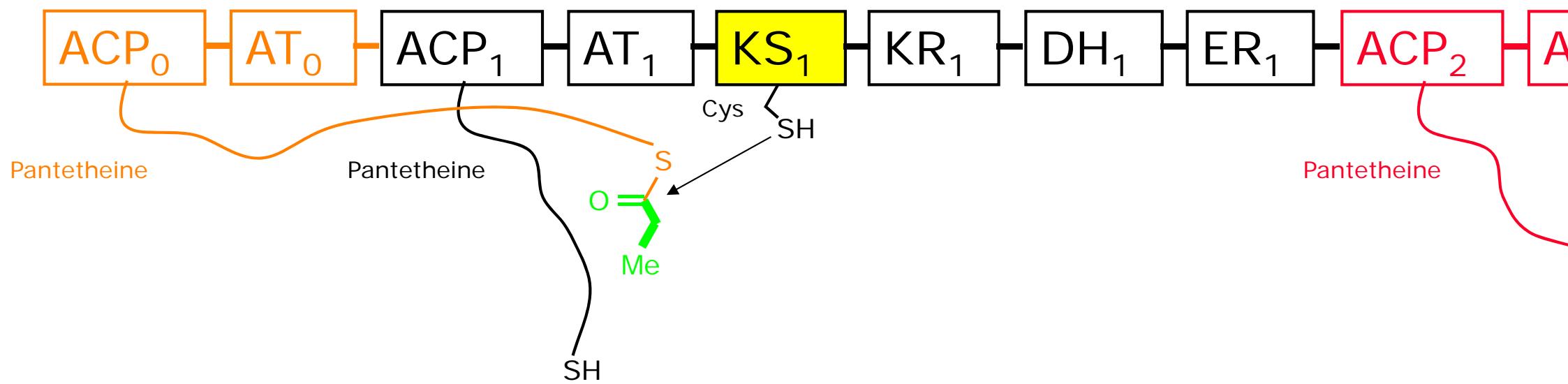


- AT_0 loads starting group (propionyl) onto ACP_0

POLYKETIDE BIOSYNTHESIS

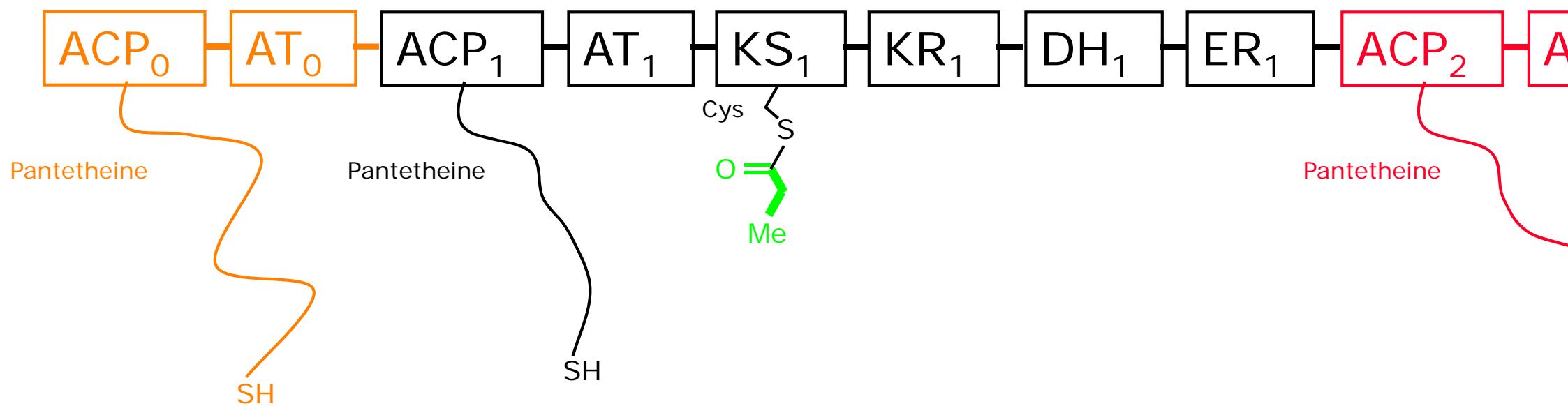


POLYKETIDE BIOSYNTHESIS

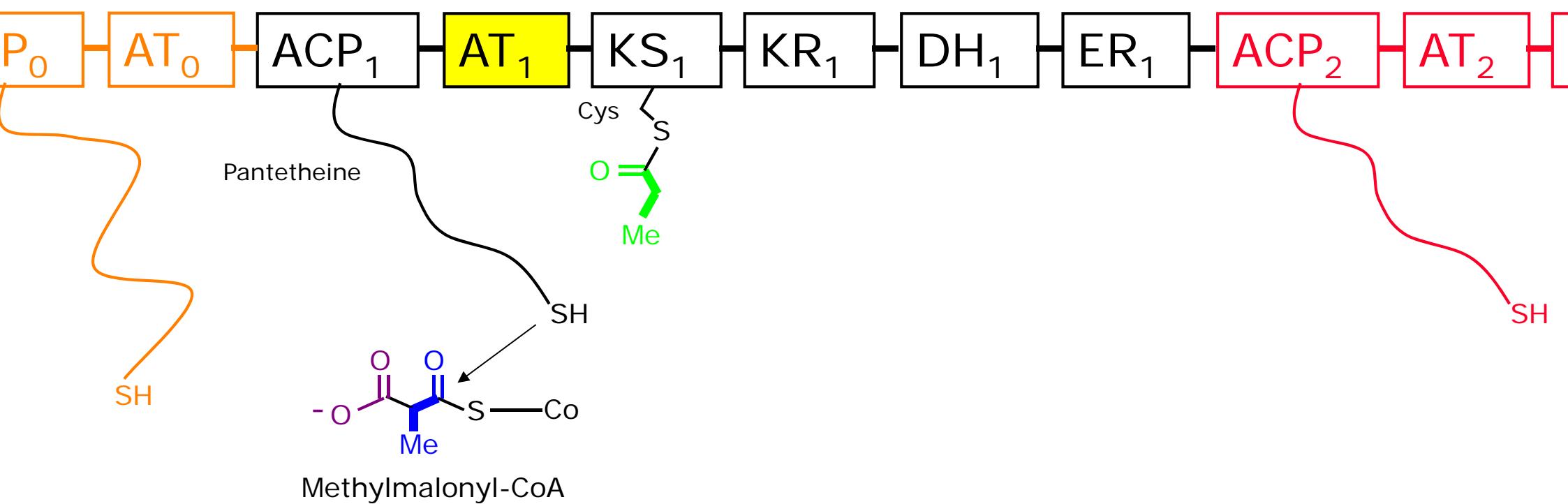


- KS₁ catalyzes translocation to module 1

POLYKETIDE BIOSYNTHESIS

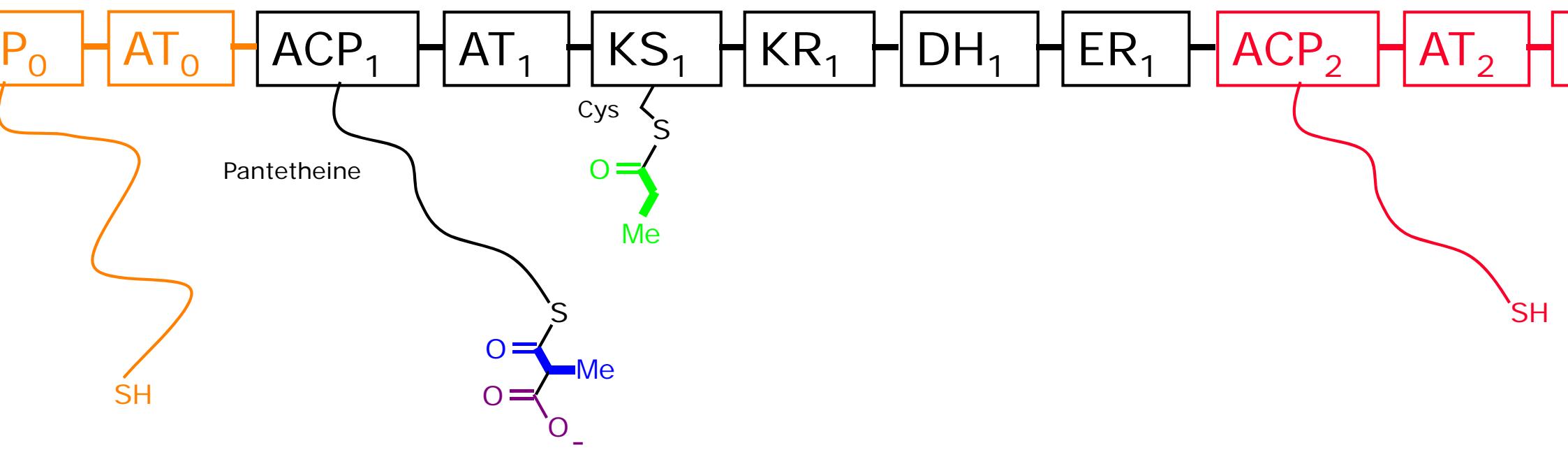


POLYKETIDE BIOSYNTHESIS

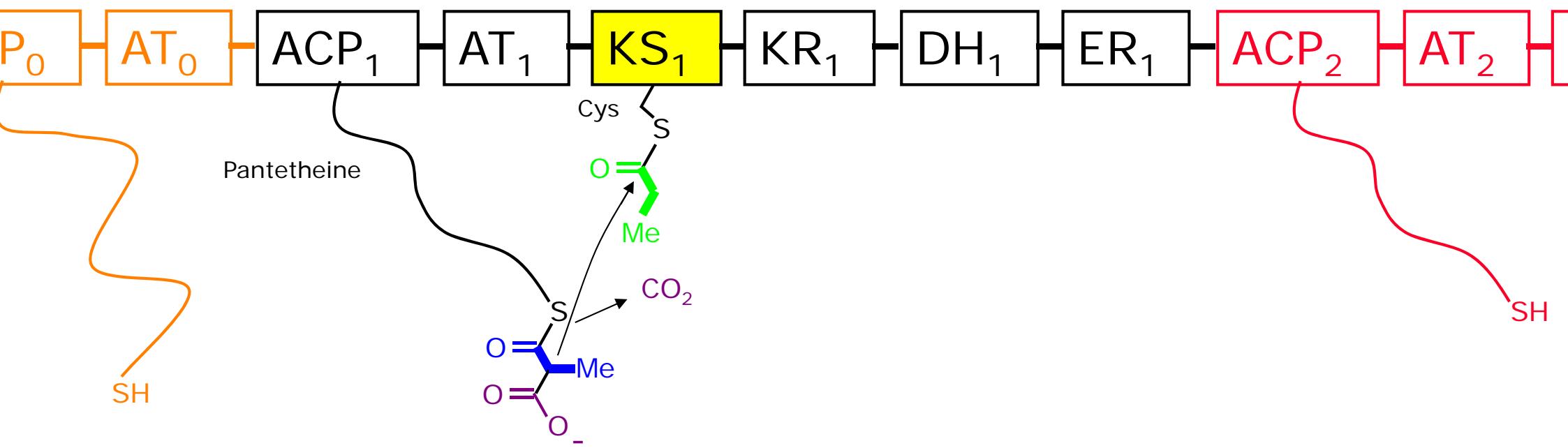


- AT_1 loads methylmalonyl group onto ACP_1

POLYKETIDE BIOSYNTHESIS

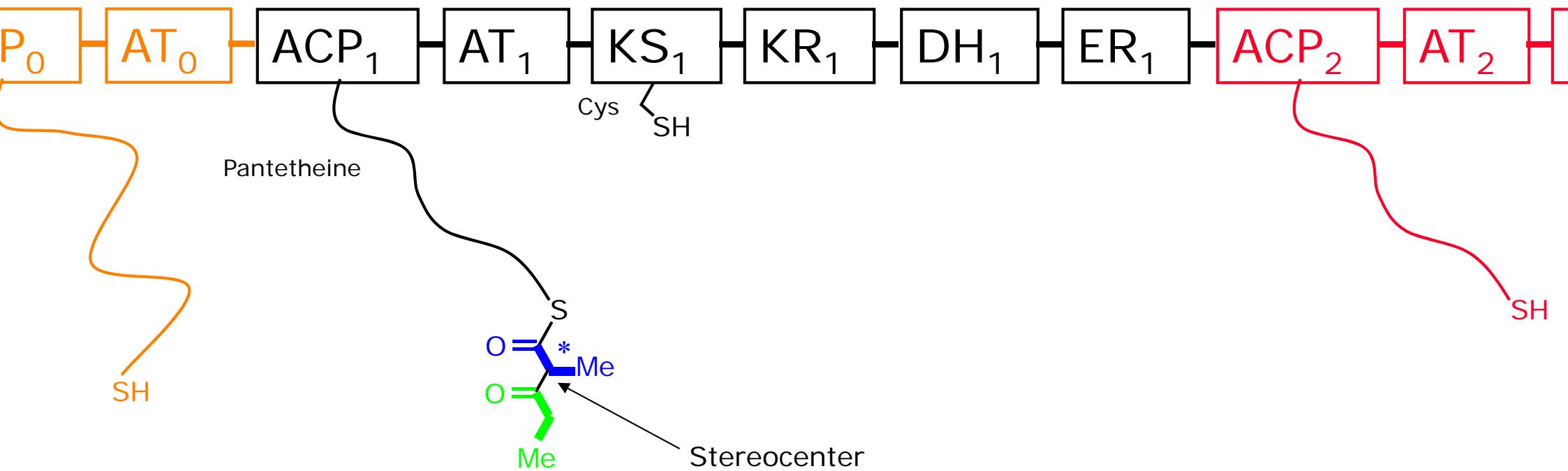


POLYKETIDE BIOSYNTHESIS

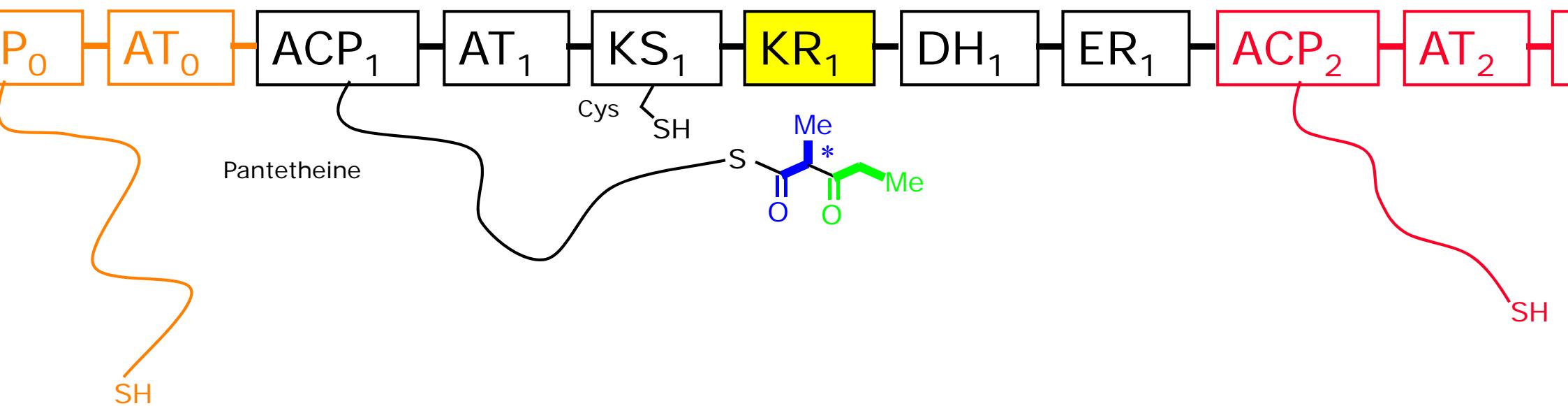


- KS_1 catalyzes Claisen condensation

POLYKETIDE BIOSYNTHESIS

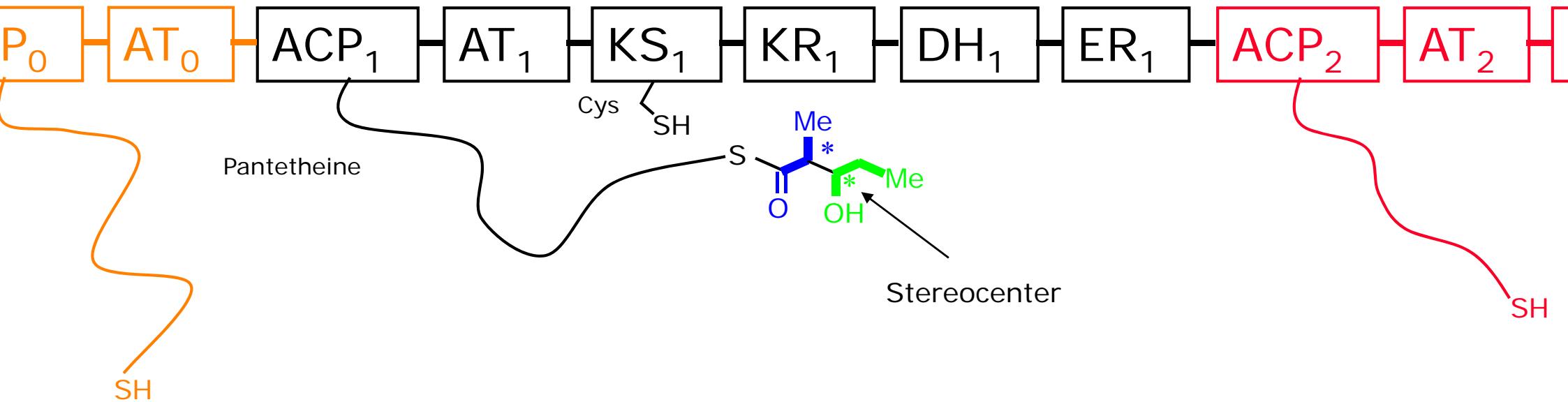


POLYKETIDE BIOSYNTHESIS

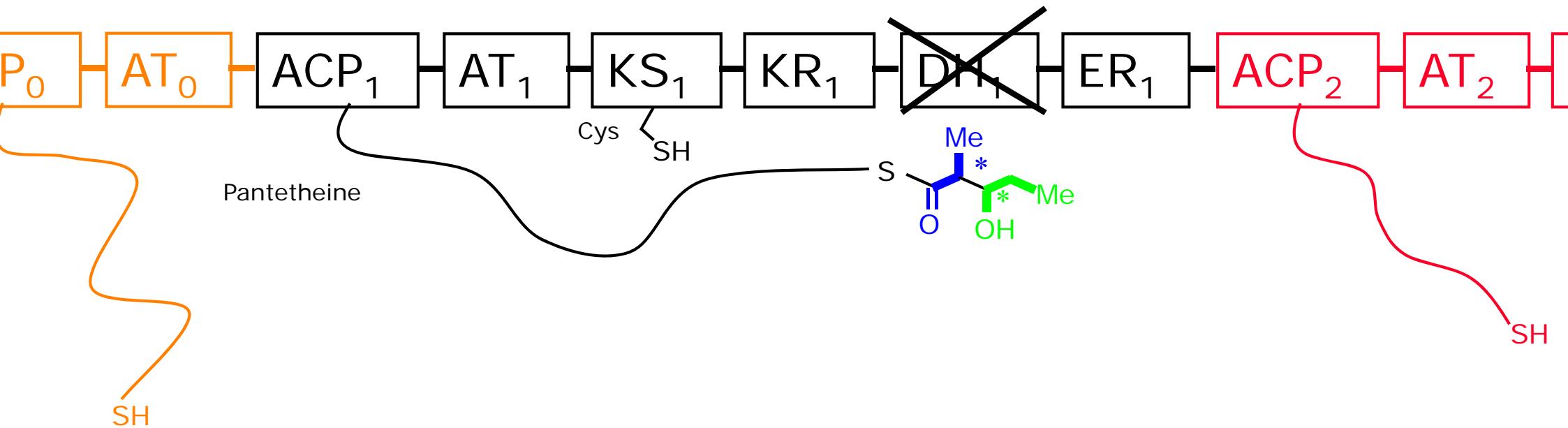


- KR_1 catalyzes reduction of ketone

POLYKETIDE BIOSYNTHESIS

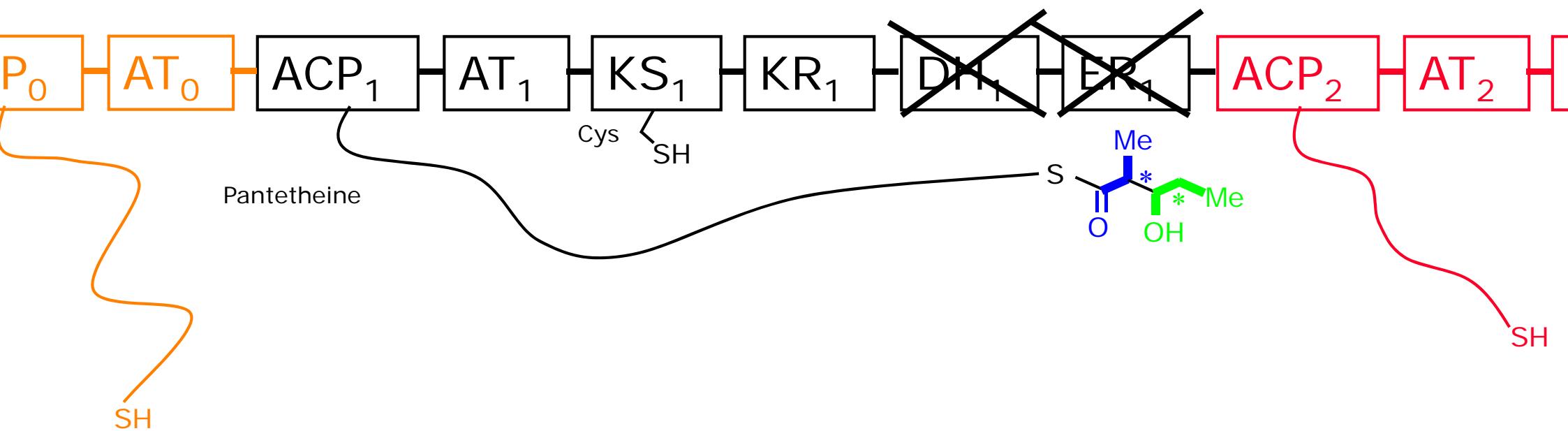


POLYKETIDE BIOSYNTHESIS



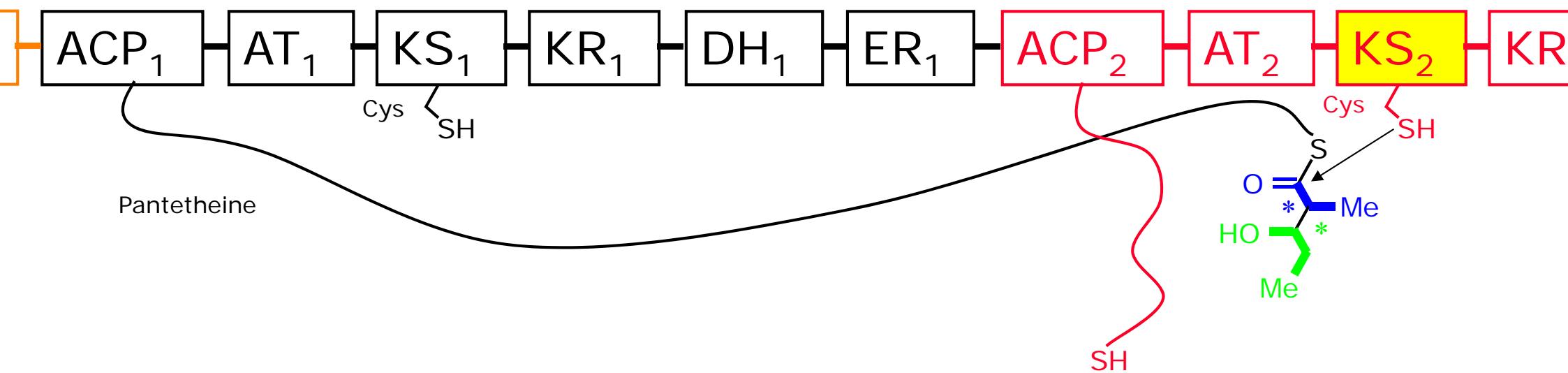
- no DH_1 activity

POLYKETIDE BIOSYNTHESIS



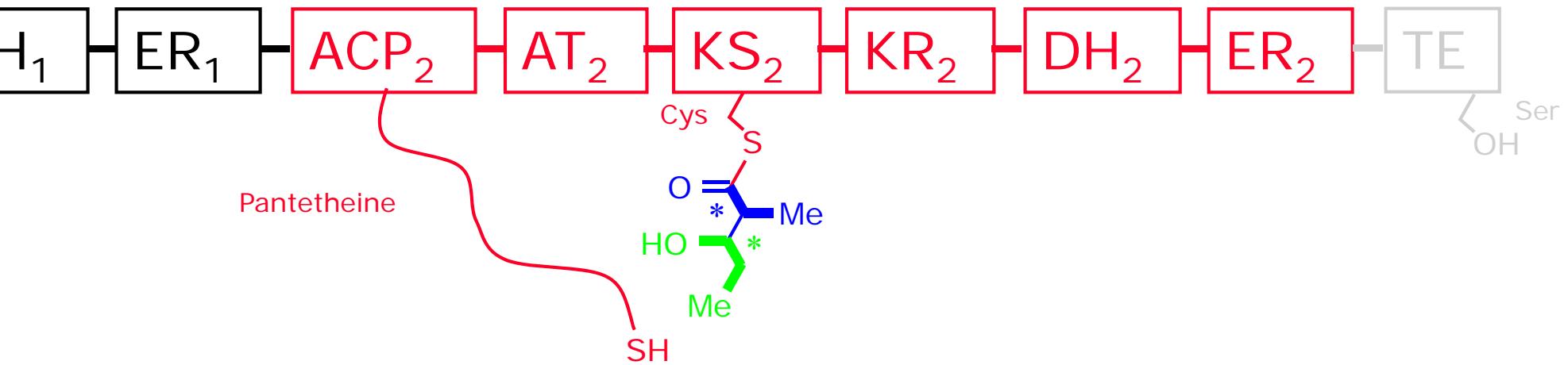
- no ER_1 activity

POLYKETIDE BIOSYNTHESIS



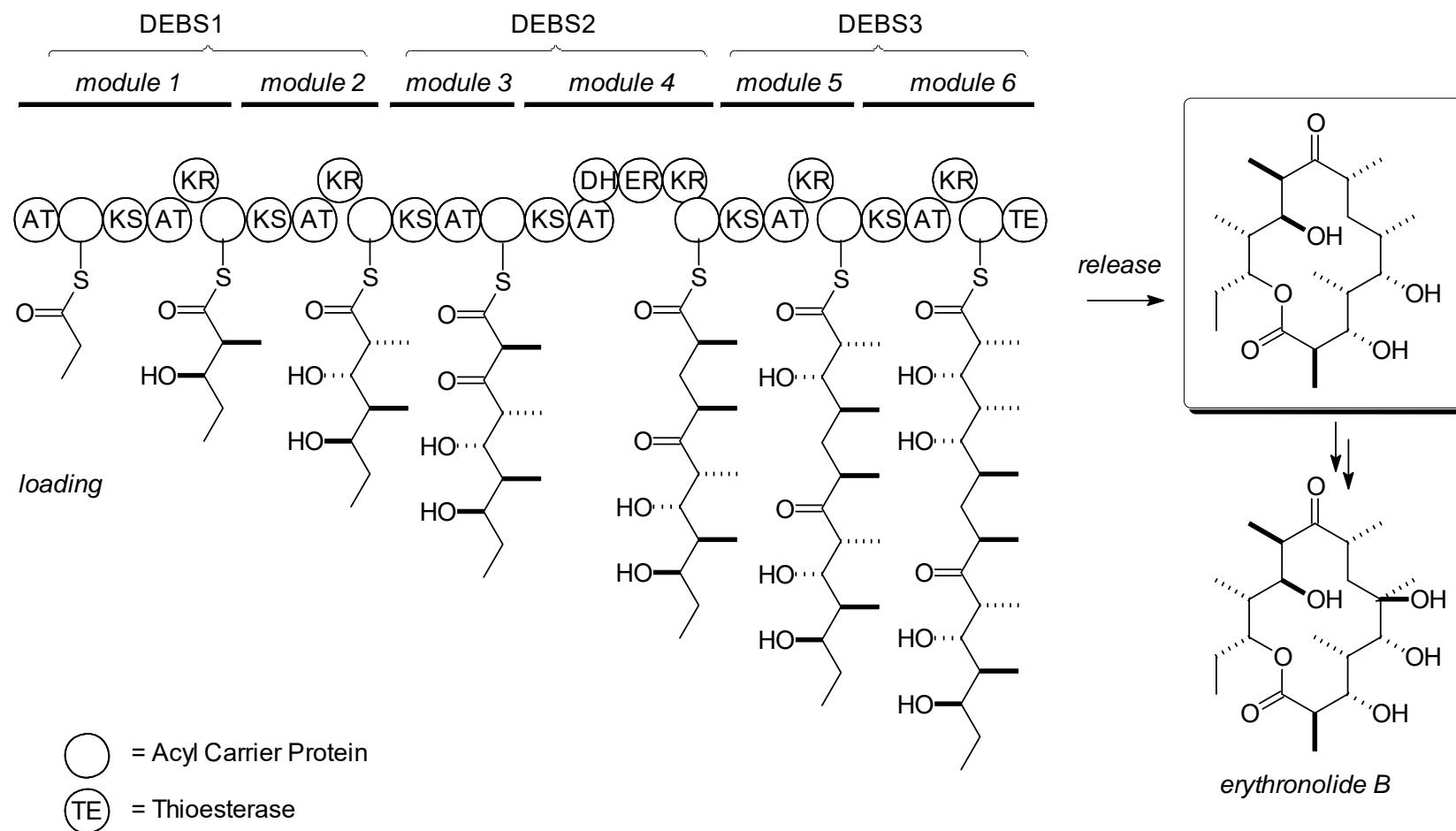
- KS₂ catalyzes translocation to module 2

POLYKETIDE BIOSYNTHESIS



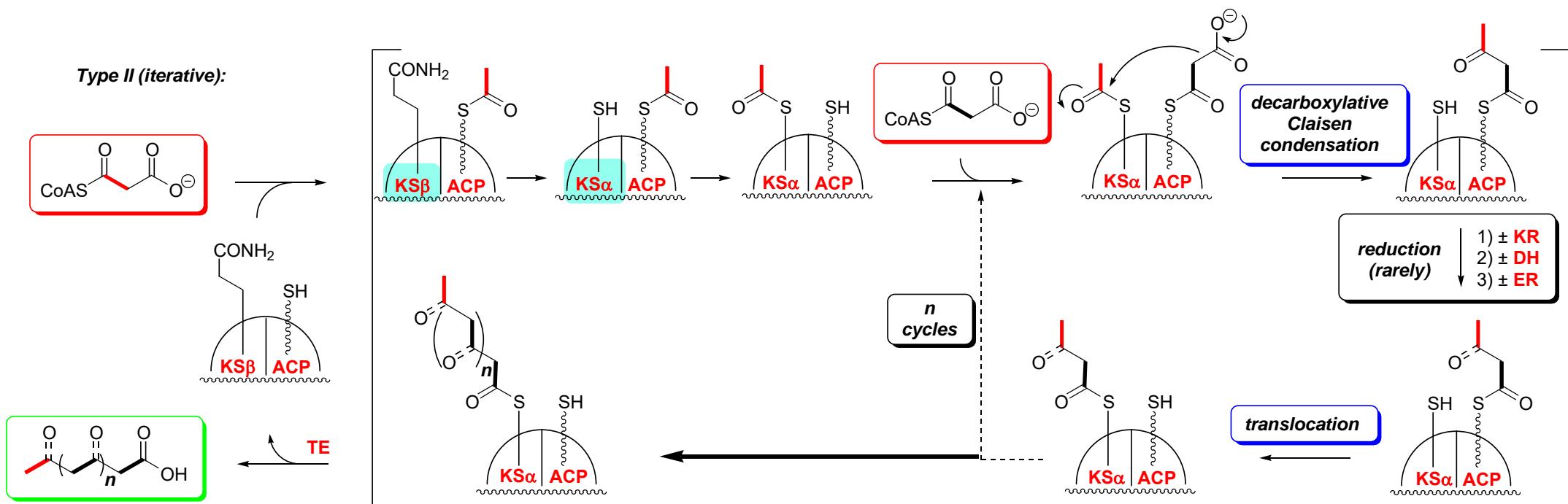
Biosynthesis of Erythromycin – Type I(modular) PKS

- **6-deoxyerthonolide** is a precursor to **erythromycin A** – **bacterial** antibiotic (*Streptomyces erythreus*):
 - propionate based **heptaketide**; 3 multifunctional polypeptides (DEBS1, DEBS2 & DEBS3, all ~350 kDa)
 - Katz et al. *Science* **1991**, 252, 675 ([DOI](#)); Staunton, Leadley et al. *Science* **1995**, 268, 1487 ([DOI](#)); Khosla et al. *J. Am. Chem. Soc.* **1995**, 9105 ([DOI](#)); **review:** Staunton & Weissman *Nat. Prod. Rep.* **2001**, 18, 380 ([DOI](#))



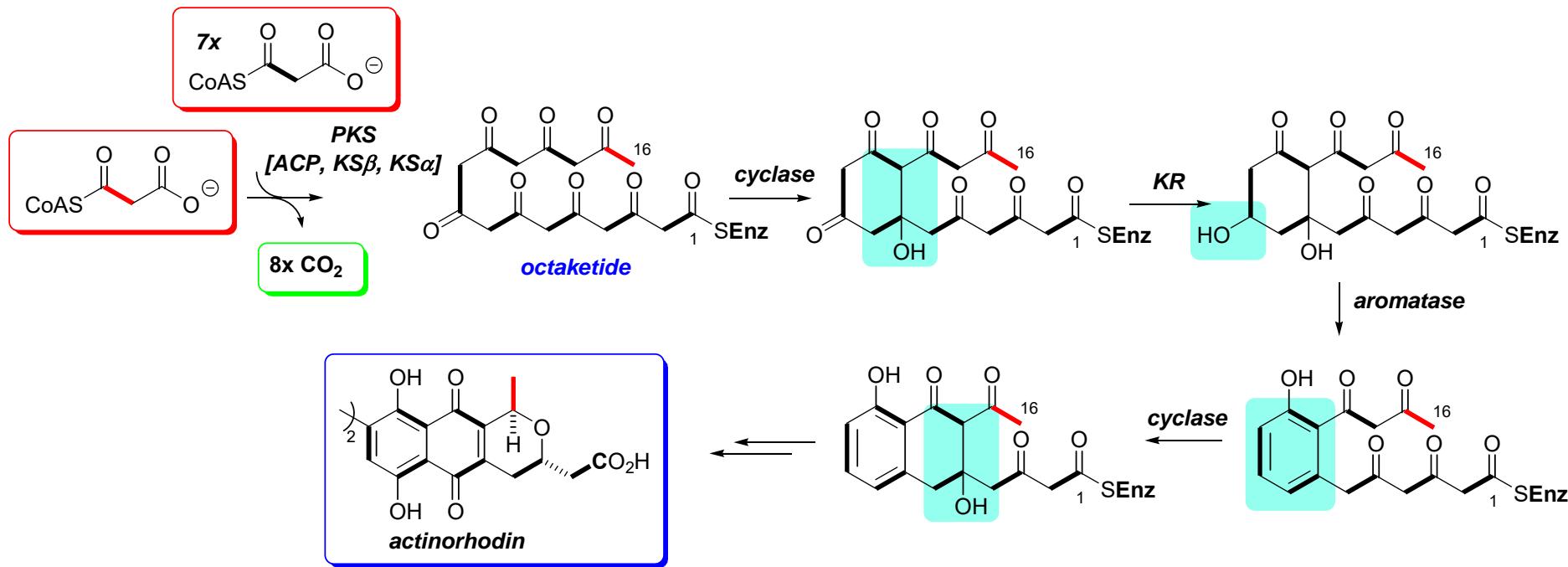
Type II PKSs – Enzyme Clusters (Microbial)

- Type II PKSs:** single set of discrete, dissociable single-function proteins (ACP & 6× catalytic functions): **ACP**, **KS_β**, **KS_α**, **[KR, DH, ER, & TE]** [NB. NO acetyl or malonyl transferases (AT, MT)]
 - **iterative** - each catalytic module *may* operate in each iteration (cf. FASs) (→ **aromatics/polyphenols**)
- these clusters (generally) use **malonate** as BOTH **starter** & **extender** unit
- their **ACP proteins** are able to load malonate direct from malonyl CoA (no MT required)
 - the **starter malonate** is **decarboxylated** by ‘**ketosynthase**’ β (**KS_β**) to give **S-acetyl-ACP**
 - the **extender malonates** undergo **decarboxylative Claisen condensations** by **ketosynthase α** (**KS_α**)
- these clusters rarely utilise **KR**, **DH** or **ER** activities and produce ‘true’ polyketides:



Biosynthesis of Actinorhodin – Type II PKS

- **actinorhodin** – octaketide **bacterial antibiotic** (*Streptomyces coelicolor*)
 - Hopwood *Chem. Rev.* **1997**, 97, 2465 ([DOI](#))

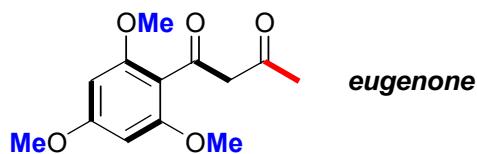


- **timing** of **1st cyclisation** and mechanism of **control of chain length** uncertain
 - **octaketide** synthesis then cyclisation? (as shown above)
 - **hexaketide** synthesis then cyclisation then two further rounds of extension?
- indications can sometimes be gleaned from **biomimetic syntheses**.

Scope of Structures - Type II PKS

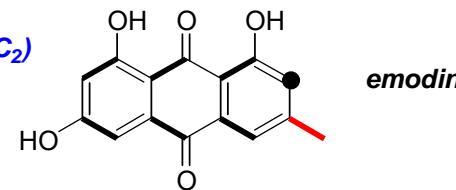
- ***microbial polyphenolic*** metabolites:

pentaketides (5x C₂)



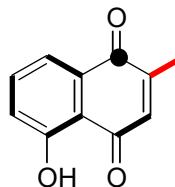
eugenone

octaketides (8x C₂)



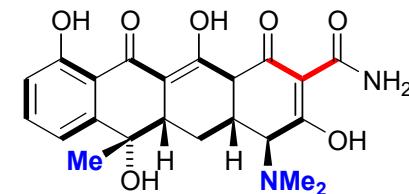
emodin

hexaketides (6x C₂)



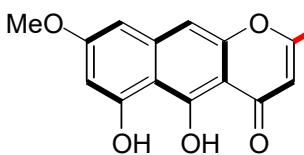
plumbagin

nonaketides (9x C₂)



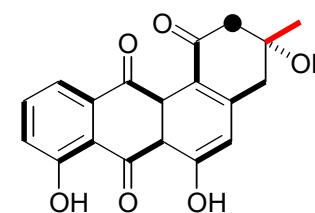
tetracycline

heptaketides (7x C₂)



rubrofusarin

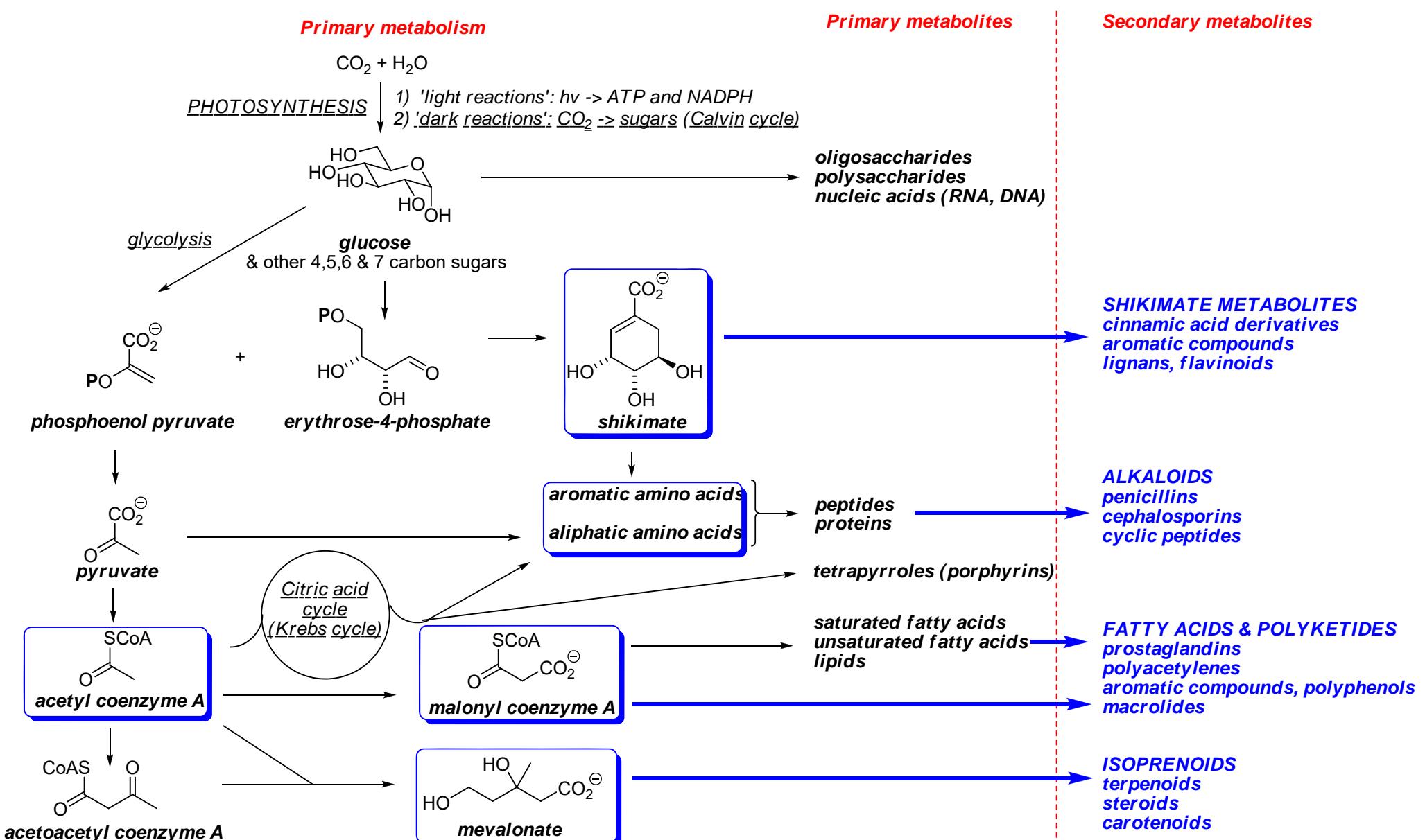
decaketides (10x C₂)



rabelomycin

- many display interesting biological activities...

Primary Metabolism - Overview

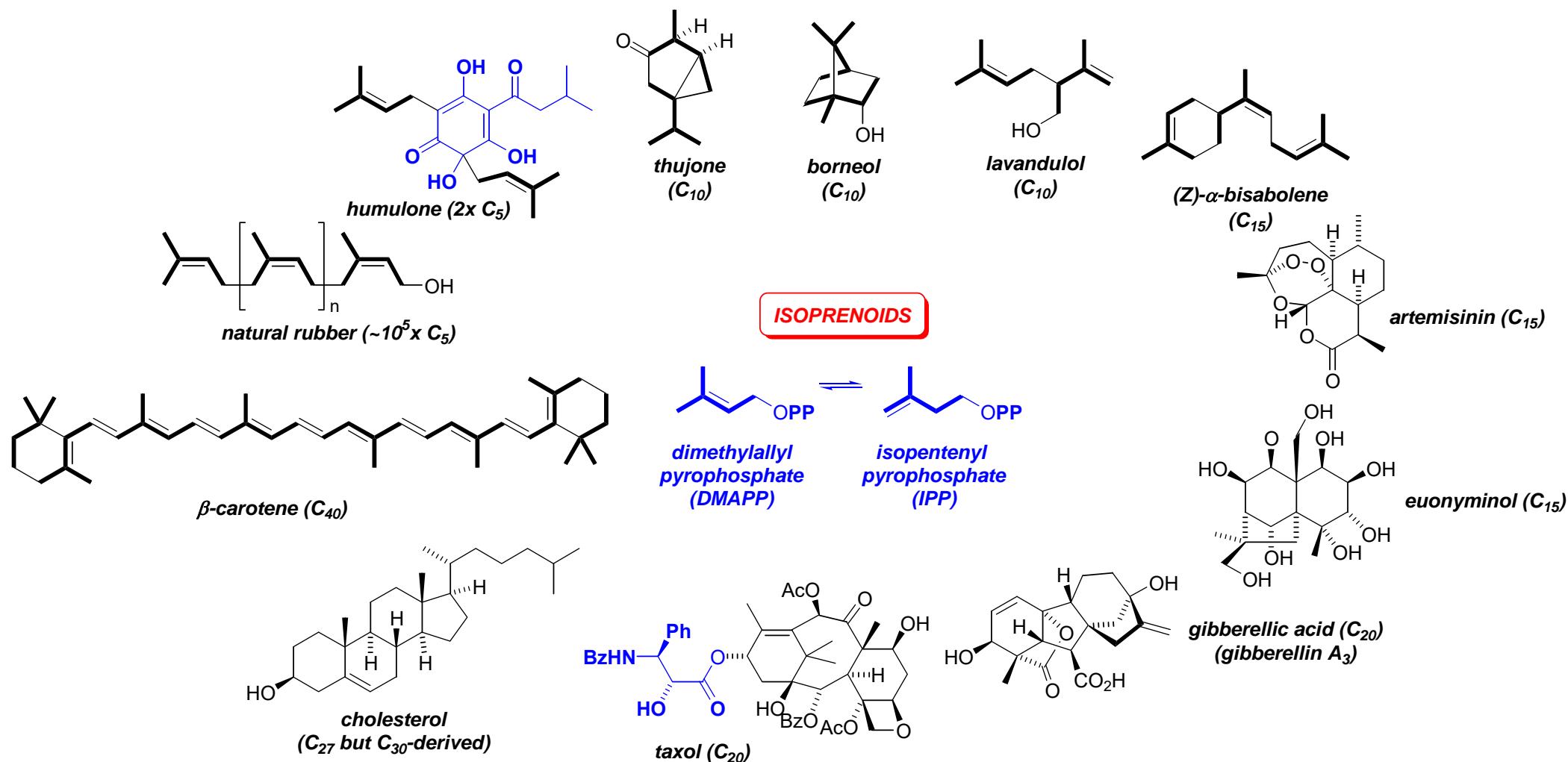


For interesting animations' of e.g. photosynthesis see: <http://www.johnkyrk.com/index.html>

Isoprenoids

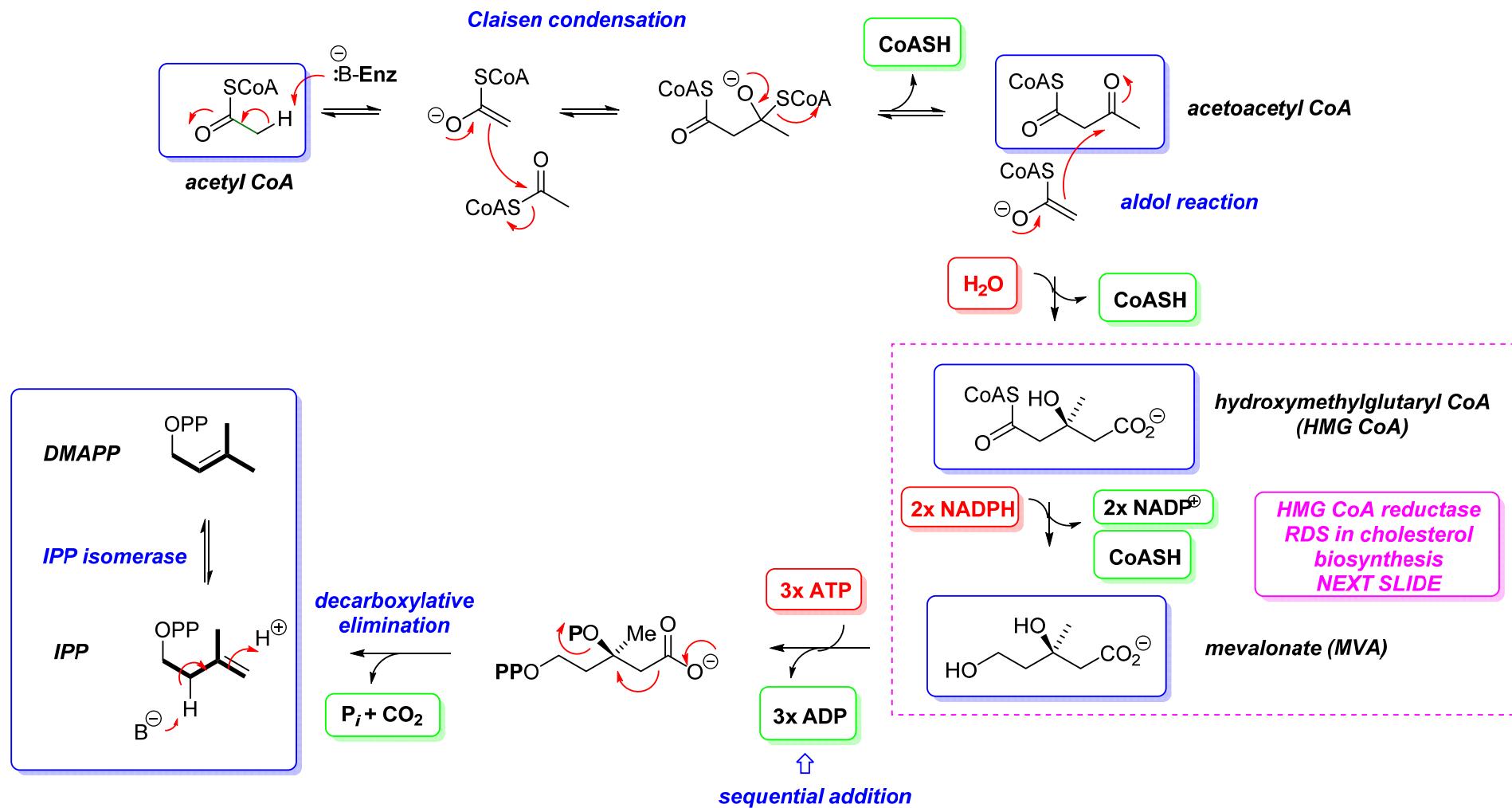
Isoprenoids

- **isoprenoids** are widely distributed in the natural world
 - particularly prevalent in plants and least common in insects; >30,000 known
 - composed of integral numbers of C₅ ‘isoprene’ units:
 - **monoterpenes** (C₁₀); **sesquiterpenes** (C₁₅); **diterpenes** (C₂₀); **sesterterpenes** (C₂₅, rare); **triterpenes** (C₃₀); **carotenoids** (C₄₀)



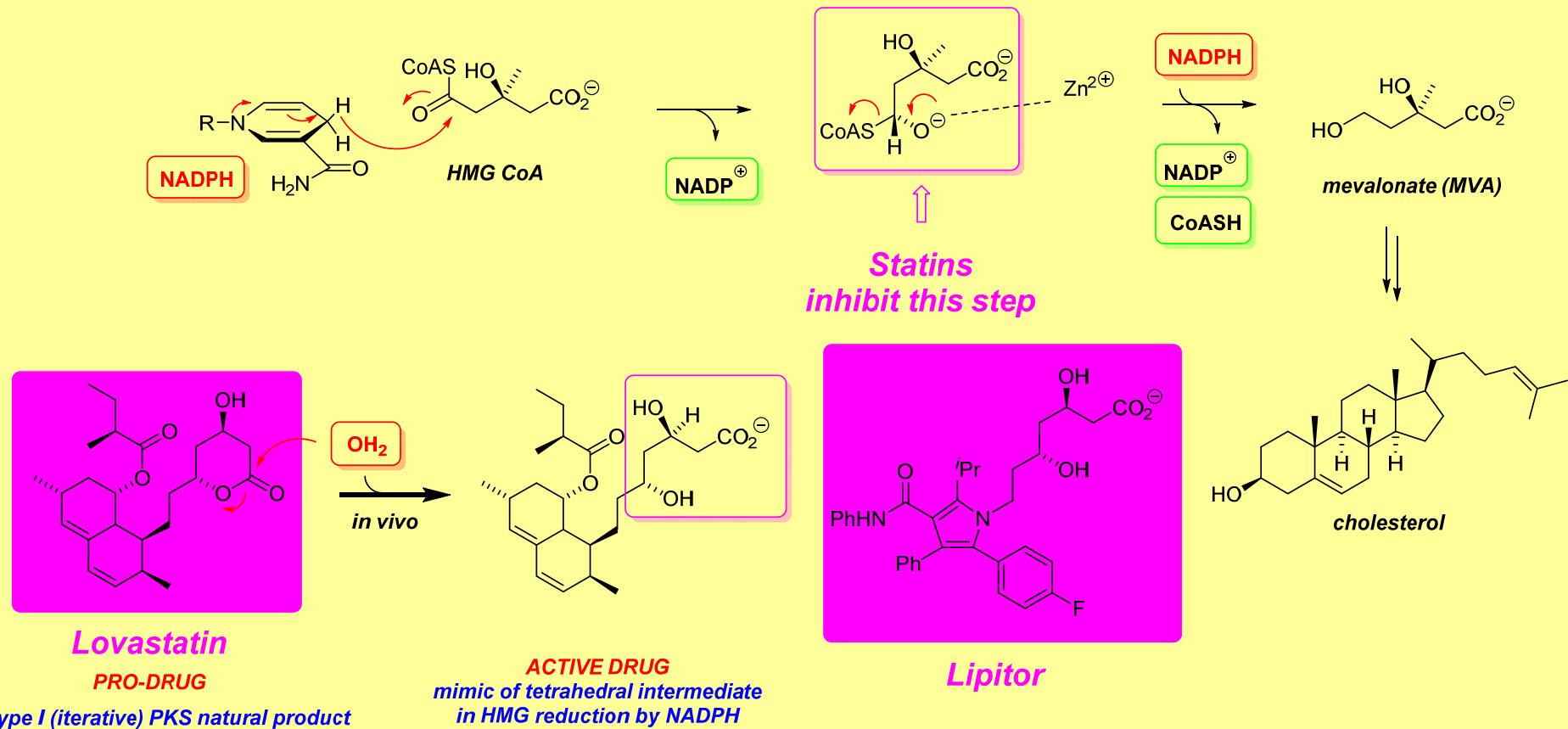
Biosynthesis of IPP & DMAPP - via Mevalonate

- **IPP & DMAPP** are the key *C₅* precursors to *all isoprenoids*
 - the *main pathway* is via: **acetyl CoA** → **acetoacetyl CoA** → **HMG CoA** → **mevalonate** → **IPP** → **DMAPP**:



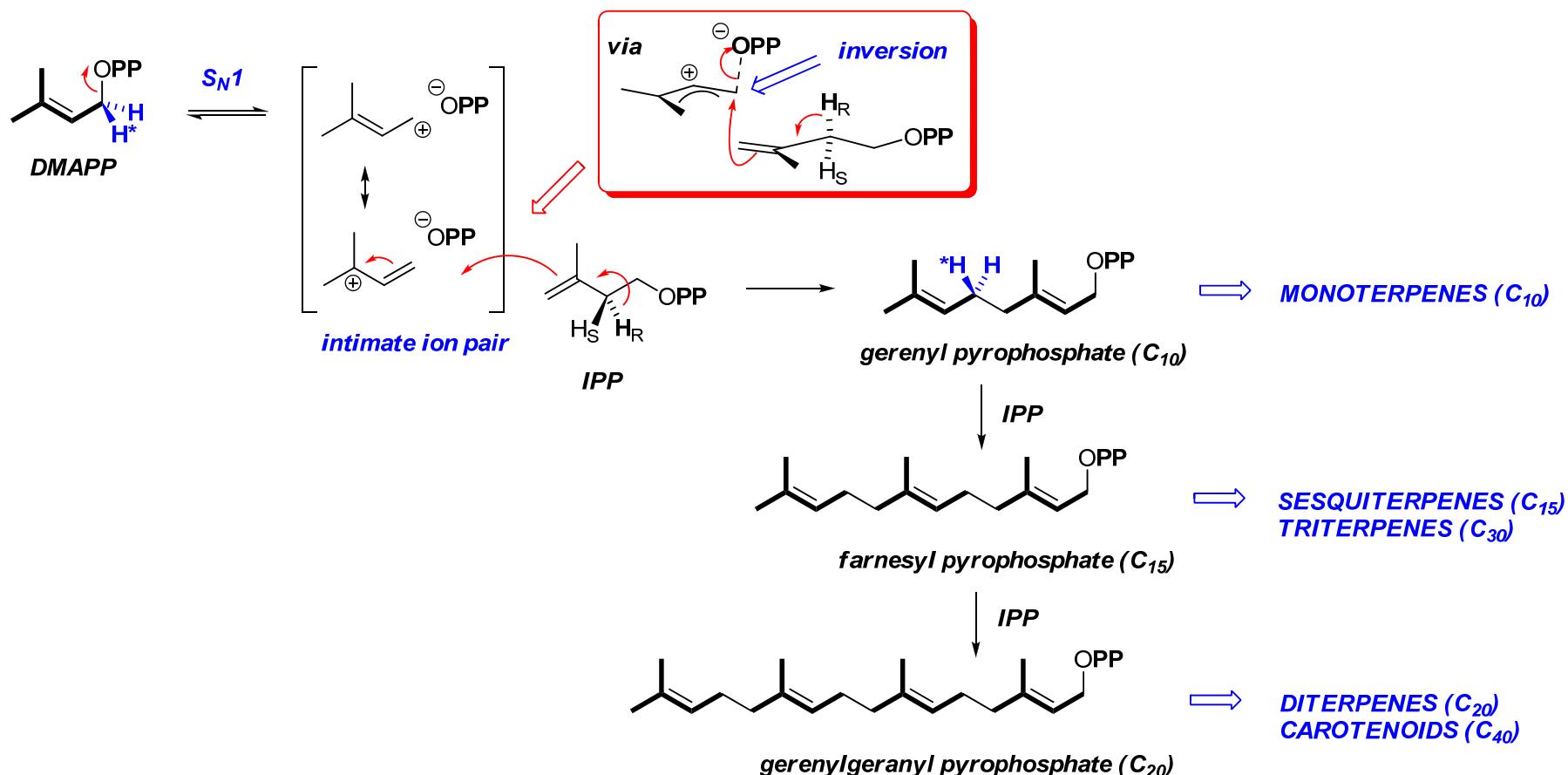
Rational Anti-cholesterol Development - *Statins*

- **HMG CoA → MVA** is the *rate determining step* in the biosynthetic pathway to **cholesterol**
- ‘**Statins**’ inhibit HMG CoA reductase and are used clinically to treat **hypercholesterolaemia** - a causative factor in **heart disease**, see: Wu et al. *Tetrahedron* 2015, 71, 8487 ([DOI](#))
 - e.g. **mevinolin** (=lovastatin®, Merck) from *Aspergillus terreus* is a competitive inhibitor of HMG-CoA reductase
 - e.g. **Lipitor** (Atorvastatin calcium, Pfizer) is also a competitive inhibitor of HMG-CoA reductase and the worlds biggest selling drug [first drug to reach \$10 billion sales (2004: \$10.8 bn)]



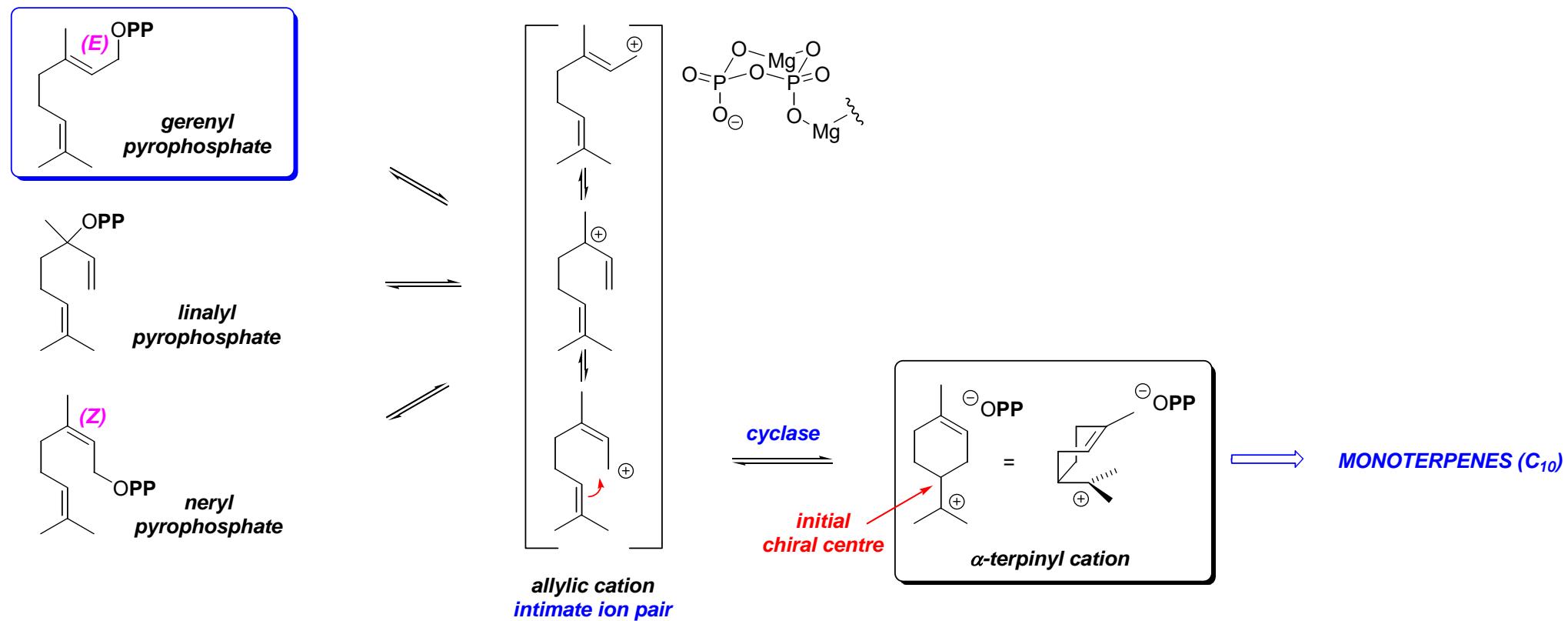
Linear C_{5n} ‘head-to-tail’ Pyrophosphates

- head-to-tail **C₅ oligomers** are the key precursors to isoprenoids
 - geranyl** pyrophosphate (C₁₀) is formed by **S_N1 alkylation** of **DMAPP** by **IPP** → **monoterpene**s
 - farnesyl** (C₁₅) & **geranylgeranyl** (C₂₀) pyrophosphates are formed by **further S_N1 alkylations** with **IPP**:



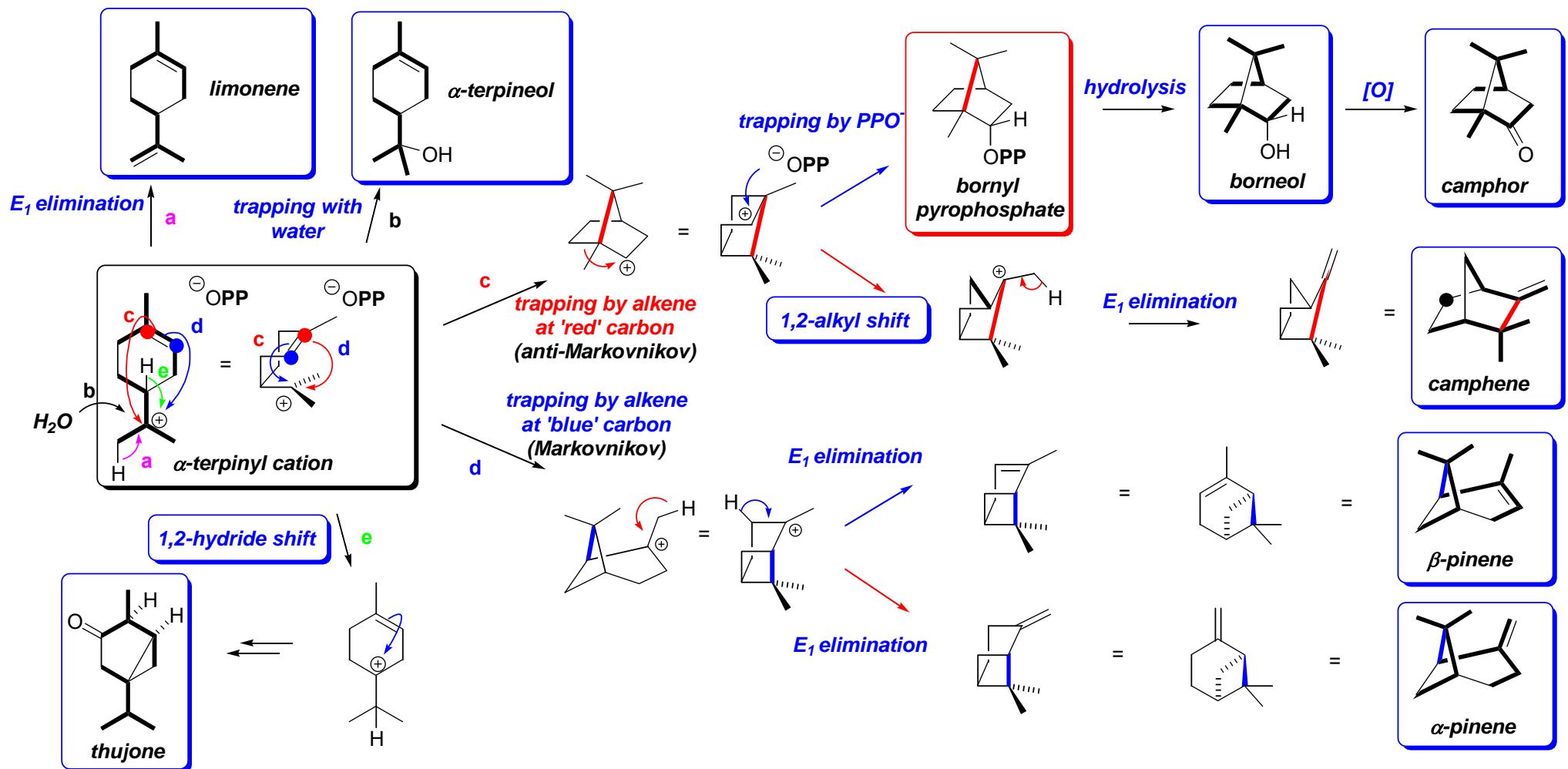
Monoterpenes – α -Terpinyl Cation Formation

- **geranyl pyrophosphate** isomerises readily via an allylic cation to **linalyl** & **neryl** pyrophosphates
 - the leaving group ability of pyrophosphate is enhanced by coordination to Mg^{2+} ions
 - all three pyrophosphates are substrates for **cyclases** via an **α -terpinyl cation**:



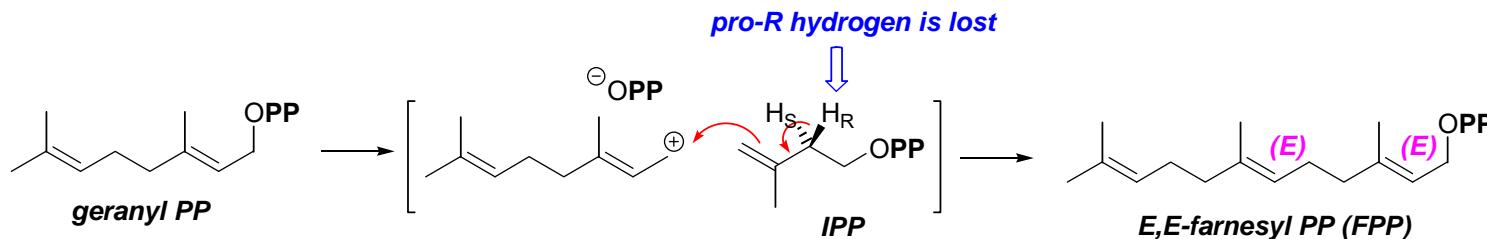
Monoterpenes – Fate of the α -Terpinyl Cation

- The α -terpinyl cation undergoes a rich variety of further chemistry to give a diverse array of monoterpenes
- Some important enzyme catalysed pathways are shown below
 - NB. intervention of Wagner-Meerwein 1,2-hydride- & 1,2-alkyl shifts

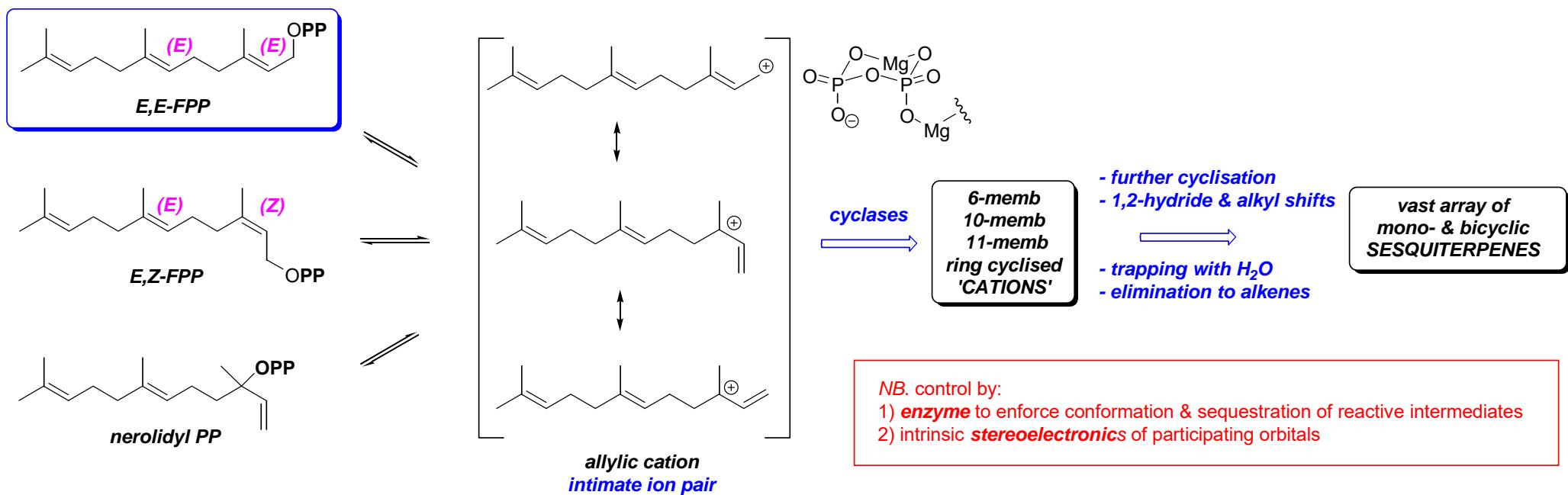


Sesquiterpenes – Farnesyl Pyrophosphate (FPP)

- ‘S_N2’-like alkylation of geranyl PP by IPP gives farnesyl PP:

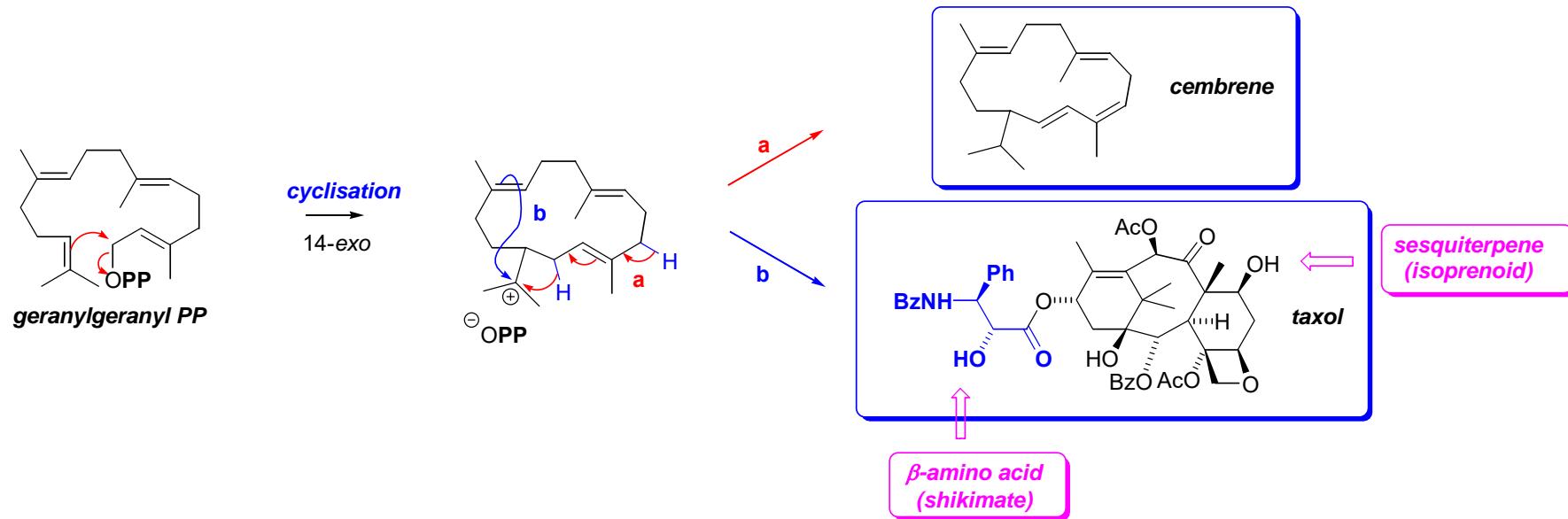


- just as **geranyl PP** readily isomerises to neryl & linaly PPs so **farnesyl PP** readily isomerises to equivalent compounds – allowing many modes of cyclisation & bicyclisation



Diterpenes – Geranylgeranyl PP → Taxol

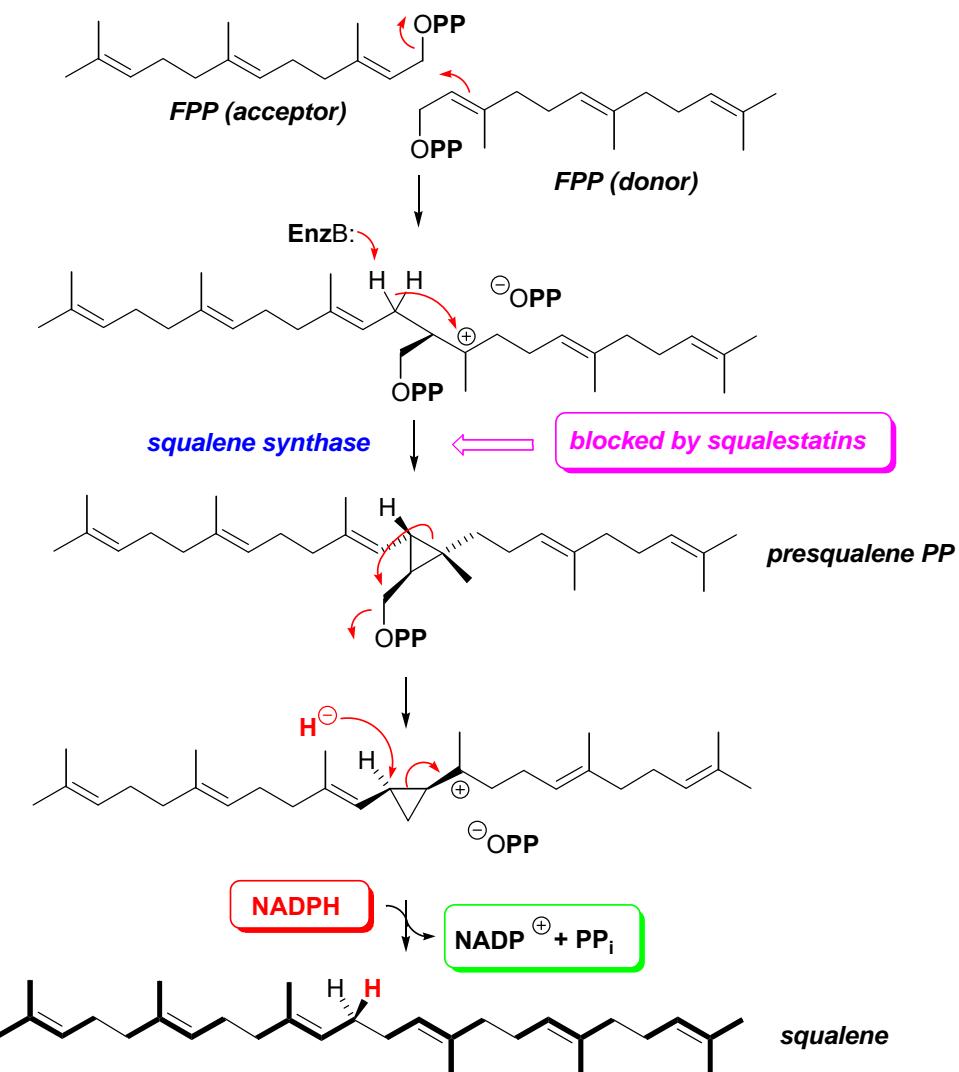
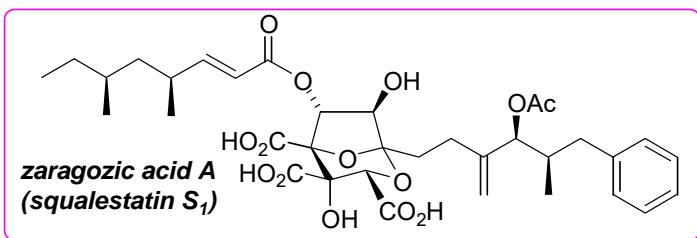
- **Taxol** is a potent **anti-cancer agent** used in the treatment of **breast & ovarian cancers**
 - comes from the bark of the **pacific yew** (*Taxus brevifolia*)
 - binds to tubulin and interferes with the assembly of microtubules
- biosynthesis is from **geranylgeranyl PP**:



- for details see: <http://www.chem.qmul.ac.uk/iubmb/enzyme/reaction/terp/taxadiene.html>
- home page is: <http://www.chem.qmul.ac.uk/iubmb/enzyme/>
 - recommendations of the Nomenclature Committee of the International Union of Biochemistry and Molecular Biology on the Nomenclature and Classification of Enzyme-Catalysed Reactions
 - based at Department of Chemistry, Queen Mary University of London

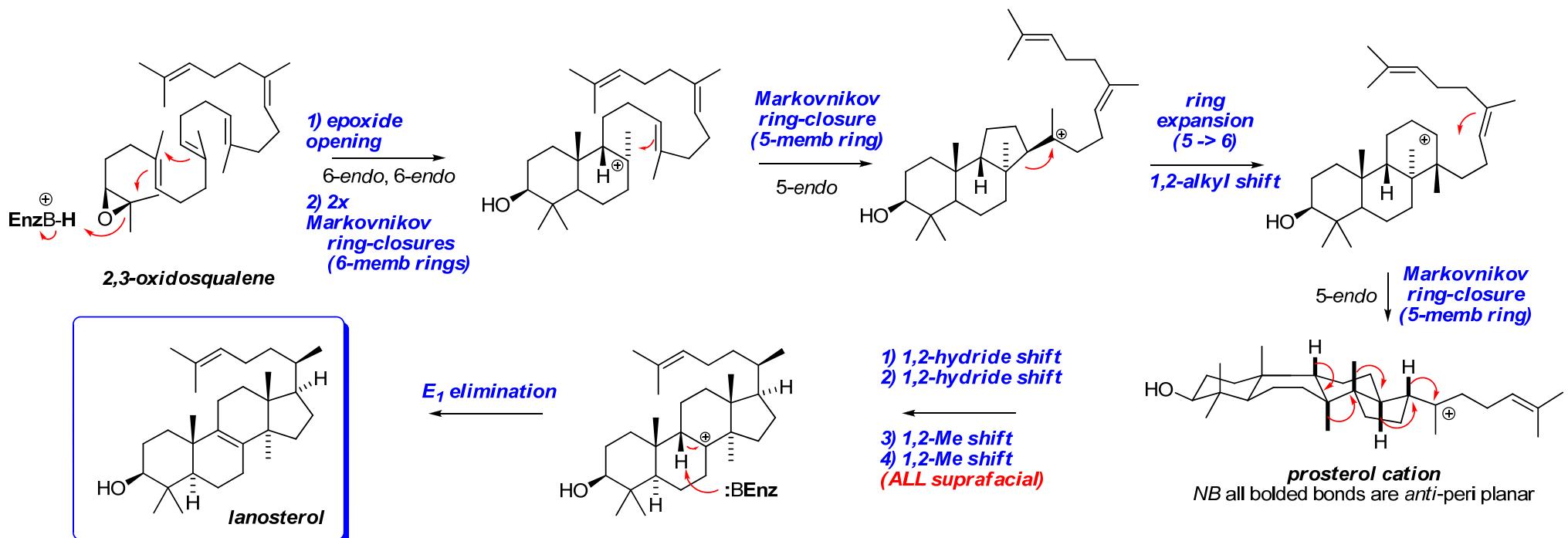
Triterpenes – $FPP \rightarrow Squalene$

- **triterpenes** (C_{30}) arise from the '**head to head**' **coupling** of **two farnesyl PP units** to give **squalene** catalysed by **squalene synthase**:
 - squalene was first identified as a steroid precursor from **shark liver oil**
 - the dimerisation proceeds *via* an unusual mechanism involving electrophilic cyclopropane formation - rearrangement to a tertiary cyclopropylmethyl cation and reductive cyclopropane ring-opening by NADPH (*NB.* exact mechanism disputed)
 - **Zaragozic acids (squalestatins)** mimic a rearrangement intermediate and inhibit squalene synthase. They constitute interesting leads for development of new treatments for **hypercholesterolaemia & heart disease** (*cf.* statins)



Oxidosqualene-Lanosterol Cyclase – Mechanism

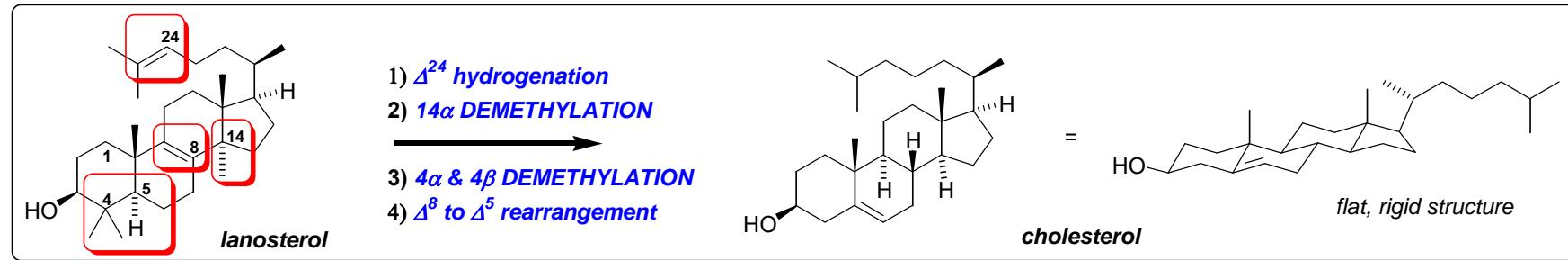
- **oxidosqualene-lanosterol cyclase** catalyses the formation of **lanosterol** from **2,3-oxidosqualene**:
 - this cascade establishes the characteristic ring system of **ALL steroids**
 - ring-expansion sequence to establish the C ring
 - the process is **NOT concerted**, discrete **cationic intermediates** are involved & **stereoelectronics dictate the regio- & stereoselectivity** although the enzyme undoubtedly lays a role in pre-organising the ~chair-boat-chair conformation



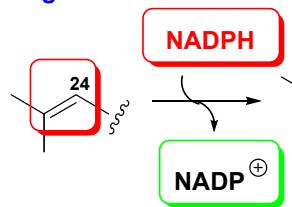
- “The enzyme’s role is most likely to shield intermediate carbocations... thereby allowing the hydride and methyl group migrations to proceed down a thermodynamically favorable and kinetically facile cascade”
 - Wendt et al. *Angew. Chem. Int. Ed.* **2000**, 39, 2812 ([DOI](#)) & Wendt *ibid* **2005**, 44, 3966 ([DOI](#))

Lanosterol → Cholesterol – Oxidative Demethylation

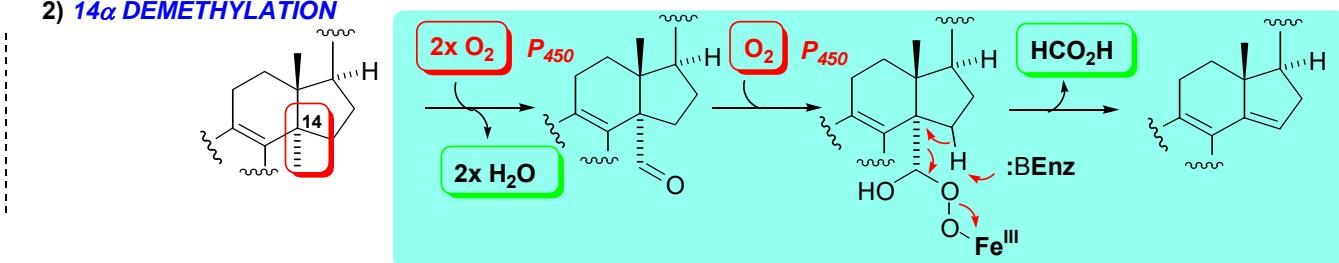
- Several steps are required for conversion of *lanosterol* to *cholesterol*:



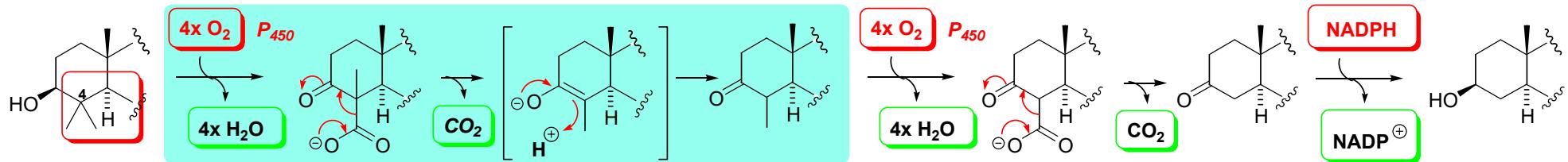
1) Δ^{24} hydrogenation



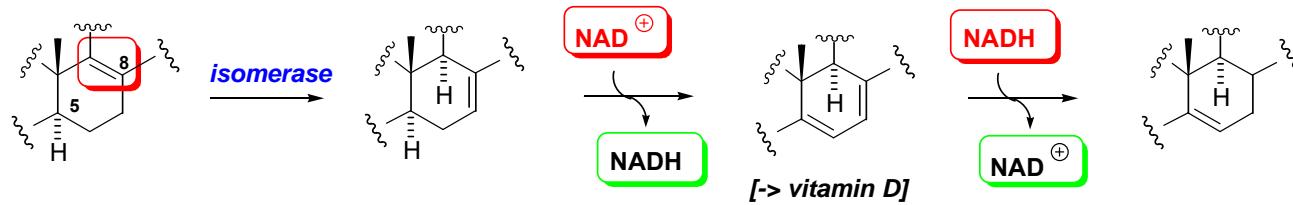
2) 14α DEMETHYLATION



3) 4α & 4β DEMETHYLATION

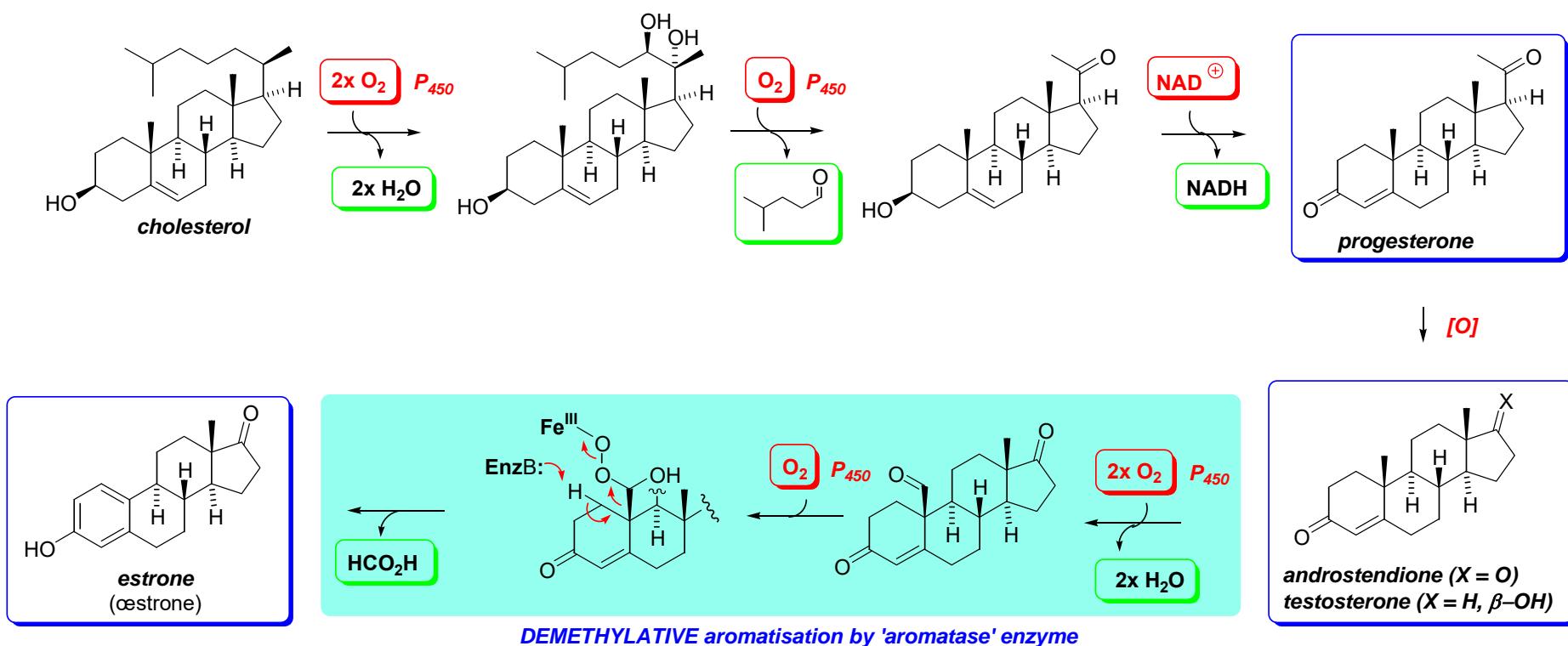


4) Δ^8 to Δ^5 rearrangement



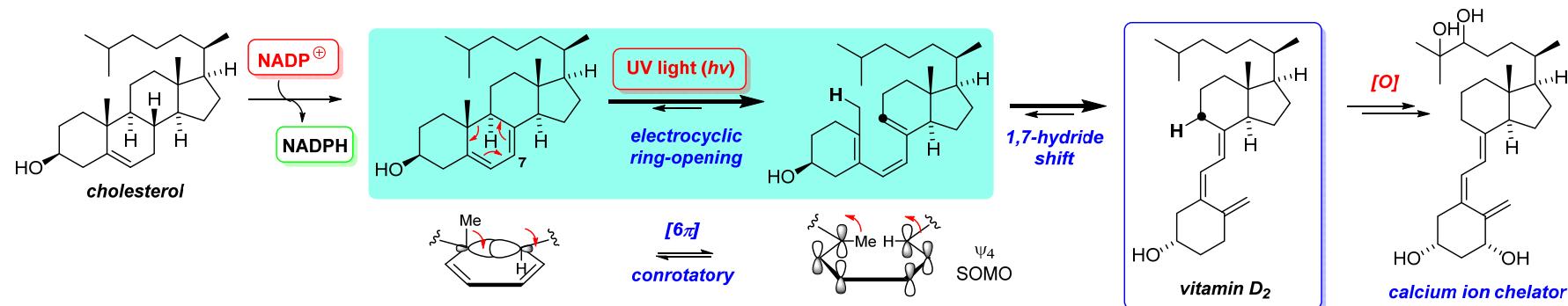
Cholesterol → Human Sex Hormones

- **cholesterol** is the precursor to the human sex hormones – **progesterone**, **testosterone** & **estrone**
 - the pathway is characterised by **extensive oxidative processing** by **P_{450}** enzymes
 - **estrone** is produced from **androstendione** by **oxidative demethylation** with **concomitant aromatisation**:

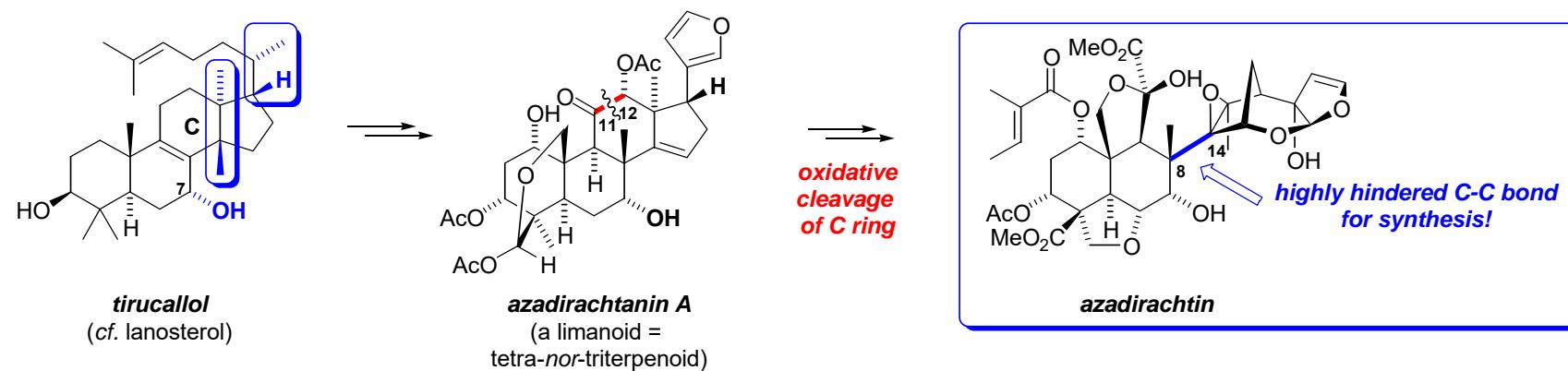


Steroid Ring Cleavage - Vitamin D & Azadirachtin

- **vitamin D₂** is biosynthesised by the **photolytic cleavage** of **Δ^7 -dehydrocholesterol** by UV light:
 - a classic example of **photo-allowed, conrotatory electrocyclic ring-opening**:



- D vitamins are involved in **calcium absorption; deficiency** leads to **rickets** (brittle/deformed bones)
- **Azadirachtin** is a potent **insect anti-feedant** from the Indian **neem tree**:
 - exact biogenesis unknown but certainly *via* steroid modification:



Summary of Presentation

- ***Metabolism & Biosynthesis***
 - some definitions, 1° & 2° metabolites
- ***Shikimate Metabolites***
 - photosynthesis & glycolysis → shikimate formation → shikimate metabolites
 - Glyphosate – a non-selective herbicide
- ***Alkaloids***
 - acetylCoA & the citric acid cycle → α-amino acids → alkaloids
 - Opioids – powerful pain killers
- ***Fatty Acids and Polyketides***
 - acetylCoA → malonylCoA → fatty acids, prostaglandins, polyketides, macrolide antibiotics
 - NSAIDs – anti-inflammatory's
- ***Isoprenoids/terpenes***
 - acetylCoA → mevalonate → isoprenoids, terpenoids, steroids, carotenoids
 - Statins – cholesterol-lowering agents