

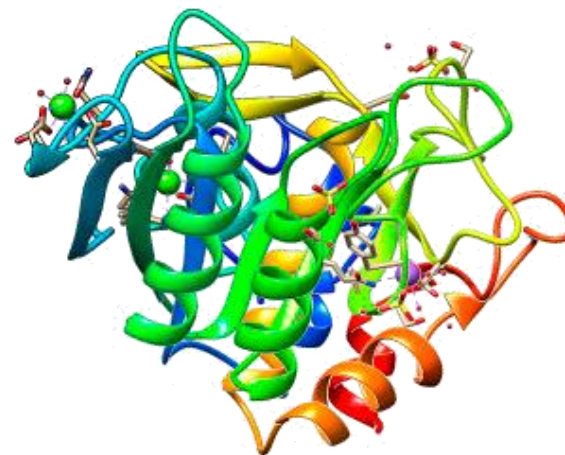


Subtilisin-like proteins and lipid signaling events: the missing links in grapevine resistance to *P. viticola*

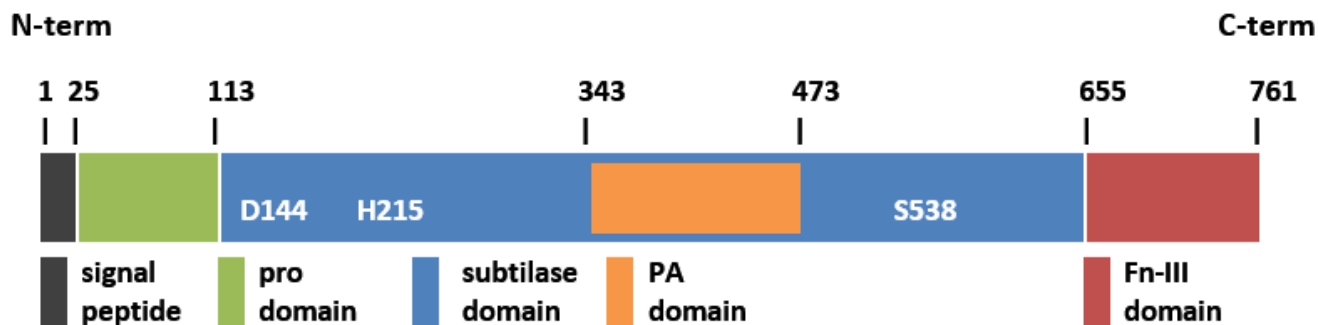
Andreia Figueiredo, Joana Figueiredo, Gonçalo Laureano, Ana Rita Cavaco, Marisa Maia, Ana Rita Matos, Marta Sousa Silva

Subtilisin-like proteases

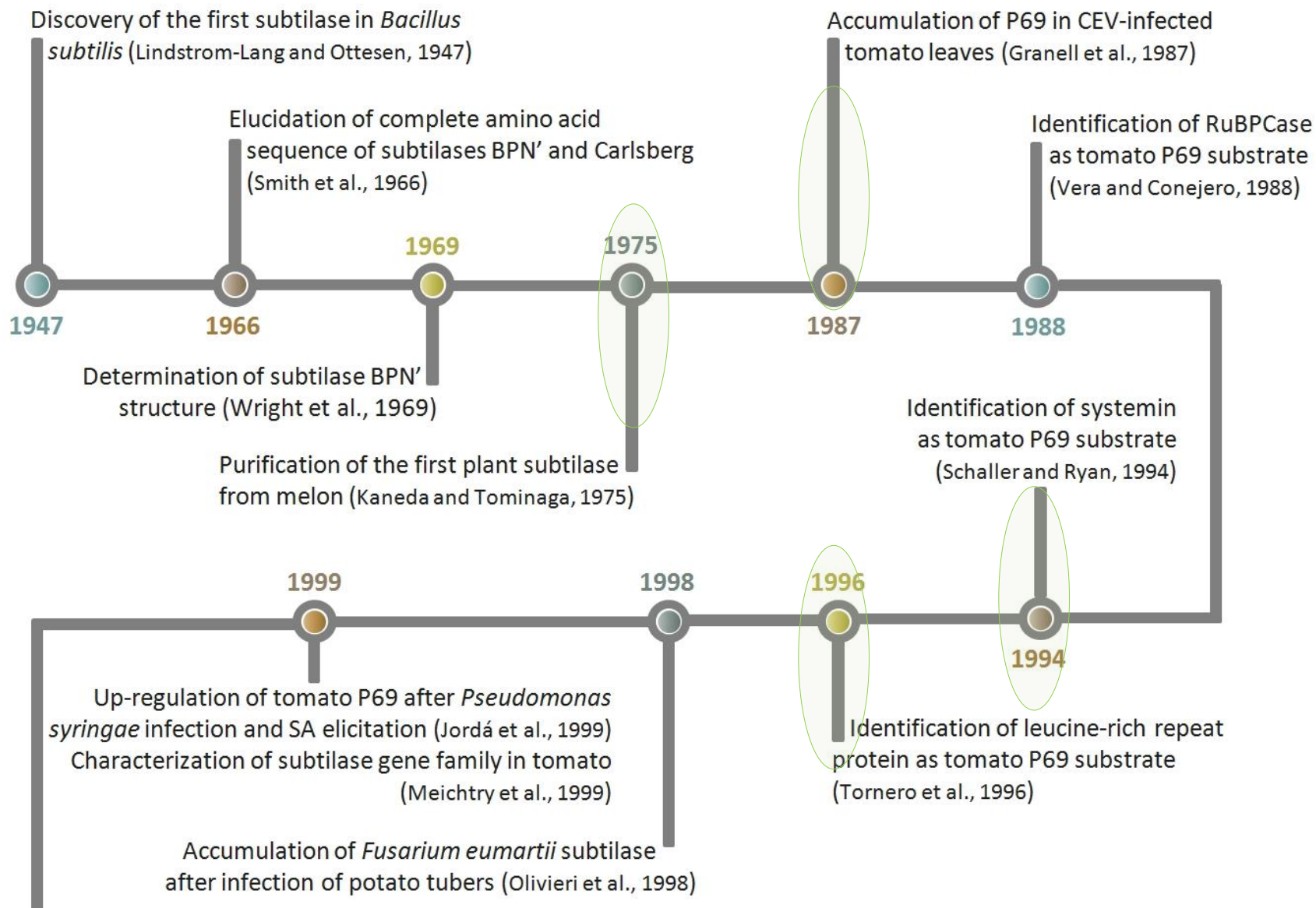
- Large group of serine peptidases
- S8 peptidase family (MEROPES)
- Broad spectrum of biological functions
- Pre-pro-proteins
 - Catalytic triad (Asp/His/Ser)
 - Pro-domain (I9 inhibitor domain)
 - Subtilase domain (S8 peptidase domain)
 - Protease-associated domain (PA domain)
 - C-terminal FnIII-like domain



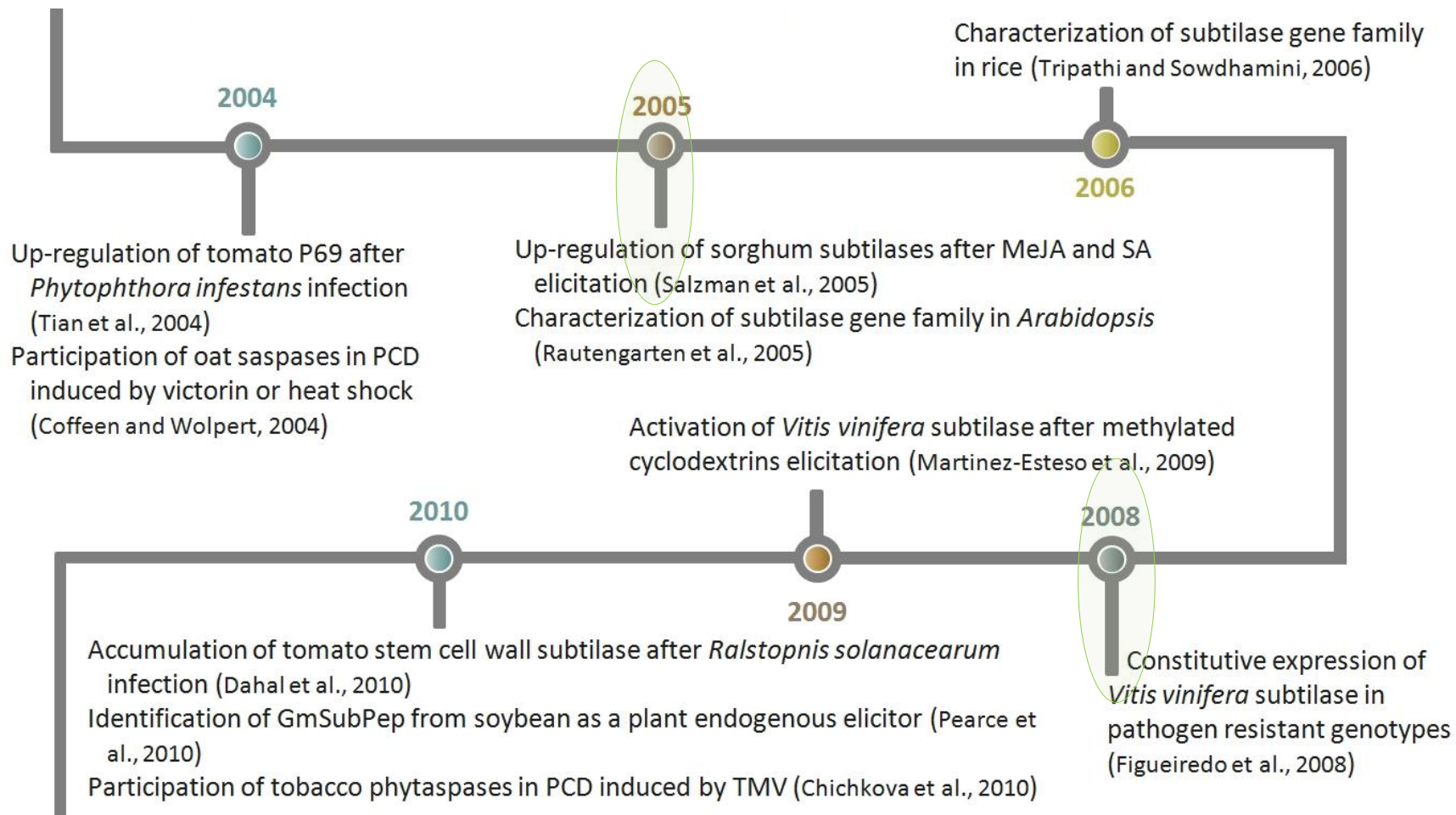
Structure prediction of the SISBt3 monomer
(PDB code:1THM)



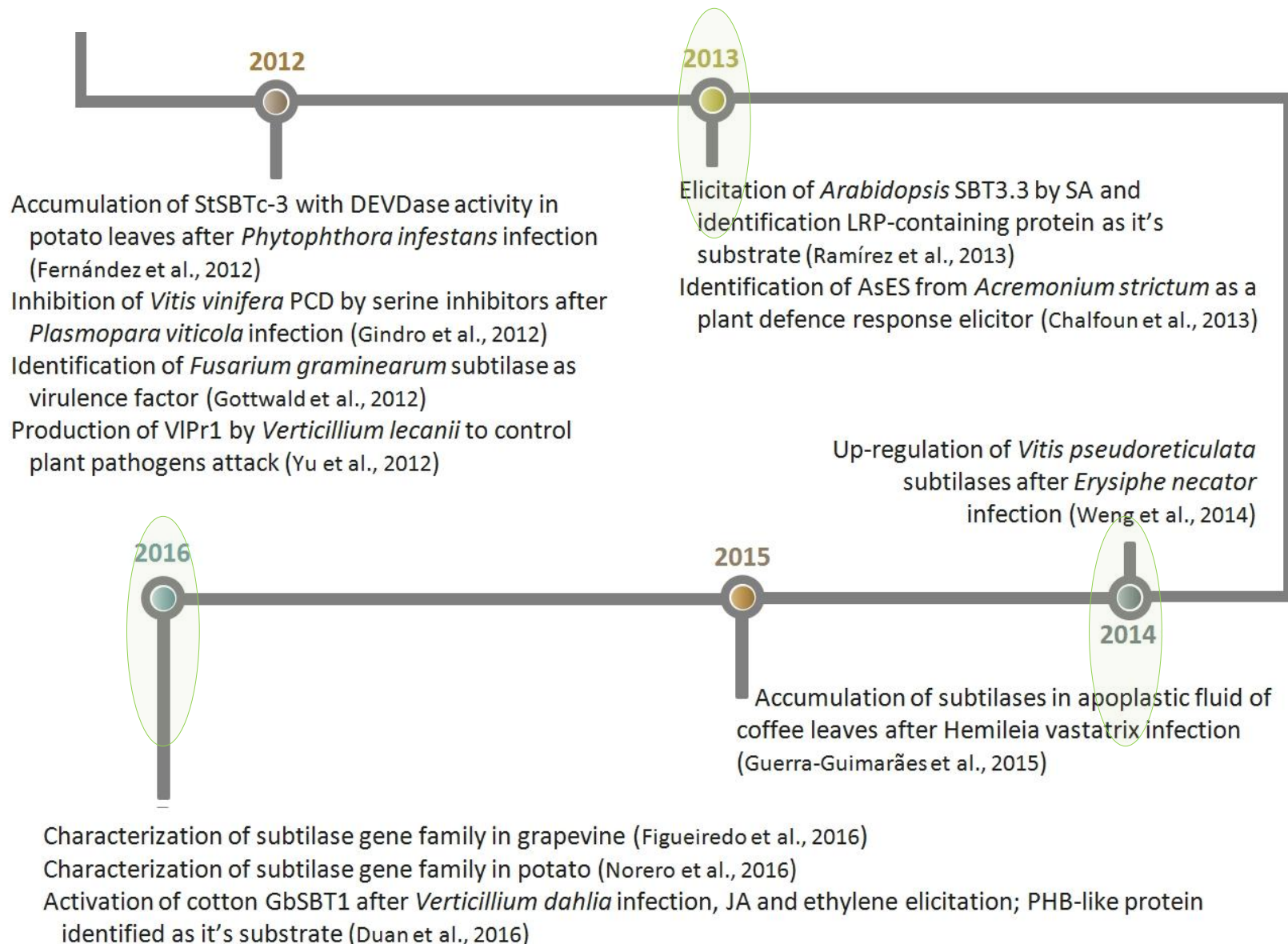
Plant subtilisin-like proteases



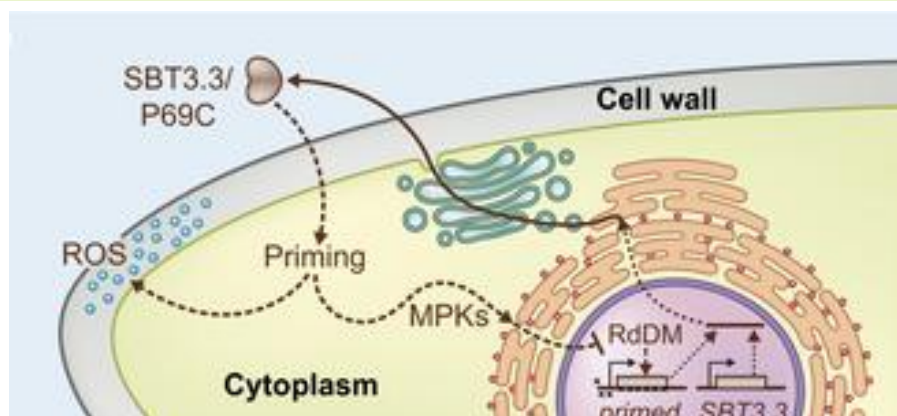
Plant subtilisin-like proteases



Plant subtilisin-like proteases

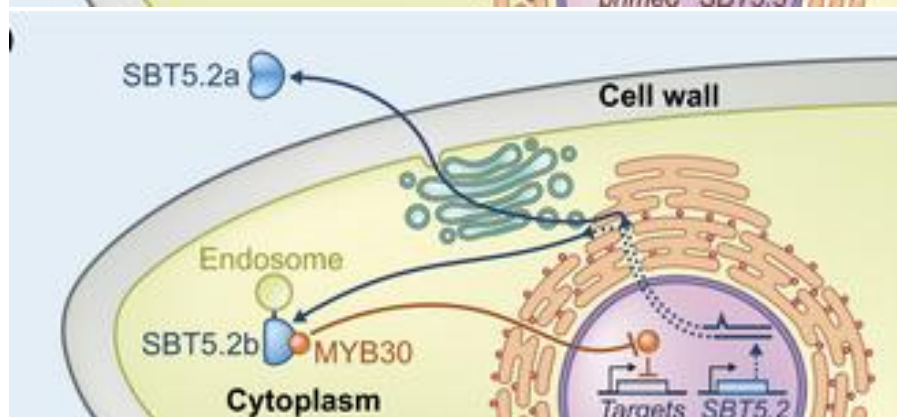


Subtilisin-like proteases in biotic interactions



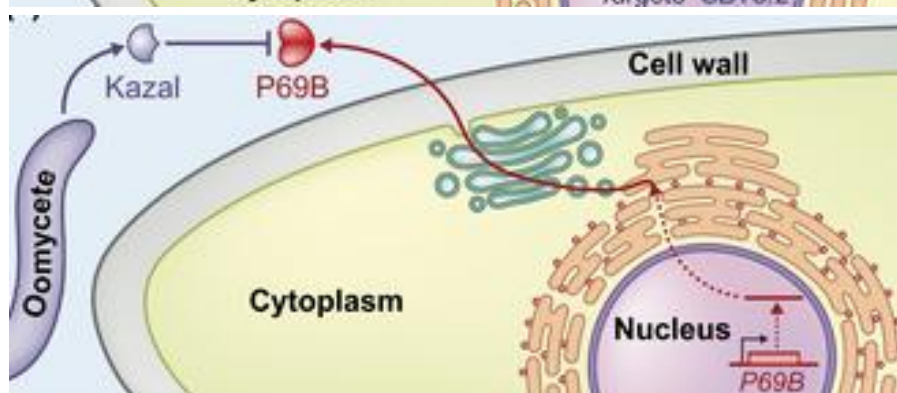
SBT3.3/P69B and immune priming

Pathogen triggers the expression of SBT3.3/P69 – negatively regulates RdDM – priming Increase of SBT3.3 expression – enhances the expression of OXI1 kinase, ROS and MAP kinase.



SBT5.2(b)

Interacts with MYB30 – nuclear exclusion – defense gene expression impairment – HR attenuation enhances the expression of OXI1 kinase, MAP kinase and ROS.

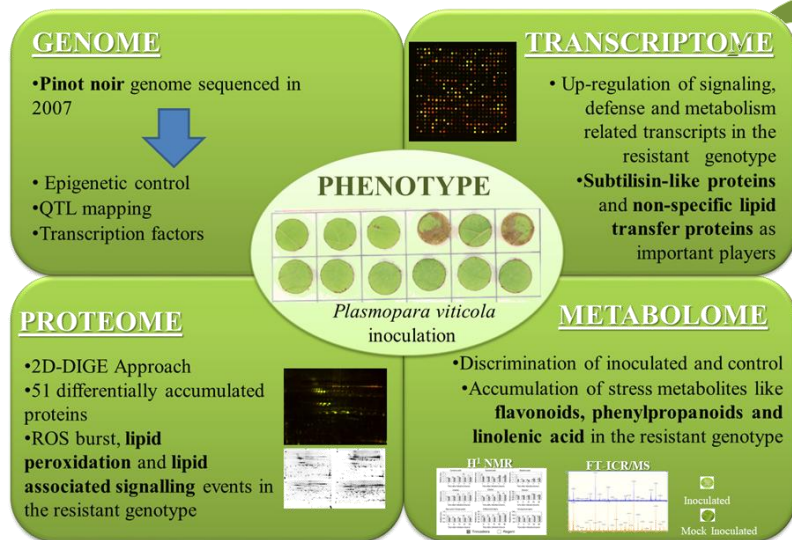


P69B(-like)

Inhibition by Kazal inhibitors from oomycetes.

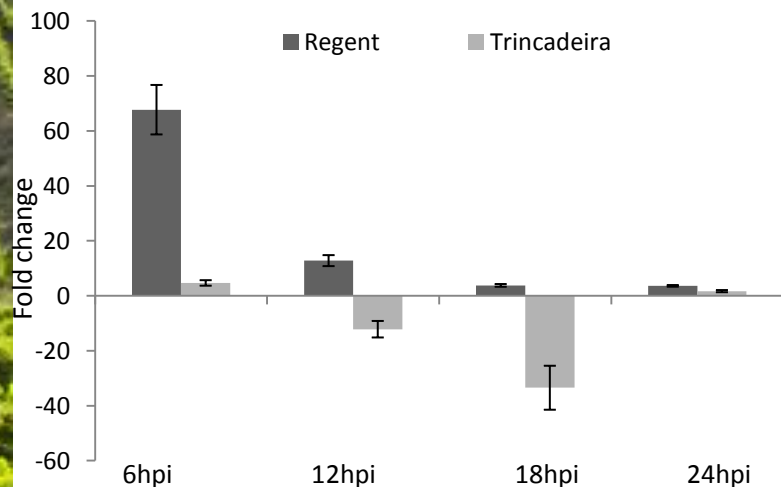
Schaller et al. (2018). New Phytologist, 218:901-915

Grapevine SBTs and *P. viticola* resistance



Grapevine SBT gene family

- Composed by 87 genes encoding 109 predicted proteins
- Divided into 5 groups
- Unevenly distributed among 15 of the 19 grapevine chromosomes
- 80% secreted, 10% mitochondrion, 9% chloroplast
- Glycosylated



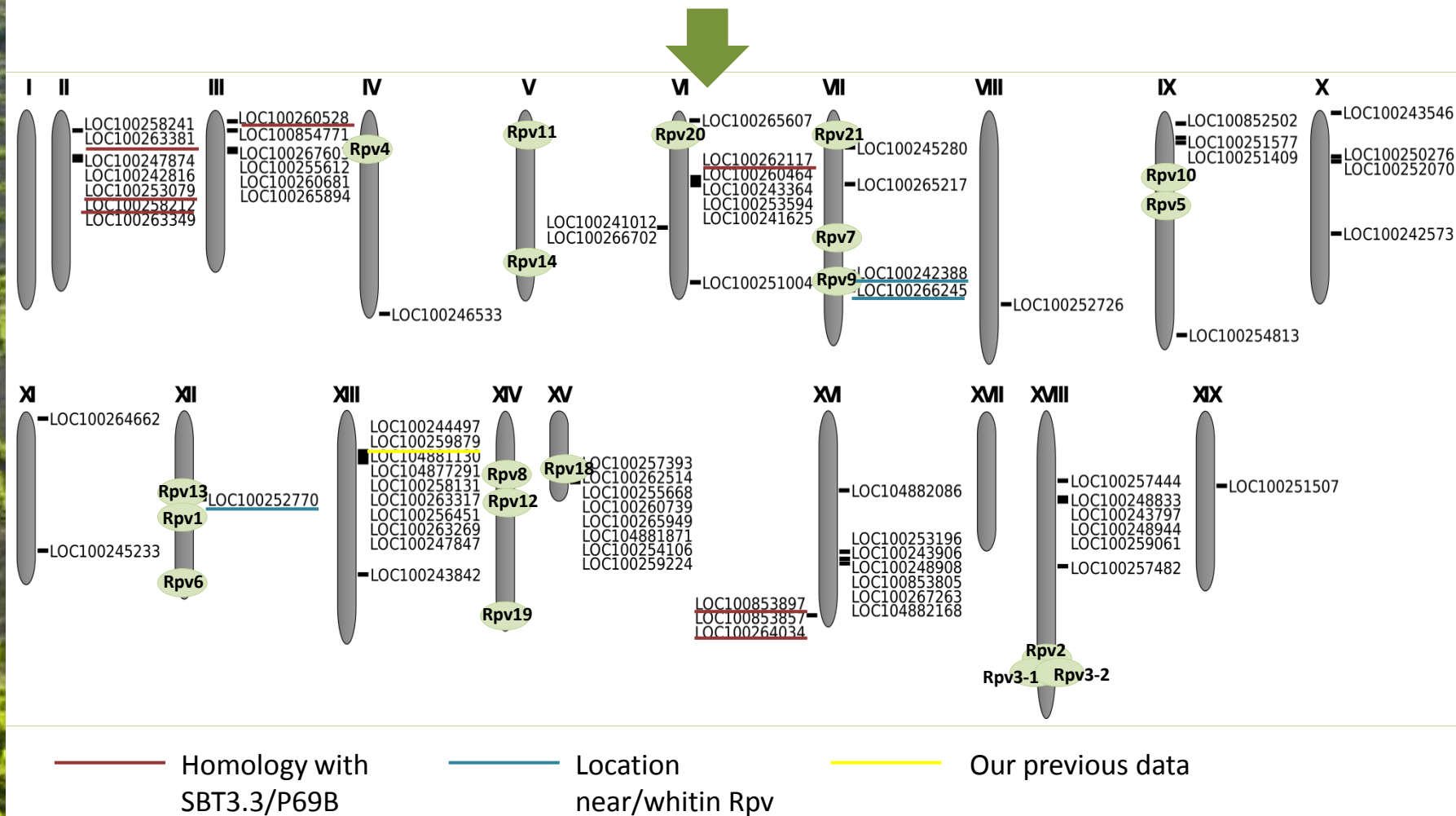
Selection of 14 grapevine subtilases:

- Previous microarray and expression data
- Sequence similarity with AtSBT3.3 and P69C
- Location near RPV loci (RPV9 in Chr7 and between RPV1 and RPV13, Chr12)

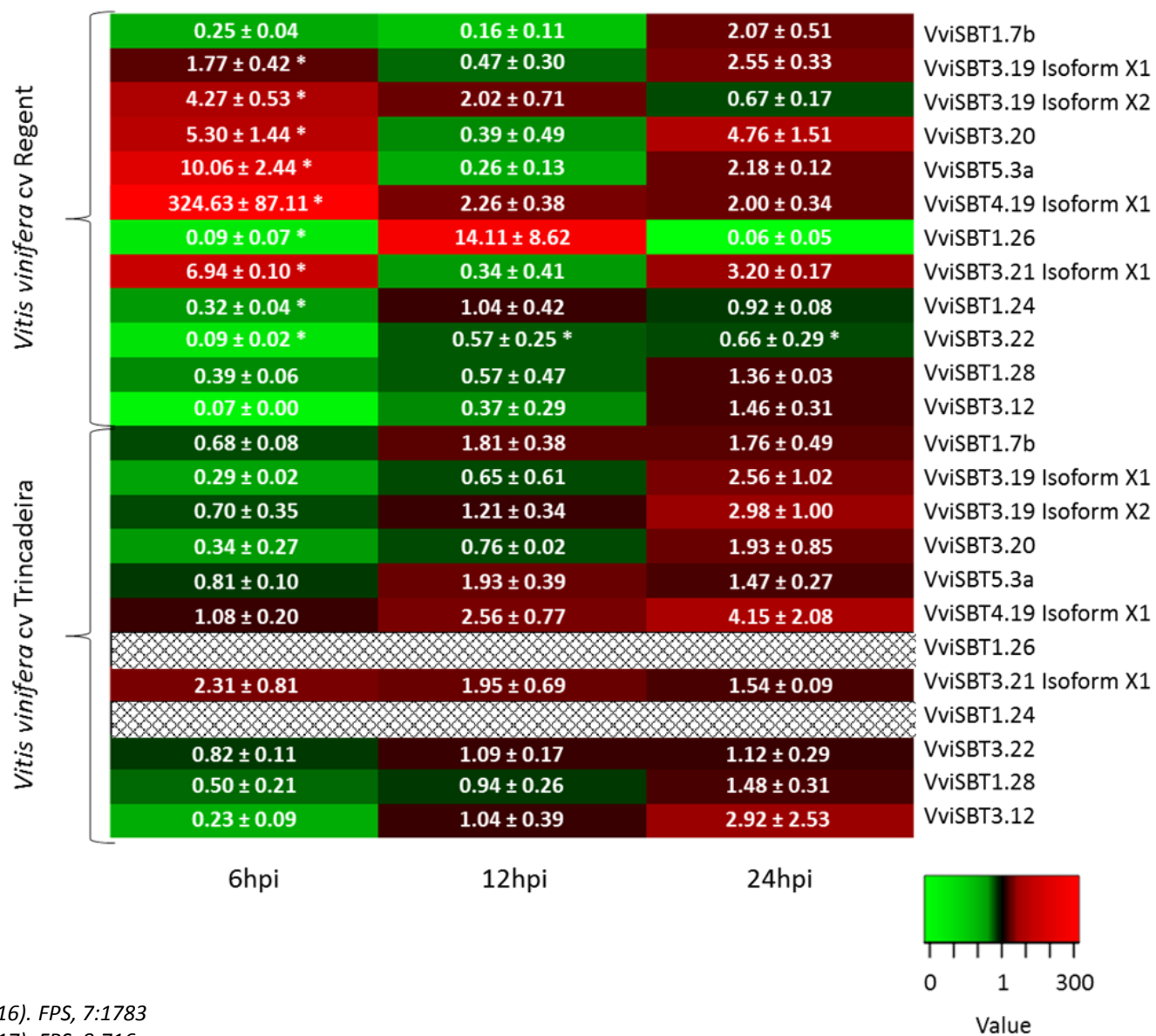
Figueiredo et al. (2016). FPS, 7:1783
Figueiredo et al. (2017), FPS, 8:716

Grapevine SBTs

New RPV identified - revision of candidates



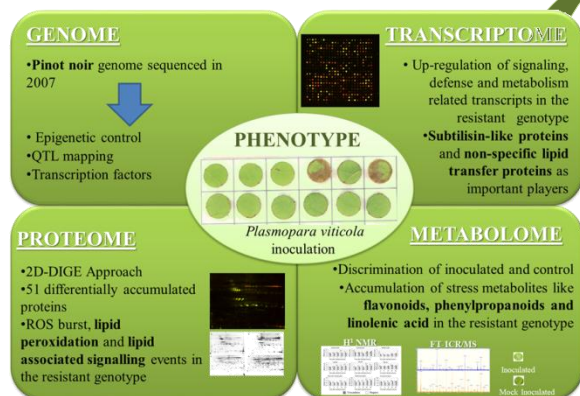
Grapevine SBTs



Figueiredo et al. (2016). FPS, 7:1783

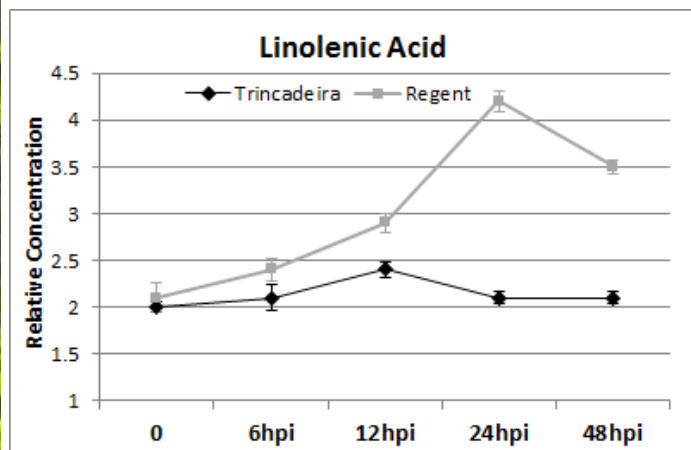
Figueiredo et al. (2017). FPS, 8:716

Lipid signaling in grapevine defense

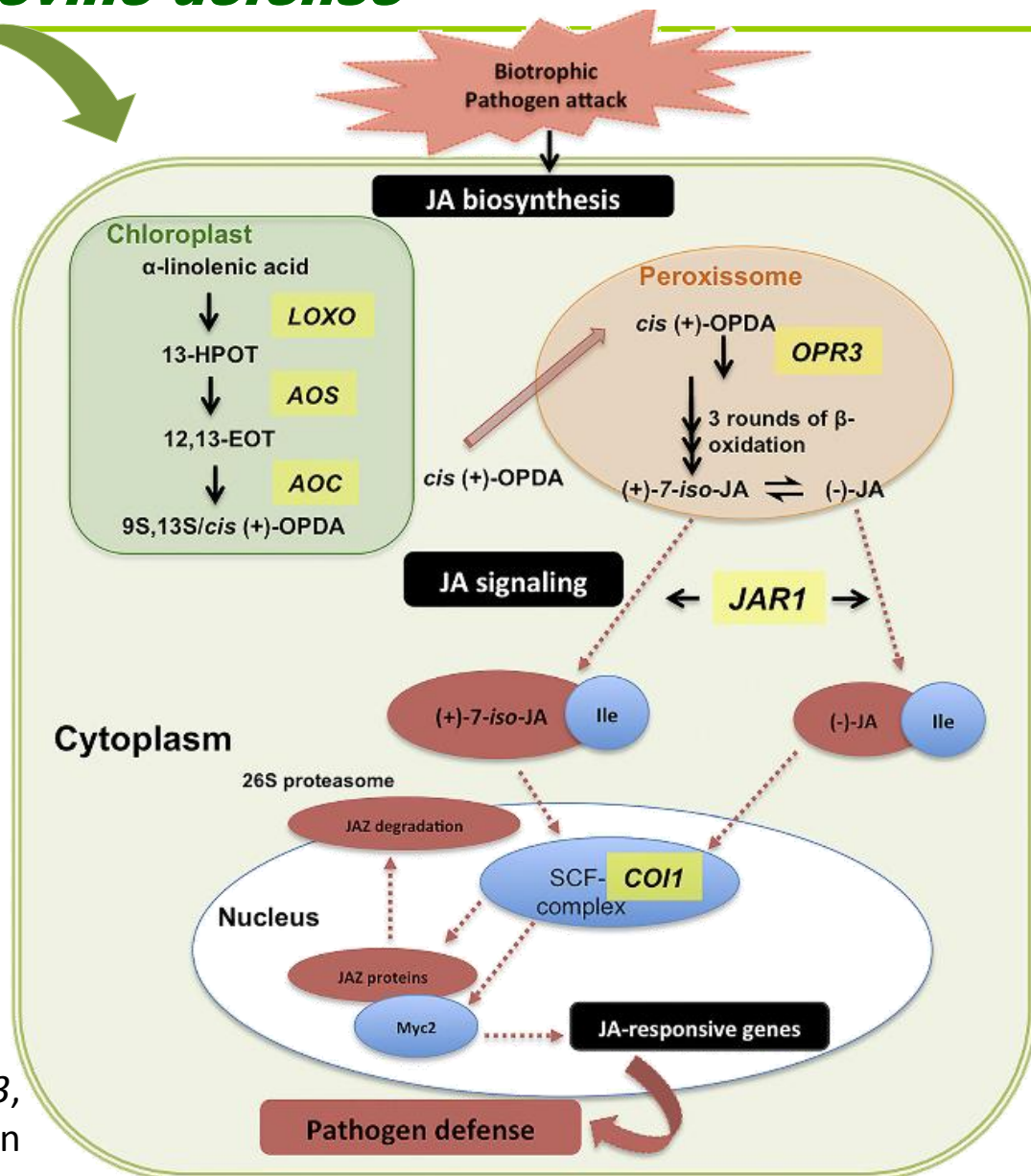


JA synthesis:

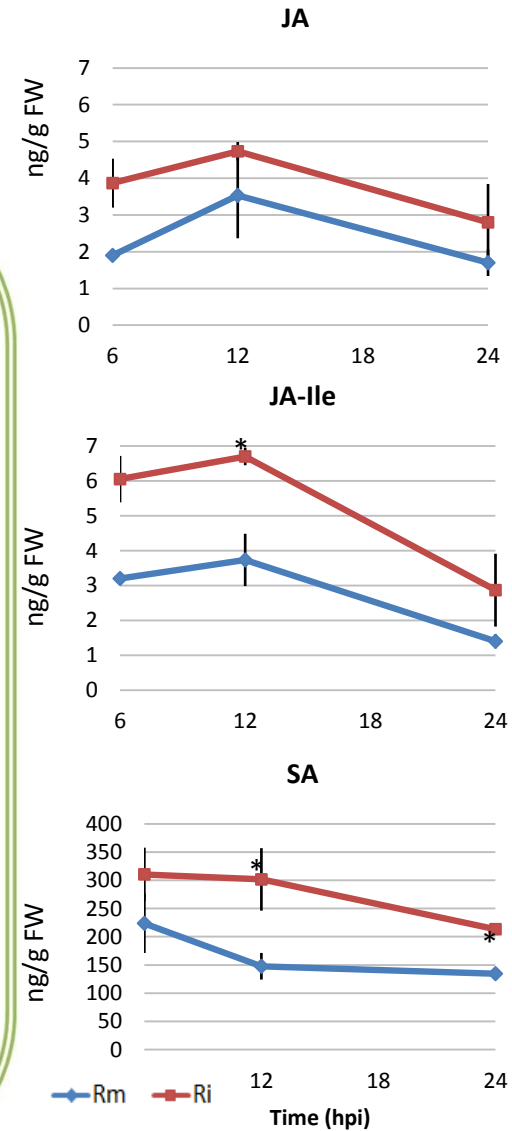
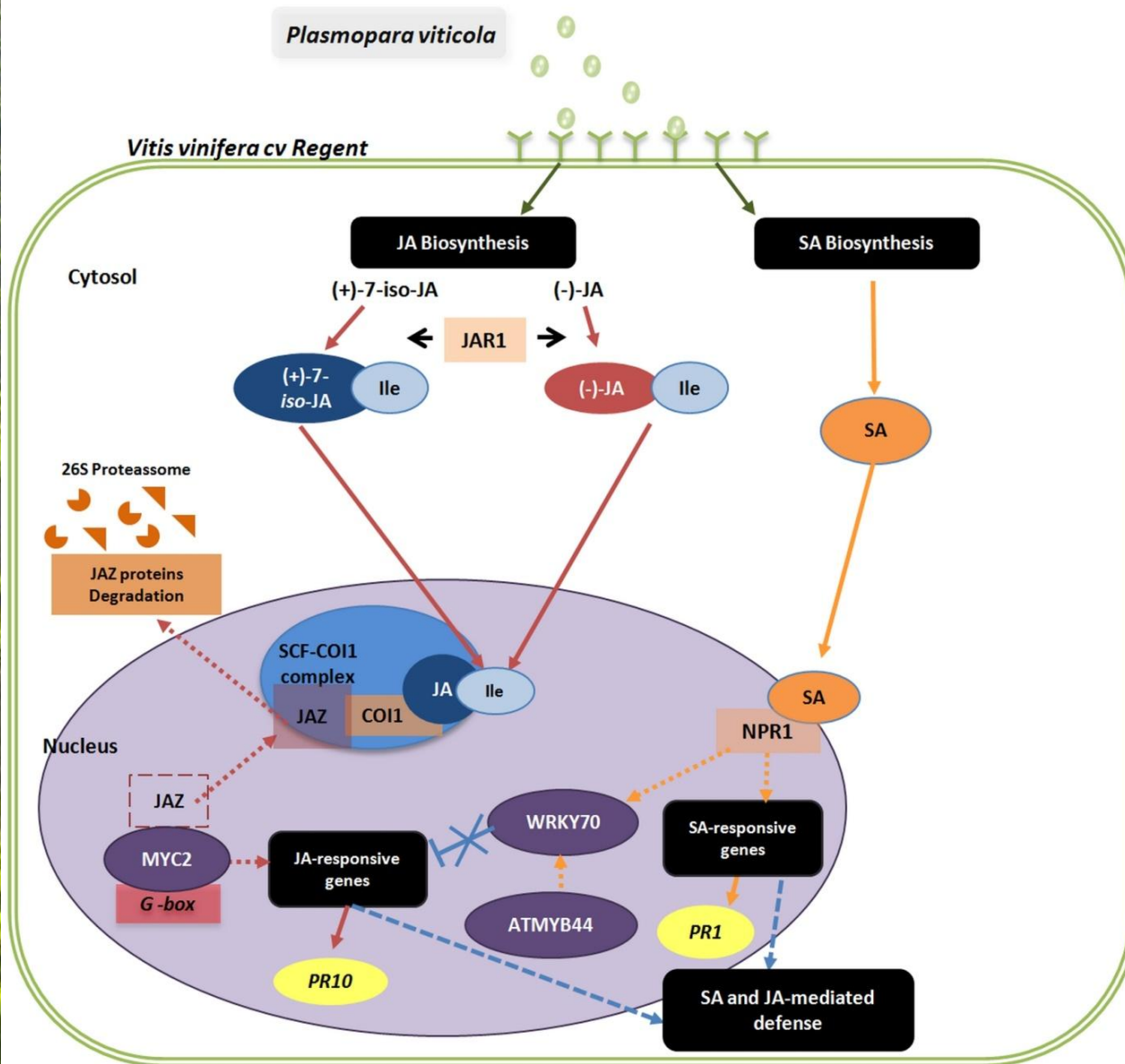
-C18:3 accumulation



-Increase of *LOX2*, *AOC*, *AOS*, *OPR3*, *JAR1*, *COI1* after *P. viticola* inoculation
 – incompatible interaction



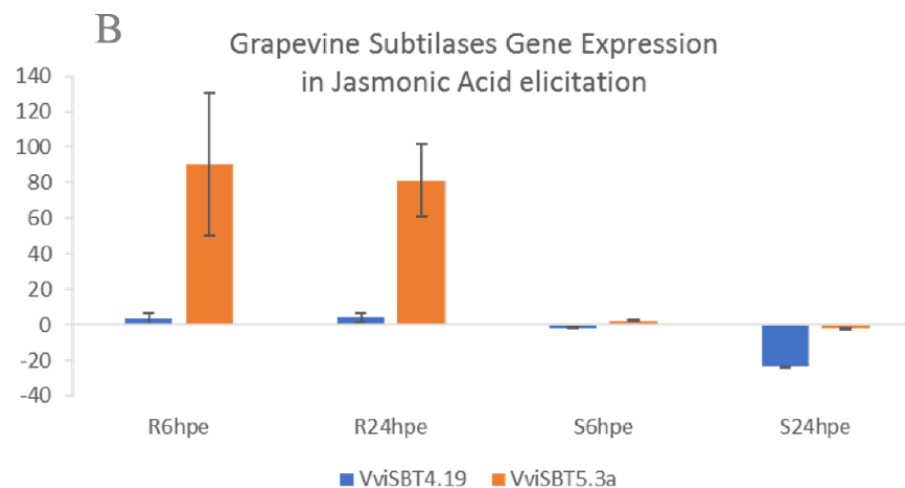
Lipid signaling in grapevine defense



Guerreiro et al. (2016). FPS, 7:565

SBTs and lipid signaling

- Systemin is derived from a precursor called prosystemin, similar to polypeptide hormones
- Systemin activates a lipid-based signaling cascade
- Wounding of tomato plants triggers the release of systemin from its precursor prosystemin by SBT3
- Systemin is perceived at cell surface by a LRR receptor-like kinase - JA biosynthesis induction – JA accumulation



SBTs and lipid signaling in *P. viticola* resistance

- Some grapevine subtilases are linked to defense mechanisms towards *P. viticola*
- Lipid signalling and jasmonic acid are also important players
- Subtilases, particularly SBT3.3 and P69C are believed to be associated to immune priming events and activation of signalling cascades
- A link between JA and SBT3.3 has been shown for other model



Are grapevine subtilisin-like proteins linked to lipid signaling events – grapevine immune priming?

Apoplast
proteomics and
lipidomics

Recombinant protein
production (structure
and substrates)

Interactome
analysis

Acknowledgments



BioISI

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Characterization of grapevine subtilisin-like proteases and their role in pathogen recognition and immune priming



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Thank you for
your attention!