

Genetic Basis for the 3-ADON and 15-ADON Trichothecene Chemotypes in *Fusarium*

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Bacterial Foodborne Pathogen and
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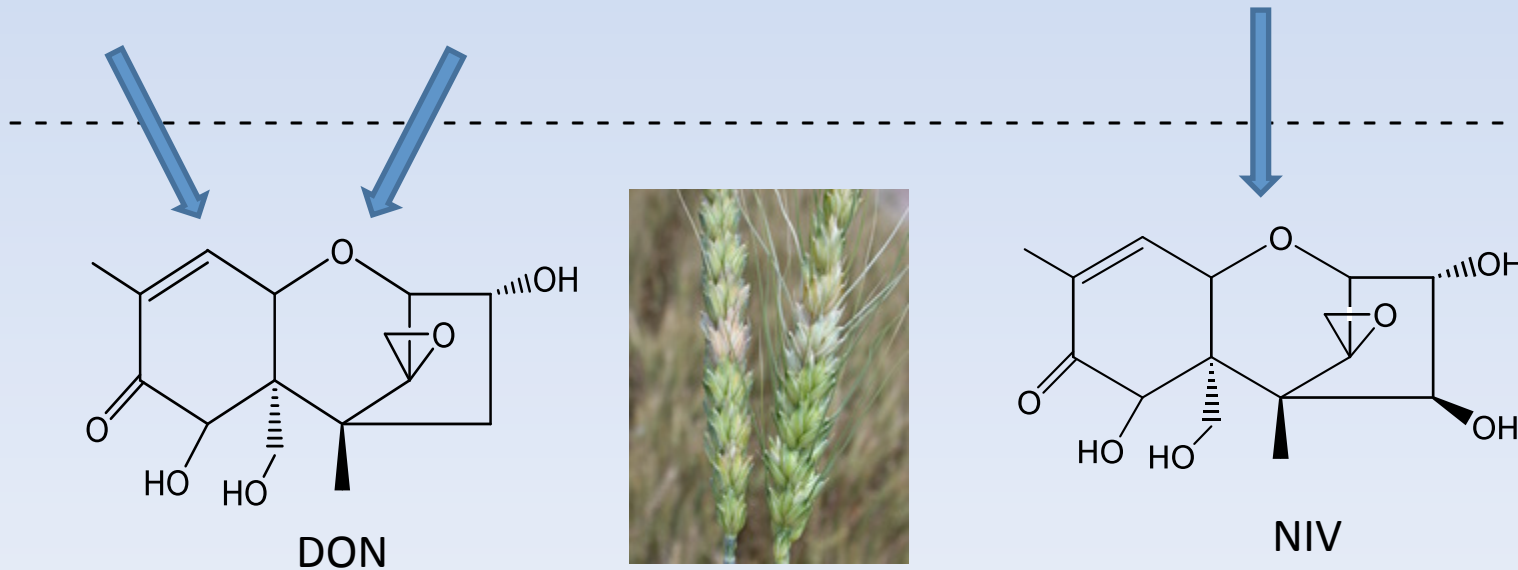
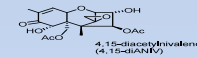
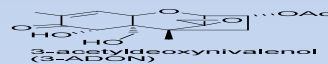
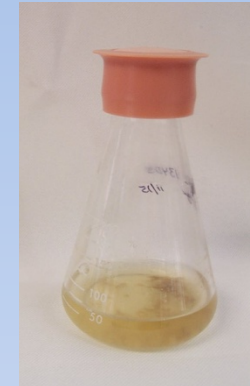
Major mycotoxins found in *Fusarium*-infected grains

DON

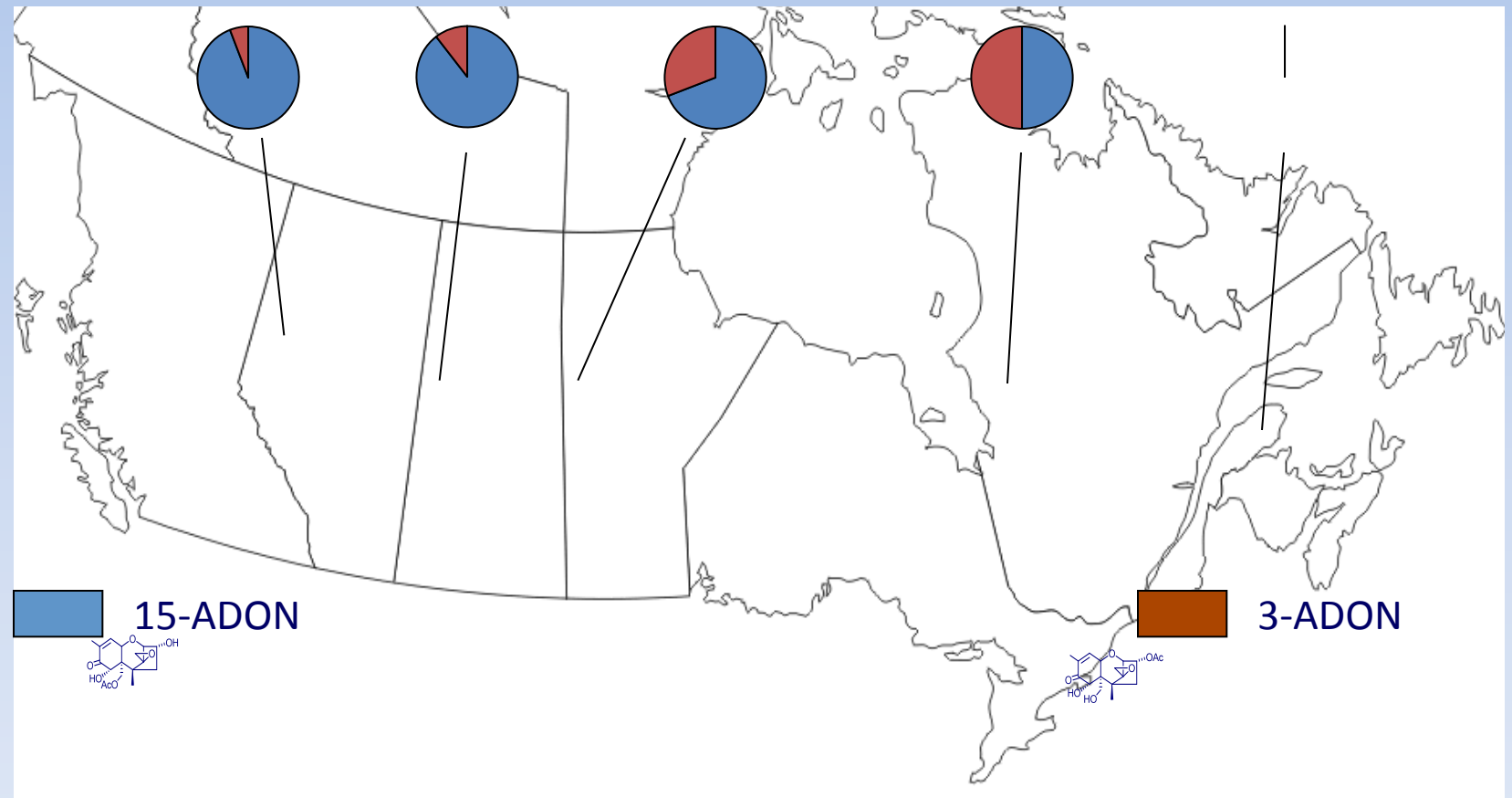
NIV

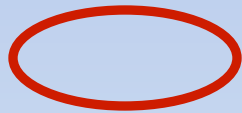
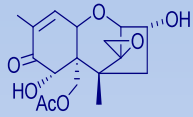


Mycotoxins produced by *Fusarium* in laboratory culture

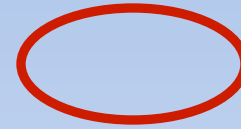
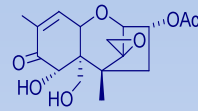


Emerging populations of *F. graminearum* 3ADON chemotypes

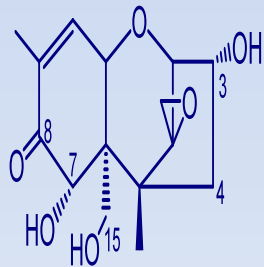




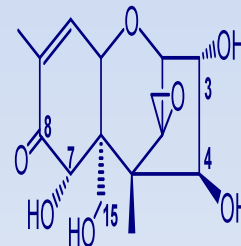
15-ADON



3-ADON

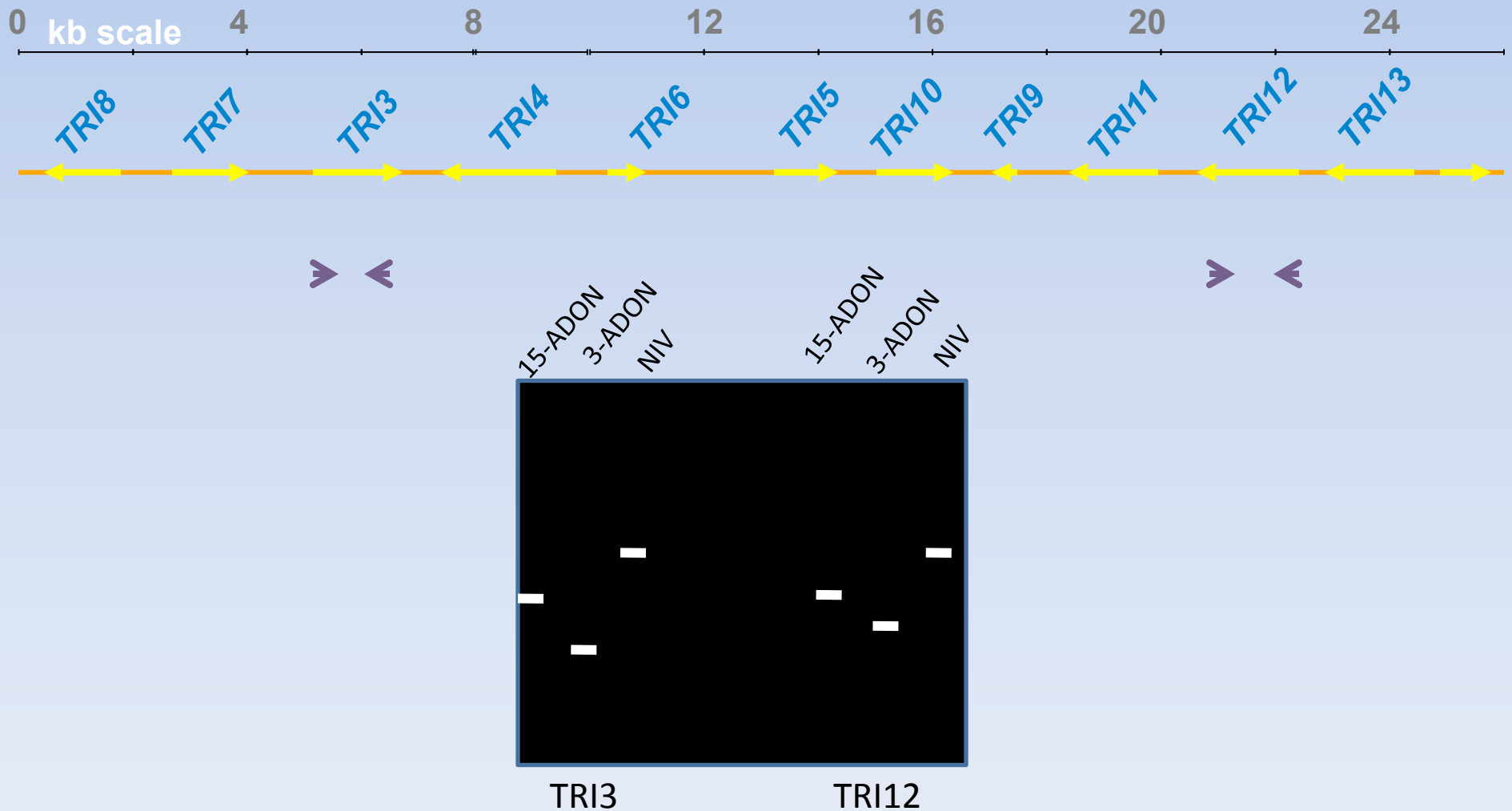


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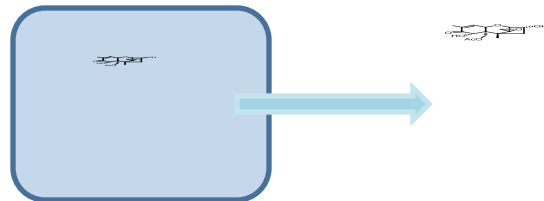


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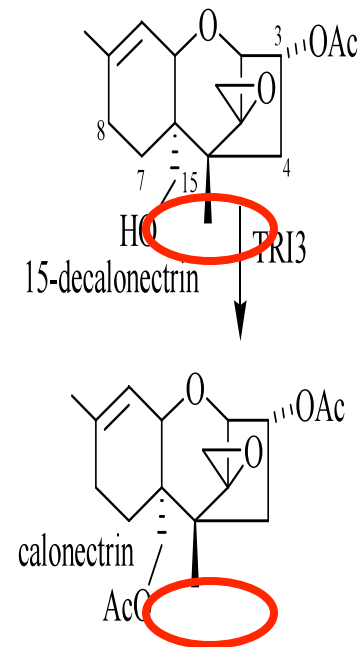
Chemotyping: using PCR analysis to correlate with chemistry



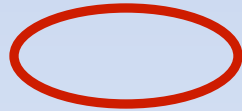
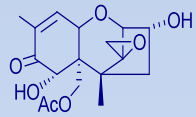
TRI12 = trichothecene efflux pump



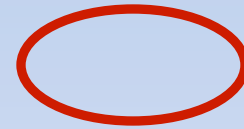
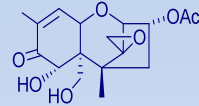
TRI3 = C15 acetyltransferase



Is *TR13* a determining factor?

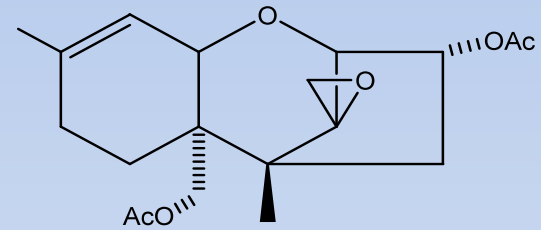
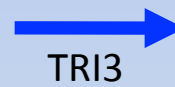
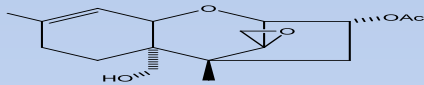


15-ADON



3-ADON

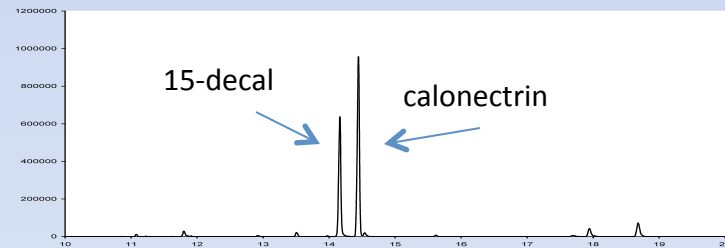
Yeast Feeding Experiments to assess TRI3 activity



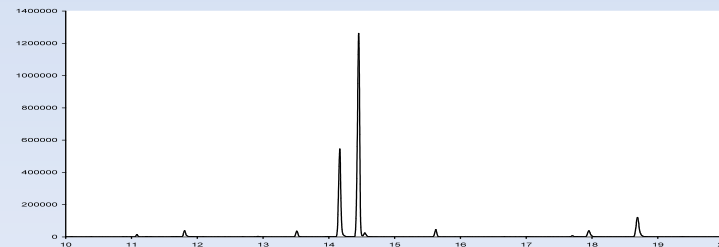
15-decal

Calonectrin

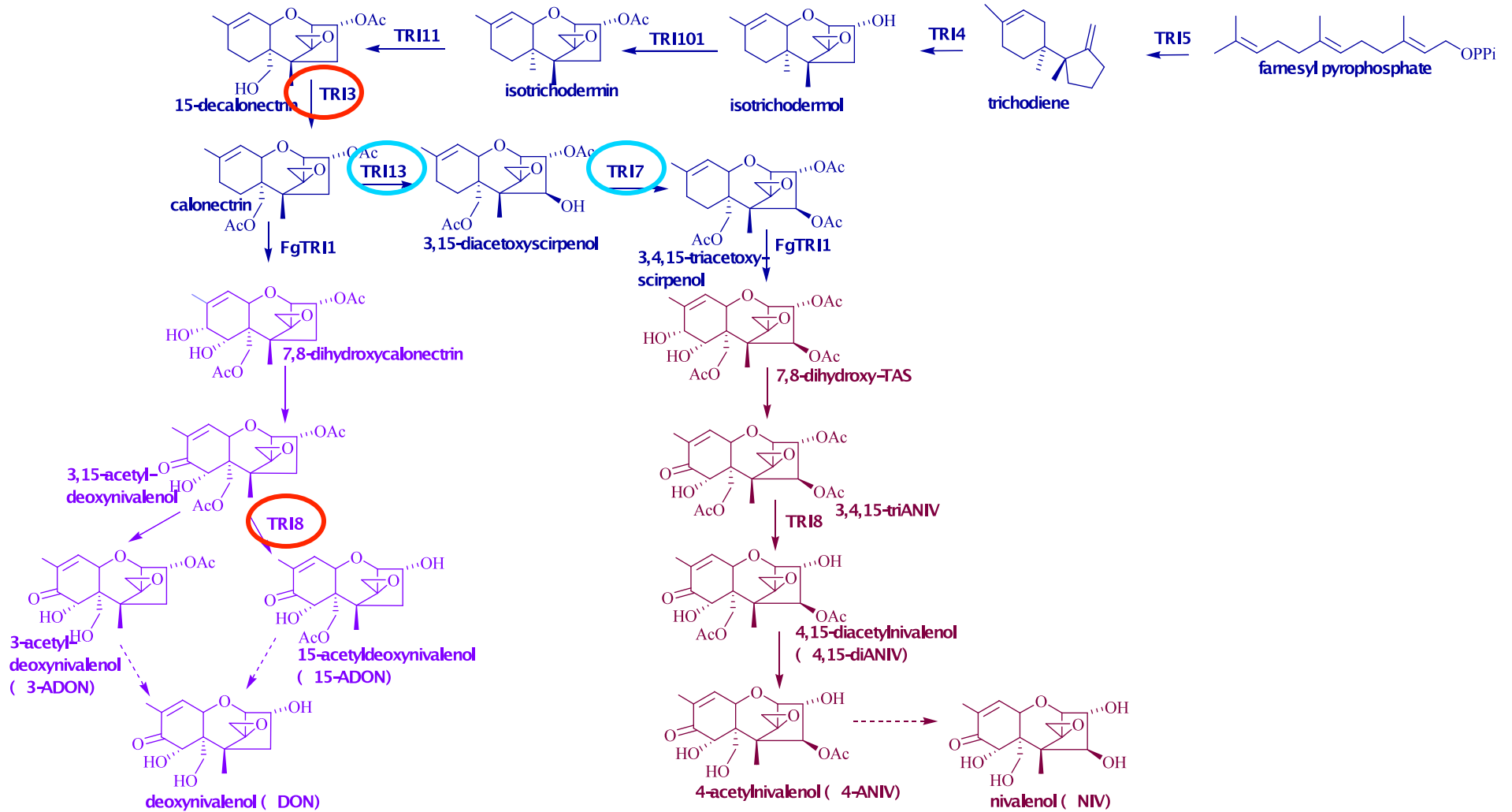
15-ADON TRI3 + 15-decal



3-ADON TRI3 + 15-decal



Trichothecene pathways



Nucleophilic
elbow



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3-ADON 1 ATGGTTCTCG ATCGTTTGTT GTTTCTTCTG AGCTTGTTGGC TGGGCTTTGT TGGCGCCACT
15-ADON 1 ATGGCTCTCG ATCGTTTGTT GTTTCTTCTG GGCTTCTGGT TGGGCTTGGT TGGCGCCGCT
61 CAAGCTGCGT TATC...AGA ACCTATTCCG CCTAGCAAGG ACCCGTGGTA TACTGCACCT
61 CAAGCTGCGT TGTCTGTCAGA ACCGCTTCCG CCCAGCAAAG ACCCGTGGTA TACTGCACCT
118 CCTGGATTTG AGAATGCCGA GCCTGGAAGT GTCTTCAGAG TCGCGCCTGC GCCTGGAAAC
121 CCTGGATTTG AGAATACCGA GCCTGGAACA GTCTTGAGAG TCGGACCTGC GCCTGGAAAC
178 TTAACCAGCG TCATCGGCAA TTGCTCTGCT TCTTACAACA TTCTCTATCG TACGACCGAC
181 TTGACCAGCG TCACCAGCAA TTGCTCTGCT TCTTACAACA TCCTCTACCG TACGACCGAT
238 AGCCATTCA AGCCTACTTG GGCAGTCACC ACTCTTTTGG TCCCCAACT GGGTCCTGAA
241 AGTCATTCA AGCCCGCTTG GGCCGTCACC ACTCTTTTGA TCCCCGAGCT GGGTCCTGAA
298 AGTCTTGCGC AGCAGAAGTA CCAACAAAGC GCGCTTTGT CATCCAAGT GCCTTATGAC
301 AGTCTTGCGC ACCAAAAGTA CCAACAAAGC GCGTAATGT CAATCCAAGT TGCCATGAC
358 TCCCCGATG TCGATGCCAG CCCTAGTAAC GCTATGTACG ACGCGAGCGA TTTTTTTTCG
361 TCTCCCGACG TCGATGCCAG CCCTAGTAAC ACTATGTACA CCGCGAGCAA TTTTCTTCC
418 AATTACTATG GAGCTGCTCT AGGTGAAGGT ATCTTCGTTT CTGTTCCGGA TTATGAGGGA
421 ATCTACTATG AAGCTGCTCT AGGTCAAGGG CTCTTCGTTT CTGTTCCAGA TTACGAGGGA
478 CCCTTGCGTG CTTTTACCGC TGGTCTCATC TCTGGATATG CTACTTTTGA CTCAATTCGC
481 CCCTTGCGTG CTTTTAGCGC TGGTGTATC TCTGGGTACG CTACTTTTGA CTCAATTCGC
538 GCTATTCTGT CGCTCGGTCT TGGCTTCAAC ACGATCGACA CGCCCAGTGT CGCCCTTTGG
541 GCTGTTCTAT CGCTGGTCT GGGCTTCAAC ATGACCAACA CGCCCAGTGT TGCCCTTTGG
598 GGCTATTCTG GCGGTGCGTT TGCCACGGAA TGGGCATCGG AACTTGCAGT ACAGTATGCA
601 GGCTACTCTG GCGGTGCGTT TGCCACGGAA TGGGCATCAG AACTTGCAGT ACAGTATGCA
658 CCCGAGCTTG TAGCAGGACC CGTTATTGGT GCTGTGATGG GAGCCCCGTT GCCCAATATC
661 CCAGAGCTGA TAACAGGACC CGTTATCGGT GCTGCGTTGG GAGCCCCGTT GCCCAATATC
718 ACTTCATGCA TCGCGATGT CAATGGAGGA CTAAGTCCG GTCTGGTGGT AAACATCTG
721 ACTTCACTCC TGTACGATGT CAATGGCAA CCTGGAGCCG GTCTGGTGGT AAATATGTTG
778 CTGGGTCTTA CGGGTCAATA TCCCGATGTC CGCAAGCACC TTGTATCCAA ACTCAACGAT
781 TTGGGTCTTA CAAGCCAATA TCCTGACGTC CGCAAGTACC TTATATCCAA ACTCAACGAT
838 GATGGCCAGT ACAACAAGGC TGGCTTTCTC GCTGCTGAGG GTTTTACTAT CAGCGAAGCG
841 GATGGCCAGT ACAACAAGAC TGGCTTTCTC GCTGCTGAGG GTTTTACCAT CAACGAAGCT
898 CTAGCACCT TTTCTGGCAA C...ATCAAC AAGTACTTCC AGAAGGGGAC TGATATTCTC
901 GGCGTCGCCT TTTATGGAAT TGATATCAAC AAGTACTTCC AGAAGGGGAC TGACATTCTC
955 AGCGACCCGA AGATTACAGC TCTTATCAAT CGGGAGGGCG TTTTGGGATA TCACGGCACC
961 AGCGACCCGA AGATTGTAGC TCTTCTTAAC CAAGAGGGCC TTCTGGGATA TAACGGTACC
1015 CCCAGATGGC CCATGTTTAT TTACCAAGCT ATCTCTGACG AGGTCACGCC GATTGCTGCT
1021 CCCAGATGGC CTTTGTTTAT TTACCAAGCT ATCCACGACG AGGTCACACC TATAGCTGAT
1075 ACCGATGCCG TAGTCGAGAG ATATTGTTCA GTGGGGGCTG ACGTTCACTT TGAAAGGAAC
1081 ACCGATGCCG TAGTCAACAG ATATTGTGCC GTGGGGGCTG ACATTCACCT CGAAAGGAAT
1135 ACTCTCGGCT CGCACGACGA GGAAGCCGGC AATAGCTATG ACGCGGCTTT CAGTGGCTT
1141 ACTATCGGCG GGCACACCA GGAAGCCGAC AATAGCTATG ACGCGGCTTT TCAGTGGCTT
1195 TTGGACATCT TCTCCGGCCA GCGTGACACA AAGGTTGTG TTATCAAGGA CGTGACGAGA
1201 TTGGACATCT TCTCCGGCCA GCGTGACACA AAGGTTGTG TTATCAAGGA ACGTACGAGA
1255 GACGTGGACG TGACGGGAGA CGTGACGGGA GACGTGACGA GAGACGTAAC ACGAGAGTTG
1261 AACATTACCG GAAGTGTCTT ACAAACACGT GAGAATGTTT AAAAGAGTGG AGTTGACTTC
1315 TGA
1321 TGGAGATCTG CCTGGTAA
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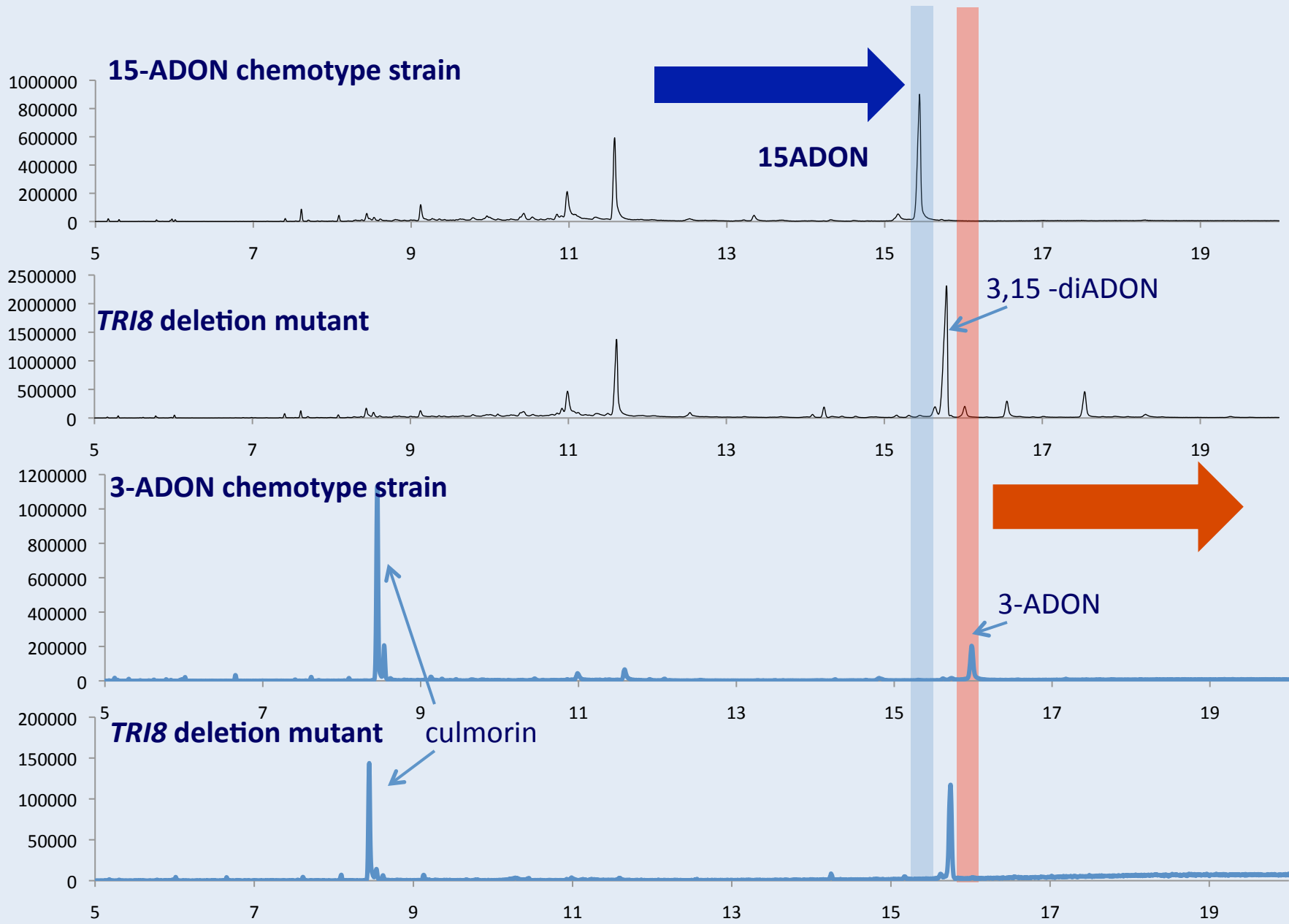
TR18 sequence comparisons

chemotype	# nucleotides	% aa ID	TR16 bs
3-ADON	1299	78	none
15-ADON	1338	85	-525 nt
NIV	1335		-525 nt

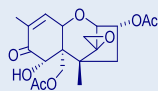
3-ADON 1255 **GACGTGGACG TGACGGGAGA CGTGACGGGA GACGTGACGA GAGACGTAAC ACGAGAGTTG**
 15-ADON 1261 **AACATTACCG GAAGTGCCT ACAAACACGT GAGAATGTC AAAAGAGTGG AGTTGACTTC**

 1315 **TGA**
 1321 **TGGAGATCTG CCTGGTAA**

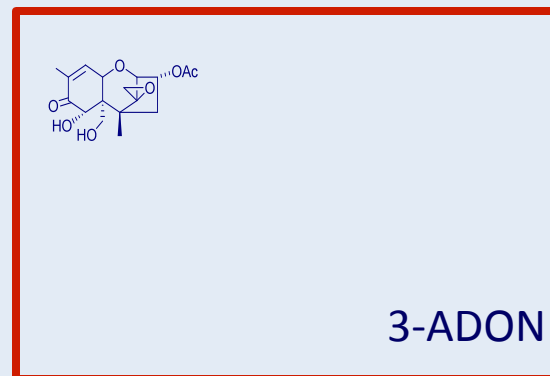
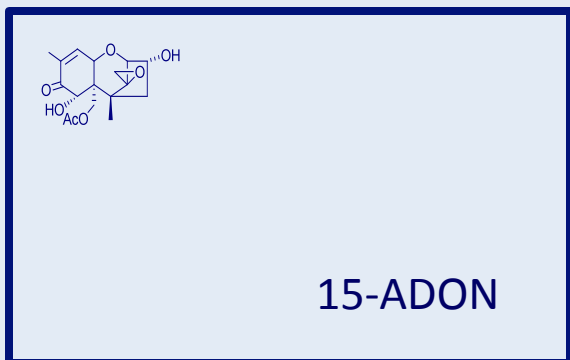
TRI8 Wild Type and deletion mutants



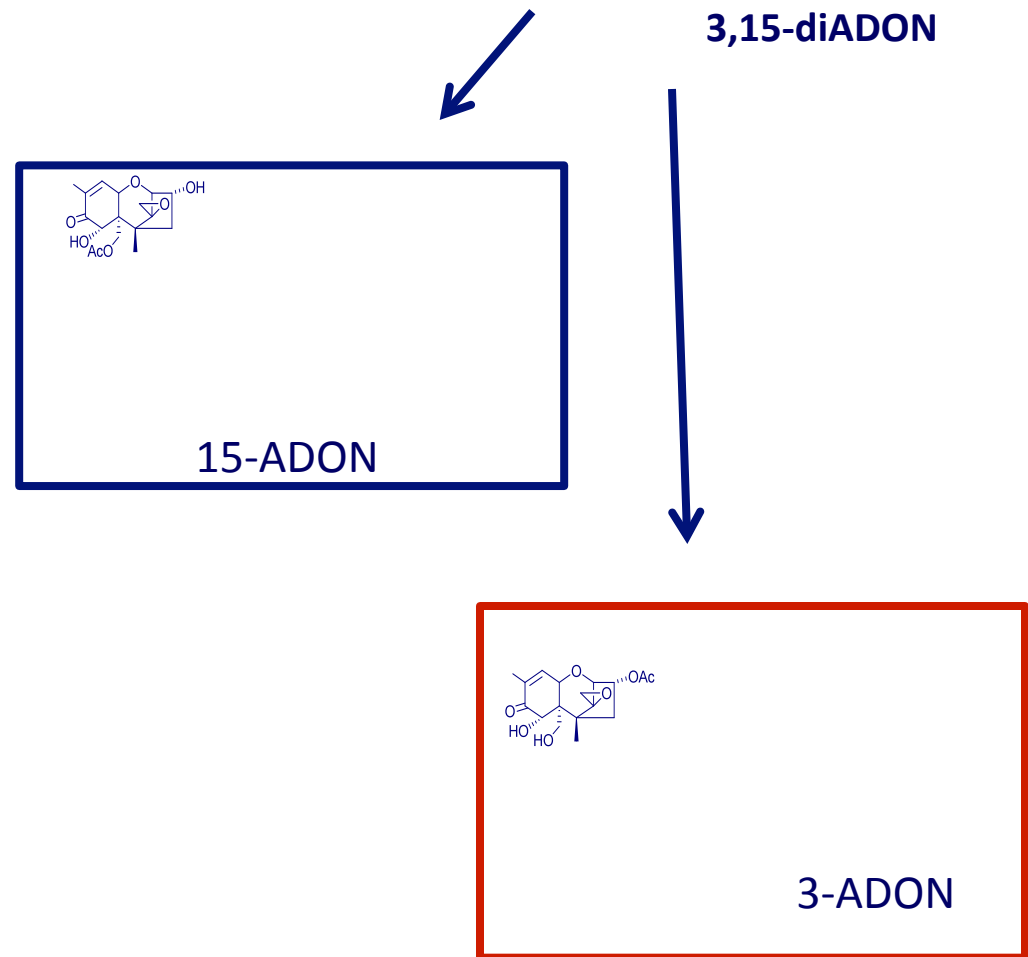
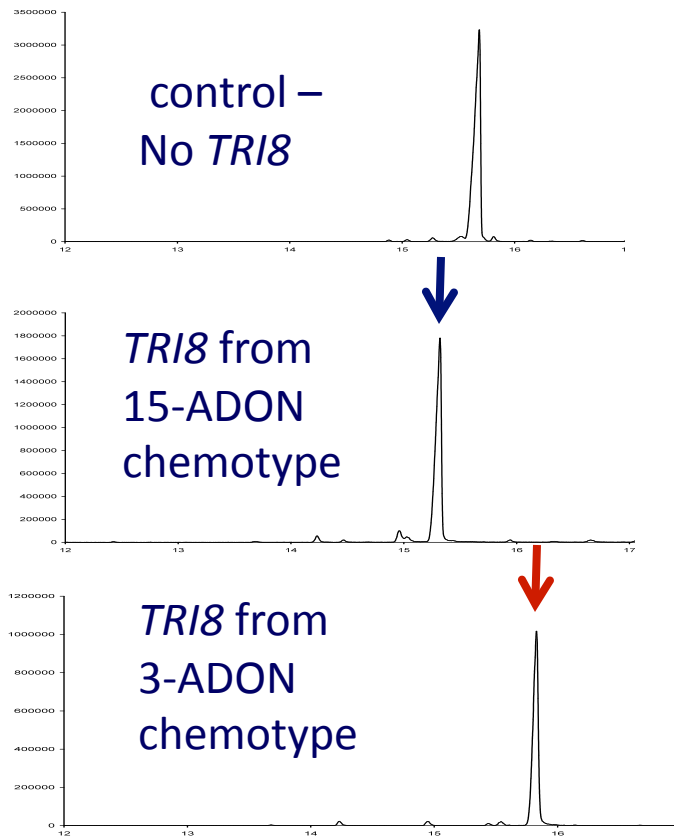
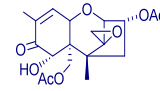
Deletion studies confirmed that 3,15-diADON was precursor for TRI8 from both 3-ADON and 15-ADON producers



3,15-diADON

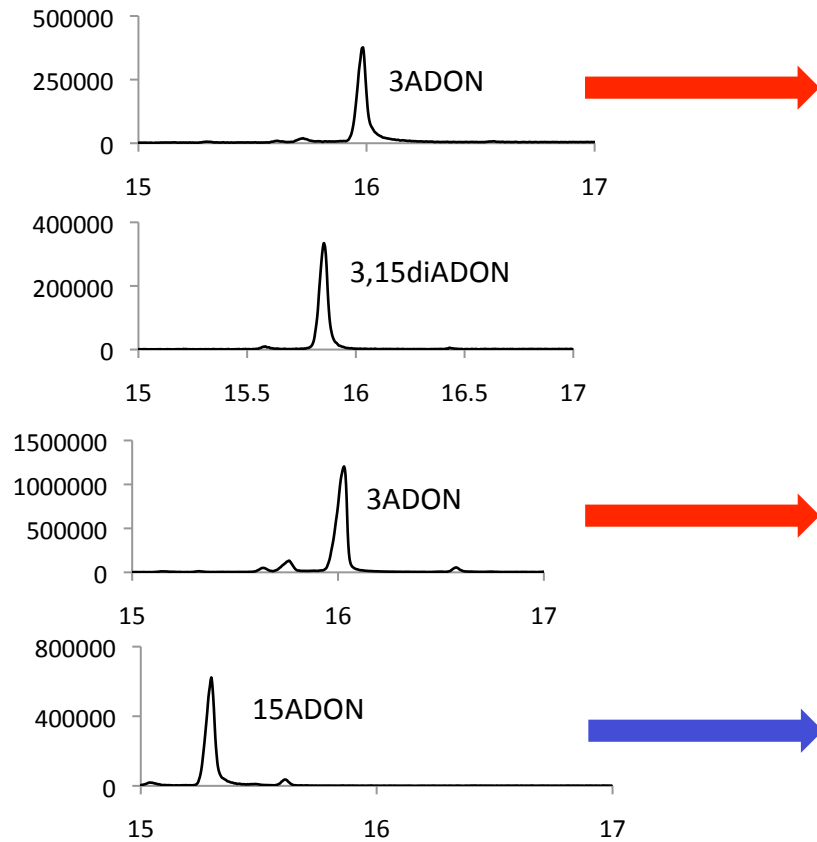


Expression of *TRI8* in yeast – feeding 3,15-diADON

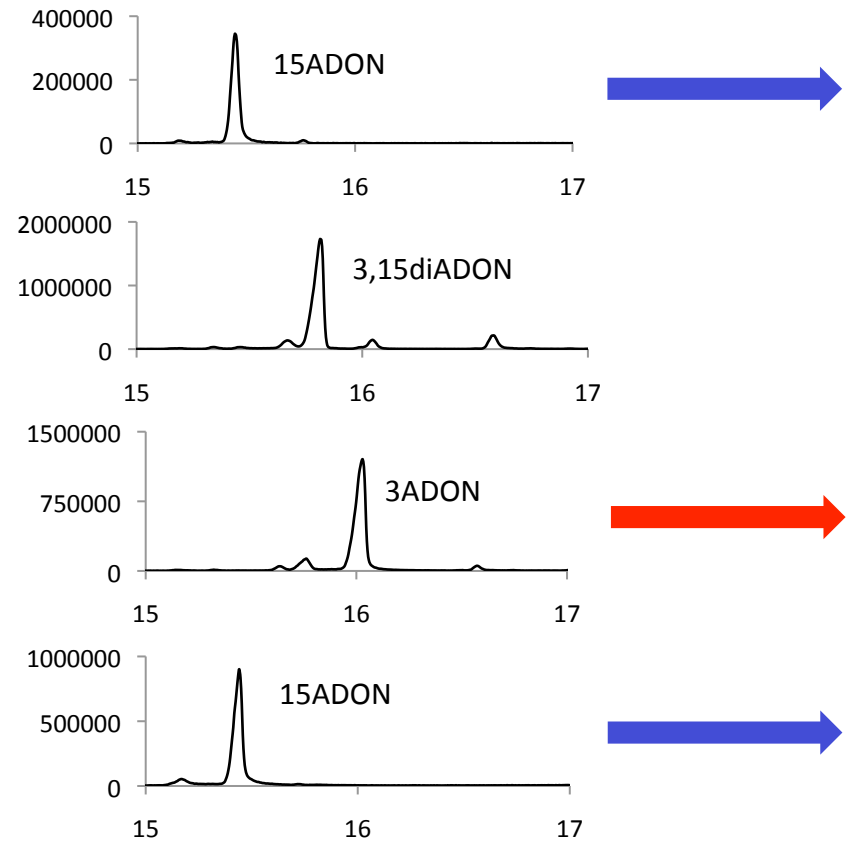


Adding back intact *TRI8* to deletion mutants

3-ADON Background

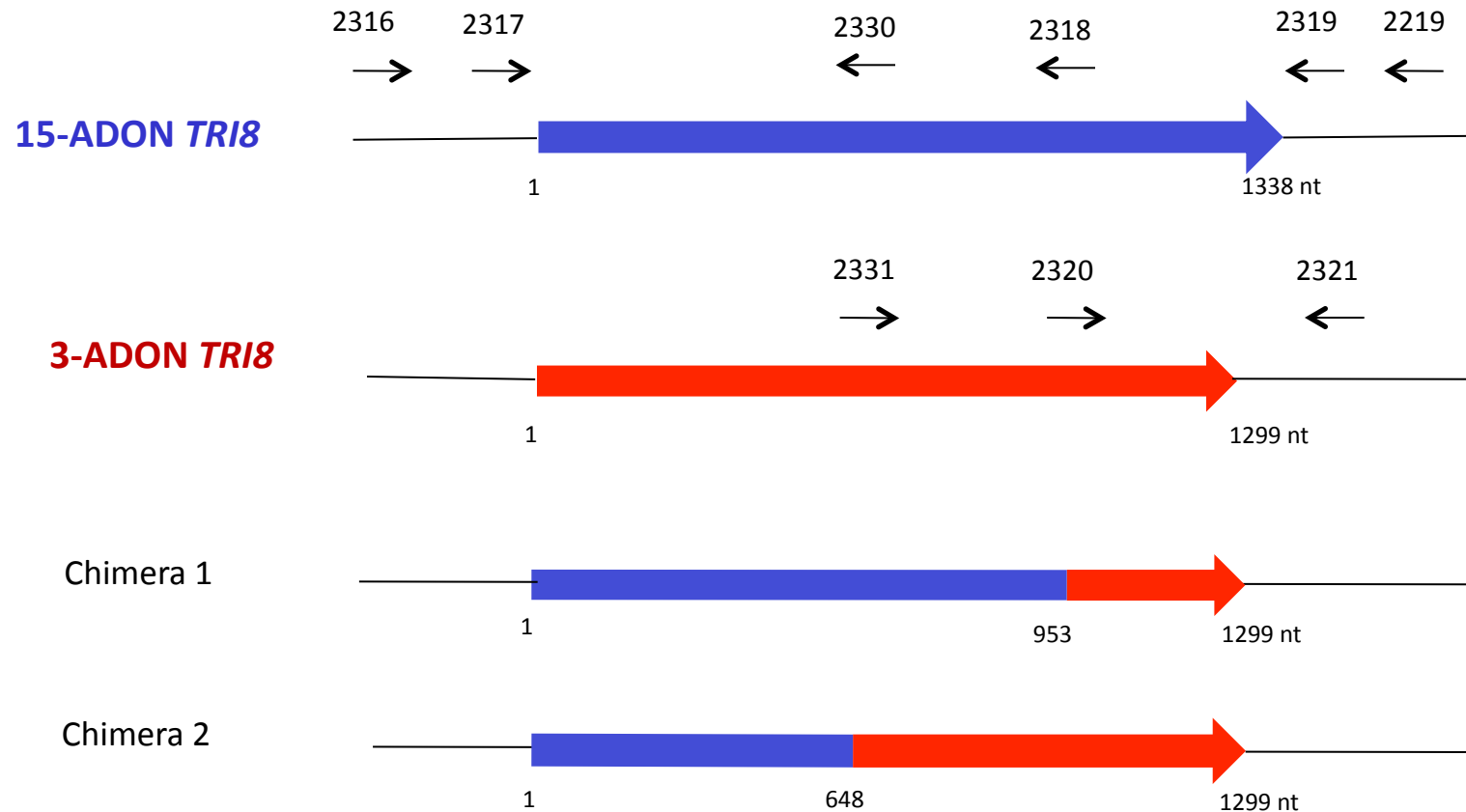


15-ADON Background



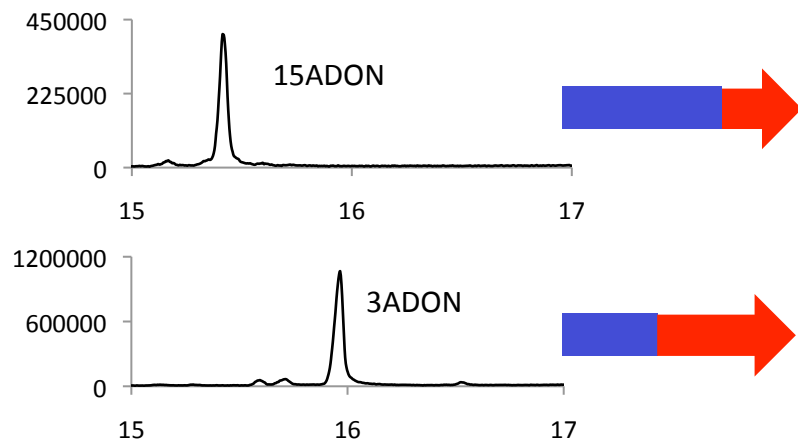
Conclusion: *TRI8* has different function

Determination of *TRI8* region responsible for different esterase activity

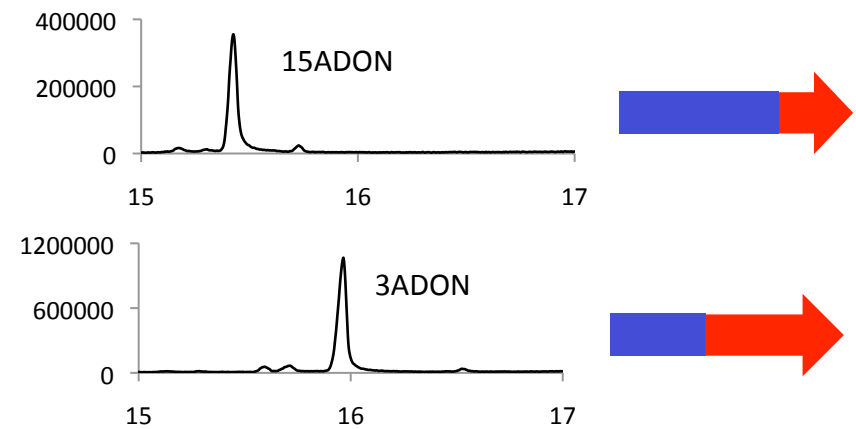


Phenotype of chimeras

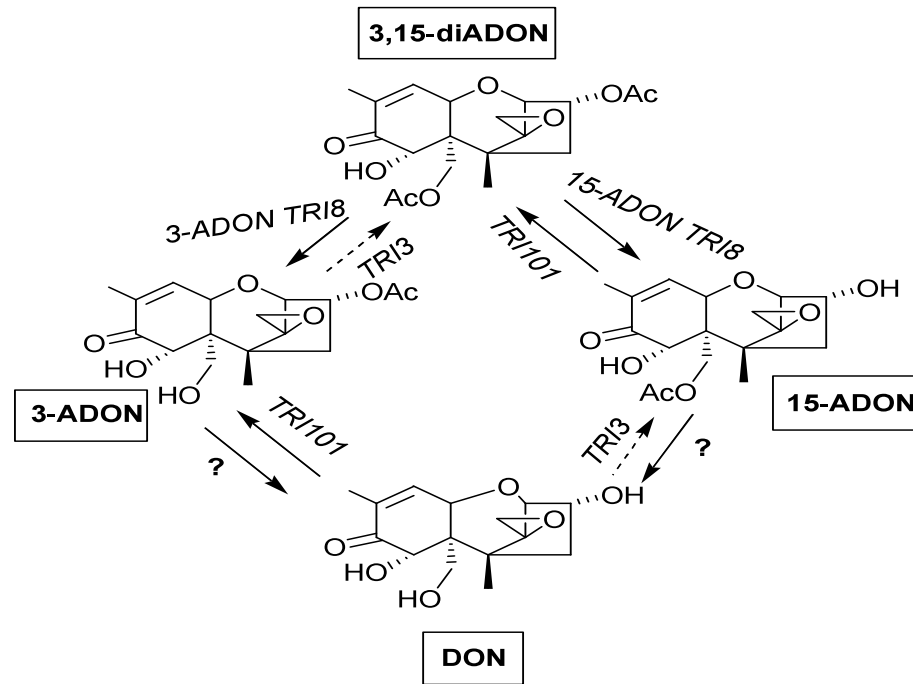
3-ADON background



15-ADON background



Proposed pathway to explain small amounts (*in planta*) of 3-ADON in a 15-ADON producer



Conclusions

- *TR18* from a 3-ADON producer encodes C-15 esterase
- *TR18* from a 15-ADON producer encodes C-3 esterase
- Middle region of *TR18* defines whether a strain produces 3-ADON or 15-ADON
- Small sequence differences can impact functionality



Acknowledgments



Jennifer
Teresi



Stephanie
Folmar



Kim
MacDonald

Visit Poster # 26 at the PBG Poster Session this afternoon.