

Combining Different Genetics-Based TSNA Reduction Strategies in Burley Tobaccos

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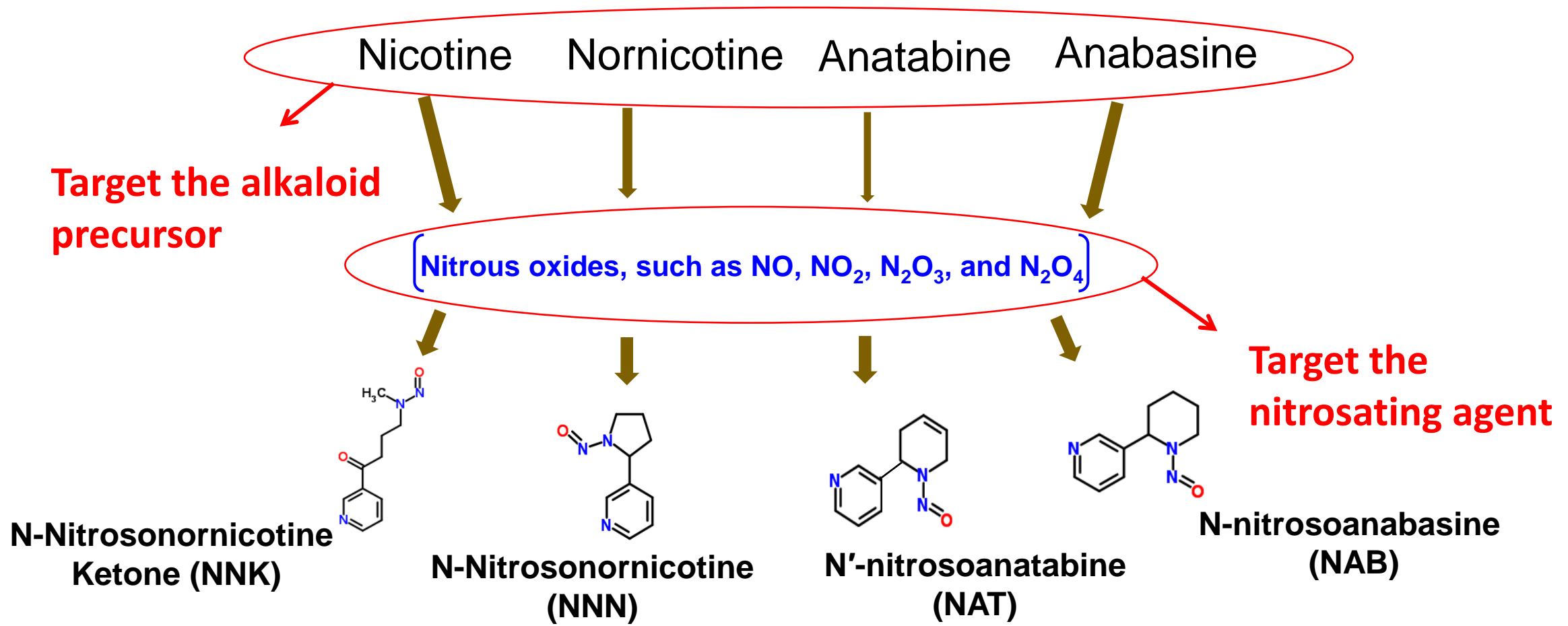
Simon Goepfert and Lucien Bovet

Philip Morris International, Neuchâtel, Switzerland

January 17, 2024

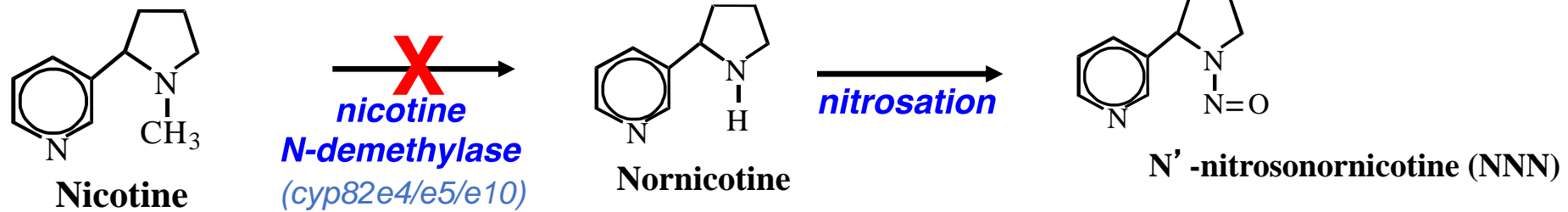
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Philip Morris International.*

Reducing Tobacco-Specific Nitrosamines (TSNA) via Genetic Modification



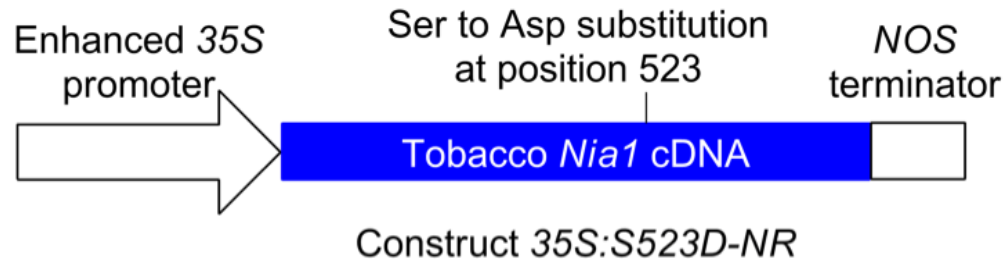
Effective Genetic-Based Strategies to Reduce TSNAs

A) Inactivation of Nicotine Demethylase Genes (Zyvert™)

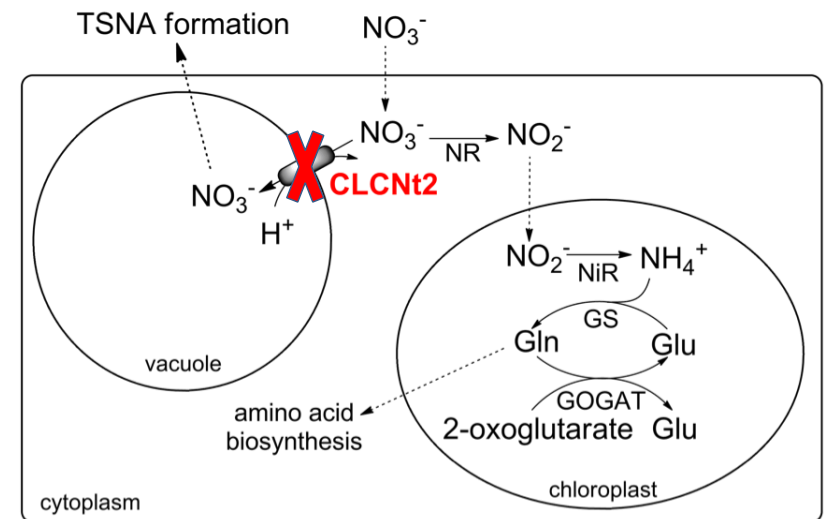


B) Reducing the Levels of Free Nitrate

1. Constitutive activation of a deregulated nitrate reductase gene



2. Inactivation of putative nitrate transporters (*CLC-Nt2* genes)



Inactivation of Nicotine Demethylase Genes (Zyvert™)

Conversion of nicotine to nornicotine in *Nicotiana tabacum* is mediated by CYP82E4, a cytochrome P450 monooxygenase

Balazs Siminszky^{*†}, Lily Gavilano^{**}, Steven W. Bowen^{*§}, and Ralph E. Dewey[§]

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RNA interference (RNAi)-induced suppression of nicotine demethylase activity reduces levels of a key carcinogen in cured tobacco leaves

Ramsey S. Lewis^{1,*}, Anne M. Jack², Jerry W. Morris³, Vincent J. M. Robert^{3,†}, Lily B. Gavilano², Balazs Siminszky², Lowell P. Bush², Alec J. Hayes³ and Ralph E. Dewey¹

Phytochemistry 71 (2010) 1988–1998

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Phytochemistry

journal homepage: www.elsevier.com/locate/phytochem

Three nicotine demethylase genes mediate nornicotine biosynthesis in *Nicotiana tabacum* L.: Functional characterization of the *CYP82E10* gene

Ramsey S. Lewis, Steven W. Bowen, Matthew R. Keogh¹, Ralph E. Dewey^{*}

Summary of Results

- Can reduce both nornicotine and NNN accumulation by ~80%
- Appears to have no significant impact on yield or flowering time

Constitutive Expression of a Deregulated Nitrate Reductase Gene

Plant Biotechnology
Journal

aab
Association of Applied Biologists

SEB
Society for
Experimental Biology

Plant Biotechnology Journal (2016) 14, pp. 1500–1510

doi: 10.1111/pbi.12510

Expression of a constitutively active nitrate reductase variant in tobacco reduces tobacco-specific nitrosamine accumulation in cured leaves and cigarette smoke

Jianli Lu¹, Leichen Zhang¹, Ramsey S. Lewis¹, Lucien Bovet², Simon Goepfert², Anne M. Jack³, James D. Crutchfield⁴, Huihua Ji³ and Ralph E. Dewey^{1,*}

scientific reports

Scientific Reports | (2021) 11:4222

| <https://doi.org/10.1038/s41598-021-83797-7>

nature portfolio

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OPEN Constitutive activation of nitrate reductase in tobacco alters flowering time and plant biomass

Jianli Lu^{1,3}, Niharika N. Chandrakanth¹, Ramsey S. Lewis¹, Karen Andres¹, Lucien Bovet², Simon Goepfert² & Ralph E. Dewey^{1,*}

Summary of Results

- Can reduce free nitrates levels by ~95% in the cured leaf/cut filler
- Can reduce total TSNA's (including NNK) by ~52% in the cut filler and ~76% in the cigarette smoke
- Can accelerate flowering by 7 to 10 days
- Can lead to reductions in biomass by 30 to 50%
- *The reductions in biomass can be overcome while retaining the low TSNA trait by crossing 35S:S523D-NR plants with late flowering burley cultivars*

Inactivation of Putative Nitrate Transporter Genes (*CLC-Nt2*)

CLCNt2 Mediates Nitrate Content in Tobacco Leaf, Impacting the Production of Tobacco-Specific Nitrosamines in Cured Leaves

Lucien Bovet^{1*}, Prisca Campanoni¹, Jian Lu², Aurore Hilfiker^{1†}, Samuel Kleinhans¹, H el ene Laparra¹, Joanne Schwaar¹, Ramsey S. Lewis², Yuki Matsuba², Hong Ma², Ralph E. Dewey² and Simon Goepfert¹

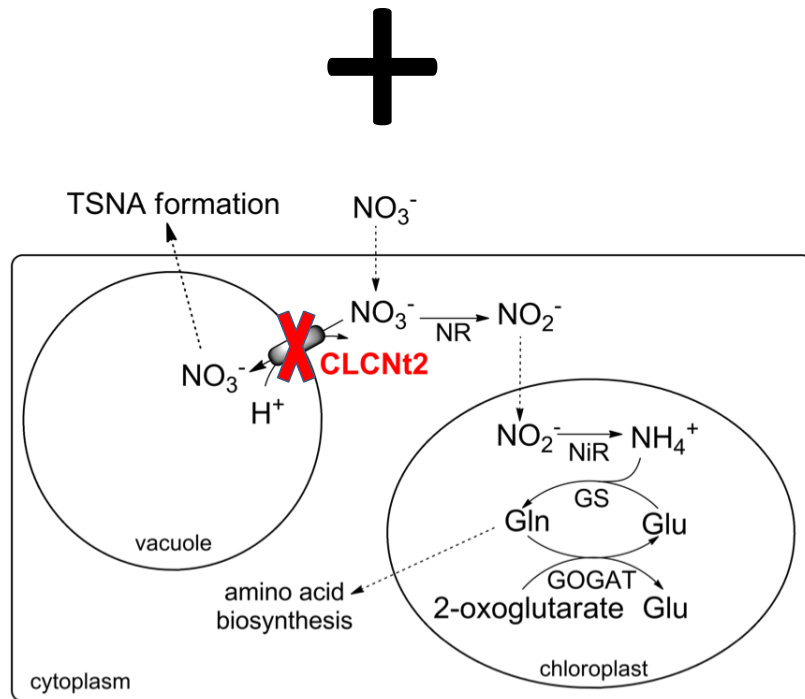
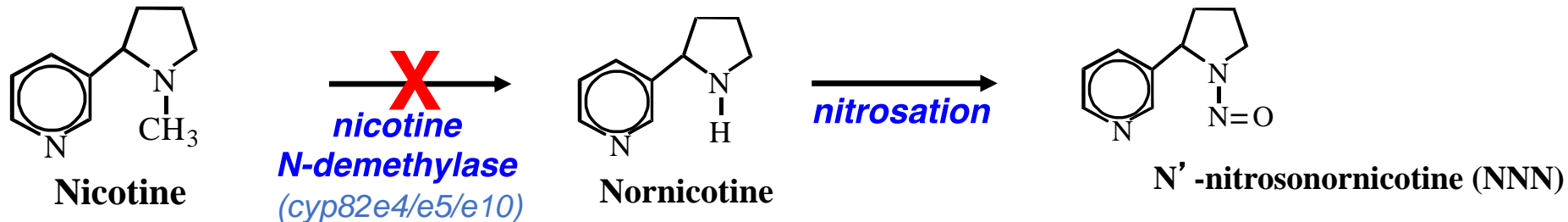
¹FMI R&D, Philip Morris Products S.A., Neuchatel, Switzerland, ²Department of Crop and Soil Sciences, North Carolina State University, Raleigh, NC, United States

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Summary of Results

- Nitrate concentration reduced by ~65%
- Total TSNA in cured leaf lamina reduced by ~40% and in mainstream cigarette smoke by ~52%
- Inactivation of *CLC-Nt2* genes was not associated with reductions in biomass but was associated with an approximate 10-day delay in average flowering time

Combining Zyvert™ Trait with *CLC-Nt2* Knockout Mutations



2022 Field Study Conducted in Clayton, NC



Line

TN90 LC

CLC-Nt2#8 5N5

TN90 Zyvert

ZYV/nt2-2

ZYV/nt2-3

Genotype

Control

clc-nt2 mutations alone

cyp82e4/e5/e10 mutations alone

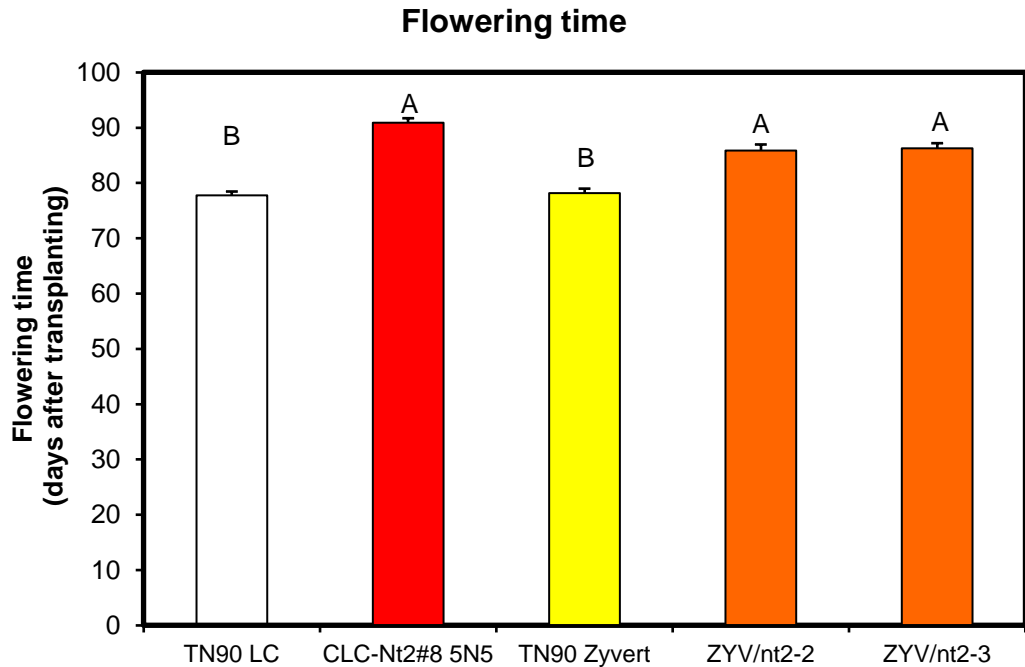
clc-nt2 + cyp82e4/e5/e10

clc-nt2 + cyp82e4/e5/e10

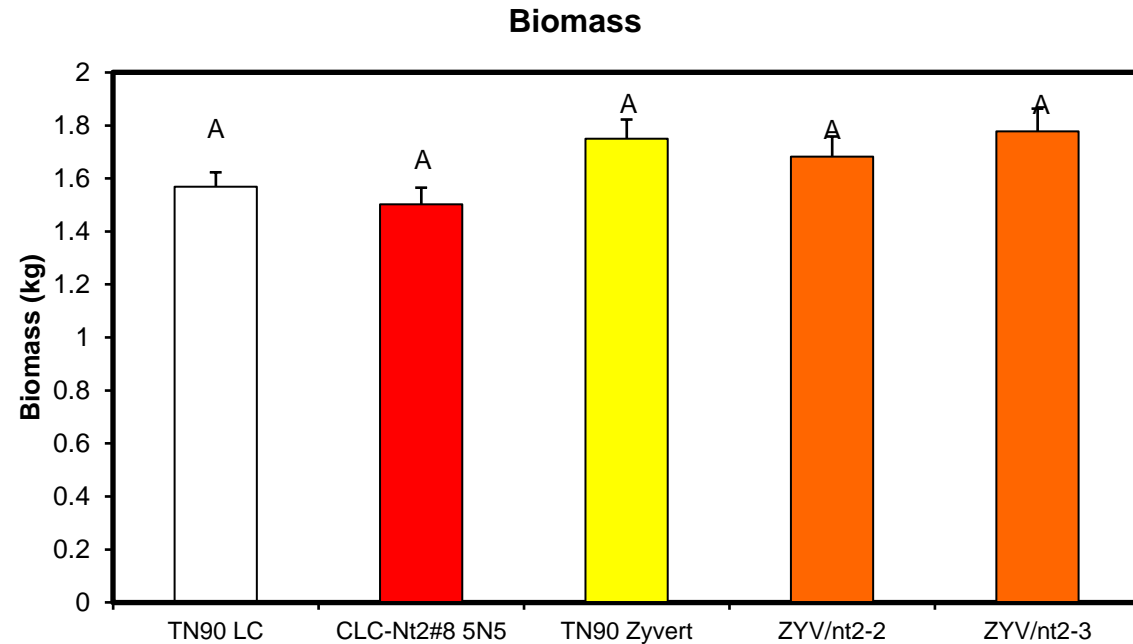
Traits Measured

- Flowering Time
- Aerial Biomass at Harvest
- Nitrate (after air-cure)
- Alkaloids (after air-cure)
- TSNA (after air-cure)

Flowering Time and Biomass Results

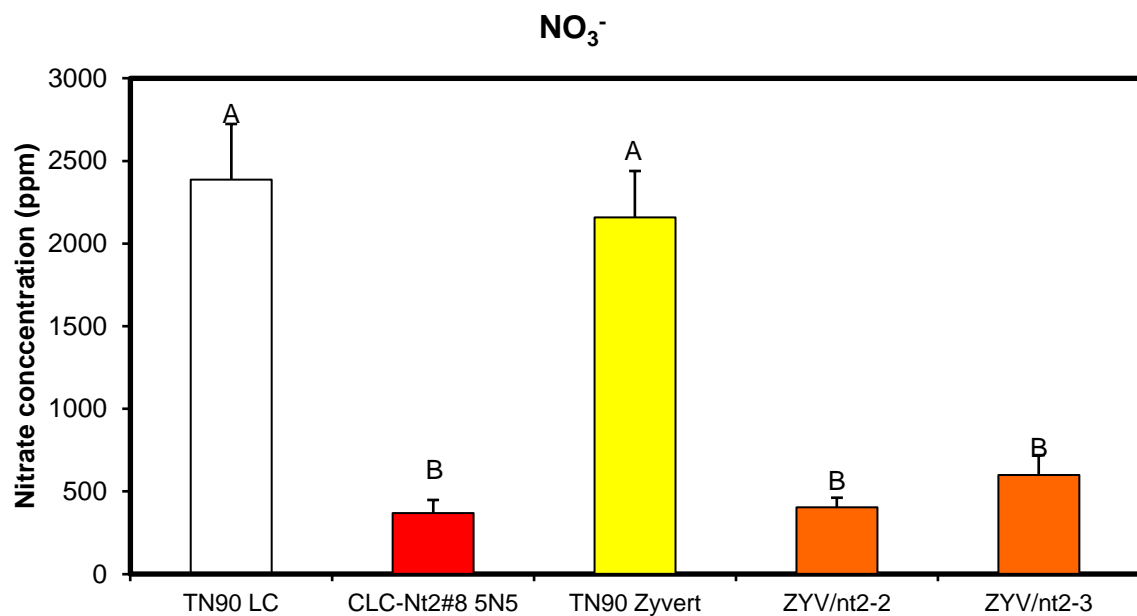


*On average, lines possessing the *clc-nt2* mutations flowered 8 – 12 days later than the TN90 LC and TN90 Zyvert lines*

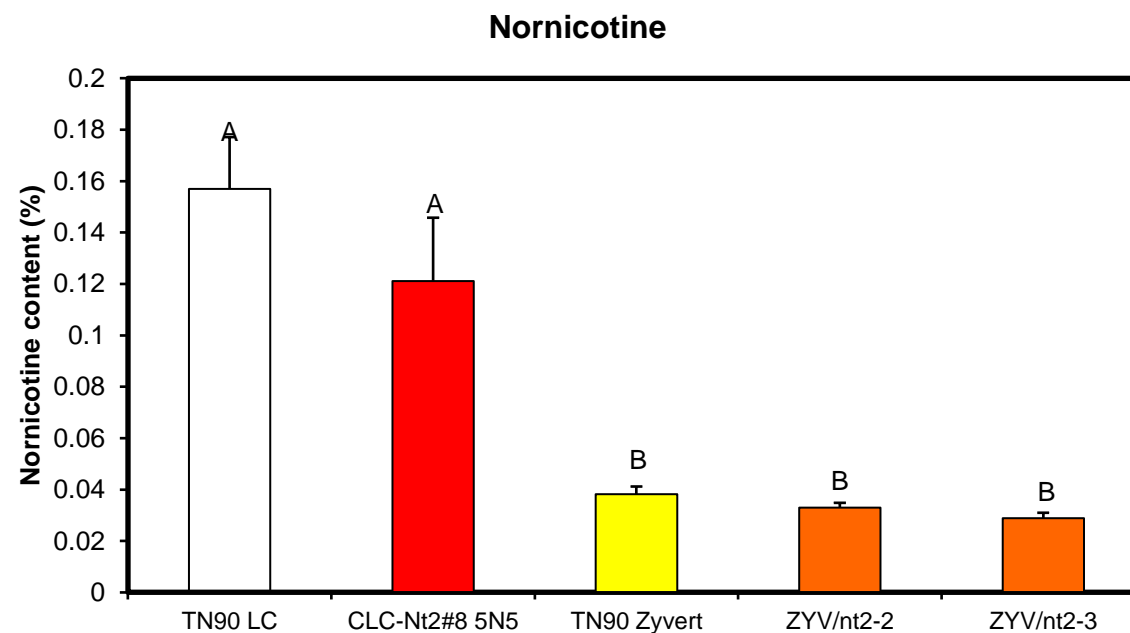


There were no significant differences in plant biomass observed among the lines

Nitrate and Nornicotine Results

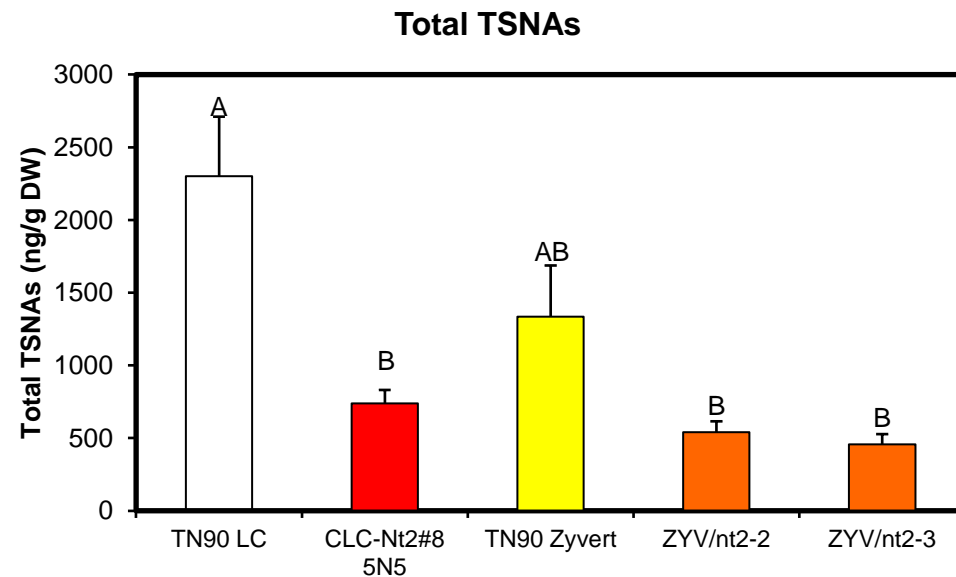
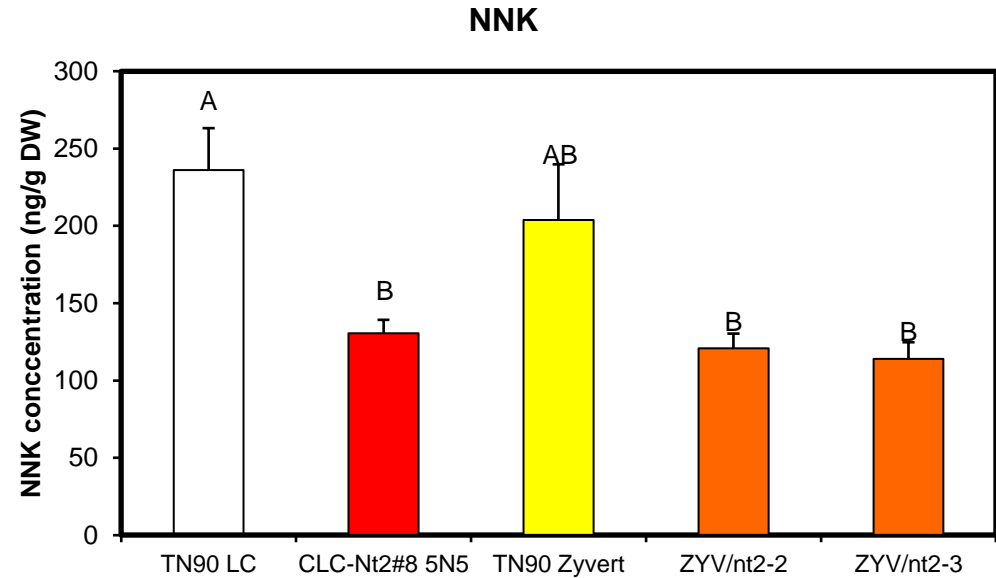
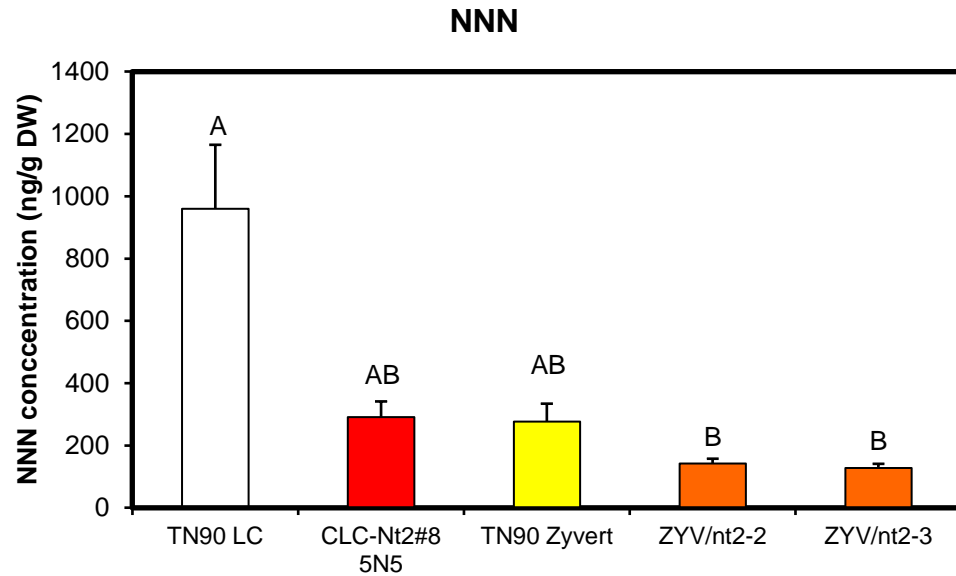


*Lines possessing the *clc-nt2* mutations accumulated 72 – 84% less nitrate in the cured leaf than the lines TN90 LC or TN90 Zyvert*



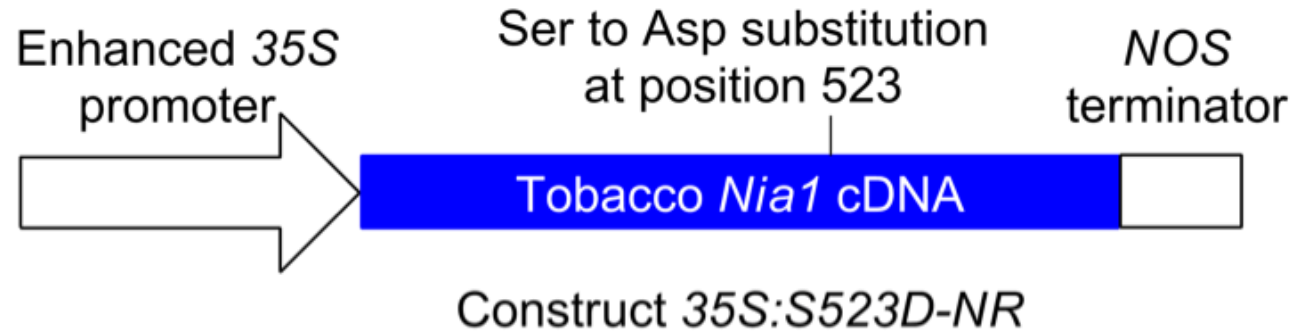
*Lines possessing the *Zyvert* mutations accumulated 69 – 82% less nornicotine than the lines without these mutations*

TSNAs

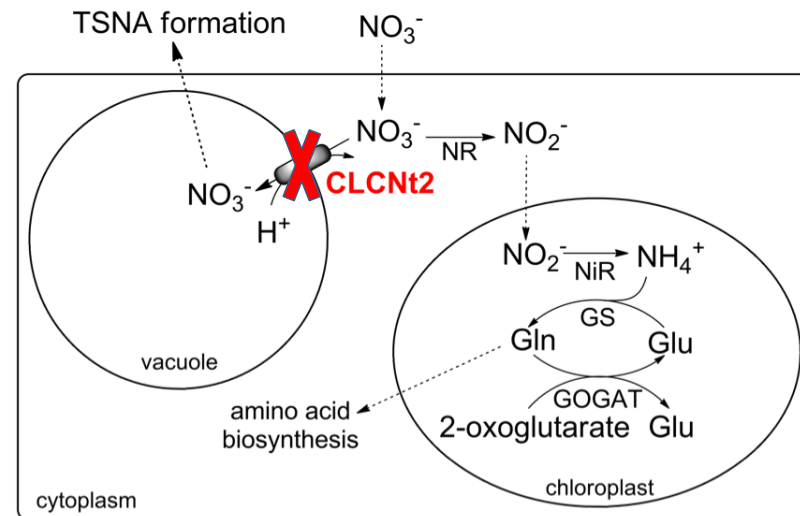


Due to the high costs of TSNA assays, only 10 plants per genotype were analyzed. The low sample size, combined with high inherent variability in TSNA formation, prevented some observations from achieving statistical significance, but the trends strongly suggest that combining the two technologies can reduce NNN levels below that achieved separately

Combining 35S:S523D-NR Transgene with *CLC-Nt2* Knockout Mutations



+



2022 Field Study Conducted in Clayton, NC

Line

Genotype

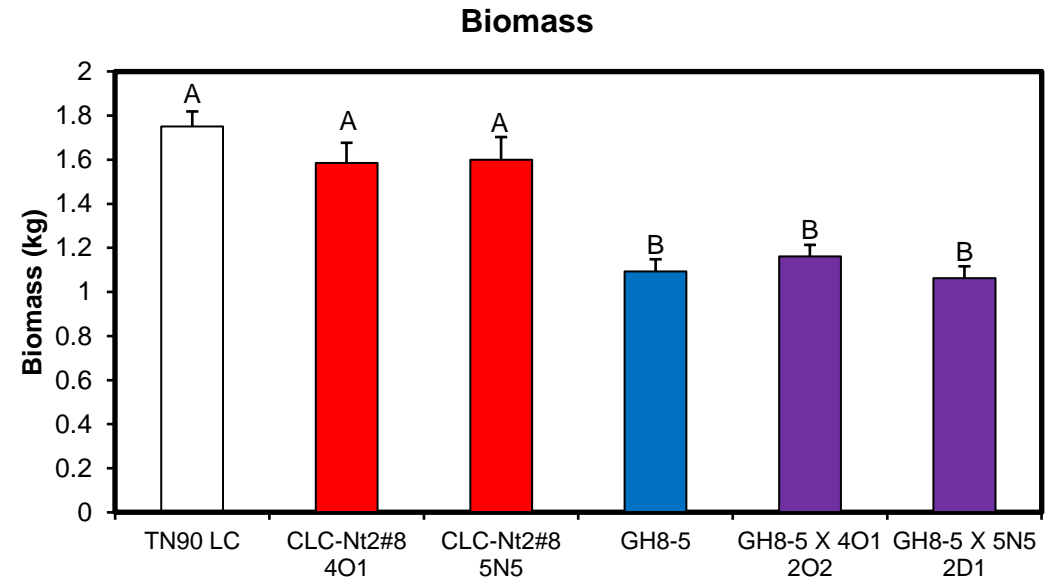
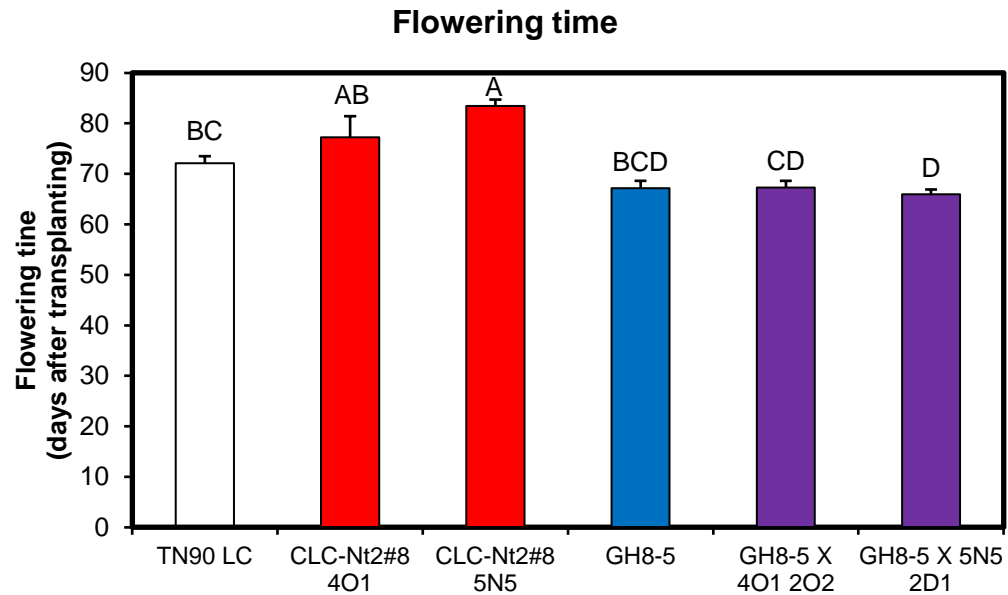
| | |
|----------------|-------------------------|
| TN90 LC | Control |
| CLC-Nt2#8 4O1 | clc-nt2 mutations alone |
| CLC-Nt2#8 5N5 | clc-nt2 mutations alone |
| GH8-5 | 35S:S523D-NR alone |
| GH8-5 4O1 2O2 | 35S:S523D-NR + clc-nt2 |
| GH8-5 5N5 2D1 | 35S:S523D-NR + clc-nt2 |
| GH8-5 5N5 2H3* | 35S:S523D-NR + clc-nt2 |

Traits Measured

- Flowering Time
- Aerial Biomass at Harvest
- Nitrate (after air-cure)
- Alkaloids (after air-cure)
- TSNA (after air-cure)

*This line converted (40% nornicotine)
so data from this line will be omitted

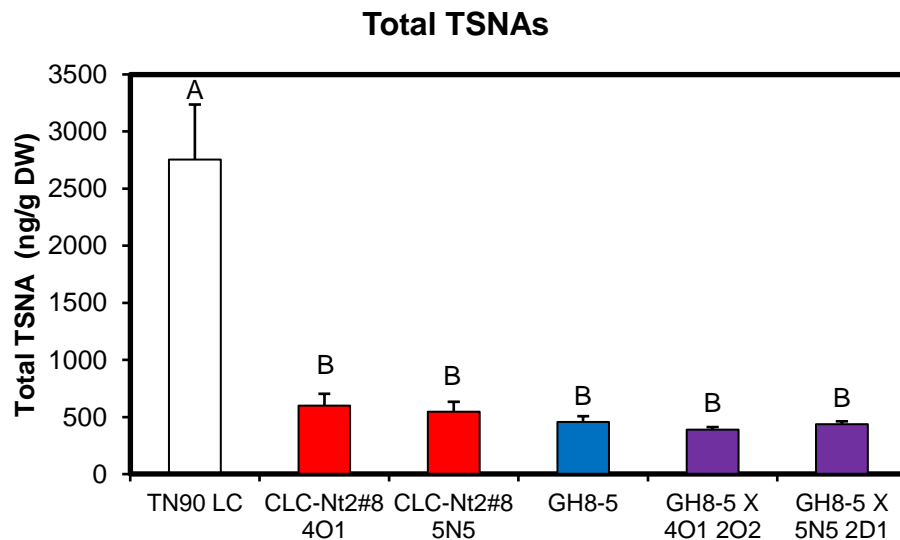
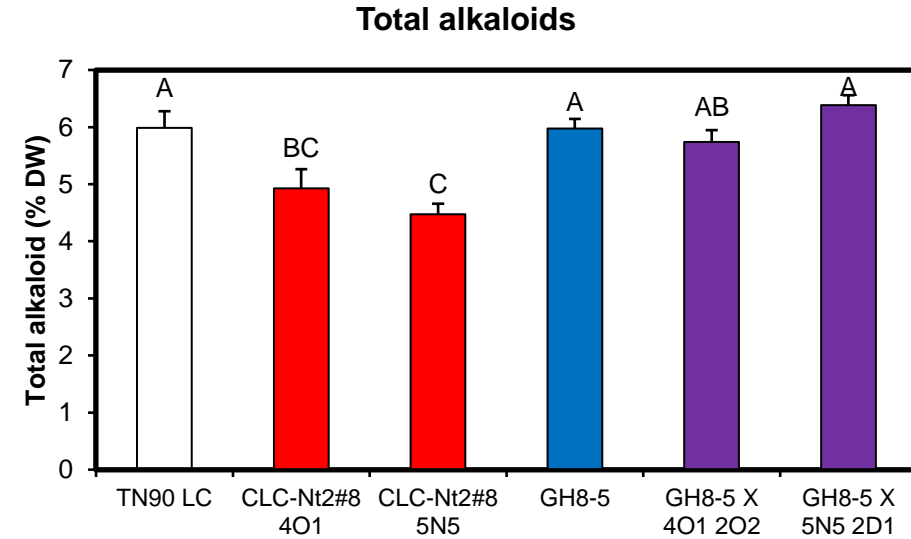
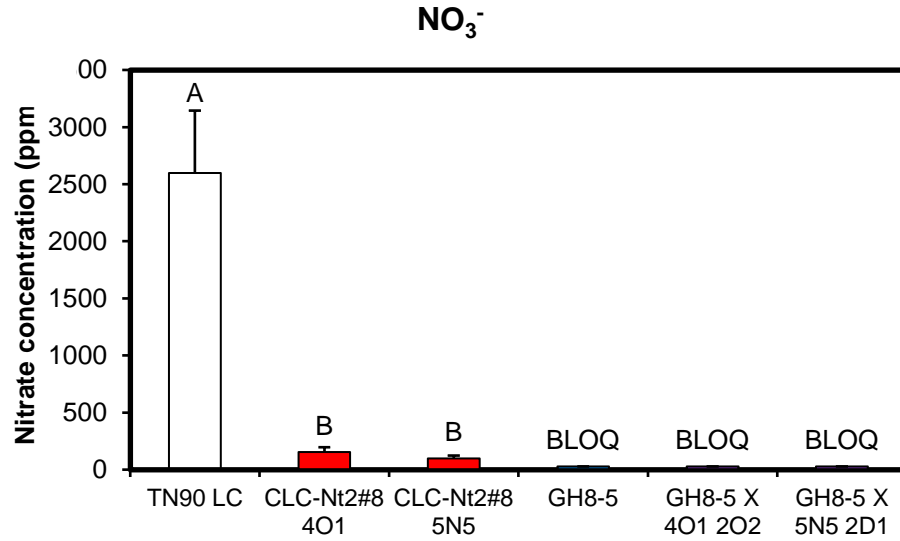
Flowering Time and Biomass Results



*Lines with only *clc-nt2* mutations tended to flower later than the rest (only one had statistical significance across all other lines), and all lines with the 35S:S523D-NR construct tended to flower earlier (though not all comparisons were statistically significant)*

All lines possessing the 35S:S523D-NR transgene showed major reductions in aerial biomass (34 – 39% decrease)

Nitrate, Alkaloids, and TSNAs



- *Both nitrate-reducing technologies greatly lowered the nitrate content*
- *The clc-nt2 mutations alone were associated with a modest reduction in alkaloids (possibly attributed to later topping dates)*
- *In 2022, both technologies led to similar overall TSNA reductions (79 – 86% decrease)*

Conclusions

- Of the two nitrate-lowering traits tested, the 35S:S523D-NR transgene was dominant to the CLC-Nt2 mutations with respect to flowering time and biomass.
- Although the CLC-Nt2 mutations alone can mediate substantial reductions in TSNA content, they would need to be deployed in combination with Zyvert mutations or LC screening because of the propensity of burley varieties to convert.
- Combining CLC-Nt2 knockout mutations with Zyvert mutations represents a promising strategy for maximally lowering TSNA content without compromising yield.