

Synthesis of three advanced biofuels from ionic liquid-pretreated switchgrass using *Escherichia coli*

Bokinsky et al, nov 2011 PNAS

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MSSB presentation, Genoscope
7 dec 11

Why did I choose this paper?

- It's new (published 4 weeks ago)
- It relates to yesterday's talk (and my research)
- Shows many basic problems in facing synthetic biologists
- JBEI is doing cutting edge research in synthetic biology



Consolidated bioprocessing for efficient conversion of biomass to fuel

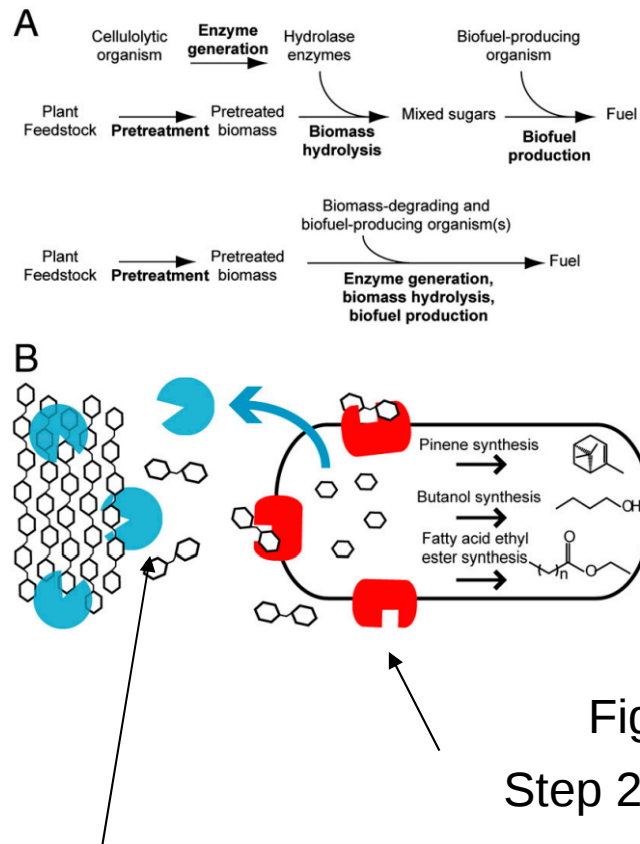
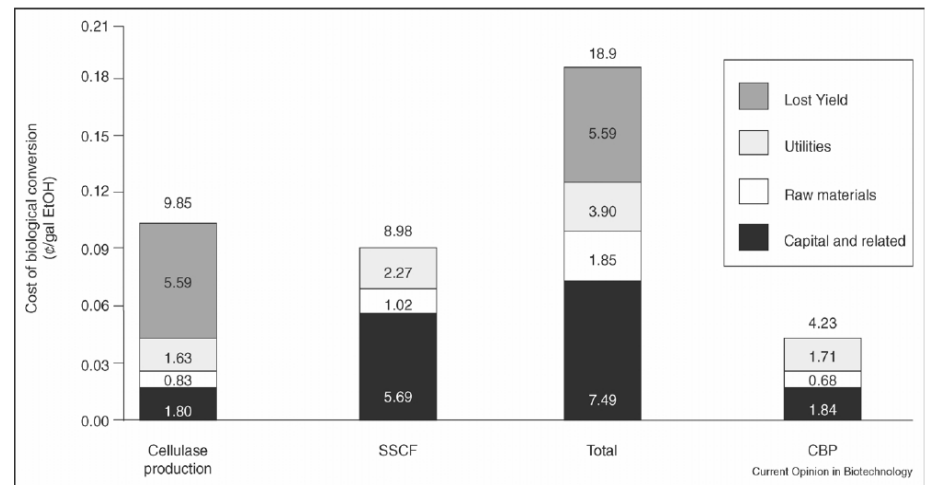


Fig 1

Step 2: glucosidase/xylobiosidase to cut short chains into sugars



Lynd et al, 2005

Step 1: cellulase/hemicellulase to cut polymers into short chains

What plant feedstocks (inputs) did they use?



switchgrass: prairie grass



yard waste



eucalyptus

What feedstocks (inputs) did they use?



switchgrass: prairie grass

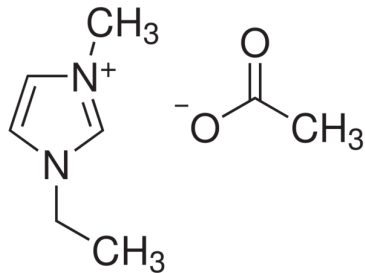


yard waste



Note: the biomass was pre-treated with an 'ionic liquid'

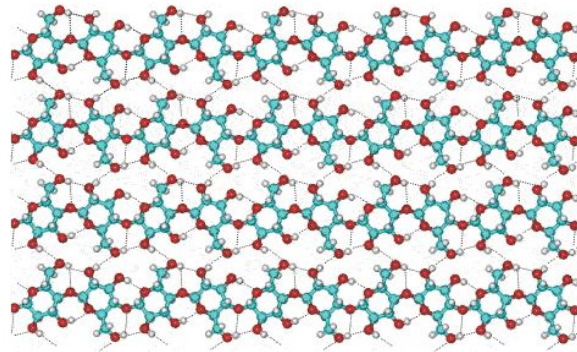
1-Ethyl-3-methylimidazolium acetate



Cost is a 'substantial barrier to commercialization' (it's expensive!)

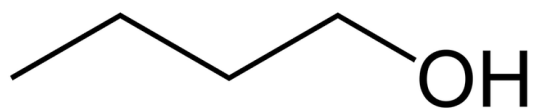
What are environmental impacts of these liquids?

On the positive side, IL 'eliminates cellulose crystallinity'

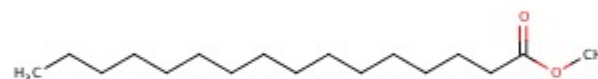


What chemicals did they produce?

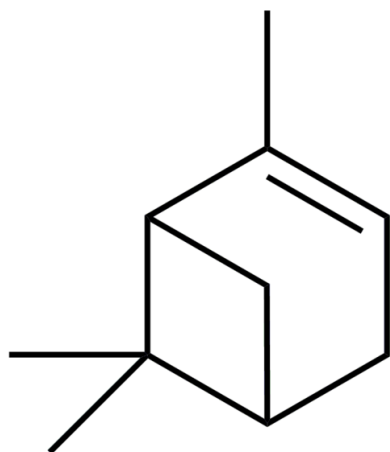
Butanol (gasoline replacement)



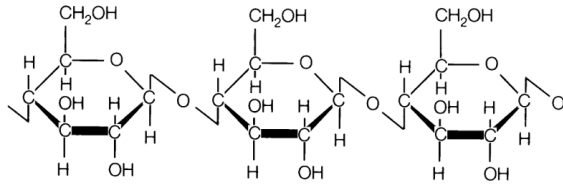
Fatty acid (biodiesel)



Pinene (jet fuel precursor)

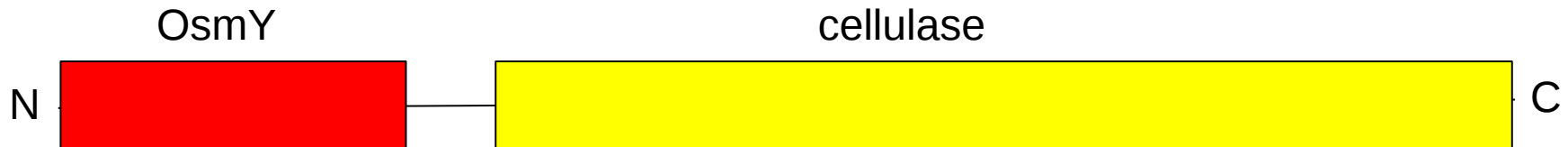


How did they make *E. coli* degrade cellulose?



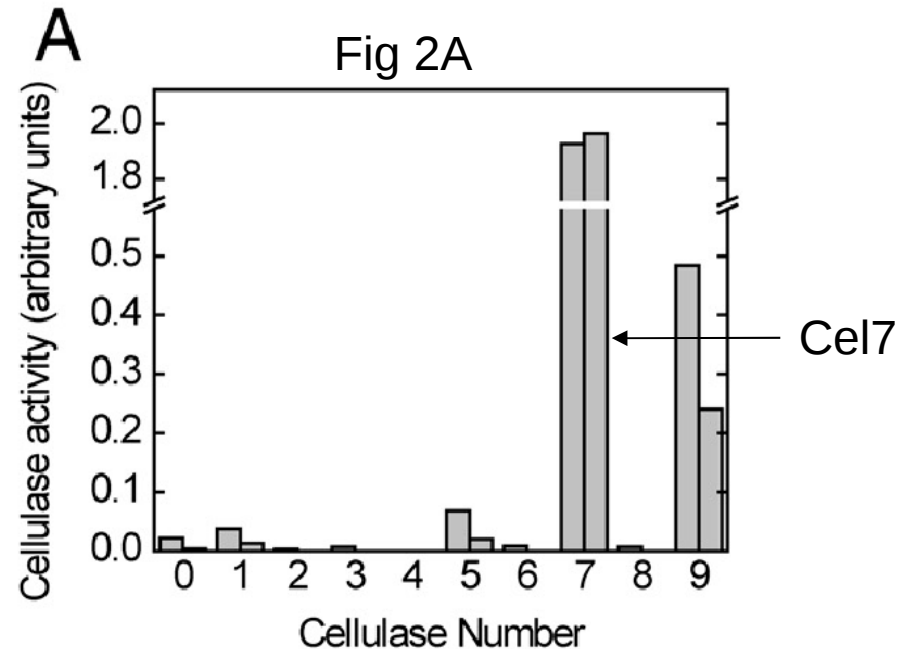
Cellulose is a polymer of glucose

Step 1: fuse cellulase genes to OsmY domain so they are secreted



How did they make *E. coli* degrade cellulose?

Step 2: find a cellulase that works in *E. coli*



Expressed 10 cellulases and tested supernatant for cellulase activity

Question:

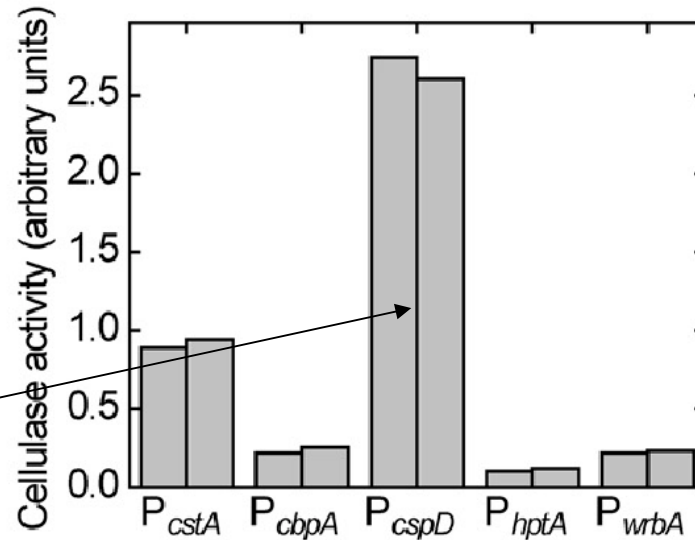
How would you select genes to test?

http://www.cazy.org/GH5_all.html

How did they make *E. coli* degrade cellulose?

Step 3: find a which promoter gives highest expression of *cel7*

Fig 2F



Chose PcspD

CstA: peptide transporter
CbpA: chaperone
CspD: DNA replication inhibitor
CspD: DNA replication inhibitor
WrbA: flavodoxin

Questions: What is a promoter?

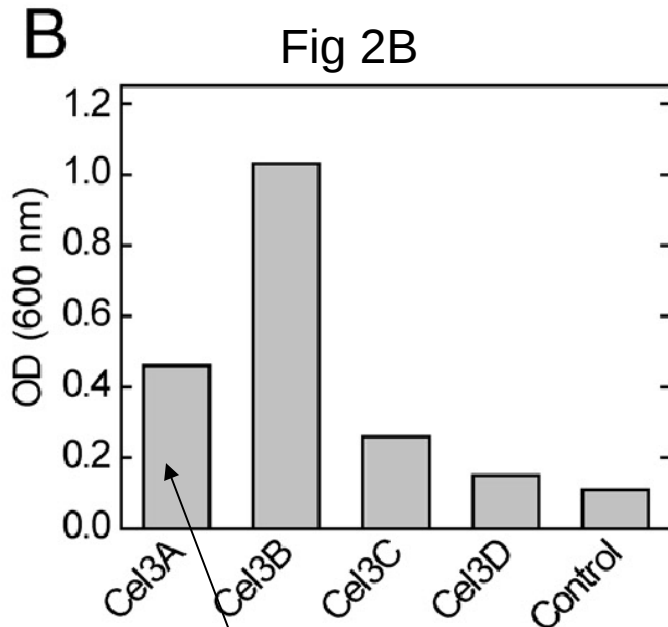
How would you choose promoters to test?

-35 Box -10 Box

```
5'-TCCCTTTGATATTGCATC---CCGCGTATATAATATGTC-3'
5'-GCCCATTTG-CATCAAGATCTGTTGAACTATAATTCCTT-3'
5'-CTGCATTGA-ATAGACTTCAACTCGCGATATAATTCAAG-3'
5'-ACACTTTGACACACCCTT-TACCCCTTATAATTAACG-3'
5'-ACCTATTGACAATTAAAGGCTAAAATGCTATAATTCCAC-3'
5'-GGCCGTTGACAGCCCAAGCAAATACTCATATAATCATAG-3'
5'-AATACTTGACATATCACTGTGATTCACATATAATATGCG-3'
5'-TGTGACTGGCGCTACAATCTTCCAAAGTCACAATTCTCA-3'
```

How did they make *E. coli* degrade cellulose?

Step 4: find a cellobiosidase that works in *E. coli*

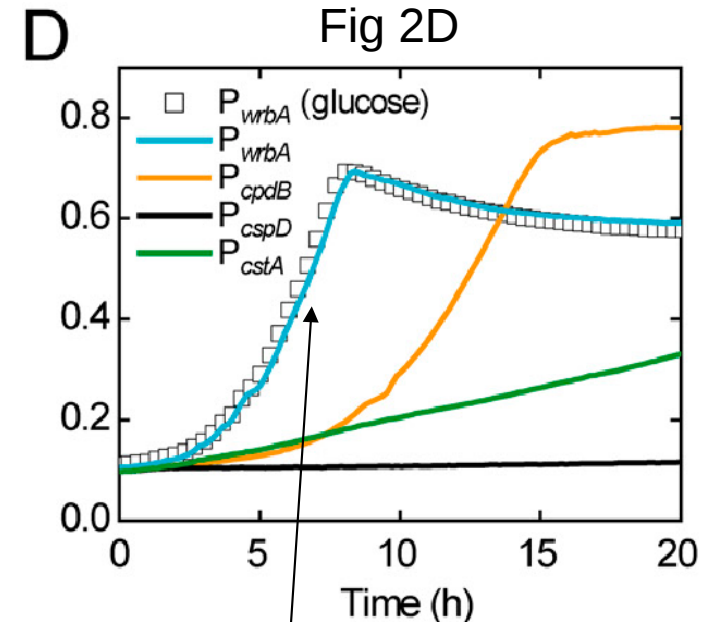


Screened 4 cellobiosidases from *Cellvibrio japonicus*. Only 4 genes?

Chose Cel3A. Why?

“*E. coli* grew best on cellobiose when expressing Either *cel3A* or *cel3B*”

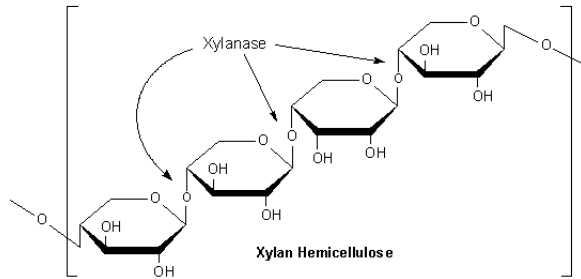
Step 5: find which promoter gives highest expression of *cel3A*



chose P_{wrbA}

How did they make *E.coli* digest hemicellulose?

hemicellulose is a xylose polymer

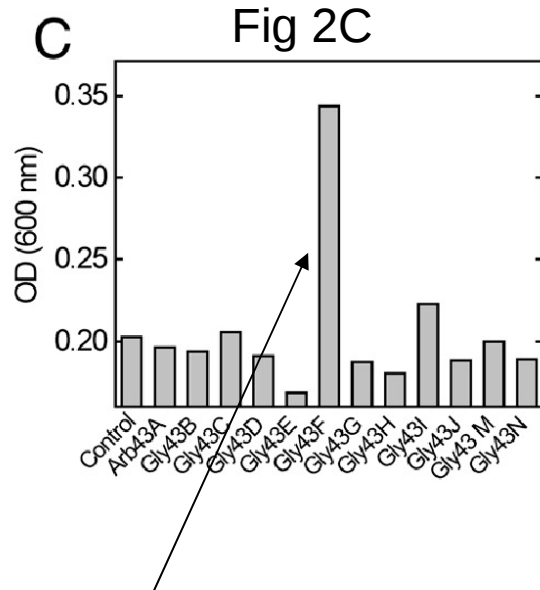


Step 1: find a hemicellulase that works in *E. coli*

This was done in another study, (Xyn10B)
but I think they used the same strategy as
for the cellulase

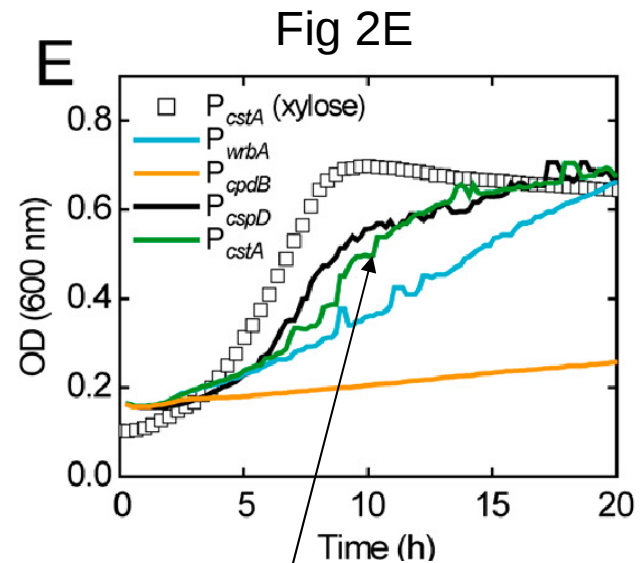
How did they make *E.coli* digest hemicellulose?

Step 2: test genes to cut short xylosaccharides by screening 12 xylobiosidases from *C. japonicus*



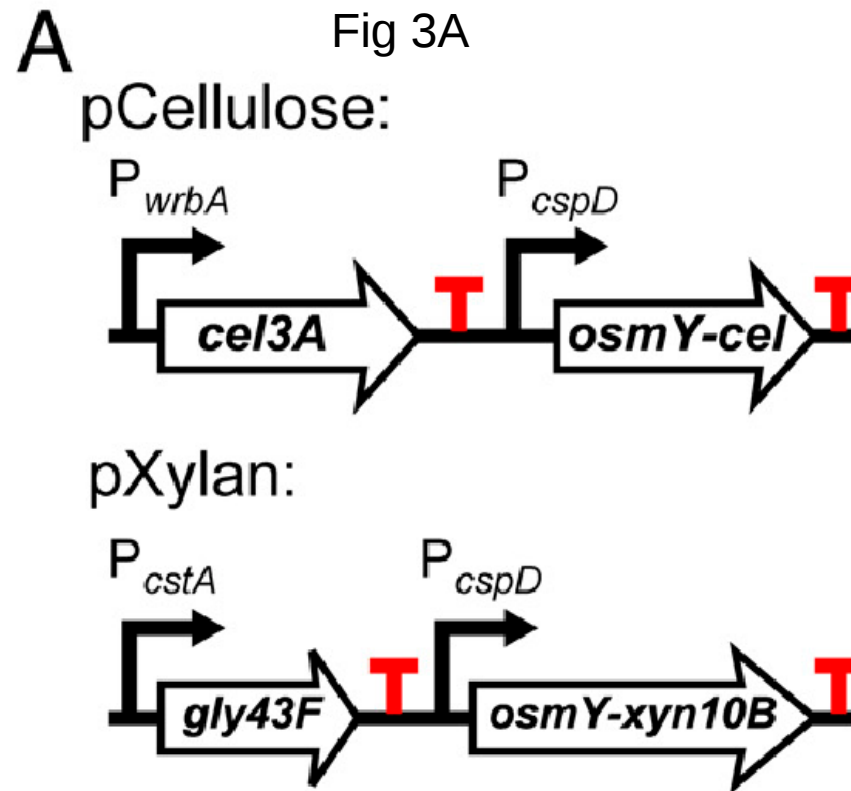
How lucky that 1 worked!
But how significant is that?

Step 3: find which promoter works best with Gly43F



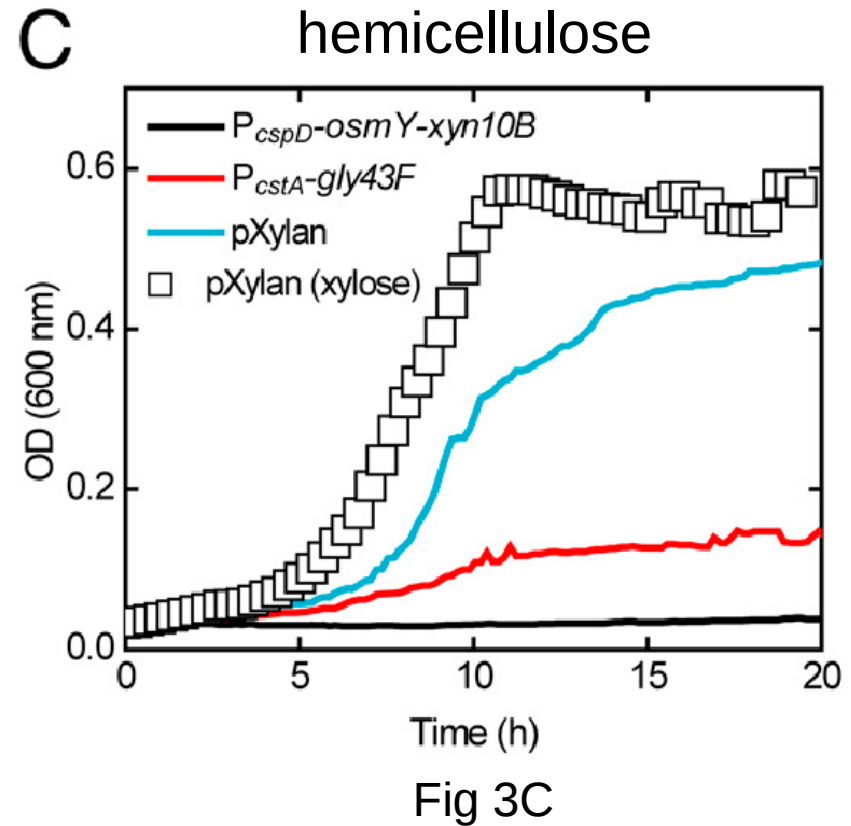
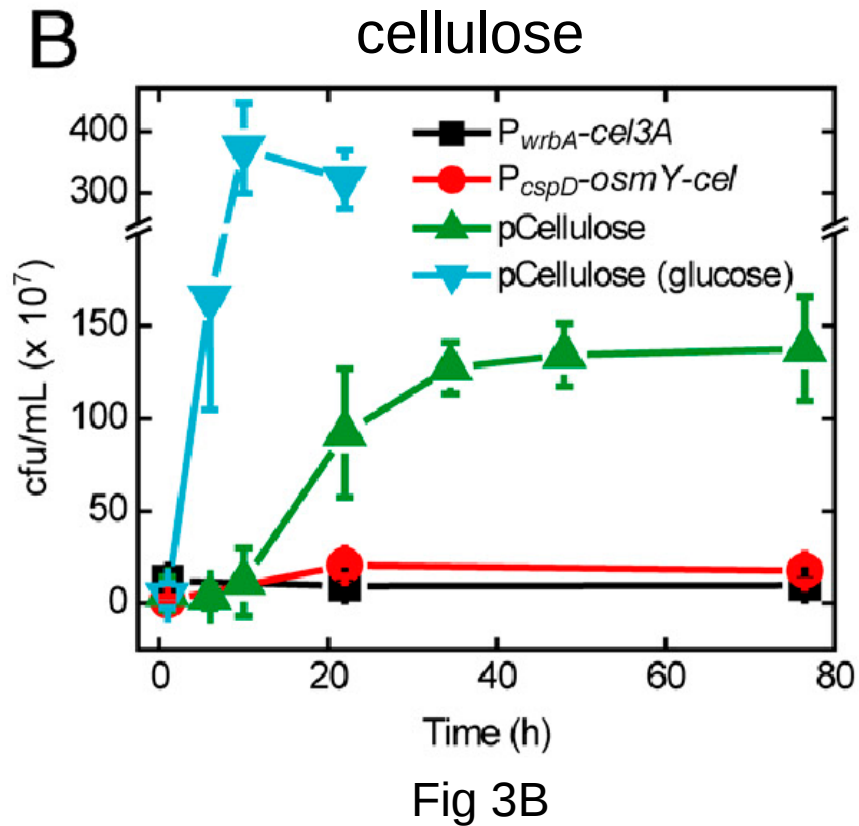
Chose P_{cstA}

clone genes onto plasmid and put them in *E. coli*

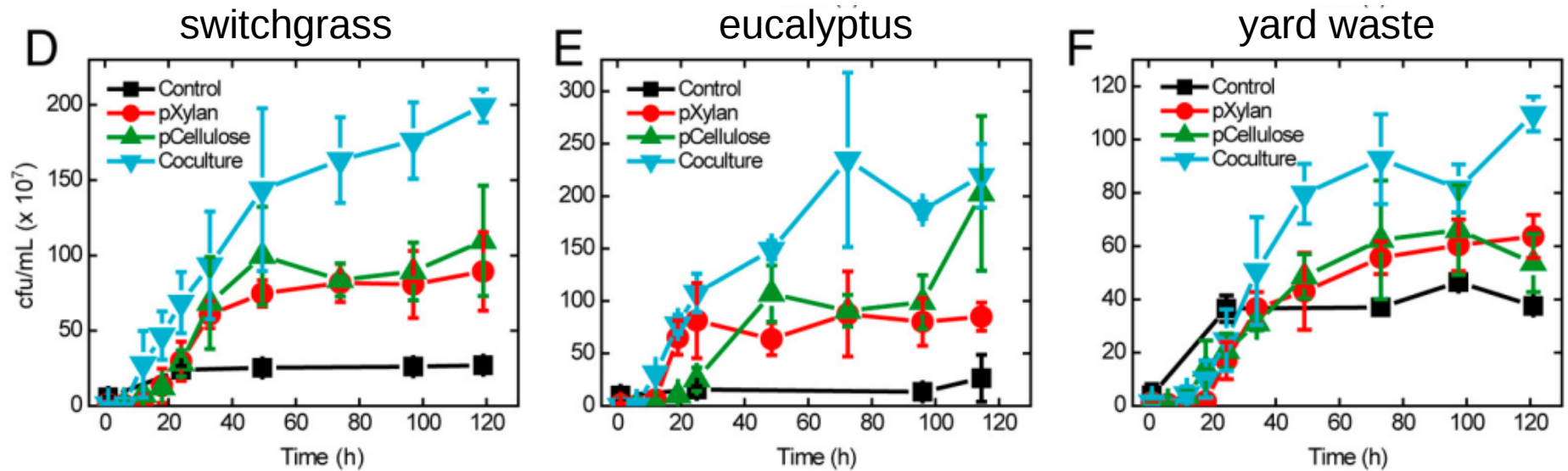


Question: why do you think different promoters worked best for different genes?

strains can grow on model substrates



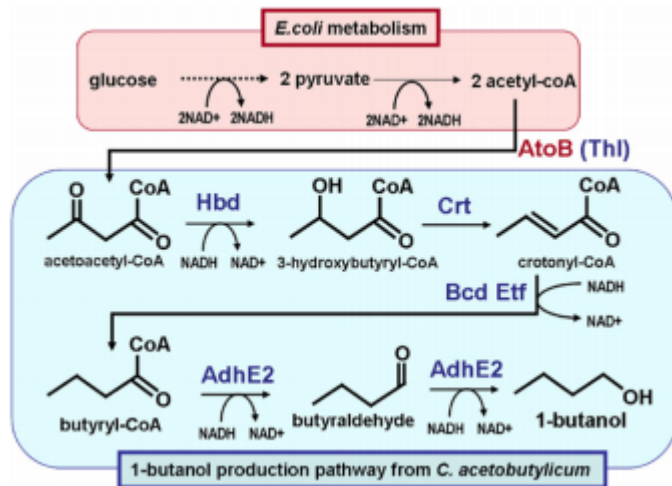
Strains can grow on biomass (sort of)



The cellulase and hemicellulases hydrolyzed 8% of sugars in switchgrass

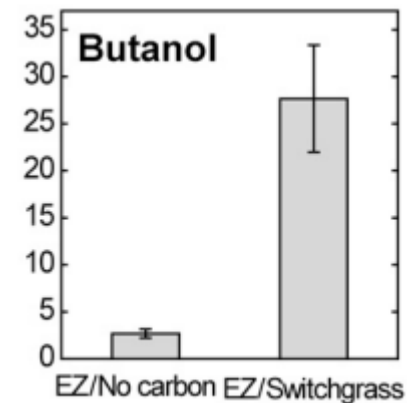
now add genes to make fuels (butanol)

Butanol=7 genes from *Clostridium acetobutylicum*



Atsumi et al, 2008

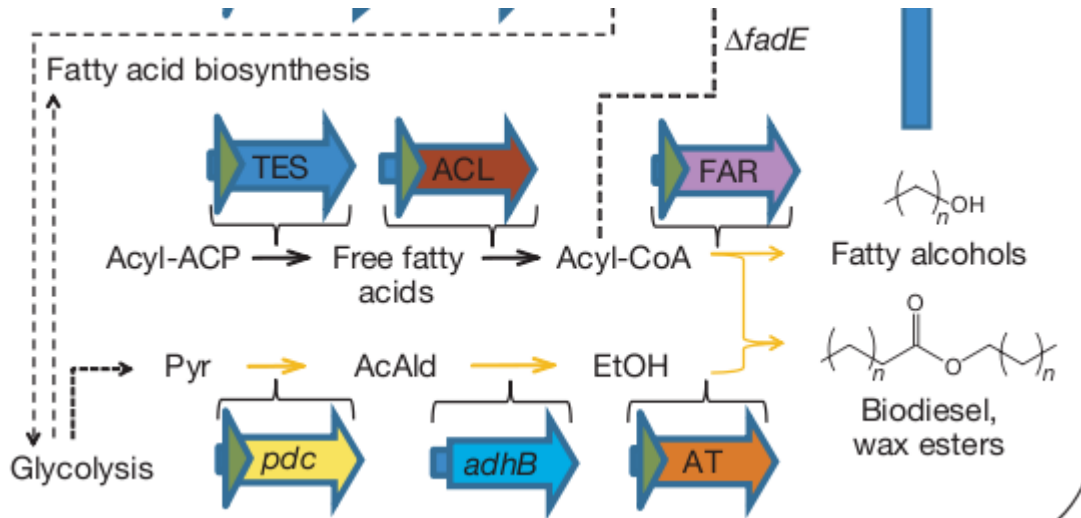
Fig 4c



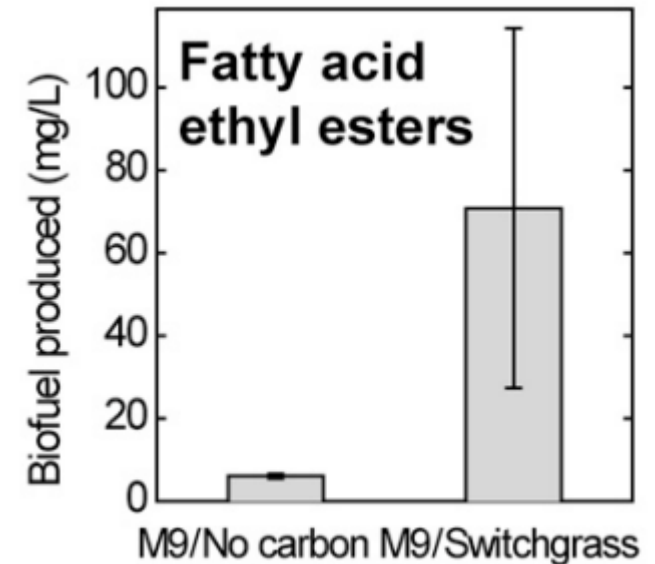
now add genes to make fuels (FAEE)

Fatty acid ethyl esters=overexpress 6 genes

Fig 4B



Steen et al, 2010



now add genes to make fuels (pinene)

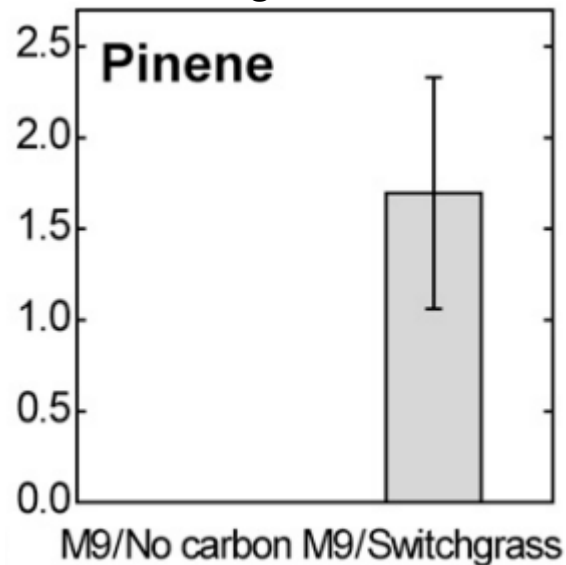
Pinene=9 genes (source unknown)

pPinene:

lacUV5



Fig 4D



Are these yields high enough to really make fuels?

butanol yield=0.028 g butanol/l of water

butanol density=0.81 g/ml

What volume of water would you need to get 1l (810g) butanol?

need 29,000 l water to get 1l butanol



What are the positives and negatives of this paper?

Positives

Made E.coli eat plants

Screen for promoters
and genes that work in
E. coli

Plant->fuel in single step

Pure fuels

Negatives

Side products?

Are other mutations
needed? Transporters?

More mutations?

Make intermediates at
higher conc?

Only 8% of plant degraded