

Bioproduct Life Cycle Analysis with the GREETTM Model

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Selection of bioproducts based on a high-level market analysis



Process simulations provided material and energy flows used in analysis



EDI: electrodeionization ³

The GREET (<u>G</u>reenhouse gases, <u>R</u>egulated <u>E</u>missions, and <u>E</u>nergy use in <u>T</u>ransportation) Model at Argonne National Lab





the fuels that can be produced from each of those feedstocks.

Carbon Accounting in the Bioproduct System Boundary



6

Bioproducts uniformly showed reductions compared to their fossil-derived counterparts



Process natural gas and feedstock consumption drive GHG emissions



PE: polyethylene; ADP: adipic acid

8



Bioproduct	Feedstock	GHG Emissions: Cradle-to-Grave		Poforoncoc
		kgCO ₂ e/kg	% Reduction	Kelerences
Propylene Glycol	Soybean & Canola	3.2	61%	ADM
	Glycerol	1.1	66%	GREET
1,3-PDO	Corn	2.7-3.5	46-71%	Urban (2009)
	Corn	1.2-2.9	37-55%	Hermann (2007)
	Sugar Cane	-1.8	62-115%	Hermann (2007)
	Glycerol	2.7	66%	GREET
	3-HP	5.3	39%	GREET
Acrylic Acid	Corn starch	2	43%	Hermann (2007)
	Corn stover	1.2	66%	Hermann (2007)
	Sugar cane	0.7	80%	Hermann (2007)
	3-HP	8.7	53%	GREET
Succinic Acid	Corn starch	0.88	90%	Cok (2014)
		1.7	81%	Cok (2014)
		1.5	83%	Cok (2014)
	Clean Sugars	1.9	86%	GREET

Conclusions and Outcomes

- Bioproducts from corn stover and algae feedstocks have the potential for life-cycle GHG emissions that are below peer fossil-derived compounds.
- Heat integration and yield increase opportunities will improve bioproduct life-cycle GHG emissions.
- The GREET bioproducts module, to be released Fall 2014, will allow the community to explore these results and generate results for bioproduct pathways of interest.
- A technical report will document data sources and methodology used to build the GREET model.
- Bioproducts module is subject to updates as additional information becomes available. It will be expanded to include additional products.

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