

Comparison of the NRC 2001 Model and DVE/OEB (Dutch) System in the prediction of protein supply to dairy cows from new co-products of bio-ethanol production

$$RUP^{DEV} = 1.1 \times CP \times \%RUP$$

$$MP = ARUP^{NRC} + AMCP^{NRC} + AECP$$

$$AMCP^{NRC} = 0.80 \times$$



$$0.75 \times MCP_{FOM}$$

$$RUP^{NRC} = CP \times \%RUP$$

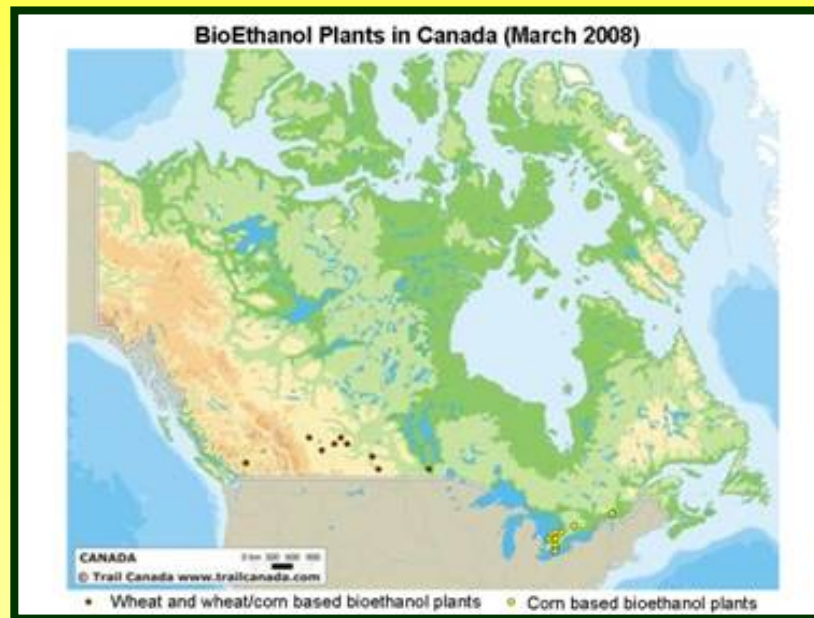
$$DVE = ARUP^{DVE} + AMCP_{FOM} - ENDP$$

Waldo Nuez Ortin



Research Justification

■ Increased ethanol production capacity in western Canada



1.4 million mt of feedstock



512 million liters of ethanol



460,000 mt of DDGS

38% wheat

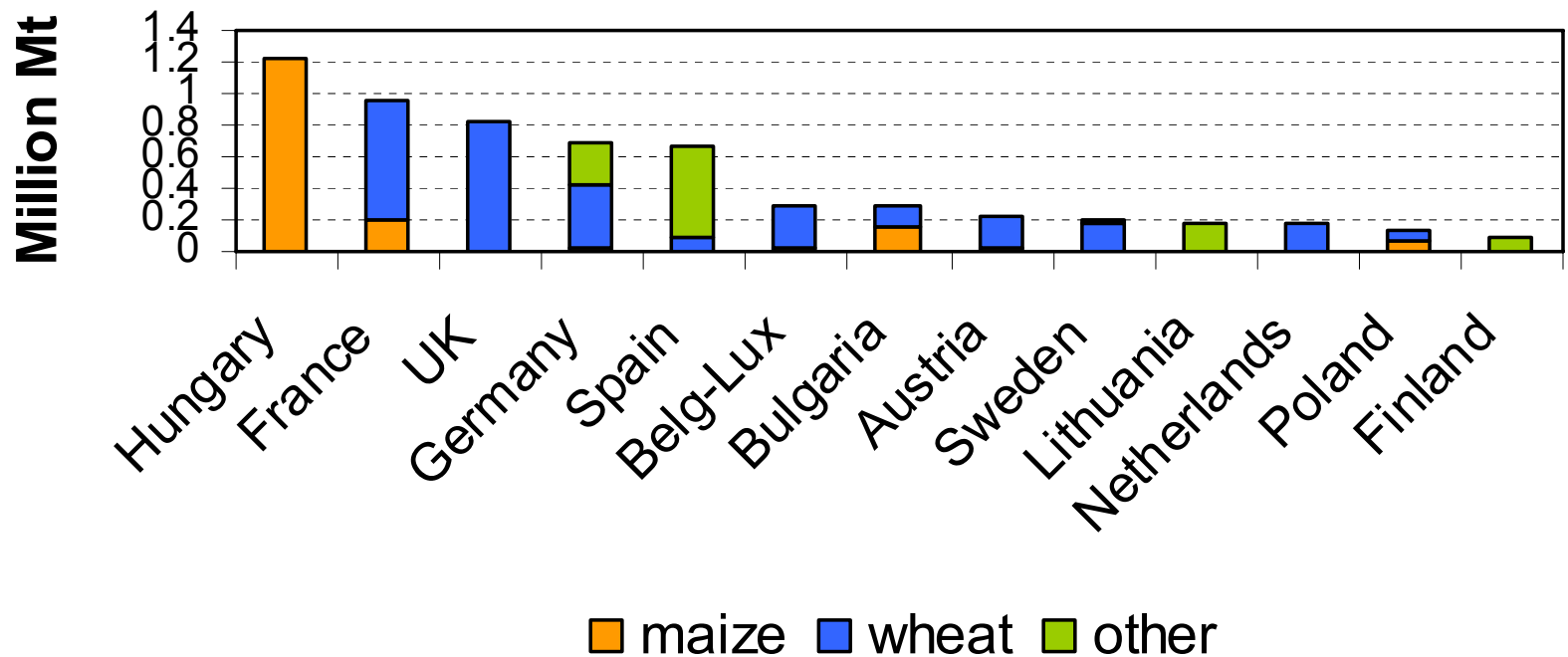
62% wheat/corn

www.ddgs.usask.ca

Research Justification

■ Europe (2.8 billion liters of ethanol in 2008)

DDGS production by country in 2010



<http://www.allaboutfeed.net>

Research Justification

■ Dietary protein evaluation of DDGS for ruminants

- **Chemical profile, CNCPS** (*Nuez Ortin and Yu, 2009*)
 - **CP (%DM): 14% (wheat) vs. 39% (wheat DDGS) vs. 32% (corn DDGS)**
 - **PB2 (%CP): 62% vs. 28% vs. 54%**
 - **PB3 (%CP): 15% vs. 51% vs. 28%**
- **In situ rumen procedure** (*Batajo and Shaver 1998, Grings et al. 1992*)
 - **RUP (%CP): 46% (corn) vs. 60% (corn DDGS)**
- **Intestinal digestibility of RUP** (*NRC 2001*)
 - **dRUP: 90% (corn) vs. 75% (corn DDGS)**

Research Justification

■ Modern protein evaluation models

- Chemical profile
- In situ rumen degradation characteristics
- Intestinal digestibility of CP



- DVE/OEB (Dutch) System *(Tamminga et al. 1994)*
- NRC 2001 Model *(NRC 2001)*



- Ruminal and postruminal availability of protein



Objective

Prediction of protein supply of DDGS to dairy cows using NRC 2001 and DVE/OEB System

1) Magnitude of difference

- **feedstock vs. wheat DDGS vs. corn DDGS vs. blend DDGS**

2) DVE/OEB System vs. NRC 2001 Model

Methodology

■ Samples

- Wheat (3)
- Corn (3)
- Wheat DDGS (5)
- Blend DDGS (70% wheat:30% corn) (3)
- Corn DDGS (3)

■ Animals

- 3 rumen fistulated Holstein cows (dry)

Methodology

Rumen incubation at 0, 2, 4, 8, 12, 24, 48 and 120h (2 runs)

Original sample: DM, Ash, CP, NDF, Starch, EE
Residue: DM, Ash, CP, NDF, Starch

*Ørskov and
McDonald, 1979*

S, D, U, T0, F0, Kd
(CP and starch)

*Calsamiglia and
Stern, 1995*

Intestinal dRUP

*Weiss et al.
1992*

TDN value

*Tamminga et
al. 1994*

Fermented OM

DVE/OEB System
Non TDN Model

NRC 2001 Model
TDN Model

Methodology

■ Availability of protein in SI

DVE/OEB System

True absorbable protein in SI (DVE)

Absorbable RUP

+

**Absorbable microbial protein
synthesized in rumen**

—

Endogenous CP losses

NRC 2001 Model

Metabolizable protein (MP)

Absorbable RUP

+

**Absorbable microbial protein
synthesized in rumen**

+

Absorbable endogenous CP

Methodology

■ Degraded protein balance

**DVE/OEB System
(Non-TDN Model)**

Degraded protein balance (OEB)

Microbial protein synthesis_{RDP}

—

Microbial protein synthesis_{FOM}

**NRC 2001 Model
(TDN Model)**

Degraded protein balance (DPB)

Microbial protein synthesis_{RDP}

—

1.18 x Microbial protein
synthesis_{TDN}

Methodology

■ Statistical analysis

■ DDGS type effect on protein availability (Objective 1)

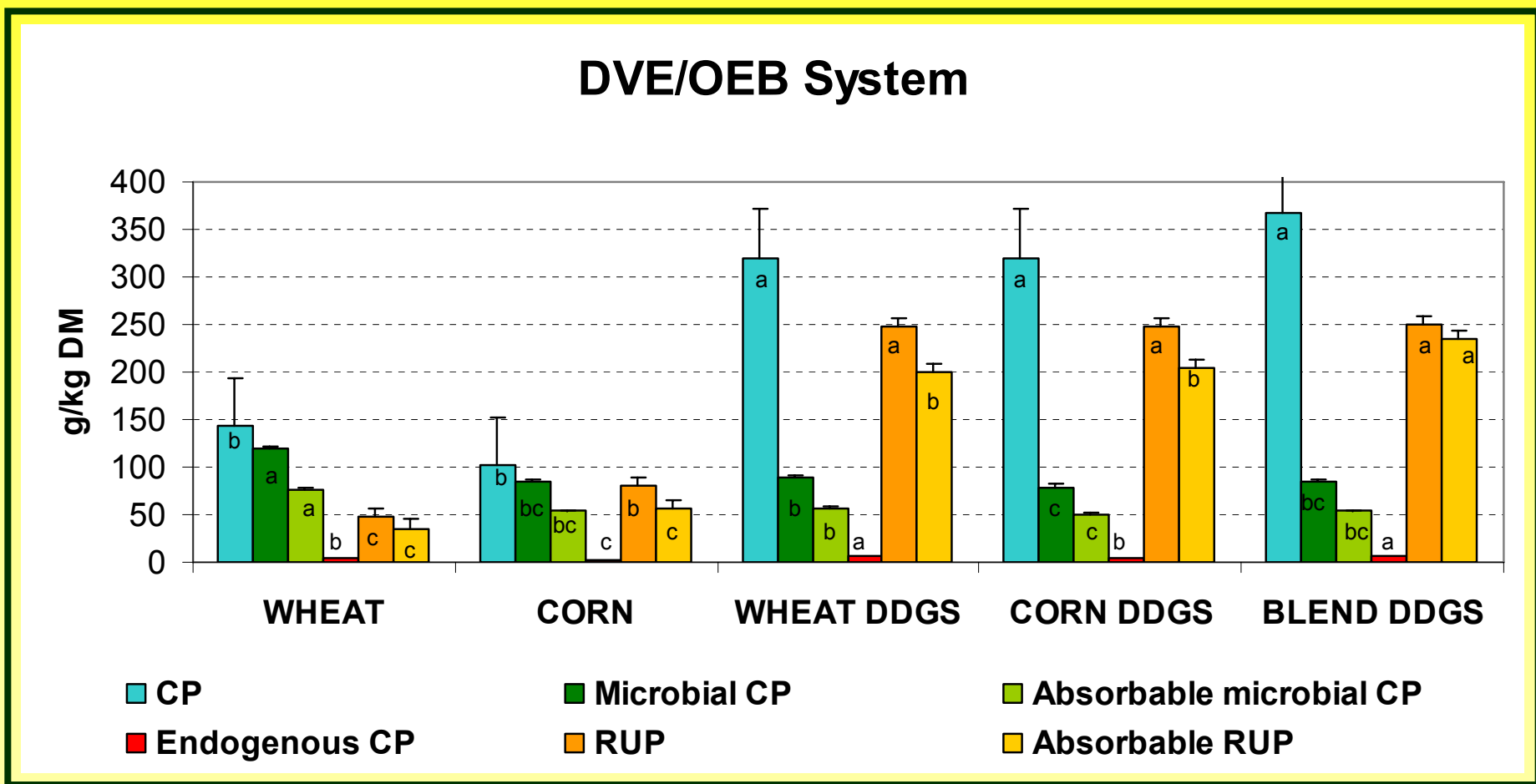
□ CRD

- Batches and runs as replicates
- SAS Mixed procedure
- Means separation using Fisher Protected LSD method ($P < 0.05$)

■ DVE/OEB System vs. NRC 2001 Model (Objective 2)

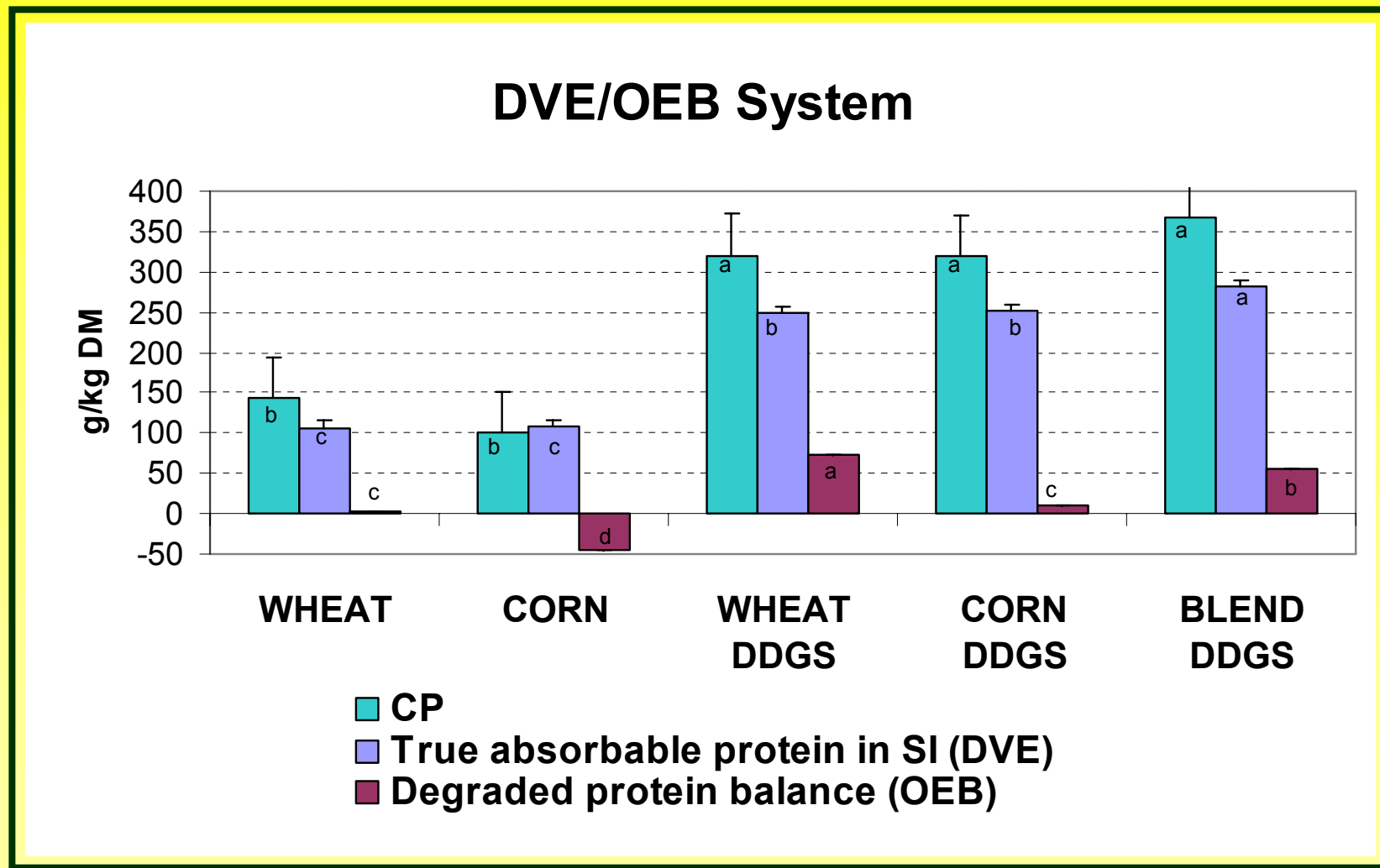
- Paired t test
- Correlation

Results (Objective 1)



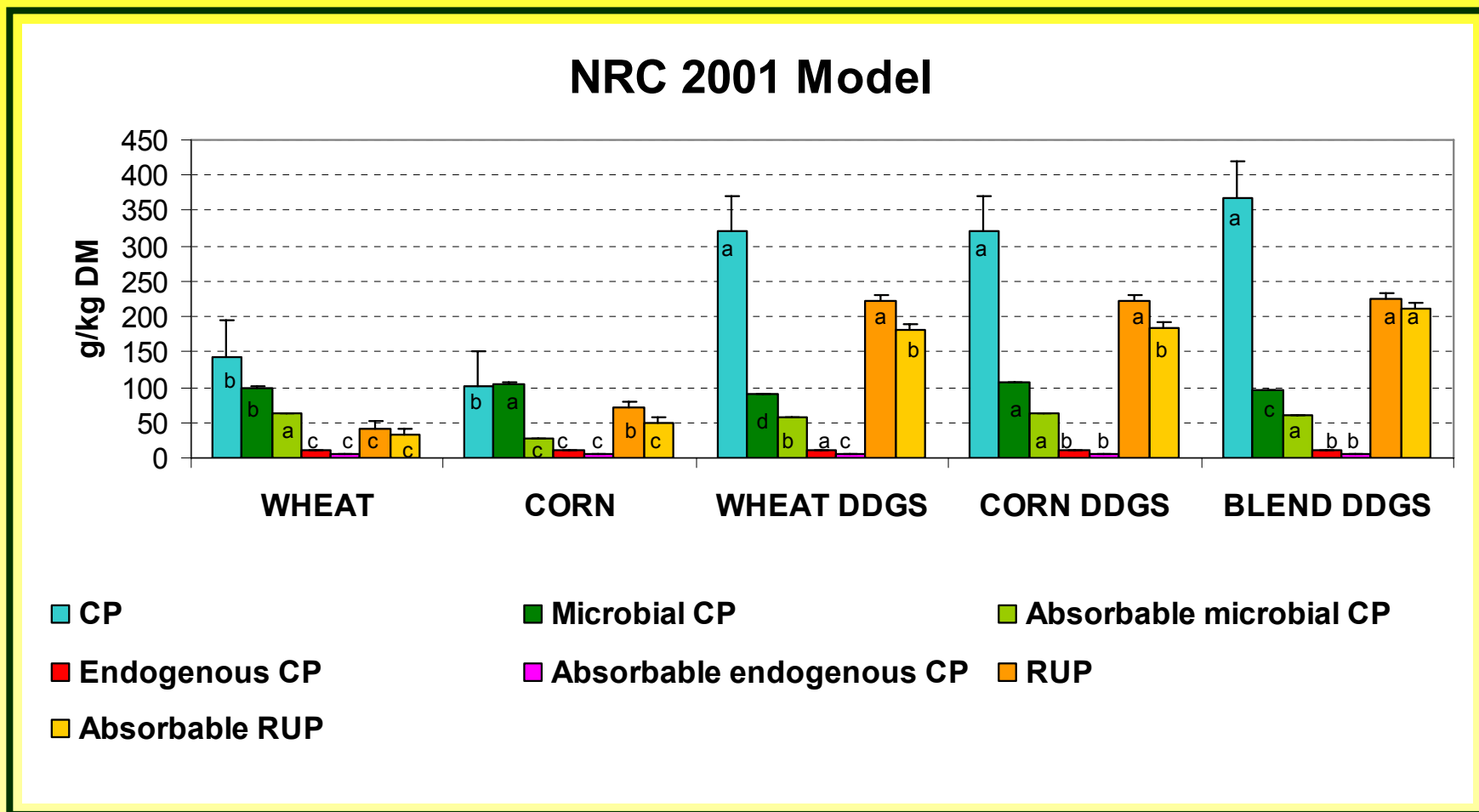
a, b, c, d Means with same letter in the same block are not significantly different ($P > 0.05$) LSD method

Results (Objective 1)



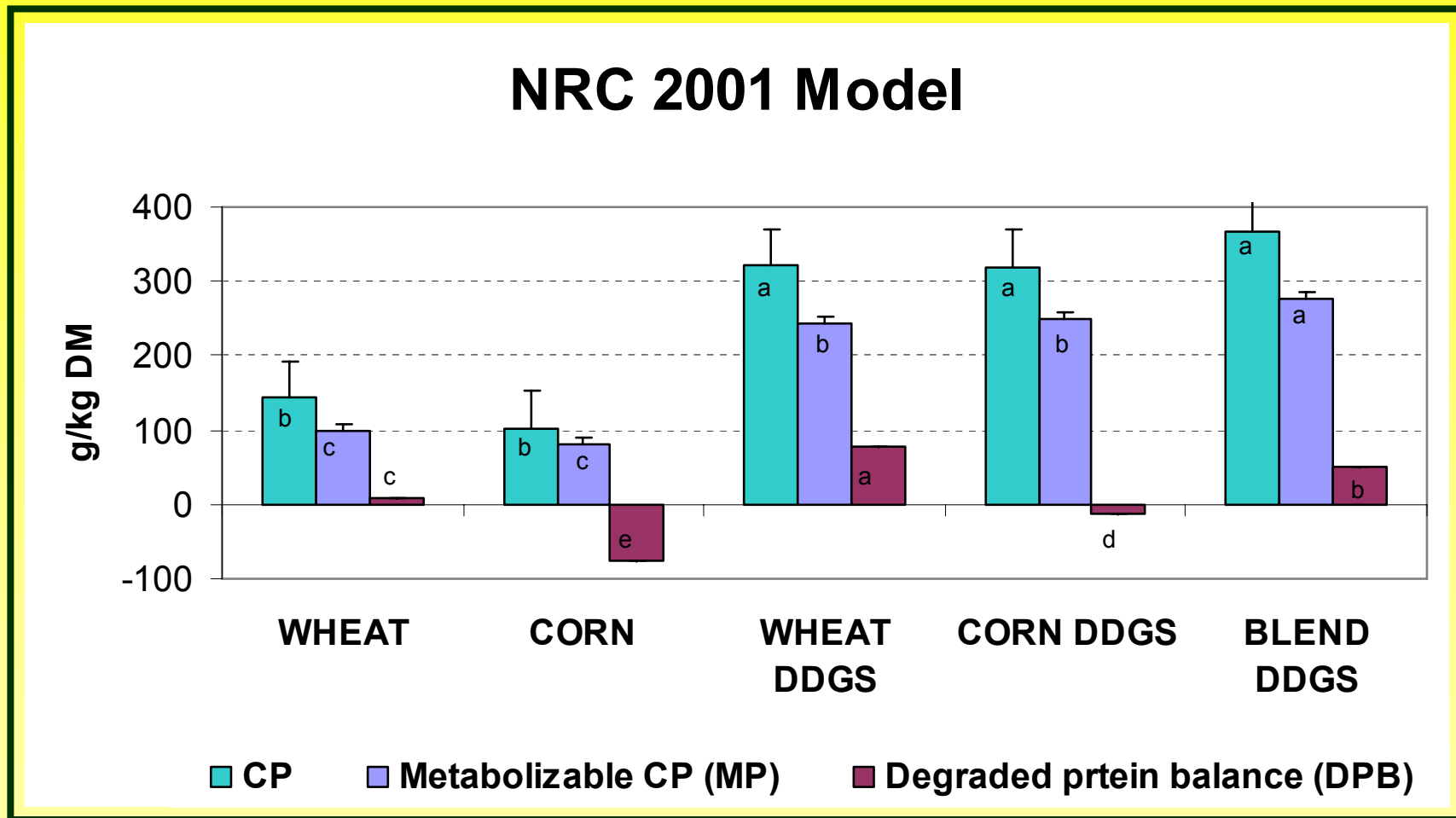
a, b, c, d Means with same letter in the same block are not significantly different ($P>0.05$) LSD method

Results (Objective 1)



a, b, c, d Means with same letter in the same block are not significantly different ($P > 0.05$) LSD method

Results (Objective 1)



a, b, c, d Means with same letter in the same block are not significantly different ($P>0.05$) LSD method

Results (Objective 2)

DVE/OEB System vs. NRC 2001 Model

Items g/kg DM	Mean DVE/OEB	Mean NRC 2001	Difference	SED	P value	R	P value
Absorbable microbial CP	53.98	60.10	-6.11	1.267	<0.0001	-0.44	0.0405
Absorbable endogenous CP	6.32	4.39	1.92	0.394	<0.0001	0.45	0.0371
Absorbable RUP	210.35	189.50	20.85	0.664	<0.0001	1.00	<0.0001
Metabolizable CP/DVE	258.01	253.99	4.01	0.793	<0.0001	0.99	<0.0001
Degraded protein balance	50.70	45.94	4.76	2.728	0.0957	0.97	<0.0001

Paired t test.
Pearson correlation.

Conclusion

■ DVE/OEB System vs. NRC 2001 Model for DDGS evaluation:



■ True absorbable protein in SI (DVE/MP):

- DDGS > feedstock grain
- Blend DDGS (70:30) > wheat DDGS = corn DDGS



■ Balance between microbial protein synthesis from available RDP and available energy (DPB):

- Corn DDGS showed the closest to zero value



■ Absorbable RUP

- MP/DVE was higher for the DVE/OEB System



Thank you!!

Advisor:

Dr. Peiqiang Yu

Advice Committee Members:

Dr. Dave Christensen

Dr. John McKinnon

Dr. Bernard Laarveld



ABIP-FOBI

Feed Chair Research