

# SPECTRUM

## Balance™

### SPECIFICATIONS



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ANIMAL NUTRITION

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Standard Analysis	% DM
Moisture	9.3
Dry Matter	90.7
Nitrogen	7.0
Crude Protein	43.7
Soluble Protein	16.7
Ammonia	0.0
NDIP	6.1
ADIP	6.1
Crude Fiber	5.0
ADF	8.4
NDF	21.6
Lignin	1.4
Total VFA	0.0
Lactic	0.0
Acetic	0.0
Sugar	3.0
Starch	0.9
Soluble Fiber	6.8
NFC	10.8

Fatty Acids	% DM
Fat	16.7
TFA	16.7
Glycerol	0.0
Pigment	0.0
C12:0	0.0
C14:0	0.1
C16:0	8.2
C16:1	0.0
C18:0	0.8
C18:1T	0.0
C18:1C	4.0
C18:2	3.2
C18:3	0.1
Other	0.2
Fat Type	1.9
Lipolysis Rate	323.3

Pool	% DM	Rate	Int dig
CHO A1	-	-	-
CHO A4	3.02	40.00	100.00
CHO B1	0.95	17.00	75.00
CHO B2	-	-	-
CHO B3	18.28	7.00	20.00
CHO C	3.36	-	-
PRO A1	-	-	-
PRO A2	16.73	7.50	100.00
PRO B1	20.91	2.09	100.00
PRO B2	-	-	-
PRO C	6.10	-	-

Protein Bio-Availability	% DM
RUP % CP	74.8
RUPd % RUP	81.4
dRUP % DM	26.6
dRUP-Lys % DM	1.7
dRUP-Met % DM	0.4
dRUP-EAA % DM	12.9

Macro Mineral	% DM
Ash	7.2
Calcium	1.3
Phosphorus	0.9
Magnesium	0.2
Potassium	0.9
Sulfur	0.7
Sodium	0.3
Chloride	0.3

Trace Mineral	ppm
Iron	564.9
Manganese	12.8
Zinc	50.6
Copper	5.5

Amino Acids	Product		Model	
	% DM	% of CP	% DM	% of CP
Arginine	2.0	4.6	1.9	4.3
Histidine	1.5	3.5	1.8	4.1
Isoleucine	1.1	2.5	0.9	2.0
Leucine	5.0	11.5	5.3	12.1
Lysine	2.3	5.3	2.7	6.2
Methionine	0.7	1.5	0.6	1.4
Phenylalanine	2.5	5.7	2.7	6.2
Threonine	1.7	3.9	1.8	4.0
Tryptophan	0.4	1.0	0.5	1.1
Valine	2.8	6.5	3.1	7.1
EAA	20.1	46.0	21.2	48.5

<sup>1</sup>When a feed component of a dairy ration consists of a blend of feed ingredients differing in rumen undegradable protein and/or post-ruminal protein digestibility, the amino acids of the resulting mixture differ in their ruminal degradability and post ruminal digestibility. Most ration formulation software allow only a single, common set of parameters to characterize ruminal degradability and post ruminal digestibility of all essential amino acids; one cannot enter parameters specific to each amino acid. Therefore, entering the amino acid profile of the whole protein of a feed mixture results in erroneous values of its metabolizable amino acid content. To circumvent this problem, users should use the 'model specifications' amino acid values in software such as CNCPs/NRC so that the correct supply of each metabolizable amino acid is calculated correctly.

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