1. Purpose of Component

The STOCK component encapsulates the GRAZPLAN animal biology model, as described in:

Freer M, Moore AD & Donnelly JR (1997). GRAZPLAN: decision support systems for Australian grazing enterprises. II. The animal biology model for feed intake, production and reproduction and the GrazFeed DSS. Agricultural Systems 54, 77-126.

Animals represented in a component instance may be of different genotypes. In particular, sheep and cattle may be represented within a single component instance.

and leave the simulation, and as physiological events such as maturation, birth or weaning take place. Animal groups that become sufficiently similar are merged into a single The animals represented by a component instance are classified into groups. The members of each animal group have the same genotype and age class, but may have a range have the same stage of pregnancy and/or lactation; the same number of suckling offspring; and occupy the same paddock. The set of animal groups changes as animals enter of ages (for example, an animal group containing mature animals may include four-year-old, five-year-old and six-year-old stock). The members of each animal group also

Each animal group has a unique, internally-assigned integer index, starting at 1. Because the set of groups present in a component instance is dynamic, the index number associated with a particular group may change over time. Each animal group is also assigned a paddock. The forage and supplementary feed available to a group of animals are determined by the paddock it occupies. Paddocks are referred to by name in the STOCK component. It is the user's responsibility to ensure that paddock names correspond to instances of the PADDOCK component or other sources of necessary driving variables. Each group also has a user-assigned tag and priority, which need not be unique. Tag values are generally used to manage distinct groups of animals in a common fashion. For example, all lactating ewes may be assigned the same tag value, which may then be used in management rules that keep them grazing together. Animal groups with different tag values are not merged even if they are otherwise similar. If tag values are assigned sequentially starting at 1, they can be used to generate summary variables. Priority values are used to allocate animals to paddocks in the draft event.

2. Initialisation Properties

The initialisation variable set is nearly completely optional. The idea is to allow the user to specify a minimal information set as well as a maximally detailed initialisation.

Description	Name of an XML file containing genotypic parameters. If the null string is specified, a default parameter set that is compiled into STOCK.DLL is used. If a file name is used, the parameters in the file modify (rather than replacing) the default parameter set.	Information about each animal genotype: Name used to refer to the genotype in management events	Maternal genotype (see notes)	• Paternal genotype (see notes)	• Number of generations of crossing: 0 denotes the pure-bred maternal genotype (in which case sire_breed is	not used), 1 a first cross, 2 a second cross (75% sire:25% dam), etc.	 Breed standard reference weight. The default value depends on dam_breed and sire_breed. 	• Expected rates of conception with 1, 2 and 3 young for mature ewes or cows in average body condition, over a mating period lasting 2.5 oestrus cycles. Only the first two elements are meaningful for cattle.	• Base rate of animal mortality. Default is 0.0.	• Breed reference fleece weight in sheep. The default value depends on dam_breed and sire_breed.	• Maximum average wool fibre diameter in sheep. The default depends on dam_breed and sire_breed.	 Clean fleece weight as a proportion of greasy fleece weight in sheep. Default is 0.70. 	• Potential maximum milk yield per head, in 4% fat-corrected milk equivalents, in cattle. Default is 20.0.
Required?	No	Yes											
Units							kg	ı	/yr	kg	mn	kg/kg	kg
Type	string	record[]	string	string	integer4		double	double[]	double	double	double	double	double
Property	param_file	genotypes . name	: dam_breed	: sire_breed	: generation		. Srw	: conception	: death rate	: ref_fleece_wt	: max_fibre_diam	: fleece_yield	: peak_milk

⇒ It is permitted to set both dam_breed and sire_breed to the null string. In this case the name field must be

The animal type (sheep or cattle) is implicit in the genotype fields.

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The dam_breed and sire_breed fields may contain the name of a genotype defined in an earlier element

of the genotypes array; multi-breed crosses may be specified in this way.

The set of valid breed names is set out below.

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a valid breed name.

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Description	Initial state of each animal group for cattle.	• Genotype of this group of animals. Must match the name field of an element of the genotypes property.	• Number of animals.	• Feasible values are 'cow', 'cows', 'heifer', 'heifers', 'steer', 'steers', 'bull', 'bulls'.	• Age of the animals.	 Unfasted live weight of the animals. 	 Highest weight recorded to date. 	• Genotype of the bulls to which pregnant or lactating animals were mated. Must match the name field of an	element of the genotypes property.	• Zero denotes not pregnant; 1 or more denotes the time since conception. Only meaningful for cows.	• Zero denotes not lactating; 1 or more denotes the time since parturition. Only meaningful for cows.	 Number of foetuses. Only meaningful for females with pregnant > 0. 	 Number of suckling calves. Only meaningful for cows with lactating > 0. 	 Condition score at parturition. Only meaningful for cows with lactating > 0. 	 Unfasted live weight of suckling calves. Only meaningful for cows with lactating > 0. 	 Paddock occupied by the animals. 	 Initial tag value for the animal group. 	• Priority accorded the animals in the <i>draft</i> event
Required?	No																	
Units				р	kg	kg	р			р			ı	kg				
Type	record[]	string	integer4	string	double	double	double	string		integer4	integer4	integer4	integer4	double	double	string	integer4	integer4
Property	cattle	: genotype	: number	: sex	: age	: weight	: max_prev_wt	: mated_to		: pregnant	: lactating	: no_foetuses	: no_suckling	: $birth_cs$	$: calf_wt$: paddock	: tag	: priority

Description	Initial state of each animal group for sheep.	• Genotype of this group of animals. Must match the name field of an element of the genotypes property.	• Number of animals.	• Feasible values are 'ewe', 'ewes', 'wether', 'wethers', 'ram', 'rams', 'crypto', 'cryptos'.	• Age of the animals.	 Unfasted live weight of the animals. 	 Highest weight recorded to date. 	 Greasy fleece weight of the animals. 	 Average wool fibre diameter of the animals. 	• Genotype of the rams to which pregnant or lactating animals were mated. Must match the name field of an	element of the <i>genotypes</i> property.	 Zero denotes not pregnant; 1 or more denotes the time since conception. Only meaningful for ewes. 	• Zero denotes not lactating; 1 or more denotes the time since parturition. Only meaningful for ewes.	 Number of foetuses or suckling lambs. Only meaningful for ewes. 	 Condition score at parturition. Only meaningful for ewes with lactating > 0. 	 Unfasted live weight of suckling lambs. Only meaningful for ewes with lactating > 0. 	 Greasy fleece weight of suckling lambs. Only meaningful for ewes with lactating > 0. 	 Paddock occupied by the animals. 	 Initial tag value for the animal group. 	• Priority accorded the animals in the <i>draft</i> event.
Required?	No																			
Units				р	kg	kg	kg	ш'n				р	p		1	kg	kg	1		
Type	record[]	string	integer4	string	double	double	double	double	double	string		integer4	integer4	integer4	integer4	double	double	double		
Property	sheep	: genotype	: number	: sex	: age	: weight	: max_prev_wt	$: fleece_wt$: fibre_diam	$: mated_to$: pregnant	: lactating	: no_young	$: birth_cs$	$: lamb_wt$: lamb_fleece_wt	: paddock	: tag	: priority

In no parameter file is specified, then permitted values for the dam_breed and sire_breed fields in the genotypes property are:

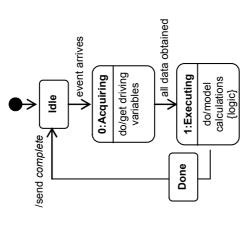
Sheep breeds	Sheep breeds	Cattle breeds	Cattle breeds
'black face x white face'	'polwarth'	'angus'	'friesian'
'border leicester'	, polypay,	'ayrshire'	'friesian x british'
'border leicester x merino'	'romney'	'beef shorthorn'	'guernsey'
'columbia'	'ryeland'	'brahman'	'hereford'
'corriedale'	'southdown'	'brahman x british'	'holstein'
'delaine-merino'	'suffolk'	'brown swiss'	'holstein x british'
'dorset x merino'	'targhee'	'charolais'	'jersey'
'dorset'	'texel'	'charolais x british'	'limousin'
'finnsheep'	'US corriedale'	'charolais x friesian'	'sahiwal'
'hampshire'	'US romney'	'charolais x holstein'	'simmental'
'large merino'	'US southdown'	'chianina'	'south devon'
'medium merino'	'US suffolk'	'dairy shorthorn'	
'small merino'			

3. Subscribed events – sequenced

3.1. do_stock

Default sequencing: 7000

Computes development, intake, growth and reproduction of all animals.



4. Subscribed events – other

4.1. buy

Causes a given number and type of animals to enter the simulation.

Parameter	Type	Units	Description
genotype	string		Genotype of the animals to be bought. Must match the <i>name</i> field of a member of the <i>genotypes</i> property.
number	integer4		Number of animals to be bought
sex	string		Sex of the animals. Feasible values are as for <i>sheep:sex</i> or <i>cattle:sex</i> , as appropriate.
age	double	months	Average age of the animals
weight	double	kg	Average unfasted live weight of the animals
$fleece_wt$	double	kg	Average greasy fleece weight of the animals. Only meaningful in sheep.
comd_score	double	1	Average condition score of the animals. If a value of zero is given, the default condition score for the weight and age will be used.
mated_to	string		Genotype of the rams or bulls with which the animals were mated prior to entry. Only meaningful if <i>pregnant</i> or <i>lactating</i> is non-zero. Must match the <i>name</i> field of a member of the <i>genotypes</i> property.
pregnant	integer4	р	Zero denotes not pregnant; 1 or more denotes the time since conception. Only meaningful for females.
lactating	integer4	р	Zero denotes not lactating; 1 or more denotes the time since parturition in lactating animals. Only meaningful for
			females.
guno_ou	integer4		Number of foetuses and/or suckling offspring.
young_wt	double	kg	Average unfasted live weight of any suckling lambs or calves.
young_fleece_wt	double	kg	Average greasy fleece weight of any suckling lambs.

4.2. castrate

Converts ram lambs to wether lambs, or bull calves to steers. If the animal group(s) denoted by group has no suckling young, has no effect.

If the number of male lambs or calves in a nominated group is greater than the number to be castrated, the animal group will be split; the sub-group with castrated offspring that were not castrated will be added at the end of the set of animal groups.

Description	Index number of the animal group, the lambs or calves of which are to be castrated. A value of zero denotes that each	animal group should be processed in turn until the nominated number of offspring has been castrated.	Number of male lambs or calves to be castrated.
Units			
Type	integer4		integer4
Parameter	group		number

4.3. draft

Assigns paddocks to animals in such a way that animal groups with the lowest tag values are placed in the paddocks with the best pasture. This event has no parameters.

Component Description – STOCK

4.4. dryoff

Ends lactation in cows that have already had their calves weaned. The event has no effect on other animals.

If the number of cows in a nominated group is greater than the number to be dried off, the animal group will be split; the sub-group that is no longer lactating will remain at the original index and the sub-group that continues lactating will be added at the end of the set of animal groups.

Description	Index number of the animal group for which lactation is to end. A value of zero denotes that each animal group should be processed in turn until the nominated number of cows has been dried off.	
Units		
Type	integer4	integer4
Parameter	group	number

4.5. join

Commences mating of a particular group of animals. If the animals are not empty females, or if they are too young, has no effect.

Description		sufficient age should be mated.	Genotype of the rams or bulls with which the animals are mated. Must match the name field of a member of the	genotypes property.	Length of the mating period.
Units					р
Type	integer4		string		integer4
Parameter	group		mate_to		mate_days

4.6. move

Changes the paddock to which an animal group is assigned.

Description	Index number of the animal group to be moved.	Name of the paddock to which the animal group is to be moved.
Units		
Type	integer4	string
Parameter	group	paddock

4.7. sell

Removes animals from the simulation.

Description	Index number of the animal group from which animals are to be removed. A value of zero denotes that each animal	group should be processed in turn until the nominated number of animals has been removed.	Number of animals to remove.
Units			
Type	integer4		integer4
Parameter	group		number

Component Description – STOCK

4.8. shear

Shears sheep. The event has no effect on cattle.

Description	Index number of the animal group to be shorn. A value of zero denotes that all animal groups should be processed. Denotes whether the main group of animals, suckling lambs, or both should be shorn. Feasible values are the null string (main group), 'adults' (main group), 'lambs' (suckling lambs), 'both' (both).
Units	
Type	integer4 string
Parameter	group sub_group

4.9. sort

Rearranges the list of animal groups in ascending order of tag value. This event has no parameters.

4.10. split

Creates two or more animal groups from the nominated group. One of these groups is placed at the end of the animal group list.

The division may only persist until the beginning of the next do_stock step, when sufficiently similar groups of animals are merged.

Description	Index number of the animal group to be split.	Feasible values are:	'age' All animals younger than <i>value</i> days are moved to a new group.	'weight' All animals with live weight less than value kg are moved to a new group.	'young' Only animals with suckling offspring are affected. Mothers with different sexes of young are divided, with	the group with all male offspring remaining in place. For mothers with twins, three groups are created; a	group with two male offspring, a group with two female offspring, and a group with one of each.	'number' value animals remain in place and the remainder form a new group	Threshold age or weight, or the number to be split, depending on the value of type. Ignored if type is 'young'.	
Units										
Type	integer4	string							double	
Parameter	group	type							value	

4.11. tag

Sets the tag value for an animal group.

Description	Index number of the animal group to be assigned a tag value.	Tag value to be assigned.
Units		
Type	integer4	integer4
Parameter	group	value

4.12. wean

Weans some or all of the lambs or calves from an animal group. The newly weaned animals are added to the end of the list of animal groups, with males and females in separate groups.

rameter Type Units Index num integer4 Index num group shou string Feasible variable	Description	ber of the animal group from which animals are to be removed. A value of zero denotes that each animal ld be processed in turn until the nominated number of lambs or calves has been weaned.	ilues are: Female and male lambs or calves are to be weaned.	Only female lambs or calves are to be weaned. Only male lambs or calves are to be weaned.	Number of lambs or calves to be weaned.
rameter Type integer4 string		Index numbe group should	Feasible valu		Number of la
rameter ir	Units				
arameter up	Type	integer4	string		integer4
Brou grou	Parameter	group	sex		number

5. Methods

None.

6. Published events

6.1. remove_herbage

Indicates the removal of herbage and seeds. This event is directed to each component instance that provides the Stock instance with a value for the plant2stock driving

Description	Mass of shoots removed in each of 5 digestibility classes.	Mass of unripe and ripe seeds removed.
Units	kg/ha	kg/ha
Type	double[]	double[]
Parameter	herbage	seed

6.2. add excreta

Indicates the excretion of faeces and urine into a paddock. Different instances of this event are directed to each component subscribing to it, with parameters depending upon the name of the paddock component to which the subscribing component belongs.

Description	faeces:	to be added.	n in faeces.	orus in faeces.	in faeces.		reted faeces:	gen in faeces.	shorus in faeces.	ur in faeces.		le.	က်	urine.	rine.	d urine.
	Organic matter in excreted faeces:	• Mass (as DM) of faeces to be added.	 Mass of organic nitrogen in faeces. 	 Mass of organic phosphorus in faeces. 	 Mass of organic sulphur in faeces. 	• Ash alkalinity in faeces.	Inorganic nutrients in excreted faeces:	 Mass of inorganic nitrogen in faeces. 	 Mass of inorganic phosphorus in faeces. 	 Mass of inorganic sulphur in faeces. 	Excreted urine:	 Volume of excreted urine. 	• Urea-N in excreted urine.	 Phosphate-P in excreted urine. 	 Sulphate-S in excreted urine. 	 Ash alkalinity in excreted urine.
Units		kg/ha	kg/ha	kg/ha	kg/ha	mol/ha		kg/ha	kg/ha	kg/ha		m³/ha	kg/ha	kg/ha	kg/ha	mol/ha
Type	record	double	double	double	double	double	record	double	double	double	record	double	double	double	double	double
Parameter	faeces_om	: weight	<i>u</i> :	d:	S:	$: ash_alk$	faeces_inorg	<i>u</i> :	d:	S:	urine	: volume	: urea	xod:	: 504	: ash alk

7. Driving properties

Description	Area of each paddock. Latitude (south is negative). Slope of each paddock.	Day length including civil twilight. Description of the pasture for use by the ruminant model.																			Consumption of supplementary feed by animals.	• Name of a paddock	 Amount of supplementary feed eaten by animals in this paddock.
Number	0+ 0+ 0+	1 0 1																			0-1		
Event: State	Initialisation Initialisation Initialisation	do_stock:0 do_stock:0																			$do_stock:0$		
Units	ha deg deg	hr	kg/ha -	kg/kg ko/ko	kg/kg	kg/kg mol/kg	1	1	•	ı		kg/ha		kg/kg	m kg/kg	m kg/kg	kg/kg	mol/kg	1				Жg
Type	double double double	double record record[]															double				5	String	double
Property	area latitude slope	daylength plant2stock · herhaoe	: dm : dmd	$: cp_conc$: s_conc	. prol_ag . ash alk	: height_ratio	: propn_green	: legume	: select_factor	: seed	: dm	: dmd	$: cp_conc$	$: p_conc$	$: s_conc$	$: prot_dg$	$: ash_alk$: height_ratio	: seed_class	supp_eaten	: радаоск	: eaten

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Description	Current time step.							Waterlogging index for each paddock.	Weather record.							
Number	1							+0								
Event:State	$do_stock:0$							$do_stock:0$	do_stock:0							
Units		р	S	S	р	S	S	ı		ွ	သွ	p/ww	p/mm	$MJ/m^2/d$	kPa	m/s
Type	record	integer4	integer4	double	integer4	integer4	double	double	record	double	double	double	double	double	double	double
Property	time	: startDay	: startSec	: startSecPart	: endDay	: endSec	: endSecPart	waterlog	weather	: maxt	: mint	: rain	: snow	: radn	pd λ :	: wind

If the following properties are not found, then alternative properties are subscribed to instead:

Description	Maximum air temperature.	Minimum air temperature.	Precipitation in all forms other than snow.	Average wind speed
Number	1	1	-	1
Event:State	$do_stock:0$	$do_stock:0$	$do_stock:0$	$do_stock:0$
Units	J _o	ွ	mm	m/s
Type	double	double	double	double
Alternative	maxt	mint	rain	wind
Property	weather	weather	weather	weather

8. Owned properties

All initialisation properties are readable. In addition, the following owned properties are available:

(a) Standard properties

Description	Fully-qualified name of the component.	Value is "Stock".	Value is "1.0".	Value is "CSIRO Plant Industry".	Denotes whether or not the component is active.	SDML description of the current state.
Units						
Type	string	string	string	string	Boolean	string
Property	пате	type	version	author	active	state

(b) Component-specific properties

Each entry in the following table describes between one and six variables: the named variable and five variants obtained by appending the texts: "_yng", "_all", "_tag", "_yng_all" and "_yng_tag'

- The variable obtained by appending "yng" is an array of the same type as the base variable. The array has one element for each animal group. Each element of the array denotes the value of the nominated quantity for unweaned lambs or calves of the corresponding animal group. If the animal group has no unweaned lambs or calves, the value is zero. For example, weight yng[4] gives the weight of unweaned lambs or calves in the fourth animal group (if any)
- The variable obtained by appending "_all" is a scalar that denotes an average or total of the quantity (as appropriate) over all animals in the component. Unweaned lambs or calves are excluded. For example, there is a weight_all variable of double type, which denotes the average weight of all animals, and number_yng_all gives the total number of unweaned lambs or calves.
- values less than or equal to zero and all unweaned lambs or calves are excluded. For example, weight tag[2] denotes the average weight of all animals with a tag value of The variable obtained by appending "_tag" is an array of the same type as the base variable. The size of this array is given by the highest tag value assigned to an animal group. Each element of the array denotes an average or total of the quantity (as appropriate) over all animals that have the corresponding tag value. Animals with tag

Note that the animal model will automatically merge and split groups of animals, so that the index position of a particular group of animals in the array variables will not necessarily remain constant.

yng	×	×	×		×	×
_tag	×	×	×	×	×	×
_all	×	×	×	×	×	×
Description		Age of animals, in months.	Fleece-free, conceptus-free weight.	Condition score at last parturition; zero if <i>lactating</i> =0	Current clean fleece weight.	Growth rate of clean fleece.
Units	p	1	kg		kg	kg/d
Lype	double[]	double[]	double[]	double[]	double[]	double[]
	age					

operty Type Units ore double[] - double[] - record[] kg/d double kg/d integer4 the integer the i	Component Description 21 Octs	1011 - 5100				I C IVIA	12 IVIAI 2000
re double[] - double[] - record[] - record[] - record[] - double kg/d double kg/d double kg/d double[] kg/d integer4[]	Property	Type	Units	Description	_all	tag	yng
double[] - record[] - record[] - record[] - record[] - record[] - record[] - kg/d double kg/d double kg/d double[] hm hm double[] hg/d do	cond_score	double[]	ı		×	×	×
double[]] record[] record[] double double double kg/d double[] m double[] m double[] m double[] kg/d double[] k	cp intake	double[]	kg/d	Crude protein intake per head.	×	×	×
record[] double kg/d double kg/d double kg/d double kg/d double kg/d double kg/d double[] m double[] m double[] record[] double[] kg/d integer4[]	dse	double[]		Dry sheep equivalents", based on potential intake.	×	×	×
double kg/d double[] m double[] trecord[] double kg/d double kg/d double kg/d double kg/d double[] trecord[] trecord[] trecord[] trecord[] trecord[] trecord[] kg/d double[] integer4[] integer4[] integer4[]	faeces	record[]		Faecal dry matter and nutrients per head.	×	×	×
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org record[] double kg/d double[]	$: ash_alk$	double	p/lom				
double kg/d double kg/d double[]	faeces_inorg	record[]		Inorganic nutrients excreted in faeces, per head.	×	×	×
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wth_diam double[]	fibre_diam	double[]	шm	Current average wool fibre diameter.	×	×	×
double[] kg record[] kg/d double kg/d double kg/d double kg/d double[] d e double[] kg double[] MJ/d double[] kg/d e double[] kg e double[] kg/d double[] kg/d integer4[] ing double[]	fibre_growth_diam	double[]	шm	Fibre diameter of the current day's wool growth.	×	×	×
record[] double kg/d double kg/d double kg/d double kg/d double[] d e double[] kg double[] MJ/d double[] MJ/d double[] kg/d integer4[] ing double[] ing double[]	fleece_wt	double[]	kg	Current greasy fleece weight.	×	×	×
double kg/d double kg/d double kg/d double kg/d double[]] d double[]] kg	intake	record[]		Total intake per head of dry matter and nutrients by each animal group.	×	×	×
double kg/d double kg/d double kg/d double[] d couble[] MJ/d double[] MJ/d double[] MJ/d double[] kg/d integer4[] integer4[] integer4[] integer4[] integer4[]	: weight	double	kg/d				
double kg/d double kg/d double kg/d double double double double double double double double mul/d double mul/d kg/d mul/d double muleger4 double muleger4 muleger	u:	double	kg/d				
double kg/d double mol/d double mol/d double double double mol/d kg mol/d double mol/d mol/d double mol/d kg mol/d mol	d:	double	kg/d				
## double mol/d double[]	S:	double	kg/d				
double[] d double[] kg kg double[] MJ/d double[] kg/d integer4[] kg/d integer4[] ss integer4[] double[] integer4[]	$: ash_alk$	double	p/lom				
e double[] kg e double[] MJ/d double[] MJ/d double[] kg/d integer4[] integer4[] integer4[] integer4[] integer4[] integer4[] integer4[] integer4[]	actating	double[]	p	If the animals are lactating, the number of days since birth of the lamb or calf, zero otherwise.	×	×	
e double[] MJ/d double[] MJ/d double[] kg/d integer4[] integer4[] integer4[] integer4[] integer4[]	nax_prev_wt	double[]	kg	Maximum previous basal weight (fleece-free, conceptus-free) attained by each animal	×	×	×
e double[] MJ/d double[] MJ/d double[] kg/d integer4[] ses double[] sinteger4 integer4[] ing double[]	,	,	:	group.			
double[] MJ/d double[] kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d	ne_intake	double[]	MJ/d	Intake per head of metabolizable energy.	×	×	×
double[] kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d kg/d k	nilk_me	double[]	MJ/d	Metabolizable energy produced in milk (per head) by each animal group	×	×	
le	nilk wt	double[]	kg/d	Weight of milk produced per head, on a 4% fat-corrected basis.	×	×	
ses double[] 25 integer4 integer4[] double[]	10 female	integer4[]		Number of female animals in each animal group.	×	×	×
integer4	no_foetuses	double[]		Number of foetuses per head in each animal group.	×	×	
integer4[] ing double[]	no_groups	integer4		Number of animal groups.			
double[]	no_male	integer4[]		Number of male animals in each animal group.	×	×	×
1040404	no_suckling	double[]		Number of unweaned lambs or calves per head in each animal group.	×	×	
	number	integer4[]		Number of animals in each animal group.	×	×	×
		, ,					

		CHIES	Description	_all	tag	yng
past_intake	record[]		Intake per head of pasture dry matter and nutrients by each animal group.	×	×	×
: weight	double	kg/d				
<i>u</i> :	double	kg/d				
d:	double	kg/d				
s:	double	kg/d				
: ash_alk	double	p/lom				
pregnant	double[]	р	If the animals are pregnant, the number of days since conception; zero otherwise.	×	×	
priority	integer4[]		Priority score assigned to each animal group; used in drafting.			
rdp intake	double	kg/d	Intake per head of rumen-degradable protein	×	×	×
rdp_reqd	double	kg/d	Requirement per head of rumen-degradable protein	×	×	×
retained n	double[]	kg/d	Nitrogen retained within the animals, on a per-head basis.	×	×	×
retained p	double[]	kg/d	Phosphorus retained within the animals, on a per-head basis.	×	×	×
retained s	double[]	kg/d	Sulphur retained within the animals, on a per-head basis.	×	×	×
- xes	string[])	See the sex field of the sheep and cattle initialisation variables. Returns "heifer" for cows			
			under two years of age.			
supp_eaten	record[]		Consumption of supplementary feed by animals.			
: paddock	string		 Name of a paddock 			
: eaten	double	kg	 Amount of supplementary feed eaten by animals in this paddock. 			
supp_intake	record[]		Intake per head of supplement dry matter and nutrients by each animal group.	×	×	×
: weight	double	kg/d				
u:	double	kg/d				
d:	double	kg/d				
S:	double	kg/d				
$: ash_alk$	double	p/lom				
tag_no	integer4[]		Tag value assigned to each animal group.			
trampling	double	kg/ha	Mass of grazers per unit area. The value returned depends on the requesting component.			
$urine_n$	double[]	kg/d	Urinary nitrogen output per head.	×	×	×
urine_p	double[]	kg/d	Urinary phosphorus output per head.	×	×	×
urine s	double[]	kg/d	Urinary sulphur output per head.	×	×	×
weight	double[]	kg	Average live weight of each animal group.	×	×	×
wt_change	double[]	kg/d	Rate of change of base weight of each animal group.	×	×	×

Component Description - STOCK

Configuration Details

Stock Component Description A.D. Moore A.D. Moore Microsoft Word 2002 15 Mar 2006

Created by: Modified by:

Processor: Printed:

Revision History

Version	Date	Changes
0.1	12 Dec 1997 First draft	First draft
0.2	17 Dec 1997 Second draft	Second draft
0.3	4 Aug 1998	Third draft
0.4		Revised to match pre-release version of component. *_tag properties added
0.5	15 Dec 2003	supp_eaten added
1.6	30 May 2005	30 May 2005 Changes to representation of genotypes described
1.7	9 March 2005	9 March 2005 Minor cleanup

Document Distribution Policy

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