Comparison of the effects of high and low milk-replacer feeding regimens on health and growth of crossbred dairy heifers

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Table S1. Descriptive data for the candidate variables tested for the multivariate models

Variables tested	Descriptive statistics	Final variable tested for inclusion
Age	Age at each examination preweaning was included in each of the repeated measures models. It was normally distributed with a mean of 32.3 ± 15 days	Continuous
Calving week	Week of birth was recorded as the calving group to give a description of where calves were born in the calving block: • Week 1 – 33 (17.6 %) • Week 2 – 17 (9.1 %) • Week 3 – 37 (19.8%) • Week 4 – 27 (14.4 %) • Week 5 – 23 (12.3 %) • Week 6 – 10 (5.3 %) • Week 7 – 20 (10.7 %) • Week 8 – 15 (8.0 %) • Week 9 – 5 (2.7 %)	Tested as an ordered factor but the raw data were found to give a significantly better fit in all models and so this was used in the model results presented.
Breed	Calf breed was recorded. Friesian n=96 (52.1 %) Jersey cross n=80 (41.7 %) Montbelliard cross n=1 (0.5%) Hereford cross n=11 (5.2 %)	Categorical
Breed Grouped Breed Data	No differences were noted between the Friesian calves and the Montbelliarde or Hereford crosses and so these animals were grouped to give Friesian type n=108 (58.3 %) Jersey cross n=80 (41.7 %)	Categorical
Passive Transfer	Total protein at recruitment was used to record passive transfer of immunity. It was normally distributed with a mean of 60.8 ± 9.5 mg/ml	Continuous

Dystocia		Assisted calvings were recorded by the farm staff. 18 heifers (9.6 %) of the cohort had calving assistance.	Binary Y/N
Size at Recruitment Week 1	Weight	Normally distributed with a mean of $41.3 \pm 8 \text{ kg}$	Continuous
	Height	Normally distributed with a mean of 74.9 ± 4.1 cm	Continuous
	Length	Normally distributed with a mean of 60 ± 3.6 cm	Continuous
	Age at Recruitment	Normally distributed with a mean of $4.5 \pm 2.7 \text{ kg}$	Continuous
Bovine Respiratory Disease	Disease pre- weaning	49/192 (26.2%) of calves had respiratory disease preweaning.	Binary Y/N
	Total score above the threshold for diagnosis pre- weaning	Data were not normally distributed, the range was 0-7 with a median of 0 and an interquartile range of 0-1.	Treating the data as an ordered factor was tested but using the raw data significantly improved the model in all cases
Diarrhoea	Disease pre- weaning	121/192 calves (64.7%) had diarrhoea pre-weaning	Binary Y/N
	Total score above the threshold for diagnosis pre- weaning	Data were not normally distributed, the range was 0-4 with a median of 0 and an interquartile range of 0-1.	Treating the data as an ordered factor was tested but using the raw data significantly improved the model in all cases.