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Genetic and environmental factors influencing milk, protein and fat yields of pasture-based dairy cows in Tasmania

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1 **Genetic and environmental factors influencing milk, protein and fat yields of pasture-**

2 **based dairy cows in Tasmania, Australia**

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- 12 +61-3-6226-2642
- 13
- 14 **Running head**: Milk yield and composition in grazing cows
- 15

16 **Abstract**. The objective of this study was to provide an update on milk production 17 performance, heritability, genetic and phenotypic correlations among production traits which 18 are valuable for management, breeding and selection decisions in pasture-based dairy systems. 19 The study utilised a total of 106,990 lactation records of Holstein-Friesian (FF), Jersey (JJ) 20 and their crossbreds (HF) from 428 Tasmanian dairy herds collected between 2000 and 2005. 21 The data were analysed using the least squares approach with a general linear model and 22 restricted maximum likelihood approach with a linear animal model. Results indicated highly 23 significant (P<0.01) effects of breed, herd size, cow's parity, season and year of calving on 24 milk, protein and fat yields. Average milk and protein yields per cow per lactation were 25 highest in the FF breed (5,212 litres and 171 kg, respectively) and lowest in the JJ breed 26 (3,713 litres and 143 kg, respectively). FF cows also produced 13.5kg more milk fat than JJ 27 and HF cows. Furthermore, milk, fat and protein yields were highest for cows calving during 28 spring and lowest for autumn-calving cows. It was also evident that cows in very large herds 29 (>1110 cows/herd) out-produced those in smaller herds. Heritability was highest for milk 30 yield and lowest for somatic cell count ranging from 0.28 to 0.41. Genetic and phenotypic 31 correlations between milk, fat and protein yields ranged from 0.41 to 0.85, and 0.66 to 0.92, 32 respectively. On the other hand, genetic and phenotypic correlations between the log of 33 somatic cell count and the production traits ranged from 0.03 to 0.09 and -0.03 to -0.05. We 34 conclude that breed, herd size, parity, season and year of calving were among the main factors 35 correlated with the productivity of dairy cows in Tasmania and adjustments for these factors 36 would be mandatory for any unbiased comparison of lactation performance within and 37 between pasture-based dairy production systems. The practical application of this information 38 would be valuable to dairy farmers for decisions related to breeding, selection and 39 management of their herds.

40 **Keywords**: Pasture-based cows, Holstein-Friesian, Genetics, Tasmania, Milk, Protein, Fat

41 **Introduction**

42 The dairy industry in Australia is very important to the agricultural sector of the economy. 43 With an ex-factory value of \$9.1 billion and farm gate value of \$3.2 billion, it is the third 44 biggest rural industry behind beef and wheat (Dairy Australia 2006). Milk production is 45 concentrated in the South-Eastern corner of Australia, with the states of Victoria, Tasmania 46 and South Australia accounting for 78% of the national output. Like in Victoria, dairying in 47 Tasmania is characterised by seasonal, low-input, pasture-based milk production reliant on 48 family labour. However, ninety percent of dairy farms use supplementary feeds such as hay, 49 silage and concentrate to augment seasonal shortages in grass production (Dairy Australia 50 2006). Deregulation of the production sector in 2000 as well as widespread drought in 2002- 51 03 led to substantial restructuring such as reduction in farm numbers, high cost of grain 52 supplements and increased herd sizes (ABARE 2003). Although the Holstein-Friesian (FF) 53 constitutes about 70% of the dairy breeds in Tasmania, there are growing numbers of other 54 breeds including the Jersey (JJ), Friesian and Jersey crossbreds (FJ), Guernsey (GG), Ilawara 55 (II) and Australian Reds (RR). Climatic factors also differ between dairy locations. Although 56 milk production from Tasmania constitutes a small proportion (7%) of the national output, 57 nearly 90% of the milk produced in Tasmania is processed for export (Dairy Australia 2006), 58 compared to 30-40% in New South Wales (NSW), Queensland and Western Australia 59 (Edward 2005). Industry statistics also indicates that the Tasmanian dairy industry is growing 60 at rates comparable to those of major dairy producing states of Australia. Tasmania's typical 61 climate offers an opportunity for year round-milk production, where precipitation is not 62 limiting. Consequently, dairy production in Tasmania will continue to play a significant role 63 in both domestic and overseas export of milk products.

64 Milk is synthesised by secretory cells in the mammary glands of lactating animals primarily 65 as nutrition for the young. Factors affecting milk production in dairy animals include genetic 66 (Tekerli *et al.* 2000), environmental and management factors, (Msanga *et al.* 2000). Several 67 studies (Madgwick and Goddard 1989, Dobos *et al.* 2001; Wales *et al.* 2006) have identified 68 factors affecting milk production of dairy cows in Victoria and other parts of Australia. 69 Tasmanian dairy farmers are one of the most efficient in pasture-based dairying (Dairy 70 Australia (2004). However, except for some performance indicators of the industry compiled 71 by the Department of Primary Industry and Water (DPIW 2005) see (Appendix 1) and Dairy 72 Australia (2005), there is paucity of recent information on the key driving factors influencing 73 dairy cattle performance in Tasmania. Therefore the objectives of this study were; to 74 characterise and quantify the milk production of pasture-based dairy cows in Tasmania as 75 influenced by breed, parity, location, herd size, season, parity, year and their interactions; to 76 identify the critical management factors underpinning milk and milk component yields and to 77 estimate the heritability of production traits and somatic cell count.

78

79 **MATERIALS AND METHODS**

80 *Site and climatic conditions*

81 Tasmania is Australia's southern-most state with a land mass of 68,000 sq km, located at 82 latitude 42° South, longitude 147° East and lies completely within the temperate zone. The 83 summer, autumn, winter and spring seasons are in the months of December-February, March– 84 May, June–August and September–November, respectively. Average maximum temperatures 85 are 21^0C (70⁰ F) and 12^0C (54⁰ F) in summer and winter respectively. Summer is warm with 86 sunny days and mild evenings, while autumn is cool with frosty nights and occasional storms. 87 Winter is mild with occasional snows on the higher mountain peaks. The annual rainfall 88 varies from 626 millimeters (25 inches) in the south to 2,400 millimeters (94 inches) on the

89 North-West Coast. The prevailing weather pattern creates a rain shadow in the west to east 90 direction leaving the East coast always warmer and drier than the rest of the state.

91

92 *Data source and editing*

93 The data used in this study were obtained from TasHerd, which is the contracted herd 94 recording agency for the Australian Dairy Herd Improvement Scheme (ADHIS) in Tasmania. 95 The data were from 428 dairy herds and consisted of 130, 366 observations on total milk, fat, 96 protein and somatic cell count (SCC) yield records of purebred Holstein-Friesian , Jersey , 97 Guernsey, Illawarra, and Australian Reds as well as crosses of Holstein-Friesian and Jersey 98 cows of different ages, season of calving, parity, and lactation length.

99

100 Two data sets were created. **DATA1** which explored non-genetic factors affecting production 101 traits used records from three breeds namely; FF, JJ and FJ, these being the predominant 102 breeds in Tasmania. Small data size and incomplete records on the other breeds in regional 103 data made genetic computation difficult, hence only records comprising of the FF breed were 104 utilised for genetic analysis in **DATA2**. For each cow, records of cow number, birth date, 105 calving date, 305-day milk yield, protein and fat yields, milksolids yield, lactation length, and 106 herd number were available. This information was used to determine cow age, calving season, 107 parity, and herd sizes. Daily milk, fat and protein yields were obtained by dividing the total 108 milk and milk component yields by the lactation length. Percentage milk fat and milk protein 109 were obtained by dividing total fat and total protein yields by total milk yield. Lactations with 110 incomplete records i.e. missing birth date, calving date, milk or milk component yields were 111 deleted. Records of cows with lactation length <100 days were also excluded from the 112 analyses. DATA1 finally consisted of 106, 990 records from 428 herds, over six production 113 years. Parities greater than four were pooled as parity5. To protect farm identities, coded herd 114 numbers and postcodes of dairy farms were used. The final data set consisting of thirty 115 postcodes was divided into 6 subsets, herein referred to as locations, based on existing local 116 council areas. For instance, herds with postcodes 7260-7265 were grouped into the North-East 117 location (Table 1). Herds were classified to correspond with the state's average herd size of 118 250 cows. Herd groups were below, similar to, twice and four times the state's average herd 119 size. The four herd size classes were; 1-210, 211-575, 576-1100 and larger than 1100 cows 120 per herd designated as small, medium, large and very large herds, respectively. For DATA2, 121 additional data omitted from the analysis were all breeds except FF, cows with parity >5 and 122 305 day milk yield <1,200 L leaving a total of 65,914 records. Parity >2 were pooled and 123 labelled as parity3. Dairy farm statistics of Tasmania is presented in Appendix 1. Least square 124 means and summary statistics of the traits for DATA1 and DATA2 are presented in Tables 1 125 and 3 respectively.

126

127 *Statistical analysis*

128 *DATA1.* General linear models (GLM) procedure in SAS (SAS 2002) was utilised to 129 compute least square means, standard errors (s.e.) and coefficient of variation (c.v.) of the 130 traits. Location, herd size, breed, calving year, calving season and parity were fitted as fixed 131 effects while age and lactation length were included as covariates. All possible interactions 132 between the fixed effects were included in the original model, but non-significant interactions 133 were dropped from the final model. The model used to describe each lactation record was:

$$
\begin{array}{c|c} \begin{array}{c} \displaystyle Y_{ijklmnopq} = & \displaystyle \mu + L_i + H_j + S_k + Y_l + B_m + P_n + (BS)_{km} + (BY)_{lm} + (SP)_{kn} + (BP)_{mn} + (PSY)_{nkl} + b_l (L\underline{L}_{ijklmno} - 136 & \displaystyle \overline{L}L \, \end{array} \end{array} \begin{array}{c} \displaystyle \sum_{ijklmnop} \displaystyle -\bar{A})^2 + e_{ijlkmnopq} \end{array} \end{array}
$$

137

134

138 where $Y_{ijklmopq}$ is the $_{ijklmopq}$ th observation of the dependent variables (milk, fat and protein 139 yields and log of somatic cell count), with fixed effects L_i of ith location (i=1,2.. 6); H_i of jth herd size (j=1,2…4); S_k of k^{th} season of calving (k=1,2… 4); Y_l of l^{th} year of calving (l=1,2… 141 6); B_m of mth breed (m=1,2... 3) and P_n of nth parity (n= 1,2.. 5). First order interaction effects 142 were (BS)_{km} of breed and season, $(BY)_{lm}$ of breed and year, $(SP)_{kn}$ of season and parity and 143 (BP)nm of breed and parity and second order interaction effects of season, year and parity 144 (PSY)_{nkl}. . There was a total of 7, 11, 13, 9, and 81 combinations of BS, BY SP, BP and PSY 145 subclasses respectively. The model terms $b_1(L\underline{L}_{ijklmn0} - \overline{L}L)^2$, and $b_2(A_{ijklmn0p} - \overline{A})^2$ represent 146 the fitting of lactation length (LL) and age of the cow (A) as covariates with b_1 and b_2 as their 147 partial regression coefficients respectively; μ is overall mean and $e_{iikmmona}$ is the random 148 sampling error with mean zero and variance σ_e^2 . Means were compared using the least 149 significant difference technique of SAS GLM procedure.

150

151 *DATA2.* In the genetic analysis of milk, fat, protein and somatic cell count, Y, P, S H and YS 152 interaction (as previously defined), were fitted as fixed effects, while cow was used as a 153 random effect. Age at calving (Age) and lactation length (LL) were included as covariates in 154 all analyses. All traits were first analysed with a univariate animal model in ASReml 155 (Gilmour *et al.* 2006) to obtain start up values for the covariance structures in subsequent 156 analyses. A single multivariate analysis using an animal model was performed in ASReml to 157 estimate heritabilities as well as phenotypic (r_P) and genetic (r_G) correlations of production 158 traits (milk, fat and protein yields) with their associated standard errors (Table 4), while the 159 log- transformed values of somatic cell count were analysed as a univariate factor.

160 The animal model is:

161
$$
\begin{vmatrix}\n\frac{1}{2} & \frac{1}{2} & \frac{1}{2
$$

163

164 where $Y_{ijklmno}$ represents the dependent variables total milk, fat, protein and log of SCC, μ is 165 the population mean, H_i , S_j , Y_k , P_l and YS_{jk} , are the fixed effects of the variables on i^{th} herd 166 (*H*=1,2…216), j^{th} calving season (*S*= (1,2…4), k^{th} calving year (*Y*=1,2 …6), l^{th} parity 167 ($P=1,2...3$), and $jkth$ first order interaction of calving year and season ($YS= 1,2...46$), 168 respectively, b_1 and b_2 are the regression coefficients of lactation length (*L*) and age at calving 169 (*A*) respectively, a_{ijklmn} is the random additive genetic effect and $e_{iklmnopj}$ is the random residual 170 error. A pedigree file tracing ancestry to the last $5th$ generation was included in the analysis of 171 DATA2.

172

173 In matrix notation the model can be written as 174

175 *y*=**X***b*+**Z***a*+**W***p*+*e*,

175
176
177 177 where y is the vector of observations, *b* is the vector of fixed effects H_i , S_j , Y_k , P_l and YS_{jk} , 178 *,* b₁*LL_i* and b₂*Age_i*, *a* is the vector of a_{ij} additive genetic effects random animal effects, *p* is the 179 vector of permanent environmental effects for cows with 305 day records, *e* is the vector of 180 random residual effects and **X**, **Z**, and **W** are the incidence matrices which relate records to 181 fixed, animal and permanent environmental effects respectively. It is assumed that the 182 permanent environment and the residual effects are independently distributed with means of 183 zero and variance σ_{pe}^2 and σ_{e}^2 respectively. The variances of the random additive, permanent 184 environment and random error effects are A σ_a^2 , I σ_{pe}^2 and I $\sigma_e^2 = R$ and variance of *y*

185 $var(y) = \mathbf{Z} \mathbf{A} \mathbf{Z'}$ $\sigma_a^2 + \mathbf{W} \mathbf{I} \sigma_{pe}^2 \mathbf{W'} + \mathbf{R}$

- 186
- 187
- 188 189

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where G is covariance matrix of the random regression coefficients, assumed to be the same 194 for all cows; *A* is additive genetic relationship matrix among the animals; ⊗ is Kronecker 195 product function (Searle 1982); *I* is identity matrix; and *U* is unstructured matrix with 196 elements that define the traits.

197

198 The mixed model equation for this model would be

199
$$
\begin{pmatrix} X^{\top}R^{-1}X & X^{\top}R^{-1}Z & X^{\top}R^{-1}W \\ Z^{\top}R^{-1}X & Z^{\top}R^{-1}Z + G^{-1} \otimes A^{-1} & Z^{\top}R^{-1}W \\ W^{\top}R^{-}X & W^{\top}R^{-1}Z & W^{\top}R^{-1}W + Ik \end{pmatrix} \begin{pmatrix} \hat{b} \\ \hat{a} \\ \hat{a} \\ \hat{a} \\ \hat{p} \end{pmatrix} = \begin{pmatrix} X^{\top}R^{-1}y \\ Z^{\top}R^{-1}y \\ W^{\top}R^{-1}y \end{pmatrix}
$$

200

201 Where $k = I/\sigma_p^2$ was assumed constant across traits.

202

203 **RESULTS**

```
204 DATA1
```
205 Least squares means estimates using Model 1 are presented in Tables 1 and 2. Milk, fat and 206 protein yields were significantly different (*P*<0.01) between cattle breeds, parity and season 207 of calving except for fat yield which did not differ between the Jerseys and crossbred cows. 208 The model fitted explained 40-43% of the variations due to environmental and management 209 factors. Lactation length accounted for about 19.3% of the total sums of squares. The other 210 factors influencing productivity were herd size, location, and cow age in order of decreasing 211 magnitude except for fat and protein yields where parity had greater influence than genotype 212 and location.

213

214 *Calving Year.* The effect of year of calving on milk yield, milk composition and the 215 log of somatic cell count of the cows is shown on Table 1. Total milk yield (L/lactation) was 216 significantly different (*P*<0.01) between all calving years except 2001 and 2003. Total milk 217 yield progressively increased annually by an average of 2.5-5.7% except for a 101L decline 218 from 2002 to 2003. Similar trends were observed for fat and protein yields. There was also a 219 general increase in annual total milk solids of 16-21 kg from 2001 to 2005 except for 2002 220 and 2003. Daily yields of milk, fat, protein and milksolids also progressively increased with 221 advances in calving year (except in 2003).

222

223 *Insert Table 1 here*

224 *Breed*. Milk yield was significantly different (*P*<0.01) between the three breeds 225 evaluated in this study (Table 1). Total milk, fat, protein and milksolids yields were highest in 226 FF cows, while JJ and FJ cows had similar fat and milk solids yields. JJ cows had the least 227 protein yield and somatic cell count. Differences between the FF and JJ in milk, fat and 228 protein yields were 1499 L, 13 kg and 28 kg respectively. JJ and FJ cows produced 29% and 229 18% less milk/lactation respectively, than FF cows. Protein yield followed a similar pattern. 230 (Table 1).

231

232 *Parity.* Parity significantly (P>0.01) influenced all milk production parameters (Table 233 1). Milk, fat, protein and milksolids yields were highest in cows with parity >4 and lowest in 234 first parity cows. There was an observed increase in milk, fat, protein and milksolids yields as 235 parity increased from 1 to >4, while the lowest SCC was observed in parity >4 cows (Table 1). 236 Milk yield differences between first vs second parity and second vs third parity cows were 237 538 L and 595 L respectively, compared with those between third vs fourth and fourth vs 238 parity>4 cows (211 L and 193 L, respectively).. Total milksolids increased from 277 kg in 239 primiparous cows, to 407 kg in cows with parities >4.

240

241 *Location.* The effect of location on yields of milk, fat, protein and milksolids are 242 shown on (Table 1). Milk and protein yields per cow were highest in FNWest, while cows in 243 KIsland produced the least milk, fat, protein and milksolids yields and had the highest SCC. 244 The difference in milk yield/cow between the dairy locations in the north-west (FNWest and 245 NWest) and the NEast was 98 L, the difference in milk yield between all the northern 246 locations and the South was 697L. Dairy herds in mainland Tasmania out-produced herds in 247 KIsland by 1274 L of milk, 48 kg of fat and 44 kg of protein per cow per year. Milk fat and 248 milksolids yields were highest in CntNorth and FNWest.

249

250 *Herd Size.* Significant variation in milk production due to differences in herd size was 251 observed. Cows in Very Large herds produced the most milk, fat, protein and milksolids 252 while Medium herds produced the least milk and protein yields (Table 1). Milk fat yield and 253 average SCC did not differ between cows in Medium and Small herds. Cows in the Small and 254 Large herds had the highest and lowest SCC respectively.

255

256 *Calving Season.* Spring-calving cows produced significantly more milk, fat, protein 257 and milksolids than Autumn and Summer-calving cows (Table 1). On the other hand, SCC 258 was highest in Summer-calving cows and lowest in Autumn-calving cows. Milk yield 259 difference between Spring and Autumn-calving cows was 676 litres, while fat and protein 260 yields between cows calving in both seasons differed by approximately 25 kg. Daily milk, fat,

261 protein and milksolids yields followed the same trend.

262

263 *Insert Table 2 here*

264

265 *Combined effect and interaction between Bbreed andby calving year interaction.* 266 Least squares means for combinations of breed and calving year are shown in Table 2. The 267 differences in total milk, fat and protein yields/cow between were significantly lower in JJ 268 compared with FF and FJ breeds. Within all breeds, total milk yield increased in all calving 269 years except 2003 when it fell. In all calving years, FF cows produced the most milk and JJ 270 the least. The general decrease in total milk yield in 2003 was lower in FF cows compared to 271 FJ and JJ breeds. The subsequent recovery and increase in milk yield in 2004 was 353L in FJ 272 compared with 288L and 149L in FF and JJ breeds, respectively. Although fat and protein 273 production patterns were similar among the breeds across calving years, gains between years 274 were slightly lower in JJ compared with the other breeds. SCC increased during the calving 275 year 2001 to 2002 in FJ and FF breeds, but declined in JJ breed. No interaction effects were 276 detected between breed and calving years for daily milk and milksolids yields. In contrast, the 277 difference in total milk yield between FF and FJ for all calving years averaged 958 L, while 278 that between FF and JJ averaged 1497 L. Similarly, the differences in protein yields between 279 FF *vs* FJ and FF *vs* JJ were 20 kg and 28 kg, respectively.

280

281 Milk yield (L) differences between the highest and lowest calving years for each breed 282 were: 818, 786, and 352 for FF, FJ and JJ breeds respectively. Milk fat yield declined by 283 between 3 and 4 kg in FF and JJ breeds from 2002 to 2003, but increased by 1 kg in FJ during 284 the same period. JJ cows produced more fat than FJ in 2000 to 2002, both produced equal

285 amounts of fat in 2003 but the FJ produced more fat than Jerseys in the subsequent calving 286 years. Of all the breeds, only the FJ produced more fat in 2003 than in 2002. Milk protein 287 yield was similar between JJ and FJ breeds in 2000-2002 but the latter produced 6-15 kg more 288 milk protein in 2003-2005. Milk protein yield was lower although not significantly (*P*>0.05) 289 in 2002 than 2003 in all breeds except FJ. The FF breed produced milk with higher SCC in 290 2000-2001 than the other breeds while FJ produced milk with the highest SCC in 2002-2005.

- 291
-

292 *Combined effect and interaction between Ccalving year andby calving season* 293 *interaction.* Milk yield/lactation for all years averaged 4845, 4463, 4249 and 4094 for spring, 294 winter, summer and autumn calving cows, respectively. Reduction in total milksolids in 295 autumn and summer-calving cows, due to the drought of 2003 were 7.5 and 18.2 kg 296 respectively, while spring and winter-calving cows produced 8 and 11 kg more milksolids 297 during the same year. With the exception of spring-calving cows in 2002, 2004 and 2005, 298 SCC was highest in summer-calving cows and lowest in autumn calvers.

299

300 *Combined effect and interaction between Bbreed by and calving season interaction.* 301 Least squares means for combinations of breed and calving season is shown in Table 2. FF 302 cows produced the most milk and milk components in all calving seasons and years. Total 303 milk yield per lactation was lowest in autumn-calving JJ cows. In contrast, milk fat yield was 304 highest in JJ and FJ cows that calved in spring and lowest in autumn calvings in all breeds. FF 305 cows that calved during autumn and spring produced the most milk protein and milk solids, 306 while autumn-calving JJ cows produced the least. SCC was highest in FJ cows that calved 307 during summer and lowest in autumn calvings.

308

309 *DATA2*

310 Unadjusted means of total milk, fat and protein yields (Table 3) were highest in third and 311 lowest in first parity FF cows. Log of SCC were similar across parities. Parity three FF cows 312 produced 651 L more milk at 305 day than their second parity counterparts while the latter 313 produced 661 L more milk than the first parity cows. Mean 305 day milk fat and protein 314 yields were highest in autumn calving FF cows. Milk yield was lowest in spring-calving cows 315 while fat and protein yields were lowest in summer calvers. Holstein Friesian cows calving in 316 autumn produced 614 L more milk than those that calved in winter. Winter-calving cows 317 produced 185 and 105 L more milk at 305d than those calving in spring and summer 318 respectively. Autumn calving FF cows produced 7% and 10% more total fat and protein 319 respectively, than their winter-calving counterparts, but the difference between summer and 320 spring calving cows was marginal.

321

322 Milk, fat and protein yields per cow increased annually from 2000 to 2005 except for a slight 323 production dip in 2002. Average annual rates of increase in yields were 3%, 3% and 4% for 324 milk, fat and protein respectively. Milk and protein yields/cow were highest in the NWest, 325 followed by FNWest and lowest in KIsland. On the other hand, FF cows in the CntNorth 326 produced the highest quantity of milk fat followed by cows in the NWest and FNWest with 327 cows in the KIsland again producing the least. The difference in milk yield/cow between the 328 highest and the lowest producing locations was 1376 L. Fat and protein yields (kg/cow) 329 differed less dramatically and ranged between 237-202 and 195–152, respectively. Somatic 330 cell counts were similar between parity groups, production years and locations.

331

332 *Insert Table 3 here*

333

Heritability of 305d milk in FF cows was highest followed by that of 305d fat yield while h^2 335 336 of SCC was lowest. Phenotypic correlations among production traits ranged from 0.66 (milk 337 *vs.* fat) to 0.92 (milk vs. protein) while that between production traits and SCC ranged from - 338 0.03 (protein vs. SCC) to -0.05 for SCC *vs.* fat (Table 4). Phenotypic correlation between fat 339 and protein was higher than that between milk and fat. Similarly, genetic correlation was 340 highest between milk and protein being 0.85 and lowest between fat and SCC at 0.03. 341 Phenotypic correlations between the milk, fat and protein yields were generally higher than 342 the corresponding genetic correlations.

343

344 *Insert Table 4 here*

345

346 **DISCUSSION**

347 *Goodness of model fitting*

348 The model fitted explained between 42-45% of the variation due to the factors tested 349 for all traits considered in the study (Table 1). This would imply that there were other 350 unaccounted explanatory variables beyond the scope of the fitted model. Such variables 351 would include temporary environmental factors like feed intake, feed quality, milking 352 frequency, housing condition, diseases and other management factors. These details about 353 herd management were not available in the data used in this study. It has been reported that 354 management factors due to individual farmer experience and openness to adoption of 355 scientific and technological tools can have tremendous impact on dairy cow productivity even 356 when animals of similar breed and production merit have been used (Chapman *et al.* 2004). 357 The results of this study should therefore be interpreted in the light of the available data and 358 tested factors.

359

360 *Calving year, calving season and their interactions*

361 Total yields of milk, fat and protein in this study were higher than the values reported for low-362 and high-bodyweight Holstein-Friesian cows in New-Zealand (Lopez *et al.* 2001), but lower 363 than the values reported by Bargo *et al.* (2002) and García *et al.* (1998; 2007) for high merit 364 cows on pasture allowance and concentrate supplementation in individual *vs*. group feeding 365 trials, respectively. Our results on total milk yield per lactation were however in agreement 366 with the findings of Grainger (1990) and García and Holmes (2001). The latter reported 367 average milk yields ranging from 4982-5409 L, s.e.=85.7 in autumn and spring-calving FF 368 cows. The higher responses in milk yield and milk component yields under experimental 369 conditions compared with aggregate data emanating from large number of cows from multiple 370 herds over diverse locations were expected. For instance, Bargo *et al.* (2002) and García *et al.* 371 (2007), offered concentrate at 1kg/4kg milk yield and 3-7kg as fed/cow/day to twenty-four 372 and fifty grazing cows respectively, whereas we evaluated data on 103,366 cows. Figures on 373 annual increases in milk yield in this study were generally lower than the national averages 374 (Dairy Australia 2006, DPIW 2005), but annual milk yields in 2000 to 2001 and 2003 to 2004 375 (Table 1) were well in agreement with published figures. The restriction of our data sets to 376 production records of only three genotypes could partly account for the discrepancy with the 377 national figures. Furthermore, differences in milk yield per cow due to higher use of 378 concentrate feeds in the states of Victoria and New South Wales (Dairy Australia 2006) may 379 also partly explain the lower milk and component yields in Tasmania. The decline in 380 production in 2002/03 calving season was attributed to feed shortages from the severe and 381 widespread drought of that season. Climatic factors such as low rainfall and adverse 382 temperatures have negative effect on milk yield in temperate cows through the 383 physiologically induced depressed feed intake (Msanga *et al.* 2000). Analysed climatic data 384 (Geography of Tasmania 2008) showed that maximum temperature was significantly lower

385 (*P*<0.05) in 2004 than in other years, while mean annual rainfall was 612.2±33.9 mm in 2002 386 to 2003 calving season compared with 780.4 mm in other years. Reduced rainfall could 387 depress pasture dry matter (DM) yield and metabolisable energy (ME) content, thus reducing 388 energy intake and productivity of pasture-based cows (Walker *et al.* 2004). Differences in 389 milk yield between calving years have been reported (Dairy Australia 2005, Msanga *et al.* 390 2000). Unlike in this study, White *et al*. (2002) found no significant interactions between 391 calving season and other factors.

392

393 Our results are also in agreement with seasonal variations in milk and milksolids yields 394 reported by García *et al.* (1998) for pasture-based cows. They reported that autumn-calving 395 FF cows produced significantly more milksolids than spring-calving FF cows due to the effect 396 of lush pasture with higher ME in spring (spring hump) and extended lactation due to greater 397 persistency. Typically, dairy cows attain peak yields between 3-8 weeks *postpartum* (Tekerli 398 *et al.* 2000). In pasture–based, winter-calving systems, it has been reported that the peak 399 month of milk production coincides with spring when production almost doubled that of the 400 lowest months between May and July (Edward 2005). Our findings revealed that lactation 401 length was significantly longer (p<0.05) in autumn-calving, compared with spring-calving FF 402 cows (306±0.34 *v.* 269±0.16 days). On the other hand, White *et al.* (2002) reported that 403 autumn and spring-calving cows had similar milksolids yields in northern Victoria. 404 Differences in production traits during different calving season reflect seasonal variations in 405 diverse calving systems practiced all over Australia aimed at minimising feed cost by 406 matching peak nutrient requirement for lactation with the period of highest availability of ME 407 from pastures (Walker *et al.* 2004). Although most farms practice split calving system, 408 percentage calving patterns of dairy herds in Tasmania were 51% and 38% for winter and 409 spring calvings, respectively.

410

412 Milk yield/lactation of FF cows reported in this study was comparatively lower than the 413 performance of the North American FF strains but similar with the milk, fat and protein yields 414 of New Zealand strains reported in the study by Horan *et al*. (2005) who also found that the 415 mean pedigree index for milk yield was significantly higher in North American strains 416 compared with the New Zealand strains. Our results were also similar to the findings of 417 Dobos *et al.* (2001) in pasture-based FF heifers over three lactations and the report of White *et* 418 *al.* (2002) in which Jersey cows produced 23.3% less milk than Holsteins.

419

420 Total milksolids yields obtained in this study was higher than the values reported for FF cows 421 in New Zealand (García *et al.* 1998, and Lopez *et al*. 2001). Their study evaluated two strains 422 of FF cows bred for low and high body weights, under low stocking rates (1.95-2.25 cows/ha) 423 supplemented with 0.4-1.20 ton DM/cow concentrate. Average stocking rate in dairy farms in 424 Australia is about 2.5 cows/ha (Dairy Australia 2005). Poor efficiency of grain supplement 425 utilisation due to higher level of substitution under low stocking rates or high pasture 426 allowance has been reported in literature (Robaina *et al.* 1998, Stockdale 1999 and Fulkerson 427 *et al.* 2000). The daily milksolids yield reported herein are in agreement with the findings of 428 Bryant *et al. (*2003) who utilised FF cows at a stocking rate of 4.4 cows/ha and supplemented 429 with 1.3-1.5 t DM per annum for three seasons.

430

431 The higher performance of FJ over the JJ cows demonstrated the beneficial effect of heterosis. 432 The rate of genetic progress in the dominant dairy breeds was evident in the annual rate of 433 increase in milk and constituent yields. Whereas increase in milk yield/lactation averaged 5% 434 and 5.3% in FF and FJ breeds respectively, it was 3% in JJ. In addition, the percentage

435 decline in lactation in 2003 was 1.7% in FF while it was 2.7% in FJ and JJ. Madgwick and 436 Goddard (1989) had highlighted the possibility of slower genetic progress which might make 437 Jersey cows less competitive in the future.

438

439 *Parity*

440 Milk production is known to increase with increase in parity and cow age due to higher body 441 weight, larger capacity for dry matter intake, increase in size of the udder and recurrence of 442 pregnancy and lactations (Capuco *et al.* 2001). Lower production in primiparous cows is 443 related to competition between tissues (e.g. mammary gland *v*. peripheral tissues) for 444 metabolites for growth and lactation in the immature animal (Tekerli *et al.* 2000). Similar 445 observations to our study on the effect of parity on dairy cow performance has also been 446 reported (Val-Arreola *et al.* 2004).

447

448 *Location*

449 Milk yield per cow in 2002/03 calving season obtained in this study was higher than reported 450 averages in other states of Australia, except Western Australia and South Australia. 451 Production per cow in Tasmania was lower than that of other states except Queensland in 452 2003/04 (Dairy Australia 2005). Concerns over low production per cow *vis-a-vis* increasing 453 use of grain concentrates to benefit from increasing genetic potential of dairy cows is a main 454 issue in pasture-based dairying (Chapman *et al.* 2004). In 2001, the Australian Dairy Herd 455 Improvement Scheme (ADHIS) adopted the Australian Profit Ranking (APR) method for sire 456 breeding value evaluation as a measure of improving long term productivity and profit per 457 cow. Pasture-based dairying in Tasmania is also focussing on production per hectare as a 458 measure of farm productivity rather than productivity per cow *per se* as evidence from 459 industry reports show that the link between profit and pasture utilisation is the strongest of all

460 performance indicators (Dairy Australia, 2004). Differences in yield traits between locations, 461 attributable to differences in the rainfall distribution pattern and geo-physical conditions, have 462 been reported extensively in literature (Msanga *et al.* 2000, Horan *et al.* 2005, Dairy Australia 463 2005). A review of climate data in the study area showed that mean annual rainfall and 464 altitude were significantly different (p<0.001) between the dairying locations in Tasmania. 465 Mean annual rainfall was lower in the South but significantly higher (p<0.01) in King Island 466 compared to the other location. Mean altitude (meters above sea level) averaged 117.7±16.4 467 m in Central North, North West, North-East and Far North-West locations compared with an 468 average of 43.6m in the South and king Island. (Geography of Tasmania 2008). In addition, 469 considerable investment undertaken in the North-West and North-East areas of the state in the 470 last decade has encouraged the emergence of corporate farmers with large herds with the 471 attendant economies of scale (DPIW 2005). It should be noted however that the relatively 472 smaller number of cows (Table 1) in the South and king Island could have some bearing on 473 some of the responses evaluated in these location.

474

475 *Herd size*

476 Herd sizes reported in this study were generally consistent with the national dairy herd 477 statistics (Dairy Australia 2006). Tasmania has the highest mean herd size of 254 cows per 478 herd, while Queensland has the smallest with 158 cows per herd. Higher performance in herds 479 with large number of cows is also in agreement with results from dairying in Victoria. A 480 benchmark study in Western Victoria that compared profitability indices of the top and 481 bottom 10 farms indicated that large herds were more profitable and gave greater returns on 482 capital than medium or small herds. As herd size increases, overhead and labour costs can be 483 spread over more units (Doyle and Kelly 1998). In addition, owners of larger herds have been 484 reported to adopt high intensity feeding systems and appeared more open to improved

485 management systems than small or medium herd owners (DRDC 1996). Smaller herds, on the 486 other hand can benefit from the flexibility in land and labour management to increase per unit 487 resource (Doyle and Kelly 1998).

488

489 *Heritability of milk, fat and protein yields.*

490 The relatively small size data used in this study could lead to an over-estimation of the h^2 491 estimates. It is known that the greater the sample size, the higher the precision of additive 492 genetic estimates (Jensen 2001). However, our h^2 estimates using both univariate and 493 multivariate approaches were within the range of values reported in literature which ranged 494 from 0.31, (Wilmink 1987) to 0.49 (Pander *et al.* 1992). Meyer *et al.* (1989) compared the 495 different methods of estimating h^2 and reported 305-day milk yield h^2 of 0.37. In another 496 study that compared alternative methods of equalizing heterogeneity of variance, Boldman 497 and Freeman (1990) reported lower h^2 estimates of 0.18, 0.22, and 0.24 for untransformed 498 milk yield in low, medium, and high producing herds, respectively. Heritability estimates of 499 fat and protein yields reported in this study were slightly higher than the values reported by 500 Visscher and Goddard (1995) from five states of Australia (excluding Tasmania), probably 501 partly because they used test-day sire models in their evaluation while we utilised an animal 502 model with 305-day records. However, our results were in agreement with the work of 503 Swalve (1995) who utilised test-day, herd-year-season animal model and reported 305d milk, 504 fat and protein h^2 estimates of 0.39, 0.32 and 0.30 respectively.

505

506 Comparison of h^2 estimation methods showed that test-day approaches generally yield lower 507 estimates, (Meyer *et al.* 1989, Pander *et al.* 1992) compared with 305-day method. Issues 508 from using aggregated 305-day milk yield and the benefits of test-day random regression 509 models have been extensively reviewed (Jensen 2001). The continuation of using 305-day 510 milk yield stems from industry tradition and the limitations imposed by computations

511 requirements until recently (Jensen 2001). There is however, a general consensus that the 512 heritabilities for fat were, in almost all cases, lower than the heritabilities for protein, and that 513 milk production has the highest heritability.

515 **Conclusions**

514

516 This study investigated the influence of genetic and environmental factors affecting dairy 517 production in Tasmania. Breed, parity, age and lactation length, were important determinants 518 of milk and milk solid yields under pasture-based dairy systems. Improved yields over the 519 years were indicative of not only improvement in dairy cow genetics, but also improvement in 520 the adoption of better management practices. Season played a significant role in the calving 521 pattern of pasture-based dairy cows in Tasmania due to the variation in micro climate across 522 the dairy regions. The inclusion of random cow effect in the animal model showed higher 523 yields attributable to additive genetic variance in the FF cows, although small data size 524 precluded the estimation of the additive genetic effects in the other breeds. This would 525 suggest the potential for genetic manipulation to increase yields. Significant herd effect also 526 suggested that there was scope to improve productivity through the adoption of improved 527 management practices. Herd size as a factor of management improved production traits in 528 very large herds thus supporting the emerging trends for larger dairy herds in Tasmania. Other 529 management factors such as access to information and market, favourable market prices, 530 technical and managerial support are very important if farmers are to cope with the challenge 531 of running profitable dairy enterprises. The heritability and genetic and phenotypic correlation 532 estimates in this study fall within the expected ranges in dairy cattle, thus confirming that 533 years of selection of the milking herd in Tasmania will likely continue to improve genetic 534 progress within the pasture-based dairy framework. A desirable future goal would be to 535 conduct a comparative economic analysis of farm profitability under high grain 536 supplementation versus grazing only to shed more light on the cost-effective benefits of

- 537 pasture-based dairying.
- 538

539

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- 545

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- 648

Table 1. Least squares means±s.e. of total milk, fat, protein yields per lactation and somatic cell count of dairy cows by breed, parity, calving year, locations, herd size, and calving season.

See materials and methods for definition of location and SCC.
Means bearing different superscripts within the same column are statistically different (p<0.01)

Table 2. Least square means of interactions of breed, calving year and calving season for milk, fat, protein and somatic cell count of three dairy cows breeds in Tasmania.

See materials and methods for definition of SCC. All tested factors were significant (P<0.01)

Table 3. Descriptive statistics of 305d milk, fat, protein and SCC of Holstein Friesian cows adjusted for terms in the animal model based on DATA2.

Category	Milk (L)				Fat (kg)				Protein (kg)				$Log of SCC^*$				
$ParityI^*$	Mean	SD	Min	Max	Mean	SD	Min	Max	Mean	SD	Min	Max	Mean	SD	Min	Max	n
	4859	1440	1209	12997	196.6	52.88	18	509	161.6	48.65	22	469	3.83	0.976	Ω	8.23	18767
\overline{c}	5520	1647	1247	13550	223.6	60.35	25	480	186.1	55.57	38	450	3.90	1.061	Ω	8.53	15677
3	6171	1746	1207	15141	250.6	66.12	27	544	207.9	57.93	16	487	4.29	1.163	Ω	8.94	31470
Calving Season																	
Winter	5648	1613	1207	14924	231.6	62.22	18	544	190.7	54.59	16	487	4.05	1.089	Ω	8.94	32998
Spring	5463	1825	1210	14946	219.1	66.37	25	517	182.3	60.62	34	468	4.12	1.134	$\left($	8.53	24600
Summer	5542	1931	1296	12944	218.1	73.53	43	460	179.7	61.9	39	414	3.89	1.535	$\overline{0}$	7.28	1230
Autumn	6262	1762	1230	15141	251.8	68.15	38	527	210.9	60.38	40	467	3.98	1.024	Ω	8.06	7086
Calving Year																	
2000	4844	1378	1229	14946	197.5	55.03	29	495	157.5	44.93	22	468	3.83	1.235	Ω	8.47	4385
2001	5436	1486	1474	14199	218.1	55.90	52	472	178.8	47.81	49	456	3.83	1.111	$\overline{0}$	7.60	6639
2002	5251	1557	1247	13112	212.6	60.81	40	507	172.9	51.27	34	432	4.08	1.014	Ω	7.82	9024
2003	5510	1608	1230	13861	222.5	60.86	18	517	185	53.01	40	438	4.01	1.079	Ω	8.15	11677
2004	5816	1699	1317	14360	236.4	63.45	27	509	197.6	56.83	26	471	4.09	1.120	Ω	8.39	14774
2005	6024	1929	1207	15141	245.1	71.05	31	544	204.8	65.06	16	487	4.21	1.109	0.69	8.94	19415
Location [#]																	
FNWest	5779	1684	1209	14924	234.6	63.73	18	544	194.9	57.44	40	487	4.02	1.068	Ω	8.53	17152
Nwest	5897	1945	1230	15141	234.9	68.15	44	509	196.9	62.42	41	469	3.97	1.231	Ω	8.35	14949
CntNorth	5749	1640	1229	13929	237.0	66.52	29	514	194.3	57.19	22	435	4.13	1.082	Ω	8.66	12376
South	5231	1298	1273	10024	209.2	55.3	46	380	174.7	43.1	44	328	4.39	0.926	2.08	8.07	1108
Neast	5365	1631	1207	14946	216.9	62.16	27	517	179.9	55.45	16	468	4.12	1.071	0.69	8.94	18596
KIsland	4581	1331	1210	9739	202.1	62.1	52	446	152.0	44.25	42	346	4.21	1.018	1.39	7.54	1733

See materials and methods for definition of locations and SCC. #Parity >2 were pooled and labelled as parity 3

Table 4. Heritability, phenotypic and genetic correlations (±se) of milk, fat, protein and somatic cell count of Holstein-Friesian cows

Heritability shown in bold (diagonal), phenotypic correlation and genetic correlation on upper and lower triangles respectively. SCC=Somatic cell count

Adapted from: DPWI 2005, Dairy Australia 2006 p=predicted, NA=Not available