

New Zealand Society of Animal Production online archive

This paper is from the New Zealand Society for Animal Production online archive. NZSAP holds a regular annual conference in June or July each year for the presentation of technical and applied topics in animal production. NZSAP plays an important role as a forum fostering research in all areas of animal production including production systems, nutrition, meat science, animal welfare, wool science, animal breeding and genetics.

An invitation is extended to all those involved in the field of animal production to apply for membership of the New Zealand Society of Animal Production at our website www.nzsap.org.nz

[View All Proceedings](#)

[Next Conference](#)

[Join NZSAP](#)

The New Zealand Society of Animal Production in publishing the conference proceedings is engaged in disseminating information, not rendering professional advice or services. The views expressed herein do not necessarily represent the views of the New Zealand Society of Animal Production and the New Zealand Society of Animal Production expressly disclaims any form of liability with respect to anything done or omitted to be done in reliance upon the contents of these proceedings.

This work is licensed under a [Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International License](http://creativecommons.org/licenses/by-nc-nd/4.0/).



You are free to:

Share— copy and redistribute the material in any medium or format

Under the following terms:

Attribution — You must give [appropriate credit](#), provide a link to the license, and [indicate if changes were made](#). You may do so in any reasonable manner, but not in any way that suggests the licensor endorses you or your use.

NonCommercial — You may not use the material for [commercial purposes](#).

NoDerivatives — If you [remix, transform, or build upon](#) the material, you may not distribute the modified material.

<http://creativecommons.org.nz/licences/licences-explained/>

be motivated by its application to non-production traits such as fertility, survival, health and product quality. There will need to be greater vertical integration and greater use of AI in these industries

to allow a financial return to entities that invest in GWS. These facets may cause a barrier to the use of GWS until genotyping costs drop sufficiently to catalyse the changes needed.

Uptake of DNA testing by the livestock industries of New Zealand

A.M. CRAWFORD, R.M. ANDERSON and K.M. McEWAN

AgResearch, Invermay Agricultural Centre, Mosgiel

ABSTRACT

This paper describes, from the perspective of the main animal genotyping service laboratory, the uptake of DNA tests by the New Zealand livestock industries. We describe a rapid increase in the use of DNA testing over the last 3-4 years in New Zealand. In particular pedigree establishment and verification has shown significant uptake by deer, cattle and sheep breeders. The highest penetration of DNA testing has been in the deer industry where we estimate 30-40% of the recorded breeding population uses DNA testing to establish and verify pedigrees. The increases have mainly been due to an increase in the number of samples per client rather than an increase in the number of clients using the service.

Keywords: DNA; pedigrees; parentage testing; gene tests.

INTRODUCTION

The last fifteen years have seen the development of microsatellite based systems to identify pedigrees in a wide range of animal species including livestock and companion animals (Dodds *et al.*, 2005). More recently a large number of QTL, and in some cases the causative gene alteration, have been discovered for an array of productive traits in livestock (Andersson & Georges, 2004). The motivation for such work is to assist farmers and breeding companies to select for improved livestock and maximise genetic gain. This paper examines, from the perspective of one major DNA genotyping service company, *Genomnz*TM, how widespread and rapidly DNA

testing in sheep, cattle and deer has been taken up by the livestock breeding industry in New Zealand.

MATERIALS AND METHODS

The commercial DNA tests offered by *Genomnz*TM are listed in Table 1. Although our data covers most non-dairy livestock DNA testing in New Zealand we are only a minor provider of DNA testing to the dairy industry where GeneMarkTM is the major provider. Two other service laboratories have been active during this time. Commercial DNA tests are provided by Lincoln University for footrot resistance and lamb survival, and between 1999 and 2003 a DNA testing service was provided by SignagenTM which has since ceased trading.

Table 1: Combined list of DNA tests offered by the *Genomnz*TM service laboratory.

Species	Test	Total number of samples tested up	
		to Jan 2007	Year test introduced
Sheep	Parentage identification	277,707	2000
	Inverdale gene test	10,642	1997
	Booroola gene test	8,747	1996
Cattle	Parentage verification and identification	15,597	1996
	Myostatin gene test	332	2001
	Calpain-1 gene test	290	2004
	BLAD gene test	232	2005
	CVM gene test	276	2005
Deer	Parentage identification	59,329	1996
	Wapiti hybrid test	~5000	1996
Salmon	Parentage verification	4,287	2003

Not included in this table is the contract R&D testing undertaken by *Genomnz*TM, in particular genome scans that are done under contract to research organisations, many of which have confidentiality provisions. The scope of this report is therefore limited to the uptake of DNA testing by commercial breeding enterprises. To our knowledge only a few single gene tests have been supplied by overseas laboratories, and all DNA parentage identification undertaken by commercial breeders in NZ has been provided by New Zealand laboratories. The thoroughbred and standardbred horse racing industries have for many years required auditing of their pedigrees. In NZ this is undertaken at Massey University for thoroughbreds and in Australia for standardbreds. The samples from all animals sent to the laboratory are stored. Most samples are either blood spots on Whatman FTA paper or hair follicles stored in envelopes. In both cases when stored in a cool dry environment these samples are likely to last for decades. This large collection of DNA samples, from many of NZ's leading livestock breeders, is likely to prove useful in tracking the genetic progress of our national herds and flock.

RESULTS

The growth of parentage testing in all three ruminant species is shown in Figures 1- 3.

In deer, cattle and sheep major growth has taken place in the last 3-4 years. Prior to 2001 relatively slow growth occurred in deer and cattle. With sheep DNA testing, which did not become available until 2000, uptake by the sheep community has shown rapid growth. Gene tests have not experienced the same rapid growth of the parentage tests (Figure 4).

This is probably due to the specialised nature of the tests and because, once a gene variant has either been fixed or eliminated in a population, testing requirements are reduced. The cause of the growth is shown in Figures 5 and 6 where the mean number of DNA samples per client has been determined for deer and cattle. In both cases the increase is due to an increase in the number of samples existing clients send to the laboratory rather than any major increase in the number of clients using the service.

DISCUSSION

It is clear from this paper that the use of DNA testing especially for parentage has become a widely used technology by the animal breeding industries for deer, sheep and cattle in New Zealand. Its adoption has primarily been by larger

scale breeders who are best able to capture the benefits of reduced labour inputs and hence justify its cost. The highly skilled labour to observe and record the birth of pastoral livestock is not required if DNA parentage is used. An additional benefit is that single sire mating is not required. This means mating groups can be larger and more efficient use of grazing can be made.

At present DNA tests for a few genes of major effect are also available for sheep and cattle. Those DNA tests for deleterious genes such as BLAD (Batt *et al.*, 1994) in cattle and spider lamb syndrome in sheep (Drogemuller *et al.*, 2005) have been widely used to screen affected populations but as carriers are removed use of the tests diminishes. Those DNA tests for traits having a beneficial effect such as DGAT1 in dairy cattle (Spelman *et al.*, 2002) and Inverdale in sheep (Galloway *et al.*, 2000) have been widely adopted by the industry and DNA testing has been used to introgress the beneficial gene variants into new populations. Once again, however, as the trait becomes fixed in the population, use of the test will diminish.

The sequencing of the cattle genome has recently been completed and an international consortium to sequence the sheep genome is in place, the goal being to complete the sheep genome within 4 years. These initiatives have or will give us over a million DNA markers. This means that for these two species DNA markers can be used directly to estimate the breeding worth of an individual. This process, called Genome wide selection (GWS), has just begun to be used in the dairy industry (Schaeffer, 2006). It will increase the rate of genetic improvement and is the logical successor to single DNA tests as it means that all the genomic contributions to a particular trait can be recognised and used to assess the breeding worth of an individual.

Unlike manufactured products where the design blueprint and assembly schedule are separate from the product itself, biological products carry the blueprint with them in the form of their DNA. We can only change many features of the biological product at breeding time. Previously this has been done using a combination of pedigree information and measuring the performance of the animal (product). Increasingly the future will see direct examination of the product blueprint (DNA) replacing trait measurement in animals. Those breeders currently collecting samples for DNA parentage will find that the samples have increasing value, over the next decade, for direct measurement of the animals DNA blueprint.

Figure 1: Annual number of Cattle parentage tests.

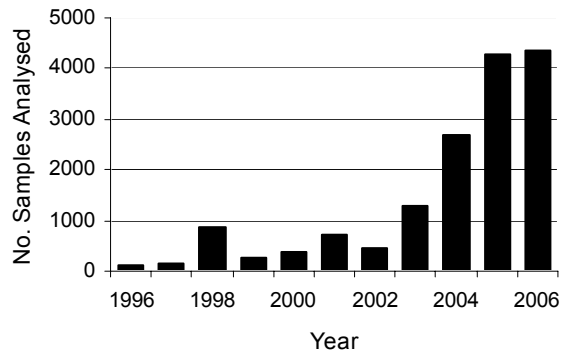


Figure 4: Annual number of gene tests in all species.

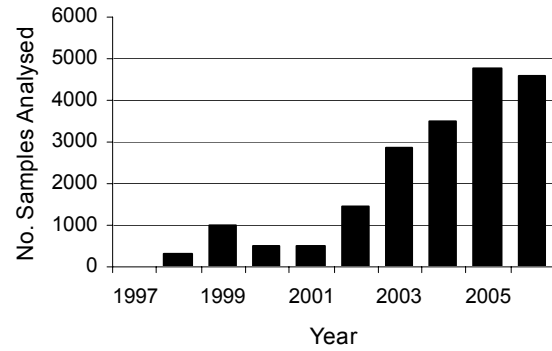


Figure 2: Annual number of Deer parentage tests.

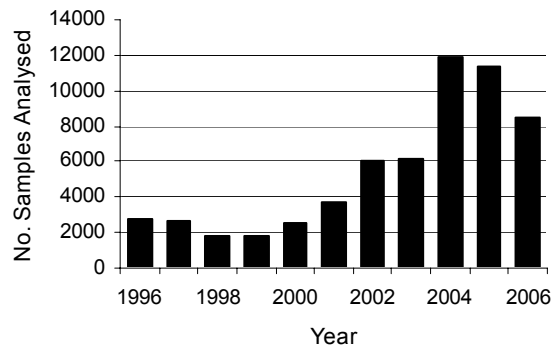


Figure 5: Mean number of Deer parentage tests per client.

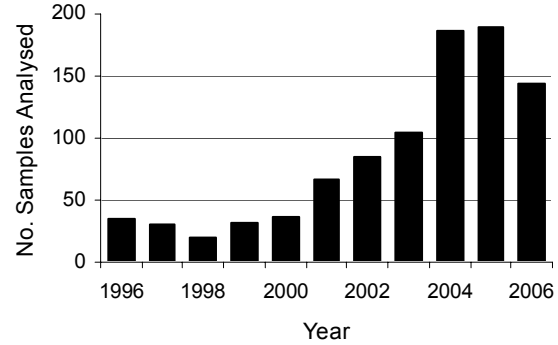


Figure 3: Annual number of Sheep parentage tests.

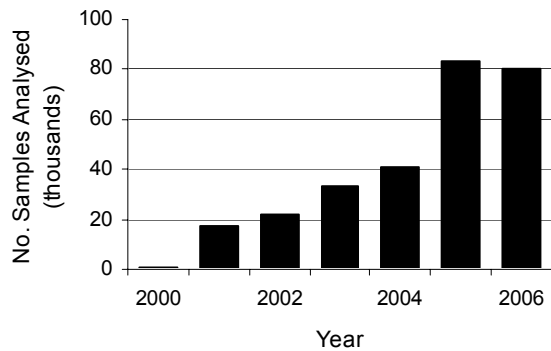
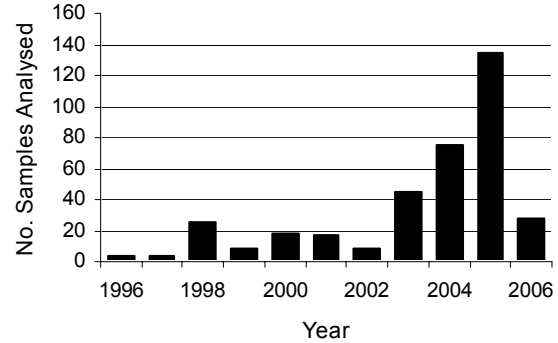


Figure 6: Mean number of Cattle parentage tests per client.



ACKNOWLEDGEMENTS

The authors wish to thank the staff of *Genomnz* for their diligence and professionalism in providing an excellent service to the NZ farming industry and for allowing us access to their records to produce this paper.

REFERENCES

Current status and future of genomic selection – combined reference list

- Amyes, N.C.; Morris, C.A.; Towers, N.R. 2002: Ryegrass staggers: genetics accounts for a six-fold difference in susceptibility between selection lines of lambs at Ruakura. *Proceedings of the New Zealand Society of Animal Production* **62**: 191-194.
- Andersson, L.; Georges, M. 2004: Domestic-animal genomics: deciphering the genetics of complex traits. *Nature Reviews Genetics* **5**: 202-212.
- Anon. 2006: Special NZ lamb for Marks & Spencer. *AgBrief* **6**: 3.
- Archer, J.A. 2003: Genetic improvement of red deer. *Proceedings of the New Zealand Society of Animal Production* **63**: 233-236.
- Asher, G.W.; Archer, J.A.; Scott, I.C.; O'Neill, K.T.; Ward, J.; Littlejohn, R.P. 2005: Reproductive performance of pubertal red deer (*Cervus elaphus*) hinds: Effects of genetic introgression of wapiti subspecies on pregnancy rates at 18 months of age. *Animal Reproduction Science* **90**: 287-306.
- Banks, R. 1997: The meat elite project: establishment and achievements of an elite meat sheep nucleus. *Proceedings of the Association for Advancement of Animal Breeding and Genetics* **12**: 598-601.
- Batt, C.A.; Wagner, P.; Wiedmann, M.; Luo, J.; Gilbert, R. 1994: Detection of bovine leukocyte adhesion deficiency by nonisotopic ligase chain reaction. *Animal Genetics* **25**: 95-98.
- Bennett, R.; Stanley, G.; McNaughton, L. 2005: Fast-forwarding dairy genetics. Accessed 25 January, 2007. http://www.lic.co.nz/pdf/grasslands_presentation.pdf
- Bissinger, P.H.; Kuchler, K. 1994: Molecular cloning and expression of the *Saccharomyces cerevisiae* STS1 gene product. A yeast ABC transporter conferring mycotoxin resistance. *Journal of Biological Chemistry* **269**: 4180-4186.
- Blott, S.; Kim, J.J.; Moisis, S.; Schmidt-Kuntzel, A.; Cornet, A.; Berzi, P.; Cambisano, N.; Ford, C.; Grisart, B.; Johnson, D.; Karim, L.; Simon, P.; Snell, R.; Spelman, R.; Wong, J.; Vilkki, J.; Georges, M.; Farnir, F.; Coppieters, W. 2003: Molecular dissection of a quantitative trait locus: a phenylalanine-to-tyrosine substitution in the transmembrane domain of the bovine growth hormone receptor is associated with a major effect on milk yield and composition. *Genetics* **163**: 253-266.
- Bodin, L.; Lecerf, F.; Bessièrè, M.; Mulsant, P. 2006: Features of major genes for ovulation in the *Lacaune* population. *Proceedings of the 8th World Congress on Genetics Applied to Livestock Production*: 04-04.
- Bonnet, A.; Thevenon, S.; Claro, F.; Gautier, M.; Hayes, H. 2001: Cytogenetic comparison between Vietnamese sika deer and cattle: R-banded karyotypes and FISH mapping. *Chromosome Research* **9**: 673-687.
- Broad, T.E.; Glass, B.C.; Greer, G.J.; Robertson, T.M.; Bain, W.E.; Lord, E.A.; McEwan, J.C. 2000: Search for a locus near to myostatin that increases muscling in Texel sheep in New Zealand. *Proceedings of the New Zealand Society of Animal Production* **60**: 110-112.
- Campbell, A.W.; McLaren, R.J. 2007: LoinMAXTM and MyoMAXTM: taking DNA marker tests from the research environment to commercial reality. *Proceedings of the New Zealand Society of Animal Production* **67**: 160-162.
- Campbell, A.G.; Meyer, H.H.; Henderson, H.V.; Wesselink, C. 1981: Breeding for facial eczema resistance – a progress report. *Proceedings of the New Zealand Society of Animal Production* **41**: 273-278.
- Campbell, A.W.; Bain, W.E.; McRae, A.F.; Broad, T.E.; Johnstone, P.D.; Dodds, K.G.; Veenvliet, B.A.; Greer, G.J.; Glass, B.C.; Beattie, A.E.; Jopson, N.B.; McEwan, J.C. 2003: Bone density in sheep: Genetic variation and quantitative trait loci localisation. *Bone* **33**: 540-548.
- Charteris, P.L.; Garrick, D.J. 1996: Characterisation of beef cattle breeding industry structure. *Proceedings of the New Zealand Society of Animal Production* **56**: 386-389.
- Crawford, A.M. 2001: A review of QTL experiments in sheep. *Proceedings of the Association for the Advancement of Animal Breeding and Genetics* **14**: 33-38.
- Crawford, A.M.; Paterson, K.A.; Dodds, K.G.; Diez Tascon, C.; Williamson, P.A.; Roberts Thomson, M.; Bisset, S.A.; Beattie, A.E.; Greer, G.J.; Green, R.S.; Wheeler, R.; Shaw, R.J.; Knowler, K.; McEwan, J.C. 2006: Discovery of quantitative trait loci for resistance to parasitic nematode infection in sheep. I. Analysis of outcross pedigrees. *BMC Genomics* **7**: 178.
- Crawford, A.M.; Anderson, R.M.; McEwan, K.M. 2007: Uptake of DNA testing by the livestock industries of New Zealand. *Proceedings of the New Zealand Society of Animal Production* **67**: 168-174.
- Cullen, N.G.; Morris, C.A.; Hickey, S.M. 2006: Genetic parameters for resistance to facial eczema in dairy cattle. *Proceedings of the New Zealand Society of Animal Production* **66**: 310-314, 319-324.
- Davis, G.H.; Dodds, K.G.; Wheeler, R.; Jay, N.P. 2001: Evidence that an imprinted gene on the X chromosome increases ovulation rate in sheep. *Biology of Reproduction* **64**: 216-221.
- Davis, G.H.; Farquhar, P.A.; O'Connell, A.R.; Everett-Hincks, J.M.; Wishart, P.J.; Galloway, S.M.; Dodds, K.G. 2006b: A putative autosomal gene increasing ovulation rate in Romney sheep. *Animal Reproduction Science* **92**: 65-73.
- Davis, G.H.; McEwan, J.C.; Dodds, K.G. 2006a: Strategies for use of molecular information in industry genetic improvement programmes for meat sheep. *Proceedings of the 8th World Congress on Genetics Applied to Livestock Production*: 04-01.

- Diez-Tascon, C.; Keane, O.M.; Wilson, T.; Zadissa, A.; Hyndman, D.L.; Baird, D.B.; McEwan, J.C.; Crawford, A.M. 2005: Microarray analysis of selection lines from outbred populations to identify genes involved with nematode parasite resistance in sheep. *Physiological Genomics* **21**: 59-69.
- Dodds, K.G.; Amer, P.R.; Spelman, R.J.; Archer, J.A.; Auvray, B. 2007: Prospects for genome wide selection in the New Zealand livestock industries. *Proceedings of the New Zealand Society of Animal Production* **67**: 162-167.
- Dodds, K.G.; Tate, M.L.; Sise, J.A. 2005: Genetic evaluation using parentage information from genetic markers. *Journal Animal Science* **83**: 2271-2279
- Drogemuller, C.; Wohlke, A.; Distl, O. 2005: Spider Lamb Syndrome (SLS) mutation frequency in German Suffolk sheep. *Animal Genetics* **36**: 539-540.
- Duncan, E.J.; Dodds, K.G.; Henry, H.M.; Thompson, M.P.; Phua, S.H. 2007: Cloning, mapping and association studies of the ovine *ABCG2* gene with facial eczema disease in sheep. *Animal Genetics* **38**: [epublished].
- Escayg, A.P.; Hickford, J.G.H.; Bullock, D.W. 1997: Association between alleles of the ovine major histocompatibility complex and resistance to footrot. *Research in Veterinary Science* **63**: 283-287.
- Forrest, R.H.; Hickford, J.G.H.; Wynyard, J.; Merrick, N.; Hogan, A.; Frampton, C. 2006: Polymorphism at the β_3 -adrenergic receptor (*ADRB3*) locus of Merino sheep and its association with lamb mortality. *Animal Genetics* **37**: 465-468.
- Galloway, S.M.; McNatty, K.P.; Cambridge, L.M.; Laitinen, M.P.E.; Juengel, J.L.; Jokiranta, T.S.; McLaren, R.J.; Luiro, K.; Dodds, K.G.; Montgomery, G.W.; Beattie, A.E.; Davis, G.H.; Ritvos, O. 2000: Mutations in an oocyte-derived growth factor gene (*BMP15*) cause increased ovulation rate and infertility in a dosage-sensitive manner. *Nature Genetics* **25**: 279-283.
- Garrick, D.J. 1997: A comparison of some challenges to genetic improvement in New Zealand industries. *Proceedings of the Association for the Advancement of Animal Breeding and Genetics* **12**: 31-39.
- Garrick, D.J.; Blair, H.T.; Clarke, J.N. 2000: Sheep industry structure and genetic improvement. *Proceedings of the New Zealand Society of Animal Production* **60**: 175-179.
- Goosen, G.J.C.; Dodds, K.G.; Tate, M.L.; Fennessy, P.F. 1999: QTL for live weight traits in Père David's x red deer interspecies hybrids. *Journal of Heredity* **90**: 643-647.
- Goosen, G.J.C.; Dodds, K.G.; Tate, M.L.; Fennessy, P.F. 2000: QTL for pubertal and seasonality traits in male Père David's x red deer interspecies hybrids. *Journal of Heredity* **91**: 397-400.
- Grisart, B.; Coppieters, W.; Farnir, F.; Karim, L.; Ford, C.; Berzi, P.; Cambisano, N.; Mni, M.; Reid, S.; Simon, P.; Spelman, R.; Georges, M.; Snell, R. 2002: Positional candidate cloning of a QTL in dairy cattle: identification of a missense mutation in the bovine *DGATI* gene with major effect on milk yield and composition. *Genome Research* **12**: 222-231.
- Haley, C.S. 1995: Livestock QTLs - bringing home the bacon. *Trends in Genetics* **11**: 488-492.
- Hanrahan, J.P.; Gregan, S.M.; Mulsant, P.; Mullen, M.; Davis, G.H.; Powell, R.; Galloway, S.M. 2004: Mutations in the genes for oocyte-derived growth factors *GDF9* and *BMP15* are associated with both increased ovulation rate and sterility in Cambridge and Belclare sheep (*Ovis aries*). *Biology of Reproduction* **70**: 900-909.
- Hayes, B.J.; Chamberlain, A.J.; Goddard, M.E. 2006: Use of markers in linkage disequilibrium with QTL in breeding programs. *Proceedings of the 8th World Congress on Genetics Applied to Livestock Production*: 30-06.
- Hayes, B.J.; Goddard, M.E. 2007: Genomic selection for accelerated genetic gain in livestock. *Proceedings of the New Zealand Society of Animal Production* **67**: 143-146.
- Heaton, M.P.; Keen, J.E.; Clawson, M.L.; Harhay, G.P.; Bauer, N.; Shultz, C.; Green, B.T.; Durso, L.; Chitko-McKown, C.G.; Laegreid, W.W. 2005: Use of bovine single nucleotide polymorphism markers to verify sample tracking in beef processing. *Journal American Veterinary Medical Association* **226**: 1311-1314.
- Henry, H.M.; Dodds, K.G.; Wuliji, T.; Jenkins, Z.A.; Beattie, A.E.; Montgomery, G.W. 1998: A genome screen for QTL for wool traits in a Merino x Romney backcross flock. *Proceedings of the 6th World Congress on Genetics Applied to Livestock Production* **26**: 433-436.
- Hohenboken, W.D.; Morris, C.A.; Munday, R.; de Nicolo, G.; Amyes, N.C.; Towers, N.R.; Phua, S.H. 2004: Antioxidants in blood from sheep lines divergently selected for facial eczema resistance. *New Zealand Journal of Agricultural Research* **47**: 119-127.
- Johnson, P.L. 2003: A directed search for QTL affecting carcass composition traits in Texel sheep. Ph.D. Thesis, Massey University, Palmerston North.
- Johnson, P.L.; McEwan, J.C.; Dodds, K.G.; Purchas, R.W.; Blair, H.T. 2005: A directed search in the region of *GDF8* for quantitative trait loci affecting carcass traits in Texel sheep. *Journal of Animal Science* **83**: 1988-2000.
- Johnson, P.L.; McEwan, J.C.; Dodds, K.G.; Purchas, R.W.; Blair, H.T. 2005: Meat quality traits were unaffected by a quantitative trait locus affecting leg composition traits in Texel sheep. *Journal of Animal Science* **83**: 2729-2735.
- Jopson, N.B.; Nicoll, G.B.; Stevenson-Barry, J.; Duncan, S.; Greer, G.J.; Bain, W.E.; Gerard, E.M.; Glass, B.C.; Broad, T.E.; McEwan, J.C. 2001: Mode of inheritance and effects on meat quality of the rib-eye muscling (*REM*) QTL in sheep. *Proceedings of the Association for the Advancement of Animal Breeding and Genetics* **14**: 111-114.
- Kappes, S.M.; Green, R.D.; Van Tassell, C.P. 2006: Sequencing the bovine genome and developing a haplotype map: approaches and opportunities. *Proceedings of the 8th World Congress on Genetics Applied to Livestock Production*: 22-01.

- Maqbool, N.J.; Tate, M.L.; Dodds, K.G.; Anderson, R.M.; McEwan, K.M.; Mathias, H.C.; McEwan, J.C.; Hall, R.J. 2007: A QTL study of growth and body shape in the inter-species hybrid of Père David's deer (*Elaphurus davidianus*) and red deer (*Cervus elaphus*). *Animal Genetics* (doi:10.1111/j.1365-2052.2007.01597.x).
- McNaughton, L.R.; Bennett, R.; Stanley, G.; Harcourt, S.; Spelman, R.J. 2005: Phenotype definition and the identification of QTL for puberty traits in crossbred dairy cattle. *Proceedings of the Association for the Advancement of Animal Breeding and Genetics* **16**: 242-245.
- Meuwissen, T.H.E.; Hayes, B.J.; Goddard, M.E. 2001: Prediction of total genetic value using genome-wide dense marker maps. *Genetics* **157**: 1819-1829.
- Montgomery, G.W.; Crawford, A.M.; Penty, J.M.; Dodds, K.G.; Ede, A.J.; Henry, H.M.; Pierson, C.A.; Lord, E.A.; Galloway, S.M.; Schmack, A.E.; Sise, J.A.; Swarbrick, P.A.; Hanrahan, V.; Buchanan, F.C.; Hill, D.F. 1993: The ovine Booroola fecundity gene (*FecB*) is linked to markers from a region of human chromosome 4q. *Nature Genetics* **4**: 410-414.
- Montgomery, G.W.; Henry, H.M.; Dodds, K.G.; Beattie, A.E.; Wuliji, T.; Crawford, A.M. 1996: Mapping the *Horns (Ho)* locus in sheep: a further locus controlling horn development in domestic animals. *Journal of Heredity* **87**: 358-363.
- Morris, C.A.; Amyes, N.C.; Cullen, N.G.; Hickey, S.M. 2006a: Carcass composition and growth in Angus cattle genetically selected for differences in pubertal traits. *New Zealand Journal of Agricultural Research* **49**: 1-11.
- Morris, C.A.; Cullen, N.G.; Geertsema, H.G. 1997: Genetic studies of bloat susceptibility in cattle. *Proceedings of the New Zealand Society of Animal Production* **57**: 19-21.
- Morris, C.A.; Campbell, A.W.; Cullen, N.G.; Davis, G.H.; Everett-Hincks, J.M.; Hall, R.J.; Henry, H.M.; Johnson, P.L.; McEwan, J.C.; Phua, S.H.; Wilson, T. 2007: Current status of QTL and association studies in New Zealand cattle, sheep and deer. *Proceedings of the New Zealand Society of Animal Production* **67**: 153-159.
- Morris, C.A.; Cullen, N.G.; Glass, B.C.; Hyndman, D.L.; Manley, T.R.; Hickey, S.M.; McEwan, J.C.; Pitchford, W.S.; Bottema, C.D.K.; Lee, M.A.H. 2007: Fatty acid synthase effects on bovine adipose fat and milk fat. *Mammalian Genome* **18**: 64-74.
- Morris, C.A.; Cullen, N.G.; Hickey, S.M.; Dobbie, P.M.; Veenvliet, B.A.; Manley, T.R.; Pitchford, W.S.; Kruk, Z.A.; Bottema, C.D.K.; Wilson, T. 2006b: Genotypic effects of *calpain 1* and *calpastatin* on the tenderness of cooked *M. longissimus dorsi* steaks from Jersey x Limousin, Angus and Hereford-cross cattle. *Animal Genetics* **37**: 411-414.
- Morris, C.A.; Daly, C.C.; Cullen, N.G.; Hickey, S.M. 2001: Correlations among beef carcass composition and meat quality traits from a genetic marker trial. *Proceedings of the New Zealand Society of Animal Production* **61**: 96-99.
- Morris, C.A.; Jordan, T.W.; Loong, P.C.; Lewis, M.H.; Towers, N.R. 1988: Associations between transferrin type and facial eczema susceptibility and some production traits in sheep. *New Zealand Journal of Agricultural Research* **31**: 301-305.
- Nicoll, G.B.; Burkin, H.R.; Broad, T.E.; Jopson, N.B.; Greer, G.J.; Bain, W.E.; Wright, C.S.; Dodds, K.G.; Fennessey, P.F.; McEwan, J.C. 1998: Genetic linkage of microsatellite markers to the Carwell locus for rib-eye muscling in sheep. *Proceedings of the 6th World Congress on Genetics Applied to Livestock Production, Armidale, Australia* **26**: 529-532.
- Page, B.T.; Casas, E.; Heaton, M.P.; Cullen, N.G.; Hyndman, D.L.; Morris, C.A.; Crawford, A.M.; Wheeler, T.L.; Koohmaraie, M.; Keele, J.W.; Smith, T.P.L. 2002: Evaluation of single-nucleotide polymorphisms in *CAPN1* for association with meat tenderness in cattle. *Journal of Animal Science* **80**: 3077-3085.
- Paterson, K.A.; McEwan, J.C.; Dodds, K.G.; Morris, C.A.; Crawford, A.M. 2001: Fine mapping a locus affecting host resistance to internal parasites in sheep. *Proceedings of the Association for the Advancement of Animal Breeding and Genetics* **14**: 91-94.
- Phua, S.H.; Dodds, K.G.; Morris, C.A.; Paterson, K.A.; McEwan, J.C.; Garmonsway, H.G.; Towers, N.R.; Crawford, A.M. 1999: Catalase gene is associated with facial eczema disease resistance in sheep. *Animal Genetics* **30**: 286-295.
- Schaeffer, L.R. 2006: Strategy for applying genome-wide selection in dairy cattle. *Journal of Animal Breeding and Genetics* **123**: 218-223.
- Sellick, G.S.; Pitchford, W.S.; Morris, C.A.; Cullen, N.G.; Crawford, A.M.; Raadsma, H.W.; Bottema, C.D.K. 2007: Major QTL affecting carcass yield in cattle: myostatin F94L. *Animal Genetics*: (in press).
- Skerman, T.M.; Johnson, D.L.; Kane, D.W.; Clarke, J.N. 1988: Clinical footscald and footrot in a New Zealand Romney flock: phenotypic and genetic parameters. *Australian Journal of Agricultural Research* **39**: 907-916.
- Slate, J.; Van Stijn, T.C.; Anderson, R.M.; McEwan, K.M.; Maqbool, N.J.; Mathias, H.C.; Bixley, M.J.; Stevens, D.R.; Molenaar, A.J.; Beever, J.E.; Galloway, S.M.; Tate, M.L. 2002: A deer (subfamily Cervinae) genetic linkage map and the evolution of ruminant genomes. *Genetics* **160**: 1587-1597.
- Spelman, R.J.; Coppieters, W.; Grisart, B.; Blott, S.; Georges, M. 2001: Review of QTL mapping in the New Zealand and Dutch dairy cattle populations. *Proceedings of the Association for the Advancement of Animal Breeding and Genetics* **14**: 11-16.
- Spelman, R.J.; Coppieters, W.; Karim, L.; van Arendonk, J.A.M.; Bovenhuis, H. 1996: Quantitative trait loci analysis for five milk production traits on chromosome six in the Dutch Holstein-Friesian population. *Genetics* **144**: 1799-1808.
- Spelman, R.J.; Ford, C.A.; McElhinney, P.; Gregory, G.C.; Snell, R.G. 2002: Characterization of the DGAT1 gene in the New Zealand dairy population. *Journal of Dairy Science* **85**: 3514-3517.

- Spelman, R.J.; Hooper, J.D.; Stanley, G.; Kayis, S.A.; Harcourt, S. 2004: Friesian Jersey crossbred trial: Generating phenotypes for the discovery of quantitative trait loci. *Proceedings of the New Zealand Society of Animal Production* **64**: 92-95.
- Spelman, R.J.; Huisman, A.E.; Singireddy, S.R.; Coppieters, W.; Arranz, J.; Georges, M.; Garrick, D.J. 1999: Quantitative trait loci analysis on 17 non-production traits in the New Zealand dairy population. *Journal of Dairy Science* **82**: 2514-2516.
- Spelman, R.J.; Sellars, M.D.; Sanders, K. 2000: Experimental design for detection of quantitative trait loci for susceptibility to facial eczema in dairy cattle. *Proceedings of the New Zealand Society of Animal Production* **60**: 281-283.
- Tate, M.L.; Goosen, G.J.; Patene, H.; Pearse, A.J.; McEwan, K.M.; Fennessy, P.F. 1997a: Genetic analysis of Père David's x red deer interspecies hybrids. *Journal of Heredity* **88**: 361-365.
- Tate, M.L.; McDonald, R.W.; Ward, J.F.; Dodds, K.G. 1998: DNA-matching in deer. *Proceedings of a deer course for veterinarians* **15**: 119-125.
- Tate, M.L.; Mathias, H.C.; McEwan, K.M.; Anderson, R.N. 1997b: Translation of information between human and livestock gene maps using a deer interspecies hybrid mapping panel. *Proceedings of the New Zealand Society of Animal Production* **57**: 301-302.
- van der Linde-Sipman, J.S.; van den Ingh, T.S.G.A.M.; Vellema, P. 2003: Morphology and morphogenesis of hereditary microphthalmia in Texel sheep. *Journal of Comparative Pathology* **128**: 269-275.
- Visser, P.M.; Medland, S.E.; Ferreira, M.A.R.; Morley, K.I.; Zhu, G.; Cornes, B.K.; Montgomery, G.W.; Martin, N.G. 2006: Assumption-free estimation of heritability from genome-wide identity-by-descent sharing between full siblings. *PLoS Genetics* **2**: e41.
- Ward, J.F.; McDonald, R.W.; Dodds, K.G.; Tate, M.L. 2001: Procedures for large scale DNA pedigree matching. *Proceedings of the Association for the Advancement of Animal Breeding and Genetics* **14**: 321-324.
- Weller, J.I.; Kashi, Y.; Soller, M. 1990: Power of daughter and granddaughter designs for determining linkage between marker loci and quantitative trait loci in dairy cattle. *Journal of Dairy Science* **73**: 2525-2537.
- Wilson, T.; Wu, X.-Y.; Juengel, J.L.; Ross, I.K.; Lumsden, J.M.; Lord, E.A.; Dodds, K.G.; Walling, G.A.; McEwan, J.C.; O'Connell, A.R.; McNatty, K.P.; Montgomery, G.W. 2001: Highly prolific Booroola sheep have a mutation in the intracellular kinase domain of bone morphogenetic protein IB receptor (ALK-6) that is expressed in both oocytes and granulosa cells. *Biology of Reproduction* **64**: 1225-1235.