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Design and implementation of the Deer Progeny Test (DPT)

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Abstract

The Deer Progeny Test (DPT) program was established in 2011, to better measure the venison production genetics available to the New Zealand deer industry and to allow commercial farmers to better utilise them to improve productivity and profitability. The DPT has five principal aims: (1) encourage and improve sire linkage across recorded breeding herds; (2) provide a platform to evaluate breeding values across the breeds of red and wapiti; (3) evaluate new traits and generate genetic correlations to allow optimisation of selection goals; (4) provide a starting point for the evaluation of maternal traits under commercial breeding environments; and (5) establish a well-phenotyped population, for future genomic tool development. The DPT is a three-year breeding program run across three different farms, progeny were born on two farms annually. Three years of artificial insemination (AI) were completed in April 2013 and involved 2417 inseminations, using 35 different sires and 1581 hinds, with an average rate of progeny weaned to hinds inseminated of 68%. All maternal (red) male and terminal (wapiti crossbred) progeny are slaughtered at approximately 11 months of age to measure carcass traits while maternal females are retained into the breeding herd to measure maternal traits. The final progeny slaughter is due late 2014. Seasonal live weights and temperament scores are collected on progeny and dams, while ultrasound eye muscle area, parasite antibody levels and foetal age are recorded on progeny. A subset of core traits is recorded by partner herds who contributed AI sires. Slaughters occur on a single date for each farm and a variety of meat yield and quality traits are assessed. DPT data is uploaded to DEERSelect from incorporation of these data is publically communicated via Sire Summary reports. The DPT has increased the percentage of linked herds recording growth on DEERSelect from 70 to 100%. This means DEERSelect breeding values and economic indexes produced from red deer can be validly compared across herds.

Keywords: deer; progeny test; DEERSelect; red deer; wapiti; sire

Introduction

In 2007 Deer Industry New Zealand (DINZ) implemented a 5-year productivity strategy (Pearse & Fung 2007). This strategy, aimed at making productivity gains for venison, growth, and reproductive traits for farmed deer, as “the deer industry has struggled to achieve any real productivity gains (in these areas) over the previous 10 years” (Pearse & Fung 2007). This strategy has since been revised and the aims of the later document included: “more deer, earlier, heavier” and “the uptake of genetics, technology and techniques by deer farmers” to enable productivity increases (DINZ 2011a).

Tools for quantitative genetic analysis which provide estimated breeding values (EBVs) for genetic selection of deer have been available to breeders since 1998 and across-herd genetic evaluation for red deer became available following the launch of DEERSelect in 2005 (Archer et al. 2005). A Sire Reference Scheme (SRS) operated at Invermay from 2003-2007 (Archer 2003), and was designed to provide sufficient linkage to allow across-herd genetic evaluations for red deer. DEERSelect, using the Sheep Improvement Limited (SIL) genetic engine with evaluation modules written specifically for deer, performed across-herd genetic evaluations for 15 linked red deer herds using the linkage provided by the SRS (Archer et al. 2005). There have been many subsequent developments in DEERSelect, including the addition of within-herd EBVs for wapiti in 2008 (Archer 2008), new red deer trait modules such as reproduction (Archer 2010), meat yield (Asher 2011) and economic indexes (Archer 2010). There have also been many other updates within the SIL database where DEERSelect resides, such as new methods for reporting genetic connectedness of herds and contemporary groups (linkage) (Visser et al. 2013).

Genetic linkage among herds needs to be maintained as historical links weaken over time. If genetic linkage is inadequate it becomes invalid to compare the genetic merit of animals from different herds (Visser et al. 2013). By 2010, there was concern that linkage established by using common sires in the SRS 2003-2005 birth cohorts had weakened due to the links established not being maintained via ongoing usage of common sires across breeding herds, as had been anticipated (Archer et al. 2011). This was confirmed, when the April 2011 DEERSelect red deer sire summaries were reported, with only 14 out of 20 herds recording growth (live weight) traits adequately connected (under present SIL criteria (Visser et al. 2013)) for across herd reporting (Table 1). Three of the six unlinked herds had contributed sires to the SRS and had been linked in 2005.

Red deer and wapiti are different subspecies with large morphological differences between them as such they are evaluated separately on DEERSelect using...
The DPT project is primarily growth and venison focussed, and therefore, both red deer (maternal sire type) and wapiti or wapiti crossbred (terminal sire type) sires were included for evaluation, as the annual contribution of wapiti crossbreds for venison production up to 50% of the annual number of animals slaughtered (Shackell et al. 2003). A range of these sires was used and their progeny evaluated under commercial conditions, such as hill and high-country farms where many commercial breeding hinds are now farmed (Netzer et al. 2009), because farmers question what the most suitable hind genotype is for such environments.

The DPT has five principal aims, to: (1) encourage and improve sire linkage across recorded breeding herds; (2) provide a platform to evaluate breeding values across the breeds of red and wapiti (maternal and terminal); (3) evaluate new traits (especially venison related) and generate genetic correlations to allow optimisation of selection goals; (4) provide a starting point for the evaluation of maternal traits under commercial breeding environments; and (5) establish a well-phenotyped, DNA resourced population, for future genomic tool development.

### Establishment of the DPT

At a meeting held at AgResearch Invermay in December 2010, involving DINZ staff, breeders, scientists, industry geneticists, farm consultants and veterinarians, a proposal to create a centralised genetics resource to achieve multiple outcomes ranging from near-term applied focus (e.g., herd genetic linkage) to longer-term research (e.g., development of resources for genomic research) was presented. At the same time, food processors Alliance Group Limited (AGL) expressed interest in supporting a progeny test for deer to evaluate meat and carcass yield and quality characteristics, similar to the Alliance Central Progeny Test (CPT) for sheep (Campbell et al. 2003). Endorsement by breeders and support from AGL and DINZ lead to establishment of a working party to guide implementation of the DPT project.

Other industry participants, particularly venison processors were invited to be involved in the industry-wide project. Following this, Landcorp Farming Limited (LFL) agreed to contribute funds alongside AGL, DEEREselect, and AgResearch (MBIE Core Funds). Breeders were asked to submit expressions of interest to become partner herds. This involved contributing semen (gratis) from sire stags used heavily in individual herds (and ideally across several others), in the year of corresponding DPT progeny conception. There was also a committed undertaking to record on DEEREselect for key growth, meat and reproduction traits.

### Goals of the DPT

The DPT project is

<table>
<thead>
<tr>
<th>Trait group</th>
<th>Herds connected Apr 2011</th>
<th>Herds connected Dec 2013</th>
<th>% Herds connected Apr 2011</th>
<th>% Herds connected Dec 2013</th>
</tr>
</thead>
<tbody>
<tr>
<td>Growth</td>
<td>14/20</td>
<td>24/24</td>
<td>70%</td>
<td>100%</td>
</tr>
<tr>
<td>Reproduction</td>
<td>4/6</td>
<td>9/12</td>
<td>67%</td>
<td>75%</td>
</tr>
<tr>
<td>Meat-yield*</td>
<td>N/A</td>
<td>9/9</td>
<td>N/A</td>
<td>100%</td>
</tr>
</tbody>
</table>

*Meat-yield traits not available on DEEREselect until October 2011

different genetic parameters (Archer et al. 2008). This means that EBVs for wapiti crossbreds and red deer are not directly comparable. Commercial deer farmers primarily use wapiti as terminal sires for venison production (Shackell et al. 2003) and the impossibility of directly comparing red and wapiti EBVs has made sire purchase decisions based on EBVs difficult or confusing for farmers.

Venison dominates the deer industry’s export revenue, accounting for greater than 75% of recent past annual revenue from 2007-2011 (DINZ 2011b). By contrast, the maximum velvet antler had contributed was 11%; less than the 13% averaged by other products associated with venison production, (e.g. co-products, hides and leather, DINZ 2011b). Popular opinion is that antler traits (velvet antler and trophy) dominate the promotion of breeding stags for sale, which is disproportionate to the revenue generated by antler across the industry.

### Operation of the DPT

The DPT has run for three breeding cycles (2011-2013). Each cycle aimed to provide 25 male and 25 female progeny per maternal sire and 25 mixed-sex progeny of terminal sires across two DPT farms (Fig. 1). It was assumed that the minimum AI conception rate of would be 70%, therefore each maternal sire would require 72 hinds and each terminal sire 36 hinds to be inseminated by trans-cervical AI (Rhodes et al. 2003). The aim was to inseminate 800 hinds across both DPT farms (400 per farm) each year, producing approximately 560 progeny from 11 sires annually, linked by one maternal and one terminal sire across the three years. Each sire was used evenly on both DPT farms where many commercial breeding hinds are now farmed (Netzer et al. 2009), because farmers question what the most suitable hind genotype is for such environments.

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partner herds used the sire(s) submitted for AI that cycle within their own herds in the same year as the DPT and recorded those and all stud progeny pedigrees and core traits on DEERSelect.

The primary selection criteria of stags submitted for use in the DPT by partner herds was their ability to improve linkage across industry (i.e. the number of recorded progeny, or planned progeny for the coming year). Genetic merit of the sires is also a consideration as the industry and DPT farm owners do expect above average performance. Table 2 shows the genetic merit of DPT red deer sires to be around 40% above the average sire recorded on DEERSelect. Dams on all farms except Invermay are commercial breeding hinds and as such had no previous individual performance data recorded on DEERSelect or otherwise.

The DPT has three categories of trait recording; core traits, venison traits and new (research) traits. Core traits include quarterly live weights, progeny fates, ultrasound-eye muscle area (EMAUS) and estimated conception date (ECD) via foetal aging (White et al. 2003) and are recorded by the partner herds and DPT farms. Venison traits are measured by as such had no previous individual performance data recorded on DEERSelect or otherwise.

Figure 1 Design of deer progeny test (DPT) for numbers of progeny generated per sire of each sire-type, traits measured and planned fates for each sex of those progeny-types.

Table 2 Mean estimated breeding values (EBVs) and economic indexes of the red deer sires of the 2011 and 2012 deer progeny test birth cohorts and mean of all red sires reported from the DEERSelect December 2013 Sire Summary.

<table>
<thead>
<tr>
<th>Red sires</th>
<th>Estimated breeding values</th>
<th>Economic indexes</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>W12 (kg)</td>
<td>CW (kg)</td>
</tr>
<tr>
<td>DPT Sires</td>
<td>+17.2</td>
<td>+9.0</td>
</tr>
<tr>
<td>All Sires</td>
<td>+10.2</td>
<td>+5.3</td>
</tr>
<tr>
<td>Difference</td>
<td>+7.0</td>
<td>+3.7</td>
</tr>
</tbody>
</table>

Where W12 = live weight at 12 months of age, CW = carcass weight adjusted for 12-month live weight, MWT = mature live weight of hinds, CD = conception date (negative is earlier conception), R-EK = replacement breeding retaining females and killing males early, and Term = terminal breeding killing all progeny (Archer & Amer 2009).
Table 3 Summary of the number of hinds artificially inseminated (AI) for the three years of deer progeny test (DPT) and the numbers and percentages of DPT progeny (calves) present at weaning conceived by DPT AI program.

<table>
<thead>
<tr>
<th>AI Year</th>
<th>DPT Farm</th>
<th>Hinds inseminated</th>
<th>Calves at weaning conceived by AI (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>2011</td>
<td>Invermay</td>
<td>399</td>
<td>258 (65%)</td>
</tr>
<tr>
<td>2011</td>
<td>Whiterock</td>
<td>417</td>
<td>258 (62%)</td>
</tr>
<tr>
<td>2012</td>
<td>Haldon</td>
<td>400</td>
<td>287 (72%)</td>
</tr>
<tr>
<td>2012</td>
<td>Whiterock</td>
<td>401</td>
<td>308 (77%)</td>
</tr>
<tr>
<td>2013</td>
<td>Haldon</td>
<td>403</td>
<td>270 (67%)</td>
</tr>
<tr>
<td>2013</td>
<td>Invermay</td>
<td>397</td>
<td>257 (65%)</td>
</tr>
</tbody>
</table>

Table 4 Summary metrics of red deer sires linked for growth traits (live weights) on DEERSelect from across-herd Sire Summary reporting (sires of last three-years-progeny cohorts) before (April 2011) and after (December 2013) the inclusion of data from deer progeny test (DPT) 2011- and 2012-born cohorts.

<table>
<thead>
<tr>
<th></th>
<th>Apr 2011</th>
<th>Dec 2013</th>
<th>Increase %</th>
</tr>
</thead>
<tbody>
<tr>
<td>Number of sires providing linkage</td>
<td>98</td>
<td>112</td>
<td>12.5%</td>
</tr>
<tr>
<td>Mean number of herds linked per sire</td>
<td>3.9</td>
<td>5.1</td>
<td>23.5%</td>
</tr>
<tr>
<td>Mean number of progeny linked per sire</td>
<td>21.3</td>
<td>24.4</td>
<td>12.7%</td>
</tr>
</tbody>
</table>

At the completion of the three breeding cycles 68% was the overall rate of progeny weaned to hinds inseminated. The highest rate was in 2012 when Haldon and Whiterock achieved better results of 72% and 77% respectively (Table 3). The rate of progeny weaned to hinds inseminated was lower and more variable for terminal than maternal sires, but this was not statistically significant (p<0.05).

Traits and communication

As of April 2014 the 2011 and 2012 DPT farm-progeny cohorts had all their growth traits (weaning, six weeks post-weaning, pre-winter, post-winter, spring and pre-slaughter live weights and ultrasound-eye-muscle area) measured. All post-slaughter yield and co-product traits have also been recorded. Meat quality and colour stability have been measured for the 2011 and 2012 cohorts, and all skin measurements are incomplete. Maternal females from 2011 have been ultrasound-scanned for estimated conception date at 18-months of age, weighed pre- and post-mating, fates recorded and they have weaned their first progeny 27-months of age. All progeny and dams in the DPT have been temperament scored on at least one occasion. CarLA samples have been collected pre-winter and pre-slaughter for 2011 and 2012, with samples for the 2011 cohort analysed as described by Mackintosh et al. (2012). The 2013 cohort has been weaned and weighed at weaning, and six weeks-post weaning. All traits accepted by DEERSelect have been uploaded.

The DPT does not publically publish any rankings of the sires used in the DPT. The intent of this is to not distract from the key aim of improving linkage. DEERSelect has been selected as the primary vehicle for communication of DPT information. Data for those traits able to be recorded on DEERSelect is uploaded to DEERSelect to strengthen herd linkage. Quarterly Sire Summary reports of red deer data are publically available via the DINZ website (http://www.deernz.org/). The first report incorporating DPT data was released in July 2012. These Sire Summaries report a range of BVs and economic indexes for all recorded breeding sires including DPT sires. Partner herds receive data summaries of DPT progeny twice a year. These are a mixture of confidential raw data e.g. live and carcass weights by sire; analysed data reported to DEEResearch e.g. carcass data by sire type; and updated linkage and genetic trend data from DEERSelect.

Linkage

In the December 2013 red deer across-herd sire summary report all herds actively recording on DEERSelect were linked for growth (n=24) and meat traits (n=9) (Table 1), compared to 70% before the DPT. There have also been increases in all metrics associated with the linkage provided by each sire (Table 4), most notably an increase in the mean number of herds linked from 3.9 to 5.1.

Across-herd analysis for wapiti (terminal sires) is not published presently by DEERSelect (as of January 2014) as there are only three breeding herds that meet the SIL connectedness criteria.
Conclusion

With the final progeny cohort weaned, the DPT is well advanced and large amounts of data have being collected. While there has been preliminary analysis of some data, a full sire genetic analysis requires the data set from all three-year’s progeny cohorts. This will not be available until June 2015 at the earliest. The DPT has already succeeded in focussing the deer industry on better performance recording (S McIntyre pers. comms.) and strengthened the linkage of data reported from DEERSelect, two of the project’s primary goals.

For the DPT to be a success and achieve maximum impact in the venison industry a number of practices still need to occur within and alongside the DPT. This includes increased numbers of commercial venison producers purchasing stags from venison-focussed breeders, who are making genetic improvement and using DEERSelect economic indexes to assist breeding stag purchase decisions. These venison-production-focussed breeders and producers need to implement clear venison-focussed breeding objectives. There needs to be on-going sire (genetic) exchange between breeders, to maintain and improve established herd linkage in the future, along with improved recording practices by those breeders. Finally utilisation of DEERSelect needs to be greatly increased across the entire NZ deer industry.

Acknowledgements

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DPT Farm Owners, Managers and staff: AgResearch Invermay, Whiterock Station and Haldon Station.

References


