

LOT	ID	PWT	YWT	YFAT	YEMD	YFD	YCFW	YFDCV	YSS	YSL	YFEC	NLW	DP+	MP+	BT	RT	SIRE	NOTES
1	259	8.9	13.7	1.5	1.6	0.4	32.0	-1.7	3.3	16.0	-44	4%	184	170	2	2	160140	
2	560	9.5	14.5	1.7	2.1	-0.2	37.4	-0.7	-0.5	11.0	-56	2%	192	179	1	1	160290	
3	351	7.1	11.2	1.2	1.7	-0.1	22.2	-0.2	-1.1	12.5	-74	7%	176	156	1	1	140474	
4	122	7.9	11.7	1.3	2.3	0.7	17.0	-2.0	5.5	10.4	-69	0%	161	144	1	1	150333	
5	1196	7.2	11.0	2.0	3.7	1.0	19.9	-2.7	5.6	16.0	-67	13%	195	160	1	1	160390 ET	
6	728	9.0	13.4	1.0	1.0	0.0	25.6	-2.9	6.0	8.7	-52	4%	175	168	2	2	160140	
7	234	6.6	10.6	1.1	2.2	0.1	16.4	-1.5	2.5	9.2	-55	2%	165	146	1	1	150048	
8	50	6.1	10.7	1.7	2.6	0.3	29.8	-1.7	2.1	12.1	-80	2%	177	160	1	1	160181	
9	108	8.2	12.7	1.8	3.1	-0.3	31.2	0.1	-4.3	11.5	-39	4%	181	160	1	1	160290	
10	463	6.7	10.3	1.5	2.1	-0.8	21.6	0.1	-3.1	5.6	-41	5%	173	157	2	2	160290	
11	321	8.6	12.2	1.1	1.6	-0.6	32.4	-0.3	-0.8	16.7	-65	1%	172	163	2	1	160210	
12	641	7.8	11.1	1.1	1.8	0.7	17.9	-0.7	3.0	11.0	-66	3%	158	142	2	2	150333	
13	781	7.2	11.0	1.0	2.2	0.1	15.4	-1.3	1.6	10.1	-53	7%	171	148	2	2	150048	
14	623	7.3	13.0	1.3	1.5	-0.6	20.4	-1.0	0.8	11.1	-71	5%	173	158	2	2	160558	
15	643	7.7	8.7	2.1	3.4	-0.3	16.4	-2.6	4.0	13.7	-76	4%	166	146	2	2	140343	
16	60	9.9	14.4	1.2	2.0	-0.1	22.6	-3.0	5.9	17.3	-70	13%	197	176	2	2	160729	
17	362	7.8	11.8	1.3	2.2	-0.5	27.7	0.1	-1.8	9.6	-47	-2%	168	155	1	1	160290	
18	279	7.6	12.2	2.1	2.9	0.8	12.5	-1.0	3.0	12.2	-68	6%	168	140	2	1	160224	
19	520	9.7	14.6	1.4	2.8	-0.7	18.7	-1.5	-0.9	8.1	-73	9%	189	162	1	1	140474	
20	862	7.6	12.1	1.3	2.4	-0.7	31.2	0.4	-2.8	11.2	-50	2%	184	168	1	1	160290	
21	1193	8.2	10.5	1.4	2.9	-0.5	22.3	-1.0	1.1	7.9	-61	0%	171	152	1	1	140474 ET	
22	1199	8.3	9.5	1.6	3.1	-0.8	14.2	-1.6	1.9	3.9	-61	0%	162	143	1	1	140474 ET	
23	123	9.5	13.4	1.6	2.6	0.6	19.3	-2.3	3.9	11.7	-55	1%	165	144	1	1	160558	
24	516	7.0	11.3	1.2	1.6	0.4	28.4	-1.9	5.2	16.8	-65	3%	176	164	1	1	160729	
25	886	7.8	11.6	1.6	2.1	0.1	16.8	-1.6	2.8	11.4	-77	5%	167	150	1	1	150266	
26	677	10.1	14.0	1.3	2.3	-0.1	16.4	-2.1	3.6	7.5	-56	9%	181	158	2	2	160558	
27	479	6.3	9.6	1.2	1.5	1.0	34.7	-0.3	0.4	21.3	-88	-3%	156	147	2	2	140178	
28	942	8.7	11.3	1.9	3.8	0.1	10.8	-2.1	2.3	7.6	-62.1	6%	170	138	2	2	140474	
29	247	6.2	9.2	2.2	2.7	0.1	20.4	-3.0	4.5	7.2	-64	0%	159	146	1	1	160558	
30	848	3.7	5.9	1.7	3.4	0.7	20.0	-2.9	7.5	13.0	-88	1%	155	138	2	2	140343	
31	204	7.3	10.4	1.0	1.7	-0.7	32.4	0.0	-1.2	12.6	-43	0%	176	168	1	1	160290	
32	917	5.5	9.2	0.8	2.0	-1.5	26.3	-0.7	0.2	-1.9	-64	-4%	167	163	1	1	160269	

33	232	9.3	11.6	1.7	3.2	-1.1	14.2	-0.8	-0.8	12.3	-66	7%	181	154	2	2	160390	
34	310	6.5	8.1	0.9	1.6	-0.6	29.8	-0.4	0.4	8.2	-60	1%	167	160	2	2	160210	
35	173	8.7	13.3	1.2	2.0	-0.1	24.5	-1.7	2.5	10.5	-62	4%	177	162	1	1	160558	
36	280	9.8	11.9	1.3	2.1	-0.3	14.4	-2.4	3.0	8.0	-62	5%	164	147	2	2	160558	
37	175	6.4	11.9	2.0	2.9	0.1	15.5	-2.9	4.9	10.3	-76	2%	168	149	1	1	160224	
38	231	8.3	11.2	1.8	3.3	-0.8	15.9	0.0	-1.9	10.5	-73	7%	179	151	2	2	160390	
39	875	7.4	12.3	2.0	2.3	-0.1	17.8	-1.4	2.6	8.7	-88	7%	177	157	1	1	160224	B/+ Booroola gene
40	464	7.8	11.7	1.8	2.7	0.1	21.7	-0.1	-1.2	3.7	-31	6%	173	151	2	2	160290	
41	408	8.2	11.5	1.3	1.9	-0.2	20.9	-0.6	1.0	11.8	-56	9%	181	160	2	2	160390	
42	721	5.1	10.0	1.3	2.5	0.1	33.6	1.1	-4.2	13.6	-40	3%	178	160	2	2	160290	
43	183	8.1	10.7	0.9	2.1	-0.7	28.8	-0.8	-0.4	7.5	-49	4%	185	169	2	2	140474	
44	436	7.3	10.5	1.7	3.1	0.7	15.2	-1.9	2.0	12.2	-64	8%	168	138	1	1	160390	
45	16	8.7	11.7	1.6	3.4	-0.3	21.9	-0.8	-0.9	11.3	-59	10%	190	159	2	2	140474	
46	414	6.6	10.6	1.5	2.6	0.0	11.1	-0.7	2.4	5.7	-76	7%	170	145	3	3	150333	
47	428	6.2	9.9	1.0	1.7	-0.4	23.8	-1.0	0.2	11.2	-75	6%	171	156	2	2	160210	
48	836	7.8	11.3	1.4	2.2	-0.9	20.8	-1.2	0.6	8.8	-78	6%	178	161	3	3	160390	
49	621	5.8	8.4	1.9	3.4	0.0	20.6	-1.5	3.6	10.9	-79	7%	177	154	2	2	140343	
50	651	7.2	11.2	1.4	1.1	0.1	26.1	-2.7	6.4	7.1	-52	2%	169	165	2	2	160140	
51	184	8.5	13.0	1.1	2.0	-0.8	18.2	-1.1	0.4	15.1	-59	5%	175	158	1	1	160729	
52	1179	6.1	10.4	1.3	1.9	0.5	23.1	-1.0	0.0	8.7	-50	0%	173	157	1	1	syndicate	
53	499	5.9	8.7	1.1	2.5	-0.4	23.0	-1.8	1.6	14.4	-71	9%	184	162	2	2	160390	
54	379	9.0	13.2	2.0	2.7	1.2	30.1	-2.4	5.4	13.5	-54	1%	176	157	2	1	160140	
55	610	6.9	10.6	1.6	2.6	-0.4	23.1	-0.7	-0.1	9.6	-54	2%	170	152	2	2	160290	
56	349	8.7	10.6	0.8	1.3	-0.2	23.2	-1.0	2.7	9.6	-61	3%	171	162	1	1	150333	
57	927	7.8	12.3	1.9	3.1	1.1	10.1	-0.9	3.0	8.7	-80	8%	167	133	2	2	160224	
58	399	8.7	11.3	1.6	1.9	0.5	21.7	-2.9	6.2	9.5	-40	1%	163	150	1	1	160140	
59	496	5.8	9.1	1.0	1.7	-0.4	24.0	-1.0	1.8	14.6	-80	3%	165	154	2	2	160210	
60	885	7.3	11.4	1.5	1.6	0.2	10.8	-2.7	4.0	10.8	-51	10%	166	147	2	2	150266	
61	177	8.7	14.2	1.7	2.3	0.1	27.0	-1.7	2.6	9.9	-41	9%	196	175	2	2	160290	
62	536	6.8	10.1	0.7	1.2	-0.5	24.8	-2.5	5.6	8.5	-51	1%	165	162	1	1	160140	
63	326	6.2	9.3	1.4	1.8	0.4	26.7	-2.1	5.9	7.8	-49	-3%	156	150	1	1	160140	
64	672	6.7	10.5	1.1	1.5	-0.9	9.1	-1.9	1.4	10.1	-60	1%	148	136	2	2	160558	
65	773	5.7	8.2	1.6	2.1	-0.1	18.9	-3.4	7.0	6.0	-44	-3%	150	143	2	2	160140	

66	686	5.3	7.0	1.1	1.8	0.5	18.7	-3.6	9.5	11.7	-53	-3%	144	139	2	2	160140	
67	374	7.4	11.0	1.7	2.8	0.7	16.1	-2.3	5.7	12.6	-53	4%	164	143	2	1	160224	
68	569	5.2	7.4	1.0	2.0	-0.2	15.7	-1.7	3.7	12.7	-58	0%	154	141	1	1	150048	
69	38	4.1	8.6	0.9	0.9	-0.8	16.9	-0.9	-1.0	11.5	-80	-2%	142	139	2	2	140178	
70	61	5.5	8.8	1.3	2.9	0.0	21.6	-0.4	0.1	15.2	-78	2%	166	145	1	1	160224	B/+ Booroola gene
71	284	4.2	6.0	1.1	2.0	-0.4	17.0	0.0	0.1	16.4	-85	2%	149	137	2	2	160210	
72	165	5.2	9.8	1.4	1.2	-0.4	20.5	-2.7	5.3	8.7	-70	-3%	155	153	1	1	160140	
73	638	5.9	7.7	0.9	0.8	-0.7	29.1	-0.7	-0.6	13.8	-74	1%	160	159	1	1	160210	
74	558	8.1	11.4	1.2	1.4	-0.8	17.9	-0.9	0.5	2.4	-60	2%	161	151	1	1	160558	
75	609	6.6	9.6	1.6	2.8	-0.2	19.0	-0.5	0.0	7.1	-48	1%	162	143	2	2	160290	
76	745	4.2	8.2	1.3	1.3	-1.1	10.7	-0.9	1.9	6.3	-62	8%	165	153	2	2	150266	
77	253	8.3	12.7	1.4	1.8	0.4	28.3	-1.0	1.6	12.0	-74	4%	180	162	2	1	160558	
78	911	7.8	11.0	0.9	1.2	-0.2	19.1	-0.5	0.2	10.1	-79	5%	162	148	2	2	160558	
79	333	6.9	8.0	1.1	2.4	-0.4	24.4	-1.1	0.5	10.4	-60	10%	184	163	2	2	140474	
80	701	5.8	9.8	1.2	2.0	0.1	18.2	-2.5	6.3	14.7	-79	4%	166	153	2	2	160729	
81	915	7.6	12.7	1.6	2.3	0.7	26.6	-2.3	6.6	15.2	-64	2%	176	160	1	1	160729	
82	413	6.3	10.7	1.1	1.2	-0.7	10.1	-1.8	1.6	8.1	-58	4%	154	142	2	2	150266	
83	233	7.3	11.9	1.3	1.8	-0.2	11.6	-2.1	3.6	9.2	-61	3%	156	141	1	1	160558	
84	139	6.1	10.7	1.8	2.8	1.4	20.9	-2.1	5.3	16.3	-86	4%	167	144	1	1	160224	B/+ Booroola gene
85	588	5.6	9.1	1.7	3.4	0.1	20.5	-0.7	-0.6	15.3	-79	11%	183	152	2	2	160194	
86	1151	4.7	8.7	1.5	2.6	0.2	19.6	-0.5	0.0	8.2	-75	0%	170	151	1	1	syndicate	
87	690	3.5	9.1	1.9	3.0	-1.1	21.3	-1.7	-0.1	10.0	-64	5%	181	161	2	1	160181	
88	553	3.7	5.3	1.6	3.1	-0.2	21.3	-1.0	1.7	11.7	-76	1%	157	141	1	1	140343	
89	294	5.4	6.6	1.1	2.7	1.1	23.1	-3.1	5.4	15.7	-65	-7%	144	131	1	1	150048	
90	115	4.6	6.9	1.0	2.3	-0.8	12.6	-1.1	1.3	6.6	-78	7%	166	147	1	1	160390	
91	896	4.1	6.4	1.5	2.4	-0.2	17.5	0.2	-1.1	9.7	-63	2%	154	137	1	1	160194	
92	475	6.9	10.1	1.3	2.7	-0.4	22.2	-1.5	1.5	9.6	-70	-1%	165	148	2	2	140474	
93	601	4.9	7.3	0.8	2.0	-1.1	24.9	0.5	-2.7	11.8	-85	-3%	152	146	1	1	160210	
94	814	5.9	9.6	1.3	2.0	-0.6	13.2	-1.7	1.5	10.1	-75	2%	158	142	2	2	140178	
95	491	6.7	11.1	1.4	1.8	-0.1	15.2	-1.7	1.6	10.5	-87	3%	159	143	2	2	140178	
96	1132	7.5	12.2	1.8	2.8	-0.3	14.8	-1.5	0.0	8.3	-56	0%	177	154	1	1	syndicate	
97	348	9.7	13.8	1.9	3.2	0.5	34.1	1.0	-3.2	14.7	-44	6%	192	165	1	1	160290	
98	679	4.7	8.6	1.2	2.0	-0.6	21.7	1.1	-4.5	12.5	-72	-1%	155	141	1	1	140178	

99	114	9.1	10.5	1.4	2.9	0.4	24.0	0.1	-1.2	15.3	-58	9%	181	151	2	2	160390	
100	722	5.7	10.7	1.4	2.4	-0.9	26.1	-0.8	-1.6	11.6	-40	3%	178	164	2	2	160290	
101	78	6.1	9.0	1.3	1.9	-0.7	23.8	-0.3	-1.4	9.9	-84	-2%	159	149	1	1	140178	
102	549	5.2	8.6	0.8	1.6	0.3	24.6	-0.8	3.0	9.9	-63	0%	157	148	1	1	160558	
103	926	6.7	8.4	1.2	1.9	-0.1	25.0	-1.8	3.8	21.6	-74	0%	159	149	1	1	160210	
104	1144	6.0	10.1	1.7	2.8	0.9	12.1	-1.5	0.0	10.3	-75	0%	159	135	1	1	syndicate	
105	545	5.7	9.0	1.2	1.6	-0.6	25.0	-1.9	2.8	10.2	-70	0%	161	156	1	1	160558	
106	583	5.8	9.9	1.7	2.3	1.1	14.4	-1.2	4.2	12.4	-67	8%	160	134	2	2	160224	
107	343	4.7	7.5	1.0	1.7	-1.1	7.3	0.1	-1.5	3.3	-62.3	9%	159	142	2	2	160194	
108	710	4.8	6.9	1.2	1.6	-0.3	11.4	-2.2	4.4	5.7	-77	5%	153	140	2	2	160194	
109	634	6.7	8.9	1.8	2.3	-0.6	16.4	-2.7	2.9	9.2	-74	4%	162	149	2	2	160558	
110	1148	5.1	8.5		3.3								156		1	1	syndicate	
111	163	6.6	9.3	1.2	2.2	-0.2	17.3	0.2	1.2	4.7	-65	-3%	151	139	1	1	150333	
112	245	6.9	8.4	1.3	2.9	-0.1	15.1	-2.0	3.1	10.0	-64	4%	164	142	1	1	140474	
113	291	7.2	12.1	1.4	2.6	-0.4	15.9	-2.4	4.9	13.6	-67	3%	171	153	1	1	160729	
114	547	5.8	8.5	1.0	1.9	0.0	27.7	0.5	-0.3	11.3	-72	-3%	154	145	1	1	160210	
sale team ave		6.9	10.4	1.4	2.2	-0.1	21.1	-1.3	1.9	10.8	-64	3%	168	152				
industry ave		3	4.6	0.1	0.3	-1	14.6	-0.6	0.5	7.7	-14	1%	138	138				

NOTES

key	
*	top 1% of animals in Merinoselect
*	top 5%
*	top 10%
*	top 20%

SIRES	all Anderson sires
PWWT	post weaning weight
YWT	yearling weight
YFAT	yearling C site fat
YEMD	yearling eye muscle depth
YFD	yearling fibre diameter
YCFW	yearling clean fleece weight
YFDCV	yearling micron CV
YSS	yearling staple strength
YSL	yearling staple length
YCURV	yearling fibre curvature
YFEC	yearling worm egg count
NLW	number of lambs weaned
INDEXES	DP+, MP+
BT	Birth type
RT	Rear type