

# ESSIE SUFFOLKS

MV Accredited: Scrapie Monitored: CLA Tested: Farm Assured

Sex	EAR No	SIRE No born	Maternal EBV	8 Wk wt EBV	Scan wt EBV	Muscle EBV	Fat depth EBV	TERMINAL Index	MATERNAL Index
M	H6:12:001	H6:N7 2	0.86	6.10	11.89	1.63	0.23	3.71	226
M	H6:12:004	H6:11:007 2	1.23	6.87	13.02	4.56	0.32	4.75	267
M	H6:12:006	Y68:L8 2	0.94	7.62	13.29	5.59	1.05	5.38	254
M	H6:12:010	239:11:004 1	0.49	6.20	13.44	3.11	-0.19	4.21	193
M	H6:12:011	H6:N7 1	1.15	6.75	11.61	2.89	0.70	4.07	252
M	H6:12:014	H6:N7 2	1.34	7.70	13.64	3.94	0.61	4.40	294
M	H6:12:017	Y68:L8 2	0.35	6.29	11.55	4.37	0.83	4.46	193
M	H6:12:018	H6:N7 1	1.23	6.48	11.06	4.74	0.01	4.09	246
M	H6:12:025	H6:N78 2	0.43	5.56	10.46	3.44	0.15	2.87	177
M	H6:12:028	H6:N6 2	0.89	6.11	10.96	4.77	0.21	4.04	237
M	H6:12:029	78X:11:055 2	0.39	7.59	14.74	6.29	0.29	5.93	206
M	H6:12:034	H6:N7 1	1.04	6.45	12.13	3.75	-0.06	4.00	236
M	H6:12:037	H6:11:007 3	0.97	6.48	13.20	3.15	-0.13	4.38	240
M	H6:12:038	H6:11:007 3	0.97	6.77	13.57	3.51	0.18	4.78	246
M	H6:12:040	239:11:004 2	0.50	7.09	12.95	6.33	0.72	5.58	215
M	H6:12:042	H6:11:007 2	0.91	7.11	14.02	4.30	0.65	4.86	261
M	H6:12:043	H6:11:007 2	0.91	6.28	11.43	3.88	-0.03	4.25	247
M	H6:12:045	78X:11:055 2	0.62	9.04	18.22	5.08	0.42	7.02	226
M	H6:12:048	239:11:004 2	0.63	5.27	9.43	3.63	-0.47	3.15	194

Breed Average

£1.51 \*

124 \*

FLOCKS

Top 25%

£2.05 \*\*

152 \*\*

H6 = Essie

Top 10%

£3.34 \*\*\*

192 \*\*\*

Y68 = Drinkstone

Top 5%

£4.03 \*\*\*\*

218 \*\*\*\*

239 = Bentley

Top 1%

£4.90\*\*\*\*\*

256 \*\*\*\*\*

2013 Terminal Sire Index

2013 Maternal Index

**Mobile 07721517969**

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M	H6:12:050	H6:N78 2	0.61	4.68	8.16	2.64	-0.51	1.85	194
M	H6:12:052	H6:11:007 2	1.12	6.31	10.75	4.50	0.37	4.56	258
M	H6:12:054	H6:N7 3	0.98	5.56	8.91	3.32	0.44	3.16	248
M	H6:12:055	H6:N7 3	0.98	5.26	9.12	2.90	-0.15	2.92	243
M	H6:12:057	78X:11:055 2	0.84	7.13	15.00	3.63	-0.29	4.30	201
M	H6:12:058	78X:11:055 2	0.84	6.99	13.59	4.07	0.30	4.71	199
M	H6:12:062	Y68:L8 2	0.63	9.07	18.11	5.27	0.44	7.05	264
M	H6:12:065	H6:N7 2	0.86	6.91	12.00	5.00	0.57	4.64	253
M	H6:12:068	239:11:004 2	0.97	6.38	11.48	4.82	-0.13	4.64	245
M	H6:12:072	H6:N78 2	0.67	6.24	11.47	3.40	0.95	3.20	227
M	H6:12:073	H6:N78 2	0.67	6.23	11.22	2.86	1.67	2.48	227
M	H6:12:075	H6:N7 1	1.32	6.95	12.26	4.67	-0.29	3.80	265
M	H6:12:080	H6:N78 2	0.67	5.93	11.53	3.70	0.07	3.69	237
M	H6:12:086	H6:11:007 3	1.13	6.13	12.67	3.69	-0.13	3.96	271
M	H6:12:087	H6:11:007 3	1.13	6.25	11.55	4.58	-0.14	4.10	273
M	H6:12:091	78X:11:055 2	0.73	7.85	15.84	4.10	0.55	5.69	237
M	H6:12:093	H6:11:007 2	0.63	7.43	14.13	4.79	-0.21	5.64	239
M	H6:12:094	H6:11:007 2	0.74	6.64	12.59	3.98	0.38	4.46	230
M	H6:12:095	H6:11:007 2	0.74	5.57	10.89	3.46	0.03	3.88	211
M	H6:12:096	H6:N7 2	0.95	5.85	10.54	3.32	-0.41	3.05	234
M	H6:12:098	H6:11:007 2	1.24	7.28	13.50	5.65	0.11	5.18	291
M	H6:12:102	Y68:L8 3	0.49	6.02	11.68	3.98	0.68	4.67	218

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M	H6:12:103	H6:N7 2	0.85	5.35	9.08	4.40	0.18	3.39	214
M	H6:12:105	H6:N7 2	0.94	5.32	8.57	3.36	0.09	3.28	233
M	H6:12:107	H6:N7 3	0.93	5.04	9.31	2.59	0.00	3.23	230
M	H6:12:109	239:11:004 2	0.72	5.59	11.99	4.88	0.05	4.52	206
M	H6:12:112	H6:N7 2	0.73	5.75	1.32	3.27	0.08	3.59	235
M	H6:12:126	H6:N7 2	1.31	6.15	10.23	3.71	0.51	3.61	253
M	H6:12:130	H6:N7 3	1.08	5.87	9.67	3.05	-0.02	3.44	249
M	H6:12:131	H6:N7 3	1.08	6.80	12.39	3.09	0.57	4.05	265
M	H6:12:133	H6:N7 3	0.76	4.30	7.61	1.11	-0.20	2.11	206
M	H6:12:137	239:11:004 2	0.38	5.81	11.21	5.83	0.21	5.17	197
M	H6:12:139	Y68:L8 2	0.81	7.81	15.70	4.24	1.04	5.33	244
M	H6:12:143	78X:11:055 2	0.03	4.84	10.58	4.24	-0.16	4.34	106
M	H6:12:145	Y68:L8 2	0.77	6.18	11.68	3.05	0.57	4.37	217
M	H6:12:150	Y68:L8 1	0.86	7.71	14.23	5.20	1.06	5.19	239
M	H6:12:151	Y68:L8 2	0.70	5.90	11.70	3.58	0.17	4.60	232
M	H6:12:155	Y68:L8 2	0.54	7.63	15.20	4.38	0.03	5.63	220
M	H6:12:163	Y68:L8 2	0.97	6.63	12.61	4.14	0.57	4.79	272

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