

## Appendix C4

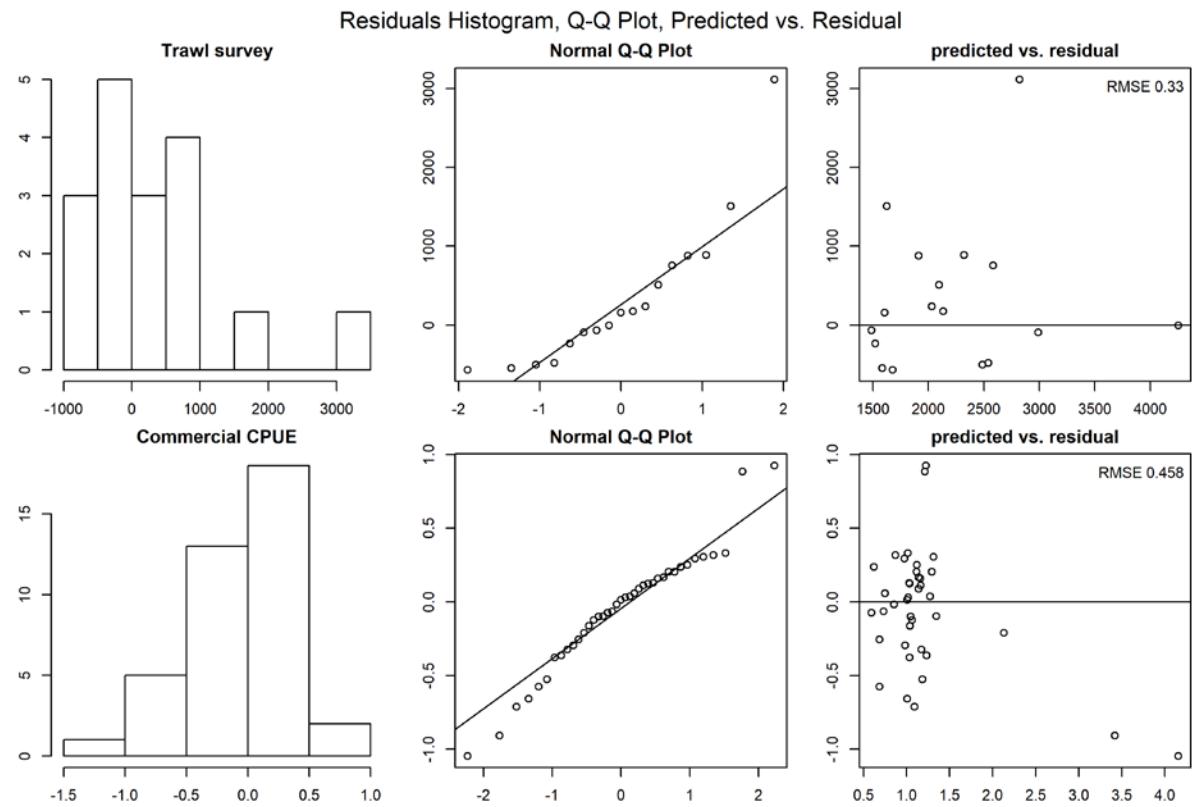


Figure C8-1. QQ Plot of Trawl survey and Commercial CPUE.

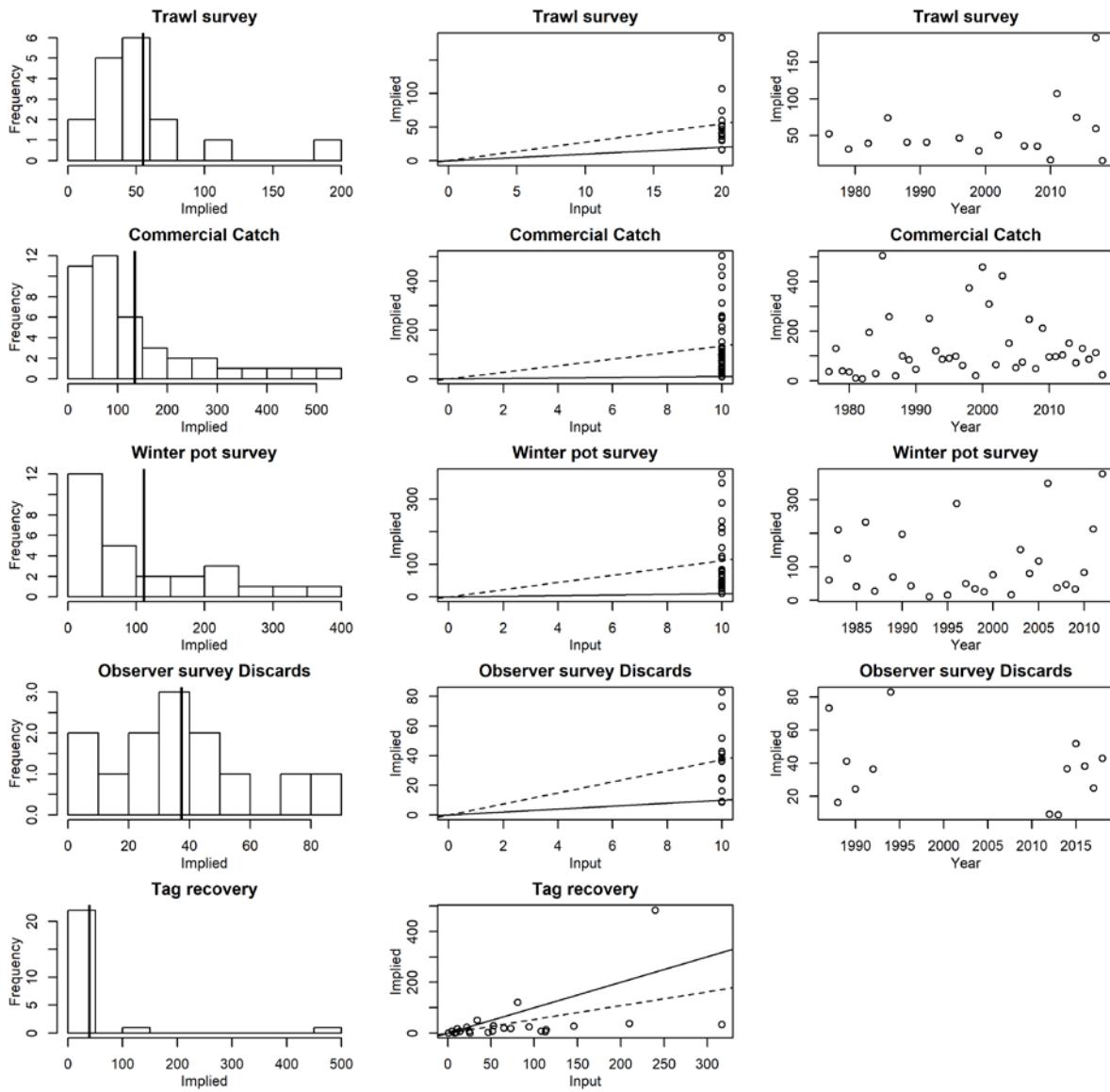


Figure C4-2: Implied effective samples. Figures in the first column show implied effective sample size (x-axis) vs. frequency (y-axis).

Vertical solid line is the mean implied effective sample size.

The second column show input sample size (x-axis) vs. implied effective sample size (y-axis). Dashed line indicates linear regression slope, and solid line is 1:1 line. The third column show year (x-axis) vs. implied effective sample size (y-axis).

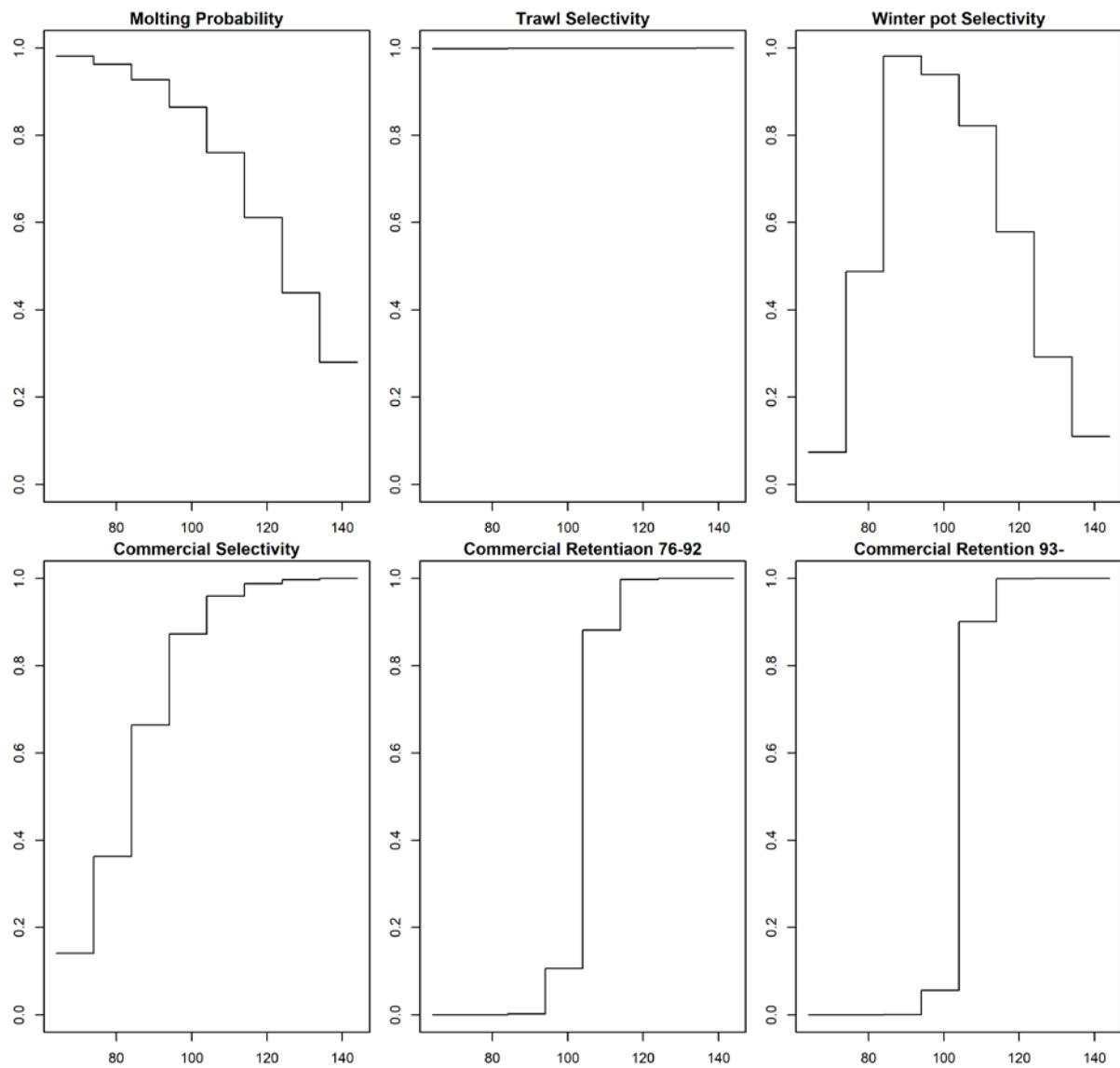


Figure C4-3. Molting probability and trawl/pot selectivity. X-axis is carapace length.

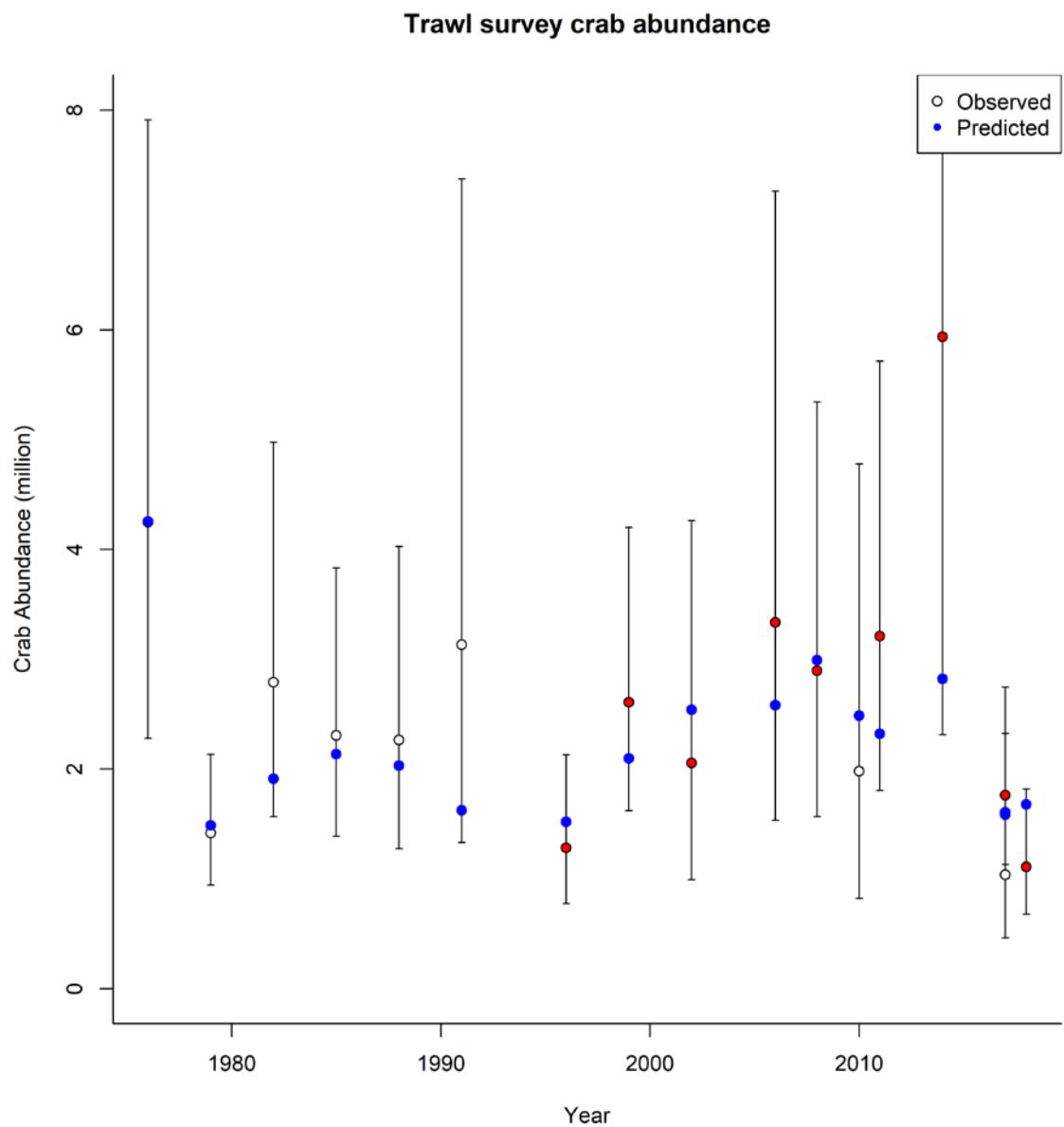


Figure C4-4. Estimated trawl survey male abundance (crab  $\geq 64$  mm CL). Observed: White: NOAA Trawl Survey, Red: ADG&G Trawl Survey

### Modeled crab abundance Feb 01

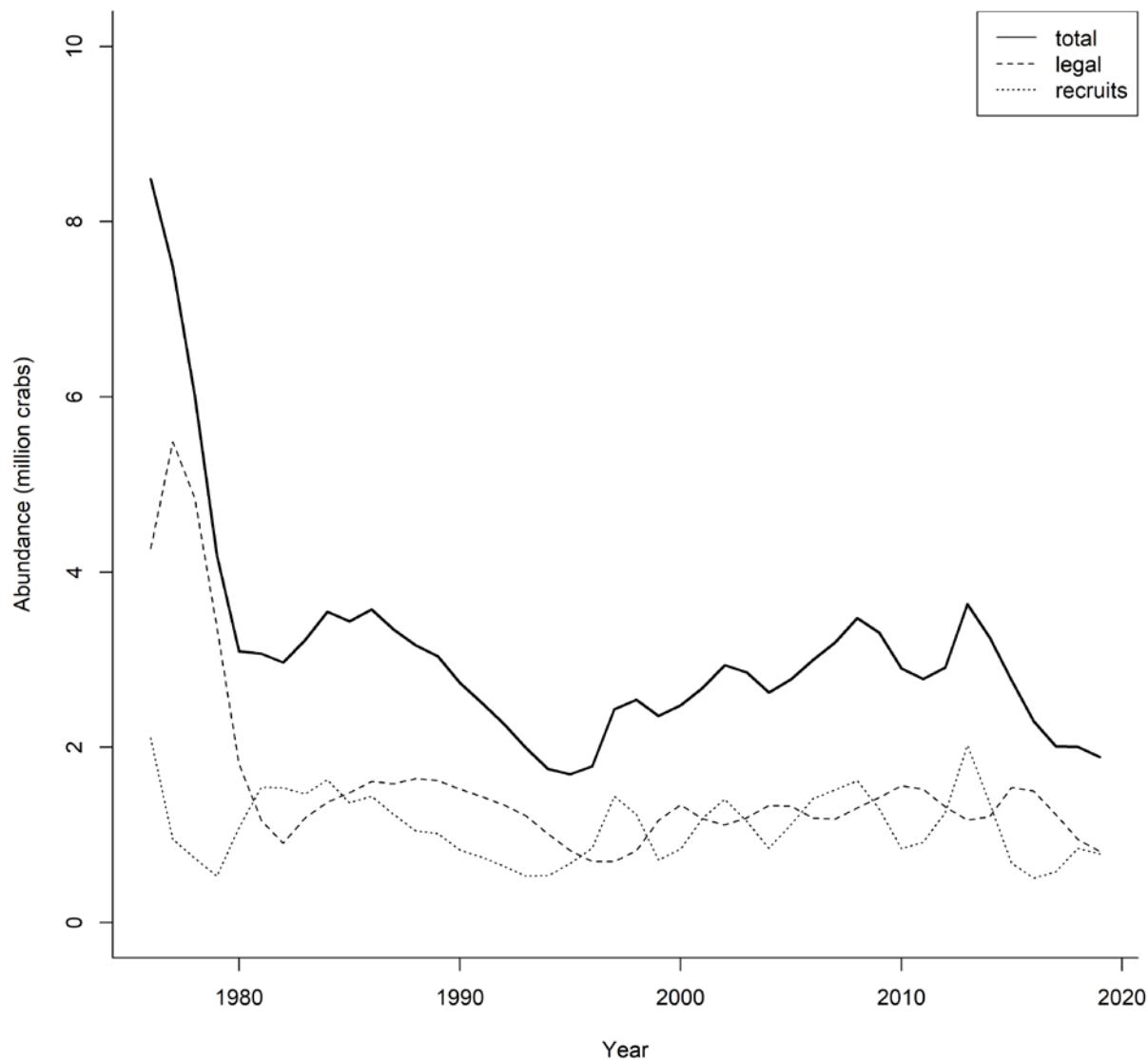


Figure C4-5. Estimated abundance of legal males from 1976-2015.

**MMB Feb 01**

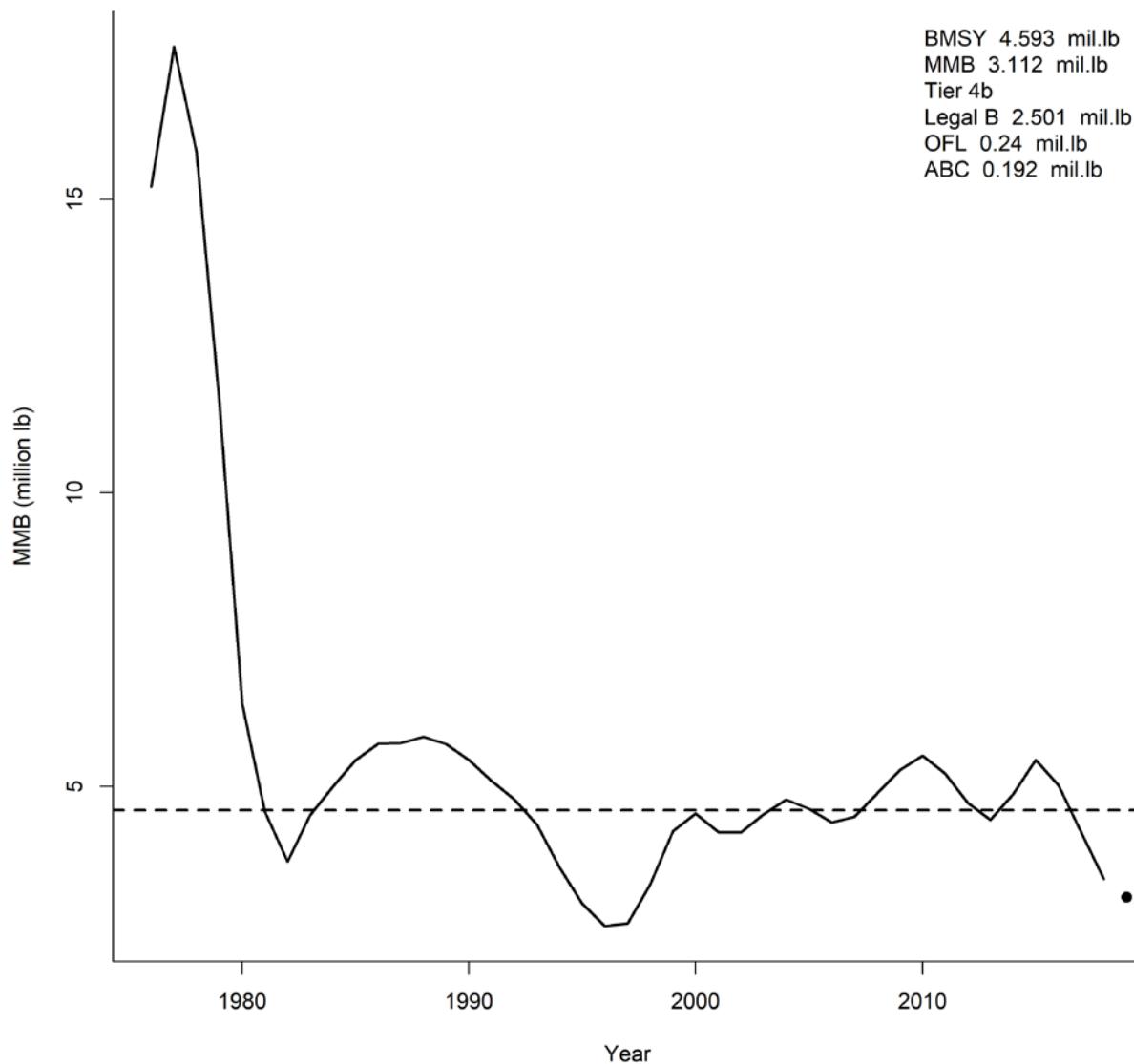


Figure C4-6. Estimated abundance of Mature Male Biomass from 1976-2019. Dash line shows Bmsy (Average MMB of 1980-2019).

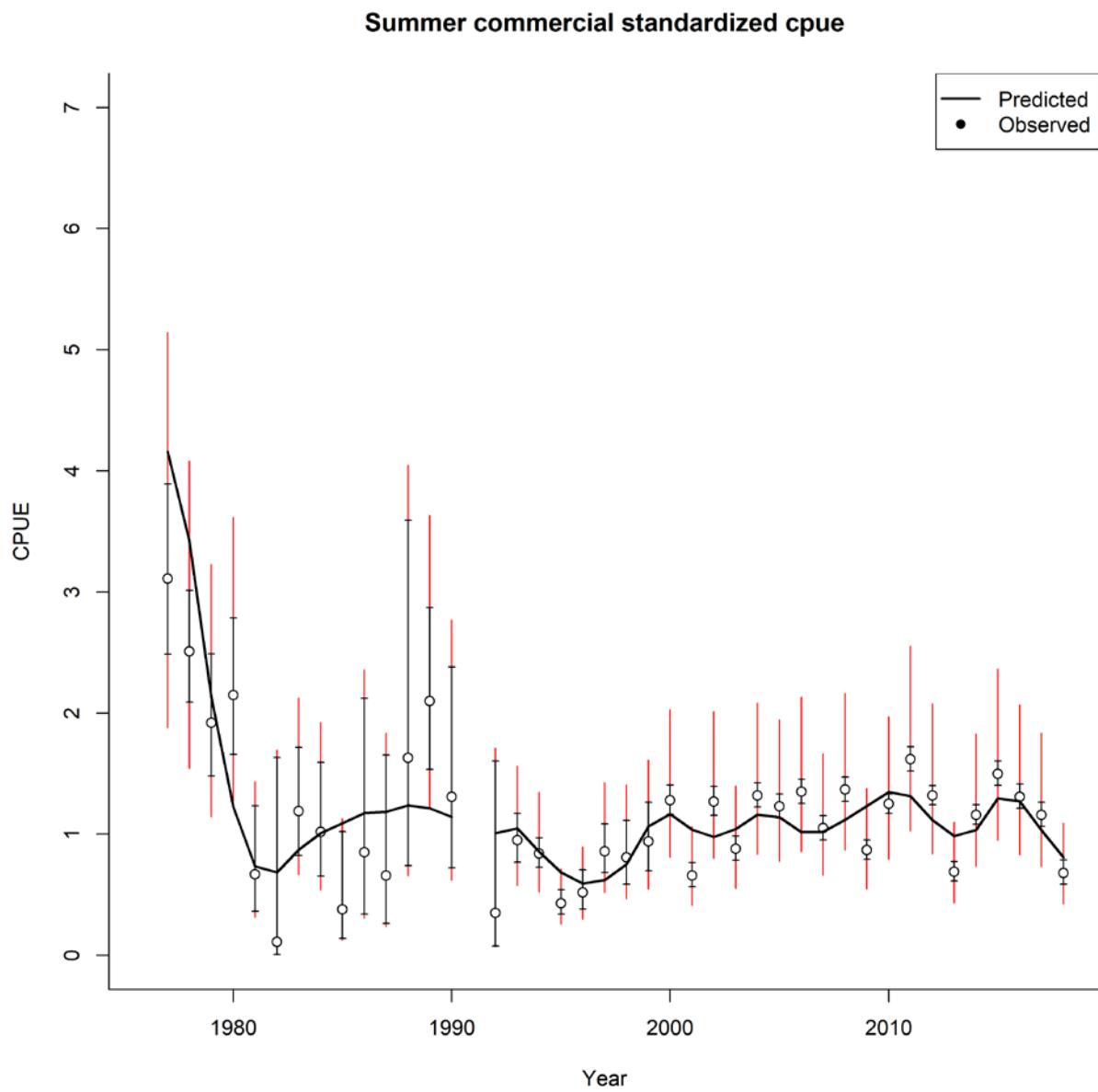


Figure C4-7. Summer commercial standardized cpue 1977-2018.

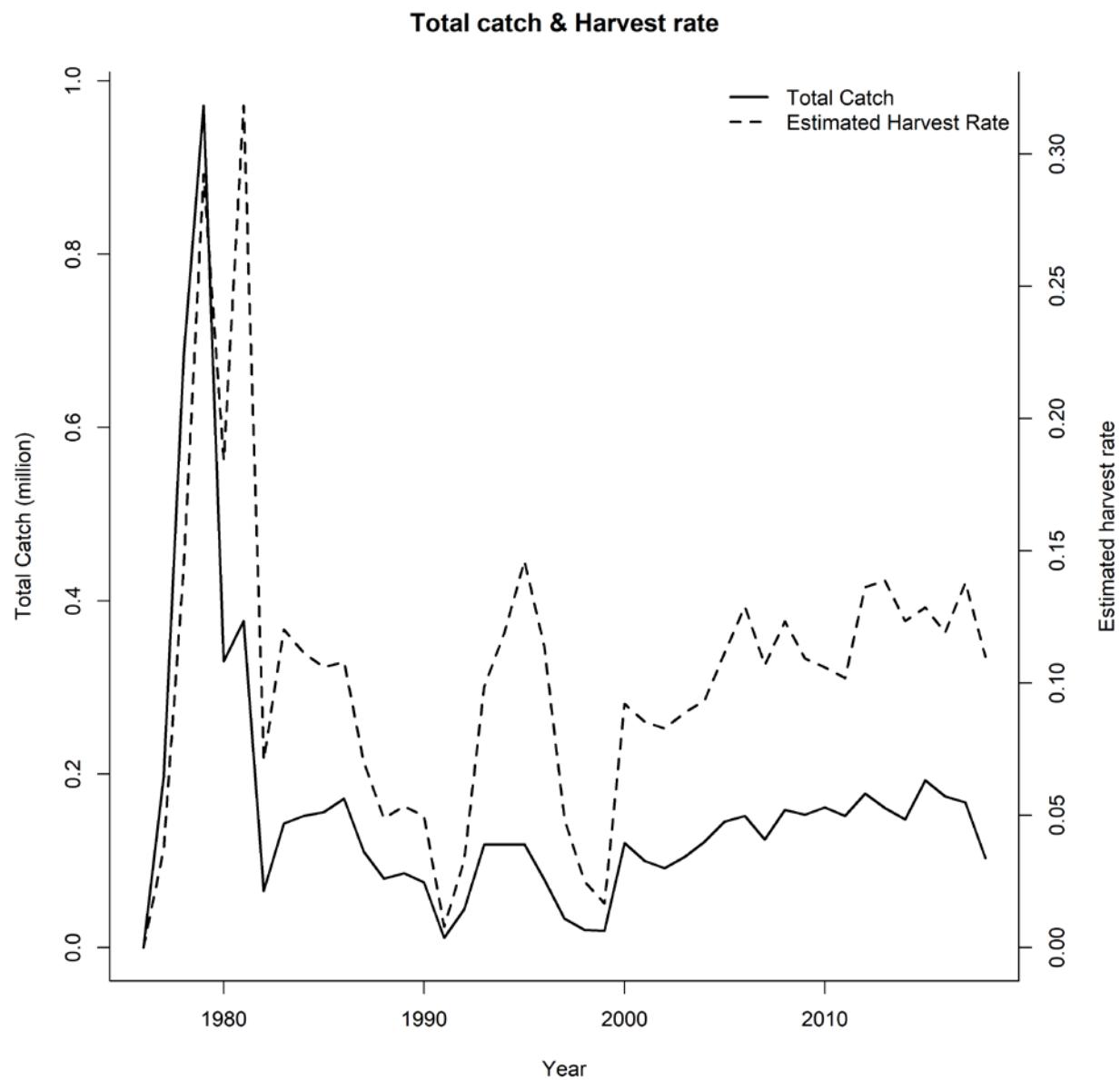


Figure C4-8. Total catch and estimated harvest rate 1976-2018.

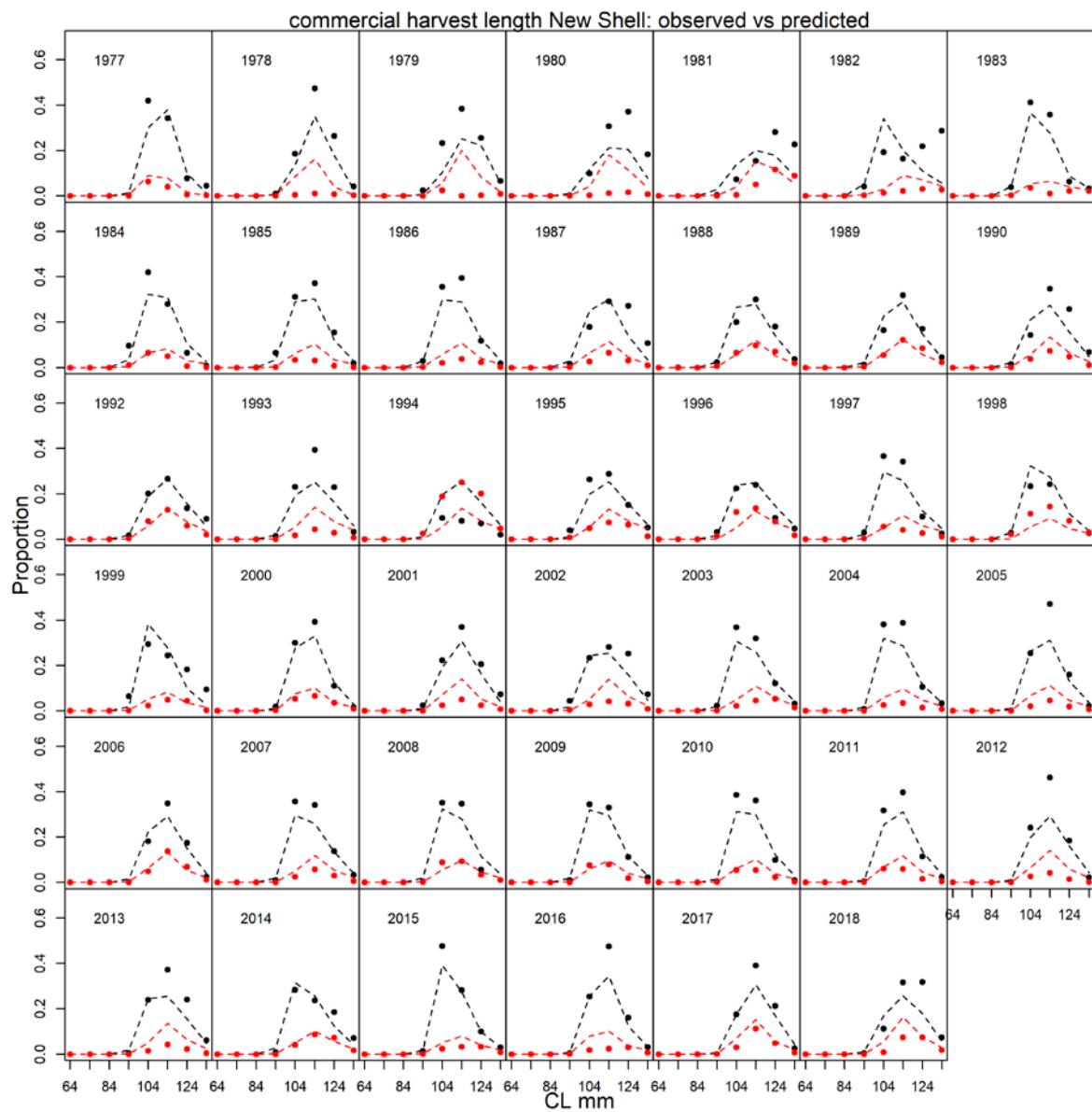


Figure C4-9. Predicted (dashed line) vs. observed (dots) length class proportions for commercial catch. Black: New Shell, Red: Old Shell

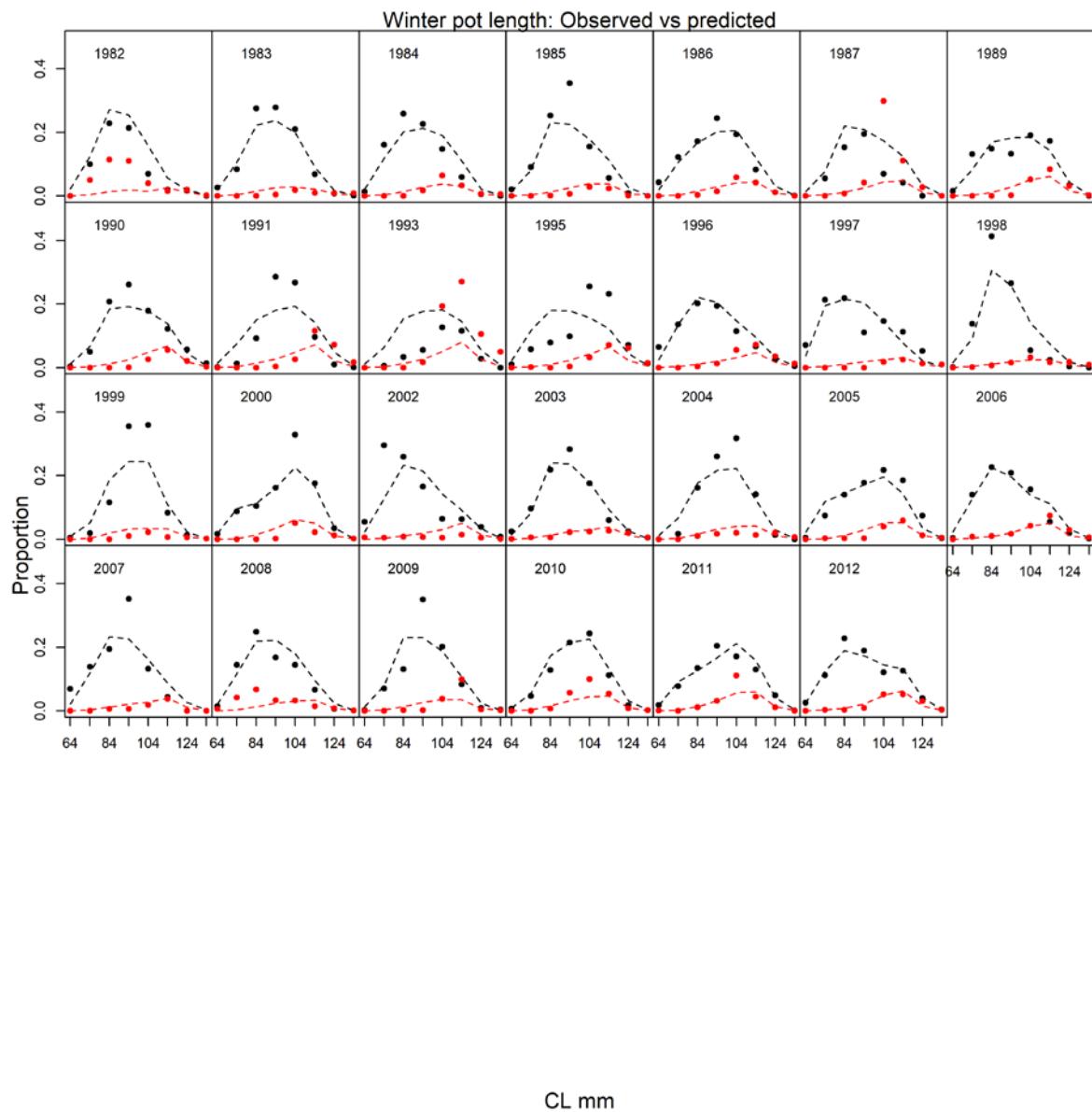


Figure C4-10. Predicted (dashed line) vs. observed (black dots) length class proportions for the winter and spring pot survey.

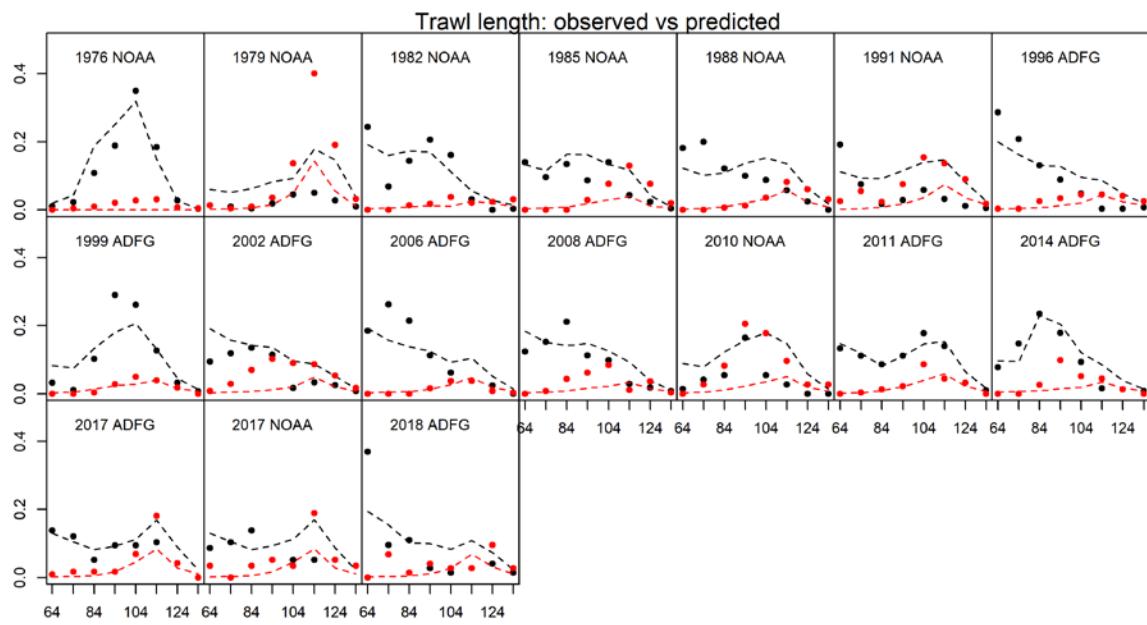


Figure C4-11. Predicted (dashed) vs. observed (dots) length class proportions for trawl survey

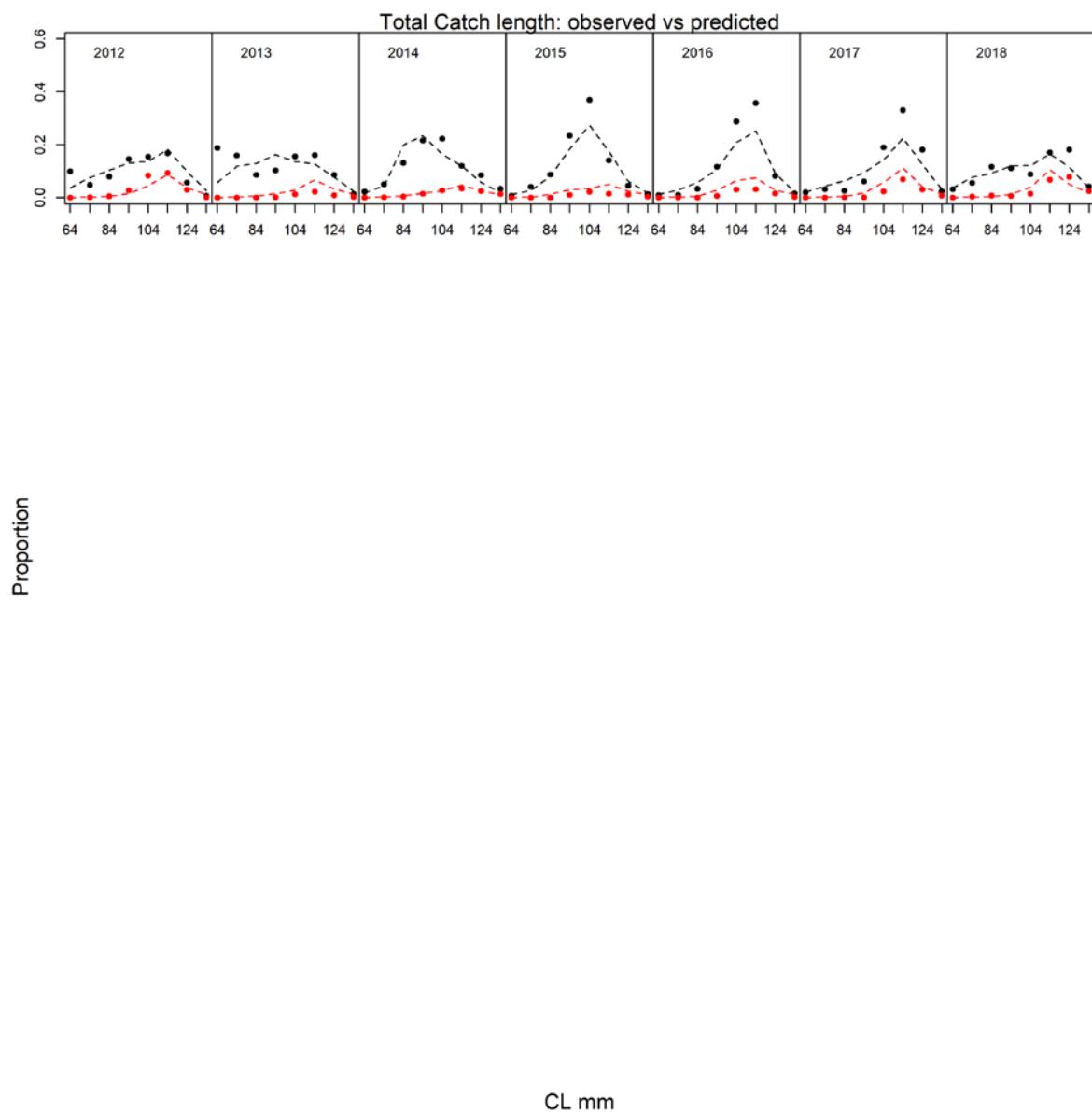
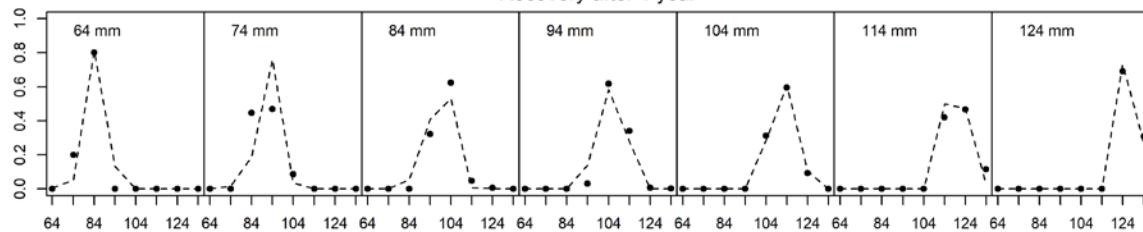
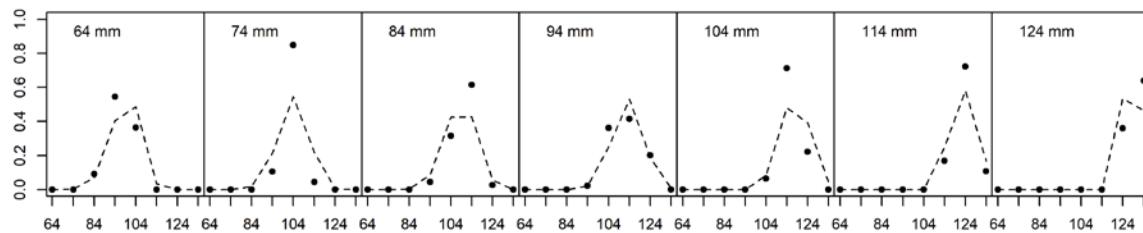


Figure C4-12. Predicted (dashed) vs. observed (dots) length class proportions for the observer survey.

Tag recovery data observed vs predicted  
Recovery after 1 year



Recovery after 2 years



Recovery after 3 years

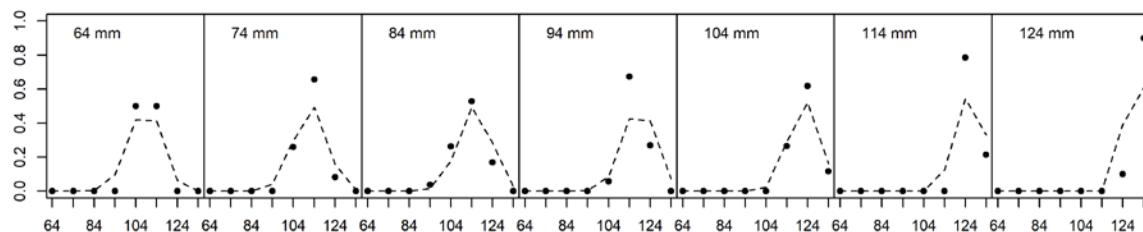


Figure C4-13. Predicted vs. observed length class proportions for tag recovery data.

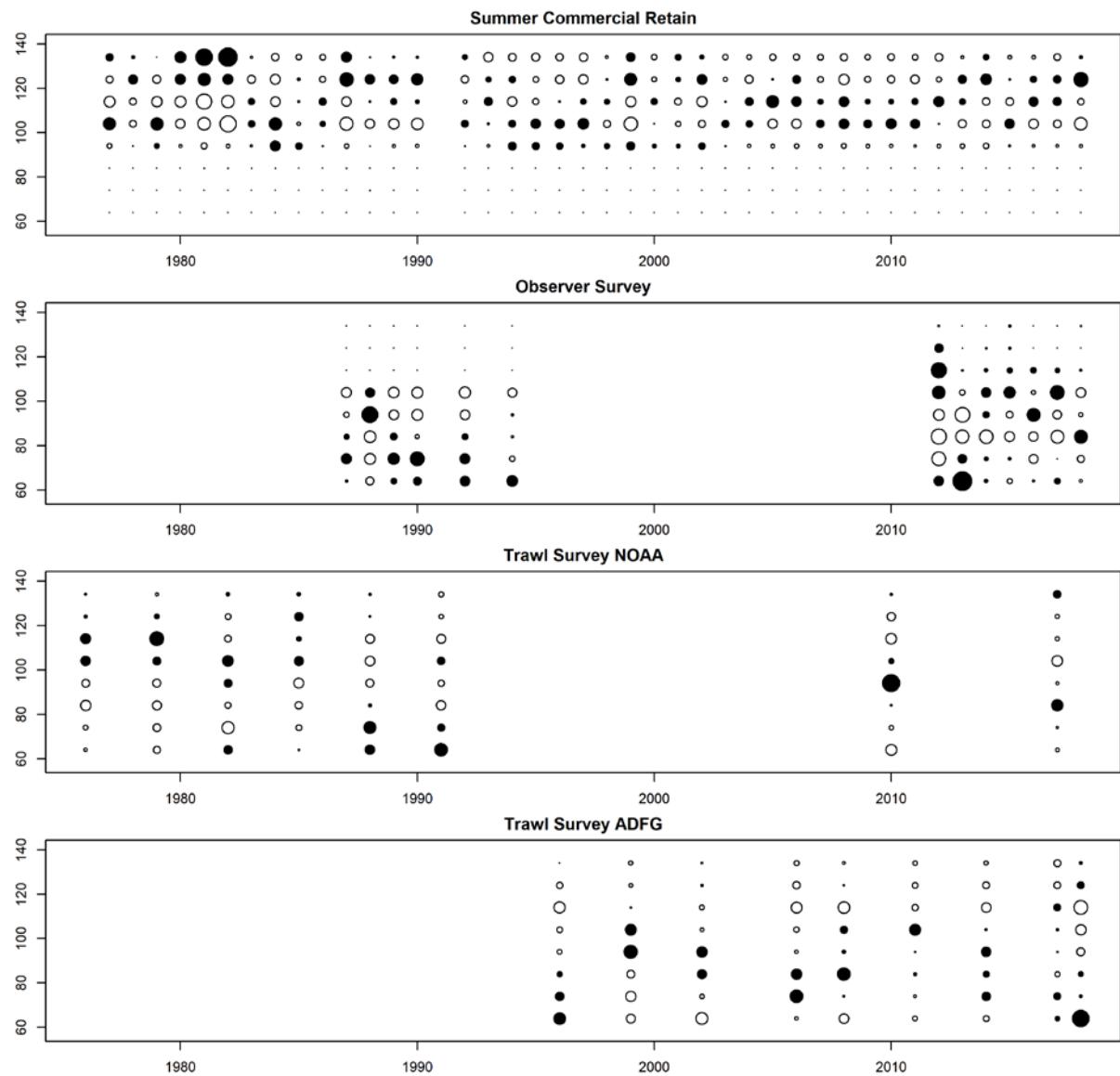


Figure C4-14. Bubble plots of predicted and observed length proportions.  
 Black circle indicates model estimates lower than observed, white circle indicates model estimates higher than observed. Size of circle indicates degree of deviance (larger circle = larger deviance).

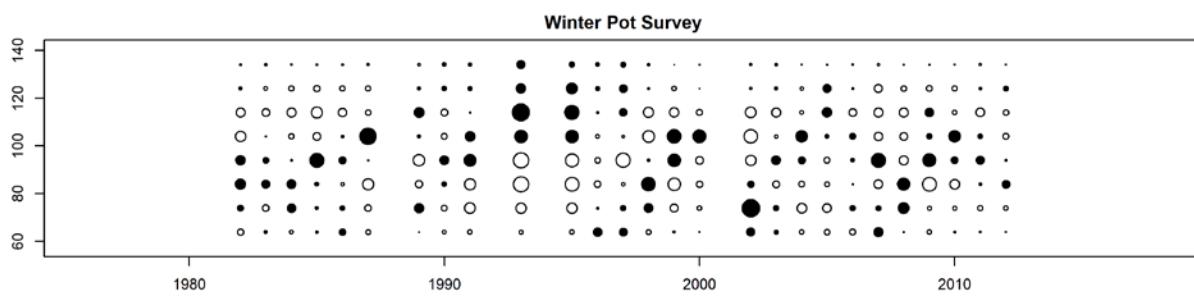


Figure C4-15. Bubble plots of predicted and observed length proportions.  
 Black circle indicates model estimates lower than observed, white circle indicates model estimates higher than observed. Size of circle indicates degree of deviance (larger circle = larger deviance).

Table C4. Summary of parameter estimates for a length-based stock synthesis population model of Norton Sound red king crab.

name	Estimate	std.dev
log_q1	-7.010	0.173
log_q2	-6.834	0.117
log_N <sub>76</sub>	9.046	0.132
R <sub>0</sub>	6.443	0.081
log_R <sub>76</sub>	0.028	0.418
log_R <sub>77</sub>	-0.530	0.370
log_R <sub>78</sub>	-0.719	0.353
log_R <sub>79</sub>	0.369	0.317
log_R <sub>80</sub>	0.516	0.282
log_R <sub>81</sub>	0.414	0.263
log_R <sub>82</sub>	0.378	0.314
log_R <sub>83</sub>	0.562	0.275
log_R <sub>84</sub>	0.171	0.291
log_R <sub>85</sub>	0.460	0.279
log_R <sub>86</sub>	0.074	0.287
log_R <sub>87</sub>	0.024	0.247
log_R <sub>88</sub>	0.026	0.260
log_R <sub>89</sub>	-0.316	0.279
log_R <sub>90</sub>	-0.286	0.256
log_R <sub>91</sub>	-0.525	0.284
log_R <sub>92</sub>	-0.705	0.306
log_R <sub>93</sub>	-0.574	0.289
log_R <sub>94</sub>	-0.287	0.256
log_R <sub>95</sub>	-0.068	0.224
log_R <sub>96</sub>	0.569	0.217
log_R <sub>97</sub>	-0.009	0.291
log_R <sub>98</sub>	-0.631	0.319
log_R <sub>99</sub>	0.000	0.308
log_R <sub>00</sub>	0.296	0.262
log_R <sub>01</sub>	0.396	0.238
log_R <sub>02</sub>	-0.006	0.312
log_R <sub>03</sub>	-0.282	0.329
log_R <sub>04</sub>	0.285	0.239
log_R <sub>05</sub>	0.416	0.219
log_R <sub>06</sub>	0.443	0.241
log_R <sub>07</sub>	0.522	0.227
log_R <sub>08</sub>	0.096	0.283

name	Estimate	std.dev
log_R <sub>09</sub>	-0.384	0.289
log_R <sub>10</sub>	0.046	0.245
log_R <sub>11</sub>	0.350	0.274
log_R <sub>12</sub>	0.888	0.190
log_R <sub>13</sub>	-0.181	0.294
log_R <sub>14</sub>	-0.651	0.312
log_R <sub>15</sub>	-0.719	0.277
log_R <sub>16</sub>	-0.448	0.239
log_R <sub>17</sub>	-0.009	0.282
a <sub>1</sub>	1.520	4.554
a <sub>2</sub>	2.300	4.242
a <sub>3</sub>	3.767	4.048
a <sub>4</sub>	4.057	4.034
a <sub>5</sub>	4.299	4.024
a <sub>6</sub>	3.536	4.054
a <sub>7</sub>	2.104	4.319
r <sub>1</sub>	10.000	0.810
r <sub>2</sub>	9.713	0.830
log_a	-2.661	0.089
log_b	4.828	0.015
log_φ <sub>st1</sub>	-5.000	0.085
log_φ <sub>wa</sub>	-2.116	0.317
log_φ <sub>wb</sub>	4.797	0.029
Sw1	0.074	0.036
Sw2	0.488	0.122
log_φ <sub>t</sub>	-2.084	0.057
log_ar	-0.884	0.178
log_br	4.641	0.011
log_ar	-0.688	0.216
log_br	4.646	0.012
w <sup>2</sup> <sub>t</sub>	0.051	0.016
q	0.753	0.130
σ	3.856	0.211
β <sub>1</sub>	12.404	0.703
β <sub>2</sub>	7.672	0.173
ms78	3.204	0.265

