**Table S2:** Maximum likelihood estimates of the overall FSC replacement rates in germaria with FSC clones of the indicated genotypes

|  |  |  |  |
| --- | --- | --- | --- |
|  | MLE (/week) | Std. Error (/week) | 95% CI (/week) |
| Wildtype | 0.146 | 0.019 | [0.116,0.182] |
| lgl(1) | 0.245 | 0.034 | [0.193,0.308] |
| Dlg(m52) | 0.280 | 0.035 | [0.224,0.350] |
| Scrib(1) | 0.024 | 0.025 | [0.004,0.060] |
| Scrib(2) | 0.156 | 0.021 | [0.123,0.196] |
| Baz | 0.148 | 0.019 | [0.119,0.179] |