Table 1.Parameters of the model describing the probability of being eaten as a function of predictors. Shown are parameters from the fixed part of the Model 1.

| Term | Estimate | SE | z-statistic | p.value |
| --- | --- | --- | --- | --- |
|  | -1.208 | 0.0924 | -13.08 | <0.0001 |
|  | 0.370 | 0.0547 | 6.76 | <0.0001 |
|  | -0.138 | 0.0631 | -2.19 | 0.0289 |
|  | -0.105 | 0.0799 | -1.32 | 0.1870 |
|  | -0.319 | 0.0852 | -3.75 | 0.0002 |
|  | -0.182 | 0.0373 | -4.88 | <0.0001 |
|  | 0.673 | 0.0942 | 7.15 | <0.0001 |

Model terms: *PMt* - a probability to be identified as Mt; *Nconsp* - a number of conspecifics for given mussel in the experimental unit; *PropT* - a proportion of mussels of T-morphotype; *Ntotal* - a number of mussels in the experimental unit in the end of exposition; *L* - mussel size; *Baster* - sea stars biomass in the end of the exposition.