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<th>strength</th>
<th>other phenotypes</th>
<th>inheritance†</th>
<th>linkage</th>
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</table>

*Columns are the total number of mutations isolated, the number isolated in the DA837 growth selection, and the number isolated in the L1 arrest escape screen.

†AD = autosomal dominant, AR = autosomal recessive, XR = X-linked recessive.

‡Since other existing eat-18 mutations suppress eat-5 but do not have an Unc phenotype, it is likely that this phenotype is caused by a second mutation irrelevant to the Sef phenotype.