A computational analysis of the dynamic roles of talin, Dok1, and PIPKI for integrin activation - Supplementary Table

Florian Geier¹,², Georgios Fengos¹, Dagmar Iber¹,*

¹ ETH Zürich, Department of Biosystems Science and Engineering (D-BSSE), Mattenstrasse 26, 4058 Basel, Switzerland; ² new address: Biozentrum, Klingelbergstrasse 70, 4056 Basel, Switzerland

* E-mail: Corresponding dagmar.iber@bsse.ethz.ch
<table>
<thead>
<tr>
<th>Parameter</th>
<th>Group 1</th>
<th>Group 2</th>
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<tbody>
<tr>
<td>k1on</td>
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<tr>
<td>k1off</td>
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<td>k2on</td>
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<td>k2off</td>
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<td>k3aon</td>
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<td>k3aoff</td>
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<td>0.014</td>
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<td>k3bon</td>
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<tr>
<td>k3boff</td>
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<td>k6on</td>
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<td>SRC&lt;sub&gt;tot&lt;/sub&gt;</td>
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</table>

**Table S1.** Mean parameter values of group 1 (high TAL:INT, Figure 3 main text) and group 2 (high DOK:INT, Figure 3 main text). On-rates are related to the KD value mentioned in the main text as $kon = koff/KD$. 