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Supplemental Information

Structural and Functional Analysis of a Talin

Triple-Domain Module Suggests an Alternative

Talin Autoinhibitory Configuration

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SUPPLEMENTAL INFORMATION

Figure S1, related to Figure 2.

A. Cross-linking of purified His-R7R8R9 protein and mutants. Cross-linking reactions were terminated at 20 min or 40 min before subject to SDS-PAGE analysis. **B.** Structural comparison of R7R8R9 and R7R8R9-3Y. The R9 domain of R7R8R9-3Y is superimposed on that of the wild-type R7R8R9. The R9:R8' interfaces in the two structures are highly similar (*left*), despite a ~30° rotation between the two R8' domains (*right*, orange for WT and lime for R7R8R9-3Y). **C.** Electron density at the R9:R8' interface in a steric view. *Upper*: electron density of R7R8R9. *Lower*: electron density of R7R8R9-3Y.

Figure S2, related to Figures 3 and 4.

A. Integrin activity analysis of talin-V1540Y. *Left*: relative integrin activity of wild-type talin and talin-V1540Y in histogram. NS: not significant compared with talin WT. *Right*: expression levels of the transfected constructs were examined by anti-GFP Western. **B. Expression levels of talin in CHO-A5 cells for integrin activity analyses.** The corresponding figures are indicated.

Figure S3, related to Figure 4.

Crystal contact prevents the alternative autoinhibitory configuration in the crystal structure of the F2F3:R9 complex. The F2 domain of a symmetrically related F2F3 molecule (F2', in lime) makes contact with several residues in the α 5 helix of the R9 domain. This crystal contact of the F2 and R9 domains likely prevents the "mode B" autoinhibitory configuration.

Figure S4, related to Figure 3.

Effect of $PI(4,5)P_2$ headgroup on the F2F3:R9 interaction. (A) *In-vitro* pull-down of His-F2F3 by GST-R9 in the presence of IP₃ at various concentrations. His-F2F3 input and GST/GST-R9 pull-down were shown by Coomassie staining. Bound His-F2F3 was shown by anti-His Western. (B) *In-vitro* pull-down of His-F2F3-KKR by GST-R9 in the presence of IP₃ at various concentrations. Input proteins and bound His-F2F3 were shown by Coomassie staining.



S1



GFP GFP Tain Tain Tain Tain FT



