Supplemental Materials Molecular Biology of the Cell

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Supplemental Figure Legends

Supplemental Figure 1. α *E-catenin binds cooperatively to filamentous actin*

(A) The α E-catenin ABD is a monomer in solution. The molecular mass of α E-catenin ABD was determined by small-angle x-ray scattering (SAXS). The calculated molecular weight of the construct is 26.5 kDa. A small amount of aggregation was apparent at the higher concentrations, but analysis of the data either assuming some aggregation or assuming no aggregation clearly showed that the ABD is a monomer at all concentrations tested.

(B) Representative images of GFP α E-catenin ABD binding to filamentous actin *in vitro*. Localization of increasing concentrations of α E-catenin ABD (25 nM to 4 μ M GFP) bound to phalloidin-stabilized F-actin (20% Cy3 labeled, also see Methods).

(C-D) Bulk actin filament pelleting assays with α E-catenin ABD **(C)** and full-length α E-catenin dimer **(D)**. Bound α E-catenin (μ M/ μ M actin) was plotted against free α E-catenin (μ M), and data were fit to a Hill equation (red line; K_d and Hill coefficient listed in red) or hyperbolic function (green line; K_d listed in green). Data shown represent one example of ≥3 experiments for α E-catenin ABD and dimer. The maximal number of α -catenin molecules bound per actin monomer (Bmax) varied between different experiments and values between 0.6-1.4 were obtained for the ABD (curve fit assuming cooperativity). For full-length α E-catenin, values ranged from 0.6-1.8 (curve fit assuming no cooperativity).

Supplemental Figure 2. α *E*-catenin ABD reduces barbed end polymerization

(A) Average barbed end polymerization rate in 2 μ M Mg-ATP-actin (10% Cy3 labeled) was measured in the presence of increasing concentrations of GFP α E-catenin ABD. Error bars indicate SD (n ≥ 30 actin filaments from ≥ 2 experiments).

(B) Kymographs of Cy3-labeled actin filaments growing in the absence or presence of 2 μ M GFP α E-catenin ABD. Arrowheads mark pauses in barbed end elongation (frequency was 0.087 min⁻¹, average lifetime was 62 ± 36 seconds, n=32 pauses).

(C) Montage of 2 μ M Mg-ATP-Actin (10% Cy3 labeled) polymerizing and bundling in the presence of 0.25 μ M GFP α E-catenin ABD. We observed a 2.9-fold increase in GFP α E-catenin ABD fluorescence per actin filament at sites of bundling. Scale bar is 10 μ m. (D) Kymograph of actin filaments bundling in the presence of 2 μ M Mg-ATP-Actin (10% Cy3 labeled) and 0.25 μ M GFP α E-catenin ABD. Note the change in BE growth during filament bundling. Vertical scale is 1 minute. Horizontal scale bar 5 μ m.

Movie Legends

Movie 1

Dynamic localization of 2 nM GFP α E-catenin ABD (green) binding to phalloidin-stabilized F-actin (20% Cy3-labeled, red). Images were acquired at 50 ms intervals. Movie plays at 20 frames per second. Scale bar 1 μ m.

Movie 2

Dynamic localization of 2 nM GFP α E-catenin ABD (green) binding to phalloidin-stabilized F-actin (20% Cy3-labeled, red) in the presence of 0.5 μ M dark α E-catenin ABD. Images were acquired every 50 ms during data acquisition. Movie plays at 20 frames per second. Scale bar 1 μ m.

Movie 3

Inhibition of Arp2/3 complex-mediated filament branching in the presence of GFP α E-catenin ABD. Actin filaments were polymerized and nucleated in the presence of 1 μ M Mg-ATP-actin (10% Cy3), 50 nM Arp2/3, 100 nM SCAR^{VCA}, and 0-1 μ M GFP α E-catenin ABD. Images were acquired every 10 sec during data acquisition. Movie plays at 15 frames per second. Scale bar 10 μ m.

Movie 4

Inhibition of Arp2/3 complex-mediated filament branching in the presence of full length GFP α E-catenin. Actin filaments were polymerized and nucleated in the presence of 1 μ M Mg-ATP-actin (10% Cy3 labeled), 50 nM Arp2/3 and 100 nM SCAR^{VCA} in the absence (right panel) or presence (left panel) of 2 μ M full length GFP α E-catenin. Images were acquired every 10 sec during data acquisition. Movie plays at 15 frames per second. Scale bar 10 μ m.

Movie 5

Inhibition of cofilin severing in the presence of α E-catenin ABD. ADP-actin filaments (20% Cy3, red) were immobilized and disassembled by 75 nM ybbr-Atto488-hCofilin (green) in the absence (right panel) or presence (left panel) of 2 μ M α E-catenin ABD. Images were acquired every 15 sec during data acquisition. Movie plays at 15 frames per second. Scale bar 5 μ m.

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Protein Concentration (mg/ml)	Rg(Å)		MW(kDa)	
	Fit assuming no aggregation	Fit assuming aggregation	Fit assuming no aggregation	Fit assuming aggregation
5.0	25.2	27.4	25.9	27.6
2.5	26.0	27.1	28.0	28.9
1.0	27.3		25.2	
0.5	28.2		21.2	



25nM

50nM

0.1µM

0.25µM

0.5µM

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2µM

4µM







10 µm