Figure S1. Quantification of the Western blot data in figure 4C. The area density of each band was measured with a UVP gel document system. The data were normalized with β-tubulin and presented as ratio relative to the area density of β-tubulin. Bars represent means ± SEM. Bars with the same letters are not significantly different from each other (p>0.05).
Figure S2. Quantification of the Western blot data in figure 5C. The area density of each band was measured with a UVP gel document system. The data were normalized with β-tubulin and presented as ratio relative to the area density of β-tubulin. Bars represent means ± SEM. Bars with the same letters are not significantly different from each other (p>0.05).
Figure S3. Quantification of the Western blot data in figure 6A. The area density of each band was measured with a UVP gel document system. The data were normalized with β-tubulin and presented as ratio relative to the area density of β-tubulin. Bars represent means ± SEM. Bars with the same letters are not significantly different from each other (p>0.05).
Figure S4. Quantification of the Western blot data in figure 6B. The area density of each band was measured with a UVP gel document system. The data were normalized with β-tubulin and presented as ratio relative to the area density of β-tubulin. Bars represent means ± SEM. Bars with the same letters are not significantly different from each other (p>0.05).