

## Microtubule Dynamic Instability Does Not Result from Stabilization of Microtubules by Tubulin-GDP-P<sub>i</sub> Subunits<sup>†</sup>

Michael Caplow\* and John Shanks

Department of Biochemistry, University of North Carolina, Chapel Hill, North Carolina 27599-7260

Received October 21, 1997; Revised Manuscript Received June 26, 1998

**ABSTRACT:** The proposal that microtubule dynamic instability results from stabilization of microtubule ends by tubulin-GDP-P<sub>i</sub> subunits (where P<sub>i</sub> is inorganic phosphate) [Melki et al. (1996) *Biochemistry* 35, 12038] was based on studies of GTP hydrolysis and microtubule assembly that showed that tubulin-GDP-P<sub>i</sub> subunits can transiently accumulate at microtubule ends. There is no direct evidence that GDP-P<sub>i</sub>-subunits can stabilize microtubules under conditions where dynamic instability is observed and this has been inferred from the observation that tubulin-GDP-BeF<sub>n</sub> subunits stabilize microtubules. To test if tubulin-GDP-P<sub>i</sub> stabilizes microtubules we sought evidence for a synergism between the effect of P<sub>i</sub> and BeF<sub>n</sub>. We found, however, that P<sub>i</sub> antagonizes the effect of BeF<sub>n</sub> by displacing it from tubulin subunits. The alternate mechanism in which P<sub>i</sub> inhibits BeF<sub>n</sub> stabilization of microtubules by displacing fluoride from beryllium was ruled out from the <sup>9</sup>Be and <sup>19</sup>F NMR spectra in the presence and absence of P<sub>i</sub>. Further evidence that tubulin-GDP-BeF<sub>n</sub> is not an analogue of tubulin-GDP-P<sub>i</sub> and that tubulin-GDP-P<sub>i</sub> is not responsible for maintaining the growth phase in microtubules manifesting dynamic instability was provided by our observation that P<sub>i</sub> did not decrease the disassembly rate under conditions where tubulin-GDP-P<sub>i</sub> subunits are expected to have formed. Results showing that BeF<sub>n</sub> binds randomly to subunits in microtubules provided evidence that P<sub>i</sub> dissociation from the tubulin-GDP-P<sub>i</sub> intermediate formed during GTP hydrolysis occurs randomly rather than processively starting at the growing microtubule tip.

The two states that allow allosteric regulation of microtubule dynamics by guanine nucleotide are not yet defined. Tubulin-GDP subunits that dissociate rapidly constitute one state. Tubulin-GTP or tubulin-GDP-P<sub>i</sub><sup>1</sup> subunits presumably constitute the second state. Understanding the role of GTP in microtubules is important because as with other G-proteins, nucleotide hydrolysis and exchange are likely to be involved in regulation of function.

The presence of an extensive tubulin-GTP cap at microtubule ends was first indicated from the observation of a 7 min discrepancy between assembly of 40 μM tubulin into microtubules and formation of 40 μM P<sub>i</sub> from GTP (1). The stabilizing effect of tubulin-GTP subunits was inferred from results with nonhydrolyzable GTP analogues, which form microtubules that are relatively stable to depolymerization by calcium and cold (2–4). However, evidence against a GTP cap came from a subsequent study showing coincidence between the GTPase and assembly rates (5); also, microtubules isolated 10–15 s after assembly did not contain measurable amounts of GTP (6). The GTP cap mechanism was, therefore, revised so that the cap is restricted to a monolayer of tubulin-GTP subunits (7–12).

An alternate model in which growing microtubules are stabilized by a cap of tubulin-GDP-P<sub>i</sub> subunits was proposed from the finding that microtubules isolated on a filter about 10 s after initiating assembly contain P<sub>i</sub> from GTP hydrolysis (11). Although these results were not confirmed (6), subsequent study using a continuous spectrophotometric assay of the release of P<sub>i</sub> from assembling microtubules (13) showed that P<sub>i</sub> would be lost from microtubules too rapidly to be detected with a filter assay in which the dead time was 10–15 s (6). The half-time for P<sub>i</sub> release was about 6 s, whereas GTP in microtubules was largely hydrolyzed in about 3 s. Since P<sub>i</sub> release is slower than GTP hydrolysis, tubulin-GDP-P<sub>i</sub> subunits accumulate in growing microtubules.

It has not been possible to determine whether tubulin-GDP-P<sub>i</sub> subunits can stabilize microtubule ends; this was inferred (13) from results with the fluoride adducts of beryllium [BeF<sub>n</sub>, in which *n* is presumably 3 (14)] and aluminum. These P<sub>i</sub> analogues bind stoichiometrically and competitively with P<sub>i</sub> (15) to tubulin-GDP subunits in microtubules and reduce the subunit dissociation rate (15, 16). The assumption that tubulin-GDP-BeF<sub>n</sub> resembles tubulin-GDP-P<sub>i</sub> is, however, unproven. In fact, X-ray analysis of the complex between myosin and MgADP-BeF<sub>x</sub> showed that this resembles the myosin–ATP species before hydrolysis (17). A similar conclusion was derived for a G-protein in which the geometry of the transducin-GDP-AlF<sub>4</sub> complex resembles a pentavalent intermediate in GTP hydrolysis, rather than a transducin-GDP-P<sub>i</sub> complex (18).

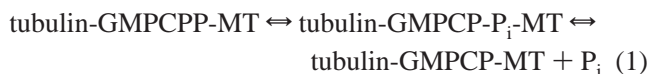
<sup>†</sup> Supported by a grant from the NIH (GM 46773).

\* Corresponding author: phone 919-966-1014; FAX 919-966-2852; E-mail caplow@med.unc.edu.

<sup>1</sup> Abbreviations: BRB buffer, 80 mM sodium 1,4-piperazinediethanesulfonate (Na-Pipes), 1 mM MgCl<sub>2</sub>, and 1 mM [ethylenebis-(oxyethylenitrilo)]tetraacetic acid (EGTA); P<sub>i</sub>, inorganic phosphate; MT, microtubule.

These structural studies suggest that tubulin-GDP-BeF<sub>n</sub> is an analogue of tubulin-GTP rather than tubulin-GDP-P<sub>i</sub>, so that results with BeF<sub>n</sub> may not be valid for estimating the stabilization from a tubulin-GDP-P<sub>i</sub> cap.

There are three additional reasons to question whether growing microtubules are stabilized by a tubulin-GDP-P<sub>i</sub> cap. First, using video microscopy to study the dynamics of individual microtubules we and others (19, 20) failed to observe stabilization of microtubules by high concentrations of P<sub>i</sub>. Second, we have measured a standard free energy change equal to +0.7 kcal for the two-step reaction sequence in which GMPCPP in microtubules hydrolyzes and P<sub>i</sub> is released (21):



On the basis of the 25 mM *K*<sub>d</sub> for dissociation of P<sub>i</sub> from microtubules (15)  $\Delta G$  is +2.18 kcal for P<sub>i</sub> dissociation in the second step in eq 1. The free energy for the first step in eq 1 in which the triphosphate is hydrolyzed is, therefore, only -1.48 kcal ( $\Delta G_{\text{overall}} = +0.7 = -1.48 + 2.18$ ). The small free energy for nucleoside triphosphate hydrolysis requires that the remaining 3.7 kcal of energy from the triphosphate bond (5.18 kcal - 1.48 kcal; 20) be stored in the microtubule lattice. The energy stored in tubulin-GMPCP-P<sub>i</sub> subunits, and presumably also in tubulin-GDP-P<sub>i</sub> subunits, is likely to be manifested as an enhanced dissociation rate so that these subunits would not be stabilizing, as suggested (11, 13, 15, 16). A third reason for questioning whether microtubules are stabilized by a tubulin-GDP-P<sub>i</sub> cap comes from an insightful analysis of the low affinity of P<sub>i</sub> for microtubules (22). It was pointed out that "... if release of P<sub>i</sub> increases the instability of microtubule ends, this would require a correspondingly less favorable dissociation of P<sub>i</sub>." The low affinity of P<sub>i</sub> (15), therefore, controverts stabilization of microtubules by GDP-P<sub>i</sub> subunits.

We have again focused on determining the effect of P<sub>i</sub> on microtubules and have used the P<sub>i</sub> analogue BeF<sub>n</sub> to determine whether failure to observe an effect of P<sub>i</sub> on dynamics (19, 20) resulted from an unmet requirement for multiple contiguous tubulin-GDP-P<sub>i</sub> subunits to generate a stable microtubule end. Also, we tested a mechanism in which P<sub>i</sub> binding is processive, starting with tubulin-GDP subunits that interface with a cap of tubulin-GTP (or tubulin-GDP-P<sub>i</sub>) subunits at the microtubule end. Studies of the kinetics of shortening of BeF<sub>n</sub>-treated microtubules showed that binding of BeF<sub>n</sub> and presumably also P<sub>i</sub> to microtubules is random rather than processive. Most important, we did not observe stabilization of microtubules by P<sub>i</sub> under conditions where P<sub>i</sub> is bound to tubulin-GDP subunits in the microtubule. P<sub>i</sub> was found to reverse the BeF<sub>n</sub> stabilization of microtubules, apparently by displacing BeF<sub>n</sub> from tubulin-GDP-BeF<sub>n</sub> subunits. The opposing effects of BeF<sub>n</sub> and P<sub>i</sub> indicates that tubulin-GDP-BeF<sub>n</sub> is not an analogue of tubulin-GDP-P<sub>i</sub> and we conclude that tubulin-GDP-P<sub>i</sub> subunits are not responsible for stabilizing microtubule ends.

## EXPERIMENTAL PROCEDURES

Beef brain tubulin was purified by two cycles of thermal-induced assembly and disassembly, followed by chromatography on phosphocellulose (23). The protein that did not

bind to the phosphocellulose was concentrated with an Amicon YM10 membrane. Protein that denatured during the several hours required for the concentration step and/or during freezing, storage, and thawing was removed immediately before the tubulin was used by centrifugation this at 13800g (Microfuge) for 10 min at 4 °C. All reactions were in BRB buffer (80 mM P<sub>i</sub>pes, pH 6.8, 1 mM EGTA, and 1 mM MgCl<sub>2</sub>) at 37 °C.

Real-time measurements of microtubule dynamics were analyzed in reactions at 37° using video-enhanced DIC (differential interference contrast) microscopy. Microtubules were assembled at the ends of sea urchin axonemes (provided by E. D. Salmon) that bound to a cover slip in a flow cell (24) with an approximately 2.5  $\mu$ L capacity. This small volume was replaced in 10–20 s by using filter paper to induce rapid flow of 20  $\mu$ L of a displacing solution. Images from a Hamamatsu video camera were recorded on VHS tape and were analyzed with a computer-based system (25).

Tetracoordinated complexes containing beryllium, fluoride, and the OH moiety from water form rapidly in aqueous solutions (26), with the fluoride stoichiometry progressively increasing from 1 to 4 with increasing concentrations of fluoride (9). Thus, BeF<sub>n</sub> was formed by including BeSO<sub>4</sub> and NaF in reaction mixtures. Studies of stabilization of microtubules by this substance were done in three stages. Microtubules were first assembled with 12–14  $\mu$ M tubulin, 3 mM MgCl<sub>2</sub>, and 1 mM GTP for 3–5 min to generate lengths that were sufficient to allow later study of their disassembly kinetics. Next, the microtubules were elongated for 10 min in the same mixture with 100  $\mu$ M BeSO<sub>4</sub> and 25 mM NaF and either 33 mM Na<sub>2</sub>SO<sub>4</sub>, 50 mM P<sub>i</sub> or 50 mM ethylphosphonate. The ionic strength is equal under these three conditions and from the dissociation constant for reaction of Mg with SO<sub>4</sub> (27) and with the phosphate dianion (28) the free Mg concentration is calculated to be 0.55 mM with SO<sub>4</sub> and 0.38 mM with P<sub>i</sub>. During the 10 min reaction the portion of the microtubule that had formed before addition of BeF<sub>n</sub> was viewed continuously to be certain that it was stable; meanwhile, most of the microtubules elongated beyond the field of view. Finally, microtubule disassembly was induced by addition of BRB buffer only or with the same solution used for the second phase of the reaction sequence, minus tubulin. Use of a diluent without BeF<sub>n</sub> assured that the last portion of the microtubule that disassembled was exposed to BeF<sub>n</sub>-free conditions for the longest time. The NaF concentration was 5 mM NaF in the experiments in Figures 2 and 3, since this condition formed microtubules that were sufficiently unstable that the rate of disassembly could be measured along their entire length. NaF at 5 mM was also used for reactions of BeF<sub>n</sub> with microtubules in glycerol since we wished to reproduce earlier results (15, 16).

An immunofluorescence assay was used to measure BeF<sub>n</sub> stabilization of microtubules in glycerol. Microtubules assembled for 10 min with 23  $\mu$ M tubulin in BRB buffer with and without 3.4 M glycerol were diluted 50-fold into a solution containing 50 mM MES, pH 6.8, 0.5 mM EGTA, 0.25 mM MgCl<sub>2</sub>, 0.1 mM GTP, 3.4 M glycerol, 100  $\mu$ M BeSO<sub>4</sub>, and 5 mM NaF. At varying times aliquots were diluted 500-fold into 1% glutaraldehyde in BRB buffer, and microtubules were isolated on a cover slip placed on a layer of polymerized rubber caulking at the bottom of a tube that

was centrifuged at 28 500 rpm in a Beckman SW41 rotor. Microtubules were visualized from the signal from a fluorescein-labeled goat anti-mouse antibody (Capel) used to detect a mouse monoclonal antibody against  $\alpha$ -tubulin (DM1- $\alpha$ , from Sigma). Images were captured with a CCD (charge coupling device) camera and microtubule lengths were measured with the digitizing program in an Ambis 10 image processor (Hamamatsu Corp.).

Nuclear magnetic resonance spectra were recorded at room temperature on a Bruker AMX500 spectrometer at 70.28 MHz, with D<sub>2</sub>O as an internal lock. Spectra were accumulated for 30 min and chemical shifts are relative to BeSO<sub>4</sub>.

## RESULTS

It has been proposed (13) that the dissociation of P<sub>i</sub> from tubulin-GDP-P<sub>i</sub> subunits in microtubules occurs by a processive path in which the reaction occurs sequentially from adjacent subunits. It is possible to test this proposal by studying the mechanism for binding of the P<sub>i</sub> analogue BeF<sub>n</sub>. Since the principle of detailed balance requires that reactions follow an identical path in their forward and reverse directions (29), the same route is followed for dissociation and binding of P<sub>i</sub> to tubulin-GDP subunits in microtubules. Thus, if BeF<sub>n</sub> is an analogue of P<sub>i</sub> and if its binding is processive, then so too is P<sub>i</sub> dissociation. We were interested in determining whether a processive mechanism holds since it predicts that exposure of microtubules to P<sub>i</sub> (and with growing microtubules to BeF<sub>n</sub>; Figure 1) will form multiple contiguous tubulin-GDP-P<sub>i</sub> subunits. We had found (12) that microtubule ends can be stabilized by 13–14 contiguous tubulin-GTP subunits; therefore, if P<sub>i</sub> binding is processive, it is expected to be highly effective in stabilizing microtubules.

Evidence for processivity was sought by determining whether the stabilization that results from a brief exposure to BeF<sub>n</sub> is localized to the region of the microtubule that grew most recently (Figure 1). Binding to this region is predicted for a processive mechanism since P<sub>i</sub> and BeF<sub>n</sub> is required to bind first to the site from which P<sub>i</sub> was most recently lost, with subsequent binding to adjacent tubulin-GDP subunits. Since the most recently formed tubulin-GDP subunits are generated from newly added tubulin-GTP, the first site for binding of BeF<sub>n</sub> is expected to be near the growing end of the microtubule. To test this model microtubules were briefly elongated in the presence of BeF<sub>n</sub> and then diluted into tubulin-free buffer to induce disassembly. Evidence that BeF<sub>n</sub> bound preferentially to the growing end was sought by looking for a reduced disassembly rate in this portion of the microtubule.

**Processivity for Binding of BeF<sub>n</sub> to Microtubules.** Although microtubule stabilization by BeF<sub>n</sub> has been carefully characterized in buffer with 3.4 M glycerol (15, 16), we wished to study this phenomenon under conditions compatible with dynamic instability and, therefore, did not include glycerol in the reactions since this inhibits dynamic instability. We found in video microscopy studies that microtubules are unstable when diluted into 100  $\mu$ M BeSO<sub>4</sub>/5 mM NaF, which was surprising since 50  $\mu$ M BeF<sub>n</sub> had been found to stabilize almost instantly microtubules in glycerol buffers (16). A field viewed 60 s after dilution had only 9

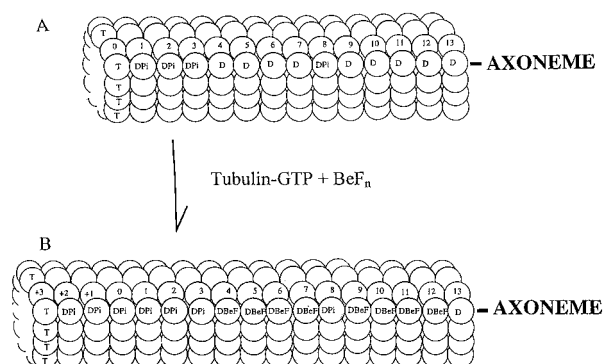


FIGURE 1: Implications of a processive mechanism for binding of BeF<sub>n</sub> and dissociation of P<sub>i</sub>. (A) Microtubules formed before addition of BeF<sub>n</sub> are presumed to have a terminal monolayer of tubulin-GTP subunits (12) and a short cap of tubulin-GDP-P<sub>i</sub> subunits (13) on a tubulin-GDP core. According to a strictly processive mechanism, P<sub>i</sub> would be lost sequentially from subunits 3, 2, 1, 0, +1, etc. (B) Subsequent elongation of the microtubule shown in panel A in the presence of BeF<sub>n</sub> first forms tubulin-GDP-BeF<sub>n</sub> in subunit 4, since this is the subunit that had most recently lost P<sub>i</sub>; subsequent binding occurs sequentially to subunits 5, 6, 7, etc. P<sub>i</sub> that would ordinarily be lost sequentially from subunits 3, 2, and 1 during addition of tubulin-GTP subunits (+1, +2, +3, etc.) is retained, since addition of BeF<sub>n</sub> to subunit 4 prevents loss of P<sub>i</sub> from subunit 3 (and 2 and 1, etc.). Thus, subunits added in the presence of BeF<sub>n</sub> will contain P<sub>i</sub>, and subunits present before BeF<sub>n</sub> will be progressively filled with this substance. P<sub>i</sub> is shown in subunit 8 both before and after elongation with BeF<sub>n</sub> because P<sub>i</sub> dissociation is not processive (see below); the concentration of tubulin-GDP-P<sub>i</sub> subunits is expected to progressively decrease toward the axoneme because these have had more time for dissociation to occur. Assuming that tubulin-GDP-P<sub>i</sub> subunits destabilize adjacent tubulin-GDP-BeF<sub>n</sub> subunits (see below), their relative absence in the "older" region of the microtubule would account for the greater stability of the portion of the microtubule near the axoneme (see Figures 2–4).

microtubules that had not fully disassembled (37  $\mu$ m cumulative length) from 97 initial microtubules (456  $\mu$ m cumulative length); equivalent results were observed in other viewing fields. Although the rate constant for BeF<sub>n</sub> binding to microtubules is only 90 M<sup>-1</sup> s<sup>-1</sup> (16), only 4 s in 100  $\mu$ M BeF<sub>n</sub> would be required to incorporate this into 4% of the tubulin-GDP subunits. This amount of substitution would be expected to increase microtubule stability since the effect of BeF<sub>n</sub> was found (16) to be described by

$$1/k_{\text{obs}} = f_-/k_- + f_+/k_+ \quad (2)$$

$f_+$  and  $f_-$  are the fraction of subunits with and without BeF<sub>n</sub> and  $k_+$  and  $k_-$  are the rates of dissociation of subunits with and without BeF<sub>n</sub>. With  $k_-/k_+$  equal to 25 (16), the rate will be reduced 2-fold when only 4% of subunits contain BeF<sub>n</sub>.

To determine whether the discrepancy from the earlier results came from the absence of glycerol in the BeF<sub>n</sub> diluent, microtubules assembled with or without 3.4 M glycerol were diluted 50-fold into glycerol-containing buffer with and without 100  $\mu$ M BeF<sub>n</sub>. Microtubules were unstable in glycerol buffer without BeF<sub>n</sub>. For example, microtubules assembled with glycerol had a mean length of 1–2  $\mu$ m and these were fully disassembled in 1 min; microtubules formed without glycerol were approximately 10-fold longer and their number concentration was reduced from an initial 22.4 microtubules/field ( $n = 11$ , SD 5.5) to 18.9 (11, 5.8) after 2 min, 15.1 (9, 2.8) after 5 min, and 3.9 (21, 2.2) after 20



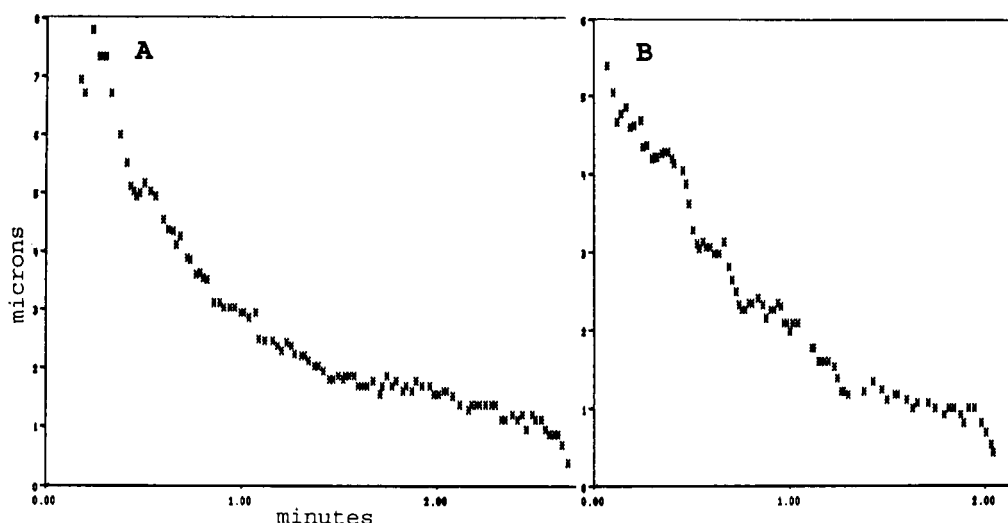


FIGURE 2: Attempt to detect processivity for binding of  $\text{BeF}_n$  to microtubules. Microtubules assembled at the end of axonemes with  $12.7 \mu\text{M}$  tubulin were reacted for approximately 30 s with  $12.7 \mu\text{M}$  tubulin subunits in 25 mM NaF and  $100 \mu\text{M}$   $\text{BeSO}_4$ ; disassembly was induced by addition of the same solution, minus tubulin. In panel A the rate during disassembly of the first half of the microtubule was  $200 \text{ s}^{-1}$ ; the rate was  $40 \text{ s}^{-1}$  for the other half. In panel B these rates were 107 and  $41 \text{ s}^{-1}$ , respectively.

min. In contrast, with microtubules assembled both with and without glycerol the number concentration was approximately the same at 1, 5, and 15 min after dilution into glycerol buffer with  $100 \mu\text{M}$   $\text{BeF}_n$ . Thus we confirmed the earlier finding (16) that microtubules in glycerol buffer are rapidly stabilized by  $\text{BeF}_n$  and deduce that  $\text{BeF}_n$  interacts with microtubules differently in the presence and absence of glycerol.

The lesser effect of  $\text{BeF}_n$  in the absence of glycerol may result because binding is weaker, so that the fraction of subunits as tubulin-GDP- $\text{BeF}_n$  (i.e.,  $f_+$  in eq 2) is lower, or because  $\text{BeF}_n$  has a lesser effect on subunit dissociation (i.e., increased  $k_-$ ). Evidence that weaker binding of  $\text{BeF}_n$  is responsible for the lesser effect in the absence of glycerol was obtained from studies in which assembled microtubules were elongated for 10 min in the presence of  $100 \mu\text{M}$   $\text{BeF}_n$ /5 mM NaF and then diluted to induce disassembly. The portion of the microtubule that was exposed to  $\text{BeF}_n$  for the entire 10 min disassembled at a rate equal to  $72 \text{ s}^{-1}$ , compared to  $621 \text{ s}^{-1}$  in a control (Table 1). Assuming a similar mechanism for microtubule stabilization by  $\text{BeF}_n$  in the presence and absence of glycerol, according to eq 2 the 9-fold reduction in rate with  $100 \mu\text{M}$   $\text{BeF}_n$  corresponds to 12.5% saturation of tubulin-GDP sites ( $f_+ = 0.125$ ). This predicts a  $K_d$  equal to  $700 \mu\text{M}$ ; a 10–13  $\mu\text{M}$   $K_d$  was determined in glycerol buffer (16).

Having found that microtubules can be stabilized by  $100 \mu\text{M}$   $\text{BeSO}_4$ , this concentration was used to determine whether the stabilization induced by a brief (30 s) exposure of microtubules to  $\text{BeF}_n$  is localized to the region near the growing end.  $\text{BeF}_n$  was formed from  $\text{BeSO}_4$  with 25 rather than 5 mM NaF since the lower concentration did not induce microtubule stabilization in 30 s. The requirement for a higher NaF concentration was surprising since the mole fraction of beryllium as  $\text{BeF}_3$ , which is presumably the reactive species for stabilizing tubulin-GDP subunits (14), only changes from 0.66 with 5 mM NaF to 0.61 with 25 mM F; however, the higher NaF increases the mole fraction as  $\text{BeF}_4$  from 0.08 to 0.34 (30). The requirement for higher NaF for rapid stabilization of microtubules suggests that in

Table 1: Rate of Disassembly of Microtubules in  $\text{BeF}_n$ <sup>a</sup>

| addition  | rate ( $\text{s}^{-1}$ ) (n, SD) |
|---|----------------------------------|
| none  | 621 (8, 217)                     |
| $\text{BeF}_n$ /5 mM NaF                            | 72 (10, 58)                      |
| $\text{P}_i$  | 684 <sup>b</sup>                 |
| $\text{Na}_2\text{SO}_4$                            | 479 (6, 105)                     |
| $\text{BeF}_n$ /25 mM NaF/ $\text{Na}_2\text{SO}_4$ | 11(10, 8)                        |
| $\text{BeF}_n$ /25 mM NaF/ $\text{P}_i$             | 248 (13, 174)                    |
| $\text{BeF}_n$ /25 mM NaF/ethylphosphonate          | 14 (8, 17)                       |

<sup>a</sup> Microtubules that had been assembled at the ends of axonemes with approximately  $15 \mu\text{M}$  tubulin in BRB buffer were treated for 10 min with a mixture of tubulin,  $100 \text{ mM}$   $\text{BeSO}_4$ , and 25 mM NaF and either 33 mM  $\text{NaSO}_4$ , 50 mM  $\text{P}_i$ , or 50 mM ethylphosphonate; control reactions lacked the  $\text{BeSO}_4$  and NaF. Disassembly was induced by addition of the same solution used for the  $\text{BeF}_n$  treatment but without tubulin. <sup>b</sup> Very rapid disassembly prevented tracing the length of individual microtubules and the rate was estimated from the reduction in length of 5 microtubules with a total length of  $41 \mu\text{m}$  to  $3.6 \mu\text{m}$  in 18 s. The calculated rate is a lower limit for the reaction since 21 other microtubules with comparable length were fully disassembled when the field was first observed.

the absence of glycerol microtubules are stabilized by  $\text{BeF}_4$  rather than by  $\text{BeF}_3$ .

The pattern for microtubule stabilization following a 30 s exposure to  $\text{BeF}_n$  was not in accord with a mechanism in which  $\text{BeF}_n$  binding is required to start near the growing end (Figure 1). Instead, the growing end was least stabilized so that the rate of disassembly progressively decreased as the microtubule disassembled (Figure 2). The ratio of the rate of disassembly of the first and second half of the microtubule was 3.2 ( $n = 7$ , SD 1.7). This is an approximate value since the decrease in the disassembly rate was progressive, so that a least-squares analysis of the data minimized the difference between the rate for the different portions of the microtubule.

It was possible that the progressive change in the disassembly rate after the 30 s exposure to  $\text{BeF}_n$  resulted because the portion of the microtubule nearest the axoneme became relatively enriched with tubulin-GDP- $\text{BeF}_n$  subunits during the time that the microtubules were disassembling in tubulin-free buffer/ $\text{BeF}_n$ . This possibility was eliminated in an

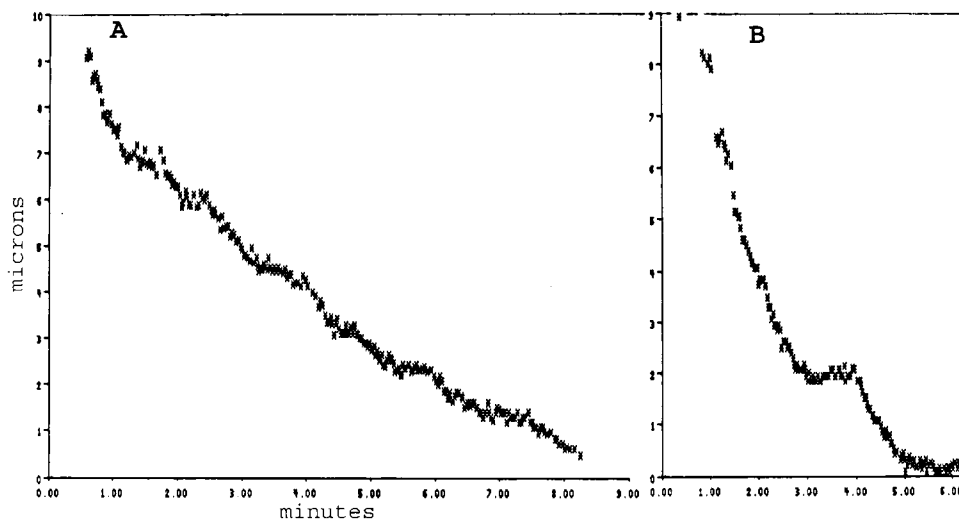


FIGURE 3: Disassembly of microtubules after reaction with  $\text{BeF}_n$  for 10 min. Kinetics are for the disassembly of microtubules that had formed at opposite ends of an axoneme in a reaction of  $14 \mu\text{M}$  tubulin without  $\text{BeF}_n$  for about 4 min and for 10 min in the presence of  $100 \mu\text{M}$  beryllium and  $5 \text{ mM}$  NaF. For the reaction in panel A, the microtubule length was  $6.4 \mu\text{m}$  before addition of  $\text{BeF}_n$  (therefore, this portion of the microtubule was in  $\text{BeF}_n$  for the entire 10 min) and  $>14.7 \mu\text{m}$  after the 10 min reaction in the presence of  $\text{BeF}_n$ . After dilution (zero time in the kinetic plot), about  $6 \mu\text{m}$  was lost during the approximately 15 s before the microtubule could be viewed. The rate of disassembly of the first and second half of the  $6.4 \mu\text{m}$  length that had been in  $\text{BeF}_n$  was  $33$  and  $20 \text{ s}^{-1}$ , respectively. The microtubule described in panel B was  $9 \mu\text{m}$  long before dilution, of which  $5 \mu\text{m}$  had formed before addition of  $\text{BeF}_n$ ; the disassembly rates were  $84$  and  $33 \text{ s}^{-1}$  before and after the pause at approximately 3 min.

experiment in which microtubules that had formed in the absence of  $\text{BeF}_n$  were further elongated for 10 min with a tubulin/ $\text{BeF}_n$  mixture, after which they were diluted into buffer without  $\text{BeF}_n$  or tubulin. NaF at  $5 \text{ mM}$  was used since this generated microtubules that were sufficiently unstable that it was possible to measure the rate of disassembly along their entire length. The predictions of a strictly processive mechanism for  $\text{P}_i$  dissociation are that maximum stabilization will be observed in the portion of the microtubule that formed in the presence of  $\text{BeF}_n$ . That is, if  $\text{P}_i$  dissociation is required to start at the interface of the GDP- $\text{P}_i$  cap and GDP core, then binding of  $\text{BeF}_n$  to subunits at this interface would lock  $\text{P}_i$  into tubulin-GDP- $\text{P}_i$  subunits in the portion of the microtubule that forms during assembly in the presence of  $\text{BeF}_n$  (Figure 1). The kinetics for dilution-induced disassembly did not support a processive mechanism since the portion of the microtubule that assembled in the presence of  $\text{BeF}_n$  was less rather than more stable than the portion that formed before addition of  $\text{BeF}_n$  (Figure 3). With almost all of the microtubules, the portion that formed in the presence of  $\text{BeF}_n$  was fully disassembled in the 15–30 s required for dilution and refocusing (Figure 3A). The low stability of the microtubule formed in the presence of  $\text{BeF}_n$  is not consistent with a processive mechanism in which binding is required to start at the interface of a tubulin-GDP- $\text{P}_i$  cap and a tubulin-GDP microtubule core (Figure 1). Also, as was observed with microtubules that had been exposed to  $\text{BeF}_n$  for 30 s (Figure 2), a 10 min exposure resulted in preferential stabilization of the microtubule that was furthest from the growing microtubule end. The half of the microtubule that was distant from the axoneme end disassembled 2.6-fold faster than the half near the axoneme ( $n = 12$ , SD 1.4). It is noted that the greater stabilization was in the portion of the microtubule that had the most time for loss of  $\text{BeF}_n$  during the disassembly process.

It was postulated that the slower disassembly of the portion of the microtubule nearest the axoneme (Figure 3) resulted

because this had a structure in which the A and B tubules of the doublet were both elongated, whereas only the A tubule was elongated in the more distal portion of the microtubule. This mechanism was tested by using ethylene glycol bis-(succinimidylsuccinate)- (EGS-) cross-linked single microtubules as seeds for elongating microtubules in the presence of  $\text{BeF}_n$ . Again, the rate was slower as the disassembly progressed toward the seed (Figure 4), ruling out the postulated mechanism: the rate for the half of the microtubule remote from the axoneme was 2.7 times ( $n = 10$ , SD 1.4) faster than the half near the axoneme.

*Can Microtubules Be Stabilized by  $\text{P}_i$ ?* Microtubules assembled on axonemes without  $\text{P}_i$  were elongated for 10 min with tubulin in  $50 \text{ mM}$   $\text{P}_i$ . When disassembly was induced by dilution into buffer containing  $\text{P}_i$ , neither the portion of the microtubule formed in the absence nor in the presence of  $\text{P}_i$  was unusually stable; most microtubules had fully disassembled in the 15–25 s required for addition of the diluent and refocusing on the field. The microtubules that could be observed to disassemble did so at a rate equal to  $684 \text{ s}^{-1}$  (Table 1), which is equivalent to that in a control reaction in which an identical ionic strength was maintained with  $\text{Na}_2\text{SO}_4$  (Table 1).

Since  $\text{P}_i$  binding is not processive so as to form multiple contiguous tubulin-GDP- $\text{P}_i$  subunits, a different approach was used to facilitate their formation. Microtubules were reacted with a mixture of  $\text{BeF}_n$  and  $\text{P}_i$ . Under these conditions the probability for forming  $n$  contiguous tubulin-GDP subunits containing either  $\text{P}_i$  or  $\text{BeF}_n$  is equal to the  $n$ th power of the sum of the mole fraction of subunits with  $\text{BeF}_n$  and the mole fraction of subunits with  $\text{P}_i$ . It was expected that binding of  $\text{P}_i$  would enhance the effect of  $\text{BeF}_n$  in regions of the microtubule in which the number of contiguous tubulin-GDP- $\text{BeF}_n$  is insufficient to stabilize maximally the microtubule end. We found, however, that  $\text{P}_i$  inhibited rather than enhanced the stabilization induced by  $\text{BeF}_n$  (Table 1, Figure 5). Representative results showing the dramatic reversal by

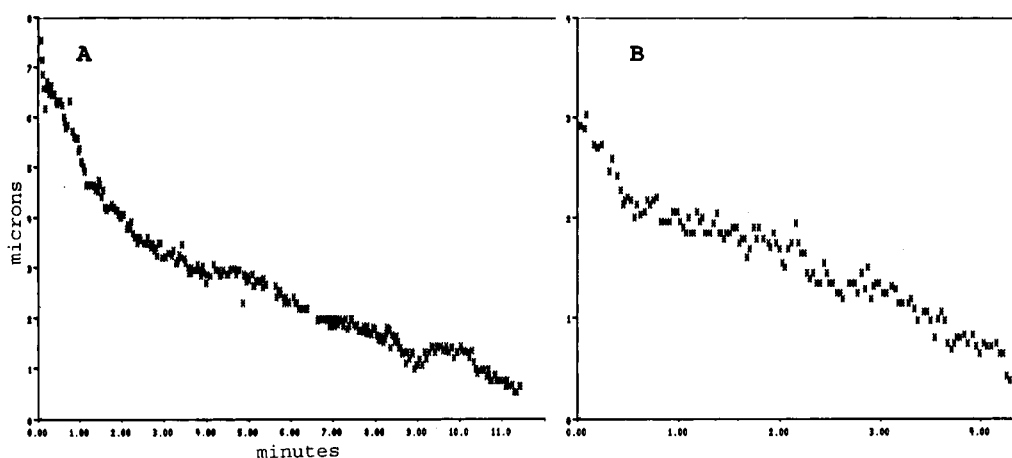


FIGURE 4: Effect of  $\text{BeF}_n$  on the stability of microtubules that had been assembled at the ends of EGS seeds. Microtubules assembled without  $\text{BeF}_n$  were elongated for an additional 10 min in 100  $\mu\text{M}$  beryllium and 5 mM NaF. For panel A the length was 8.2  $\mu\text{m}$  before dilution with BRB buffer, 4.1  $\mu\text{m}$  of which had been formed before addition of beryllium fluoride; in panel B the corresponding lengths were 4.5 and 2.2  $\mu\text{m}$ , respectively. The rates in panel A started at  $53 \text{ s}^{-1}$  and progressively slowed to  $8 \text{ s}^{-1}$  after 6 min. In panel B the rates ranged from 53 to  $12 \text{ s}^{-1}$ . Zero time in both reactions corresponds to 20–30 s after dilution with buffer.

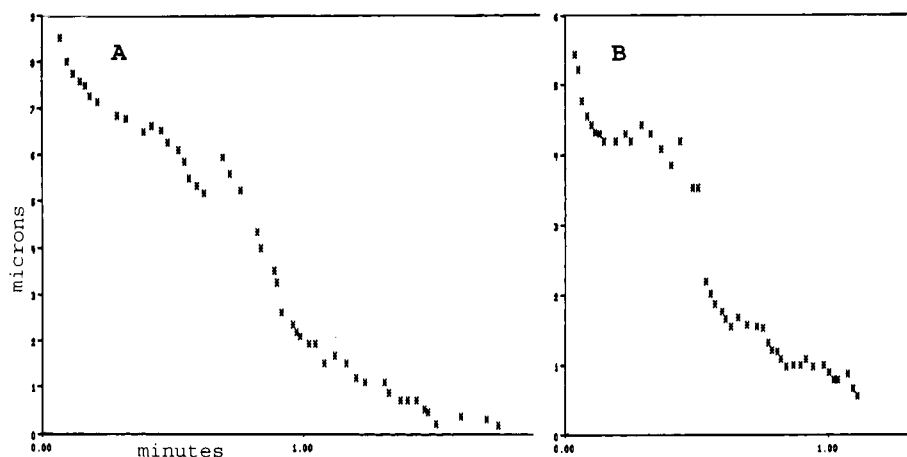


FIGURE 5: Effect of  $\text{P}_i$  on  $\text{BeF}_n$  stabilization of microtubules. Microtubules assembled without  $\text{BeF}_n$  were further elongated for 10 min in 25 mM NaF/100  $\mu\text{M}$   $\text{BeSO}_4$  and 50 mM  $\text{P}_i$  and then diluted with the same solution, minus tubulin. The predilution length of the microtubule described in panel A was 11.3  $\mu\text{m}$ , of which 3.5  $\mu\text{m}$  had formed before addition of  $\text{BeF}_n$ . The rate during the first 0.7 min was  $111 \text{ s}^{-1}$  and this increased to  $294 \text{ s}^{-1}$  for about 0.3 min and then decreased to  $84 \text{ s}^{-1}$ . The tip of the microtubule described in panel B was out the range of view for 45 s after dilution with buffer; the rate was 300 and  $452 \text{ s}^{-1}$  before and after a brief pause.

$\text{P}_i$  of the microtubules stabilization by  $\text{BeF}_n$  are shown in Figure 6. The  $\text{P}_i$  analogue ethylphosphonate did not similarly reverse the stabilization of microtubules by  $\text{BeF}_n$  (Table 1).

The possibility that  $\text{P}_i$  inhibits the stabilization of microtubules by displacing fluorine from  $\text{BeF}_n$  was ruled out from the  $^9\text{Be}$  NMR spectrum in the presence and absence of  $\text{P}_i$ . The fraction of beryllium with 1–4 coordinated fluorides can be determined from the NMR spectrum (31) and the high sensitivity of this method is illustrated from our results with 2.5 and 5 mM NaF. It was possible (Figure 7) to detect the change in the percent  $\text{BeF}_n$  ( $n = 2, 3$ , and 4) from 35, 53, and 3.5 with 2.5 mM NaF to 21, 66, and 9 with 5 mM NaF (30). Results from the  $^9\text{Be}$  NMR analysis were confirmed with  $^{19}\text{F}$  NMR. In BRB buffer (pD 6.8) containing 1 mM  $\text{BeSO}_4$  and 25 mM NaF, the integrated intensity of the signals (normalized relative to the signal from NaF) from  $\text{BeF}_3$  and  $\text{BeF}_4$  were 1.28 and 0.58, respectively, in the absence of  $\text{P}_i$ . In the presence of an equimolar mixture of 25 mM  $\text{Na}_2\text{HPO}_4$  and  $\text{NaH}_2\text{PO}_4$  the relative intensities were 1.08 and 0.57, respectively. Identical spectra were observed when these were recorded with a 1 or 5 s relaxation delay.

## DISCUSSION

There is no direct evidence that microtubules can be stabilized by a cap of tubulin-GDP- $\text{P}_i$  subunits. Instead, slow dissociation of tubulin-GDP- $\text{P}_i$  has been inferred from studies with  $\text{BeF}_n$  (11, 16), which forms tubulin-GDP- $\text{BeF}_n$  in microtubules and decreases the rate of subunit dissociation about 25-fold compared to tubulin-GDP. The conclusion that tubulin-GDP- $\text{P}_i$  stabilizes microtubules is, however, not in accord with studies of the dynamics of individual microtubules (19, 20), showing that high concentrations of  $\text{P}_i$  did not decrease the rate of subunit dissociation or the catastrophe frequency.

On the basis of results obtained in glycerol buffer, it would appear that microtubules contained tubulin-GDP- $\text{P}_i$  subunits under the conditions where  $\text{P}_i$  was found not to stabilize microtubules (Table 1; 19). A rate constant for  $\text{P}_i$  binding equal to  $5 \text{ M}^{-1} \text{ s}^{-1}$  is calculated from the ratio of the association and dissociation rate constants (equal to the  $K_d$ ), with  $K_d$  equal to 25 mM (16) and the dissociation rate constant equal to  $0.126 \text{ s}^{-1}$  (13). Thus, in reactions with 0.05 M  $\text{P}_i$  the expected half-time for binding is 2.8 s; a 10

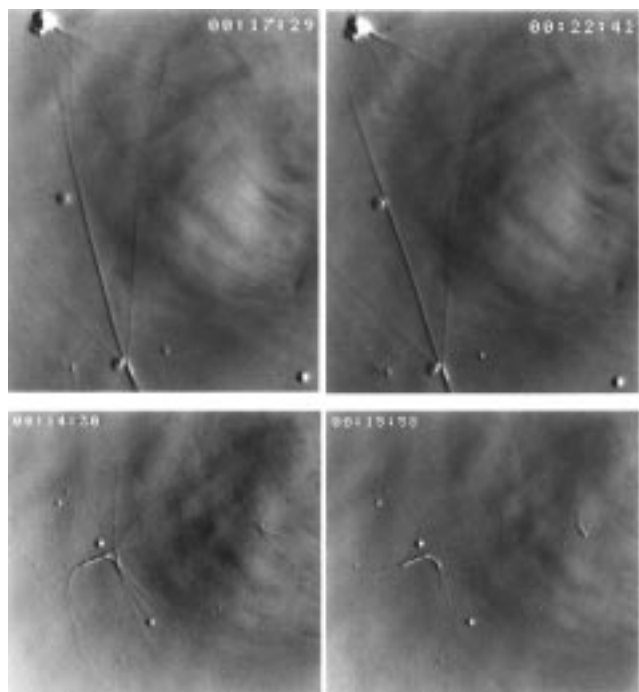


FIGURE 6: Dilution-induced disassembly of the microtubules that had been reacted with  $\text{BeF}_n$  in the presence (top) and absence (bottom) of  $\text{P}_i$ ; conditions were as described for Figure 5. Dilution was induced at 15:00 min for the reaction with  $\text{P}_i$  and at 14:05 min for the reaction without  $\text{P}_i$ .

min incubation with  $\text{P}_i$  was used in our studies. The lack of stabilization by  $\text{P}_i$  indicates that in the absence of glycerol the 25 mM  $K_d$  for  $\text{P}_i$  and/or the  $0.126 \text{ s}^{-1}$  rate constant for  $\text{P}_i$  dissociation does not hold, and/or that microtubules are not stabilized by tubulin-GDP- $\text{P}_i$  subunits.

The proposed processive path for GTP hydrolysis and  $\text{P}_i$  dissociation (13) would be expected to enhance the stabilizing

effect of added  $\text{P}_i$ , if it were true that tubulin-GDP- $\text{P}_i$  subunits can stabilize microtubule ends. According to a processive mechanism for GTP hydrolysis, this occurs predominantly at the interface of a GTP cap and the microtubule's GDP-core (32–34). The dissociation (and binding) of  $\text{P}_i$  would be similarly processive, starting at the interface of the tubulin-GTP cap and proceeding into adjacent tubulin-GDP subunits (Figure 1). This path would generate multiple contiguous tubulin-GDP- $\text{P}_i$  subunits and these are expected to have an enhanced stabilizing effect on microtubule ends since it was found (16) that the reduction in the subunit dissociation rate is proportional to more than the first power of the mole fraction of microtubule subunits containing  $\text{BeF}_n$ . The probability of  $n$  contiguous tubulin-GDP- $\text{BeF}_n$  subunits is proportional to the  $n$ th power of the mole fraction of tubulins containing  $\text{BeF}_n$ . Our finding (12) that 13 or 14 contiguous tubulin-GMPCPP subunits are necessary and sufficient to stabilize a microtubule end further supports our postulate that processive binding of  $\text{P}_i$  would be highly effective in stabilizing microtubules.

Binding of  $\text{BeF}_n$  and presumably  $\text{P}_i$  is not processive since the stabilization induced by exposure of microtubules to  $\text{BeF}_n$  was not greatest in the region of the growing end of the microtubule (Figures 2–4). Instead, maximal stabilization was in the region most distal to the growing end. Thus, the path for  $\text{P}_i$  binding is not expected to generate exclusively contiguous tubulin-GDP- $\text{P}_i$  subunits. It is nevertheless possible to form multiple contiguous tubulin-GDP- $\text{P}_i$  subunits, but as described above this is statistically likely only when microtubules contain a significant mole fraction of subunits with  $\text{P}_i$ .

The lack of microtubule stabilization by  $\text{P}_i$  did not result from failure to form sufficient contiguous tubulin-GDP- $\text{P}_i$  subunits but instead resulted because the microtubule lattice is not stabilized by binding of  $\text{P}_i$  to tubulin-GDP subunits.

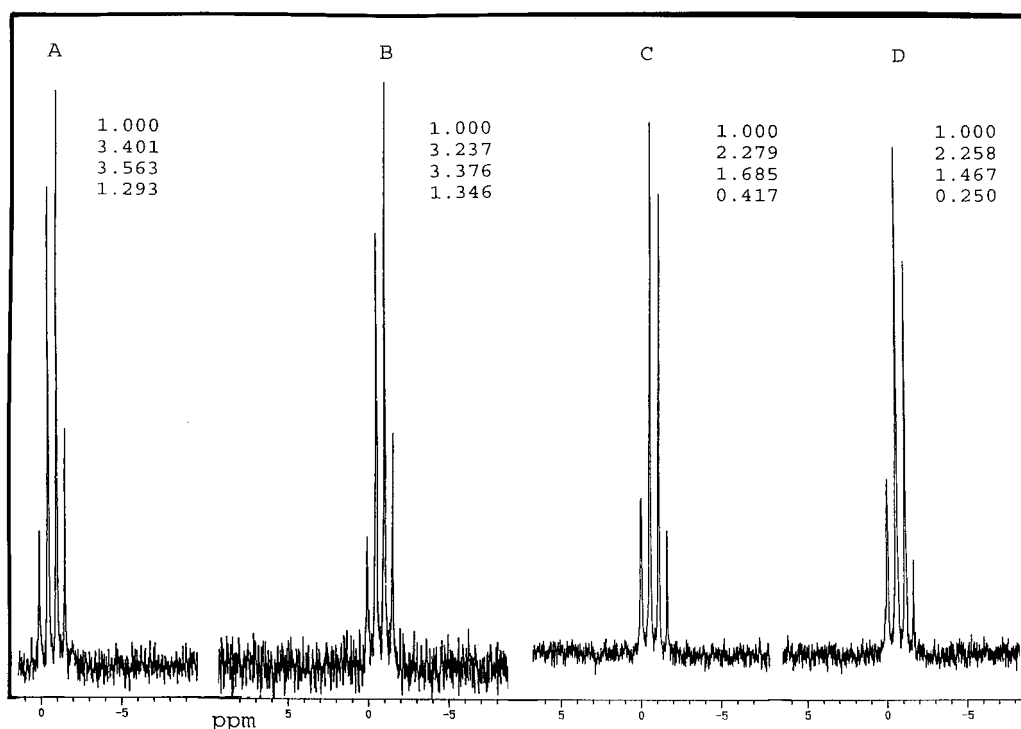


FIGURE 7: Nuclear magnetic resonance spectra of  $\text{BeF}_n$  in the presence and absence of  $\text{P}_i$ . All spectra are from  $100 \mu\text{M}$   $\text{BeSO}_4$  in BRB buffer, with 25 mM NaF (A), 25 mM NaF and 50 mM  $\text{P}_i$  (B), 5 mM NaF (C), and 2.5 mM NaF (D).



This conclusion is derived from our observation that  $P_i$  and  $BeF_n$  did not act synergistically in stabilizing microtubules. Synergism would result if binding of both of these substances to GDP subunits stabilizes the microtubule lattice and if, as with GMPCPP (12), microtubule stabilization is enhanced by multiple contiguous stabilizing subunits. Synergism is predicted because the probability for forming  $n$  contiguous tubulin-GDP subunits containing either  $P_i$  or  $BeF_n$  is proportional to the  $n$ th power of the sum of the mole fraction of tubulin-GDP subunits containing either  $P_i$  or  $BeF_n$ . In contrast with this prediction,  $P_i$  antagonized rather than enhanced  $BeF_n$  stabilization of microtubules (Figures 5 and 6). Destabilization by  $P_i$  resulted because it displaced  $BeF_n$  from the tubulin-GDP subunits and because tubulin-GDP- $P_i$  subunits are unstable in the microtubule lattice. Finally, our finding that  $P_i$  reversed the effect of  $BeF_n$  requires that  $P_i$  is bound to microtubules when this substance is present; greater binding of  $P_i$  would occur when  $BeF_n$  is not present to compete for tubulin-GDP subunits. Therefore, our finding that  $P_i$  did not increase the stability of microtubules in the absence of  $BeF_n$  means that tubulin-GDP- $P_i$  subunits do not stabilize microtubule ends.

Evidence that binding of  $P_i$  further decreases the stability of tubulin-GDP subunits comes from earlier results showing that  $P_i$  dramatically increased the catastrophe and disassembly rate and decreased the rescue rate with microtubules formed with chick erythrocyte and pig brain tubulin (20). The greater effect of  $P_i$  seen with erythrocyte microtubules (20) suggests that erythrocyte-specific  $\beta$ -6/ $\alpha$ -1 tubulin (35) has enhanced affinity for  $P_i$ . Our observation that tubulin-GDP- $P_i$  subunits destabilize the microtubule lattice provides evidence that the increased dynamics seen with this substance (20) did not result from a medium effect but instead resulted from  $P_i$  binding to the microtubule.

Our finding that the  $P_i$  analogue ethylphosphonate did not inhibit  $BeF_n$ -induced microtubule stabilization (Table 1) provides evidence against a mechanism in which  $P_i$  reduces microtubule stabilization by  $BeF_n$  by displacing fluoride from beryllium, to reduce the concentration of  $BeF_n$ . If  $P_i$  acted by reacting with beryllium then the greater basicity of ethylphosphonate compared to  $P_i$  would be expected to make it more reactive with beryllium and thus more effective in inhibiting the effect of  $BeF_n$ . Also, the monodentate complex of pyrophosphate and beryllium was completely displaced when the fluoride concentration was only 20 mM (26); therefore, it is not expected that  $P_i$  would be bound to beryllium in the presence of the 25 mM fluoride used in our experiments. Finally,  $^9Be$  and  $^{19}F$  NMR analysis of  $BeF_n$  in the presence and absence of  $P_i$  showed that the inhibitory effect of  $P_i$  on  $BeF_n$  stabilization did not result from its displacing fluoride from beryllium (Figure 7).

We do not have an unambiguous interpretation of the progressive decrease in the rate of disassembly of microtubules exposed to  $BeF_n$  (Figures 2–4). It was previously found that the disassembly rate can vary along the microtubule length (36–38), but this variability was random. The greater stabilization near the nucleating axoneme or EGS seed did not result because of longer exposure to  $BeF_n$ . Rather, because the dilution was into  $BeF_n$ -free buffer, the last-disassembling portion of the microtubule had the most time for dissociation of  $BeF_n$ . It is suggested that the greater stabilization of the portion of the microtubule near the seed

results because this first-formed portion of the microtubule has had more time to release  $P_i$  from tubulin-GDP- $P_i$  subunits (Figure 1). If tubulin-GDP- $P_i$  subunits can vitiate the stabilizing effect of adjacent tubulin-GDP- $BeF_n$  subunits, the microtubule will be most stable where the probability is greatest that  $P_i$  has been lost. The very low stability of the portion of the microtubule that forms in the presence of  $BeF_n$  may result because this region is enriched with tubulin-GDP- $P_i$  subunits, with dissociation of  $P_i$  inhibited because of binding of  $BeF_n$  to subunits that have lost  $P_i$ .

In summary, we have obtained evidence that tubulin-GDP- $P_i$  subunits destabilize the microtubule lattice so that their presence at microtubule ends cannot be responsible for microtubule dynamic instability as previously proposed (12, 15). Our results indicate that tubulin-GDP- $BeF_n$  is an analogue of tubulin-GTP rather than of tubulin-GDP- $P_i$ . This conclusion is supported by results with G-proteins and myosin where  $BeF_n$  and bind to the site normally occupied by the  $\gamma$ -phosphate moiety of GTP and ATP (39–41) and mimics the effect of the nucleoside triphosphate. The lack of microtubule stabilization by tubulin-GDP- $P_i$  and the failure to observe a large tubulin-GTP cap suggests that growing microtubules are stabilized by a monolayer of tubulin-GTP subunits (12).

## ACKNOWLEDGMENT

We are grateful to Ted Salmon for critical comments on the manuscript.

## REFERENCES

1. Carlier, M.-F., and Pantaloni, D. (1981) *Biochemistry* 20, 1918–1924.
2. Arai, T., and Kaziyo, Y. (1976) *Biochem. Biophys. Res. Commun.* 69, 369–376.
3. Weisenberg, R. C., Deery, W. J., and Dickinson, P. J. (1976) *Biochemistry* 15, 4248–4254.
4. Weisenberg, R. C., and Deery, W. J. (1976) *Nature* 263, 792–793.
5. O'Brien, E. T., Voter, W. A., and Erickson, H. P. (1987) *Biochemistry* 26, 4148–4156.
6. Stewart, R. J., Farrell, W. W., and Wilson, L. (1990) *Biochemistry* 29, 6489–6498.
7. Bayley, P. M., Schilstra, M. J., and Martin, S. R. (1990) *J. Cell Sci.* 95, 33–48.
8. Erickson, H. P., and O'Brien, E. T. (1992) *Annu. Rev. Biophys. Biomol. Struct.* 21, 145–166.
9. Martin, S. R., Schilstra, M. J., and Bayley, P. M. (1993) *Biophys. J.* 65, 578–596.
10. Drechsel, D. N., and Kirschner, M. W. (1994) *Curr. Biol.* 4, 1053–1061.
11. Melki, R., Carlier, M.-F., and Pantaloni, D. (1990) *Biochemistry* 29, 8921–8932.
12. Caplow, M., and Shanks, J. (1996) *Mol. Biol. Cell* 7, 663–675.
13. Melki, R., Fievez, S., and Carlier, M.-F. (1996) *Biochemistry* 35, 12038–12045.
14. Combeau, C., and Carlier, M.-F. (1989) *J. Biol. Chem.* 264, 19017–19021.
15. Carlier, M. F., Didry, D., Melki, R., Chabre, M., and Pantaloni, D. (1988) *Biochemistry* 27, 3555–3559.
16. Carlier, M. F., Didry, D., Simon, C., and Pantaloni, D. (1989) *Biochemistry* 28, 1783–1791.
17. Fisher, A. J., Smith, C. A., Thoden, J. B., Smith, R., Sutoh, K., Holden, H. M., and Raayment, I. (1995) *Biochemistry* 34, 8960–8972.
18. Sondek, J., Lambright, D. G., Noel, J. P., Hamm, M., and Sigler, P. B. (1994) *Nature* 372, 276–279.



19. Caplow, M., Ruhlen, R., Shanks, J., Walker, R. A., and Salmon E. D. (1989) *Biochemistry* 28, 8136–8141.
20. Trinczek, B., Marx, A., Mandelkow, E.-M., Murphy D. B., and Mandelkow, E. (1993) *Mol. Biol. Cell* 4, 323–335.
21. Caplow, M., Ruhlen, R. L., and Shanks, J. (1994) *J. Cell Biol.* 127, 779–788.
22. Purich, D. L., and Angelastro, J. M. (1994) *Adv. Enzymol. Relat. Areas Mol. Biol.* 69, 121–154.
23. Weingarten, M. D., Lockwood, A. H., Hwo, S., and Kirschner, M. W. (1975) *Proc. Natl. Acad. Sci. U.S.A.* 72, 1858–1862.
24. Vale, R. D. (1991) *Cell* 64, 827–839.
25. Glikzman, N. S., Parson, S. F., and Salmon, E. D. (1992) *J. Cell Biol.* 119, 1271–1276.
26. Feeney, J., Haque, R., Reeves, L. W., and Yue, C. P. (1967) *Can. J. Chem.* 46, 1389–1398.
27. Greenwald, I., Redish, J., and Kibrick, A. C. (1940) *J. Biol. Chem.* 135, 65–76.
28. Sullivan, J. C., and Hindman, J. C. (1953) *J. Am. Chem. Soc.* 74, 6091–6096.
29. Jencks, W. P. (1969) *Catalysis in Chemistry and Enzymology*, p 613, McGraw-Hill, New York.
30. Martin, R. B. (1988) *Biochem. Biophys. Res. Commun.* 155, 1194–1200.
31. Issartel, J.-P., Dupuis, A., Morat, C., and Giradet, J.-L. (1991) *Eur. Biophys. J.* 20, 115–126.
32. Caplow, M., Shanks, J., and Brylawski, B. P. (1984) *Can. J. Biochem. Cell Biol.* 63, 422–429.
33. Caplow, M., and Reid, R. (1985) *Proc. Natl. Acad. Sci. U.S.A.* 82, 3267–3271.
34. Carlier, M.-F. (1989) *Int. Rev. Cytol.* 115, 139–170.
35. Murphy, D. B., Wallis, K. T., Machlin, P., Patrie, H., and Cleveland, D. (1987) *J. Biol. Chem.* 262, 14305–14312.
36. O'Brien, E. T., Salmon, E. D., Walker, R. A., and Erickson, H. P. (1990) *Biochemistry* 29, 6648–6656.
37. Billger, M. A., Bhattacharjee, G., and Williams, R. C., Jr. (1996) *Biochemistry* 35, 13656–13663.
38. Dye, R. B., and Williams, R. C., Jr. (1996) *Biochemistry* 35, 14331–14339.
39. Antonny, B., Bigay, J., and Chabre, M. (1990) *FEBS Lett.* 268, 277–280.
40. Antonny, B., and Chabre, M. (1992) *J. Biol. Chem.* 267, 6710–6718.
41. Bigay, J., Deterre, P., Pfister, C., and Chabre, M. (1987) *EMBO J.* 6, 2907–2913.

BI972602+