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Relaxation of Loaded ESCRT-III Spiral Springs Drives Membrane Deformation

Graphical Abstract

Highlights

- Snf7 forms highly flexible filaments that spontaneously curl
- Snf7 filaments form spirals at the surface of lipid membranes
- Snf7 spirals are springs as they can deform under lateral compression
- Relaxation of compressed Snf7 spirals leads to membrane deformation

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In Brief
A component of the ESCRT-III membrane fission machinery self-organizes into spiral springs that trigger membrane deformation when released.
Relaxation of Loaded ESCRT-III Spiral Springs Drives Membrane Deformation

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SUMMARY

ESCRT-III is required for lipid membrane remodeling in many cellular processes, from abscission to viral budding and multi-vesicular body biogenesis. However, how ESCRT-III polymerization generates membrane curvature remains debated. Here, we show that Snf7, the main component of ESCRT-III, polymerizes into spirals at the surface of lipid bilayers. When covering the entire membrane surface, these spirals stopped growing when densely packed: they had a polygonal shape, suggesting that lateral compression could deform them. We reasoned that Snf7 spirals could function as spiral springs. By measuring the polymerization energy and the rigidity of Snf7 filaments, we showed that they were deformed while growing in a confined area. Furthermore, we observed that the elastic expansion of compressed Snf7 spirals generated an area difference between the two sides of the membrane and thus curvature. This spring-like activity underlies the driving force by which ESCRT-III could mediate membrane deformation and fission.

INTRODUCTION

ESCRT-III (endosomal sorting complex required for transport) has been implicated in the formation of intralumenal vesicles (ILVs) during biogenesis of multi-vesicular bodies (MVBs) by genetic (Babst et al., 2002; Coonrod and Stevens, 2010) and biochemical assays (Adell et al., 2014; Henne et al., 2012; Saksena et al., 2009; Wollert and Hurley, 2010; Wollert et al., 2009). ESCRT-III budding occurs in an opposite direction than in endocytosis: the limiting membrane is pushed outward from the cytoplasm instead of curving inward. ESCRT-III has been proposed to play a role in membrane deformation (Hanson et al., 2008) and fission of ILVs (Adell et al., 2014). Consistent with this, ESCRT-III is also required for geometrically similar fission reactions such as viral budding (von Schwedler et al., 2003) and abscission during cytokinesis (Carlton et al., 2008; Elia et al., 2011; Guizetti et al., 2011). ESCRT-III nucleation is promoted by ESCRT-II and its disassembly by the ATPase Vps4 (Lata et al., 2008).

It is unclear how ESCRT-III deforms lipid membranes. Because of their polymerization abilities, ESCRT-III proteins (Vps20, Snf7, Vps2, Vps24) have been proposed to generate membrane curvature by scaffolding (Cashikar et al., 2009; Fabrikant et al., 2009; Hanson et al., 2008; Lata et al., 2008). In this mode, polymers coating the membrane usually adopt a single specific shape, or, at least, a set of geometrically similar shapes. ESCRT-III filaments adopt instead a wide variety of shapes in vivo and in vitro: concentric circles, rings, spirals, helices, or linear filaments have been observed (Hanson et al., 2008; Henne et al., 2012; Pires et al., 2009). Furthermore, no unique shape for the assembly of ESCRT-III proteins arises from the molecular structure of ESCRT-III proteins (McCullough et al., 2013). Instead, curvature could be generated by other mechanisms: for example, it has been proposed that the amphipathic insertion of the N-terminal part of Snf7 could participate in the generation of membrane curvature (Buchkovich et al., 2013). We were thus interested in studying how ESCRT-III polymerization could drive membrane curvature.

RESULTS

Growth of Snf7 Patches on Supported Bilayers

To study the polymerization of ESCRT-III, we reconstituted ESCRT-III polymerization by adding purified yeast Snf7 onto supported lipid membranes. Supported membranes were obtained by bursting giant unilamellar vesicles (GUVs) composed of 40% di-oleoyl-phosphatidylserine (DOPS) and 60% of di-oleoyl-phosphatidylcholine (DOPC) on cleaned glass coverslips (Figure S1A, Movie S1). These coverslips were built into a flow chamber, allowing sequential addition and exchange of solutions.

First, Snf7 labeled with Alexa488 (Snf7-Alexa488) was flushed into the chamber, and its association to the membrane was imaged by time-lapse spinning-disk confocal microscopy (SDC). At 400 nM, Snf7 formed patches evenly distributed on...
Figure 1. Nucleation and Growth of Snf7 Patches on Supported Membranes
Lipid composition is DOPC 60% / DOPS 40% + Rhodamine PE 0.1%.
(A) Time-lapse images of Snf7-Alexa488 patches growth (green) at [Snf7] = 400 nM on supported membrane (gray).
(B) Time-lapse images (every 10 min) of a single Snf7-Alexa488 patch (green) growing at [Snf7] = 200 nM.

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the membrane surface (Figure 1A). These circular patches (Figure 1B) grew over the course of an hour until becoming confluent, eventually covering the membrane completely.

The nucleation rate of the patches depended on Snf7 bulk concentration (Figure 1C). Patch formation was not observed below 200 nM. Above 1 μM, patch formation and growth was so fast that individual patches could hardly be discriminated. Between these limiting concentrations, we were able to follow micron-sized patches individually over several tens of minutes (Movie S2). Once formed, patches disassembled with a half time of approximately 15 hr upon Snf7 washout (Figure S1B; Movie S3).

We termed “patch nucleation” this nucleation of patches in the absence of a previous Snf7 structure. The patch nucleation rate was very low, less than 1 seed, μm⁻².hour⁻¹, and depended on the amount of negatively charged lipids in the membrane (Figure S1C), revealing the critical role of these lipids in promoting Snf7 polymerization. We did not observe Snf7 assemblies in the absence of membranes.

The periphery of the patches showed dimmer Snf7 fluorescence than the center: fluorescence decayed radially at the rim over the outer 4 μm. This gradient of fluorescence was the same as the patch grew in radius (Figure 1D) and independent on bulk concentration of Snf7 (Figure 1E). The central part of the patches had the same intensity, constant over time. We postulated that the patch could be made of two parts: a central part where Snf7 entirely covers the membrane and cannot further assemble, and a rim, representing a growing front. To study whether Snf7 was assembled only in the front region, we generated patches with a solution of Snf7-Alexa488, which we then replaced with Snf7-Atto647N (Figure 1F and Movie S4). As postulated, Snf7-Atto647N fluorescence appeared only at the border of the growing patches. These observations confirmed that Snf7 patches were growing by a traveling circular front.

The front propagated at constant speed (Figure 1G). The speed was linear with Snf7 concentration (Figure 1H). As a result, and because the fluorescence gradient was independent from Snf7 concentration, fluorescence intensity curves with time at a given point (Figure S1E) could be merged into a single one (Figure S1F). The amount of negatively charged lipids also affected the front speed (Figure S1D). In summary, the growth of the Snf7 patches reflected a nucleation/growth process, with a nucleation rate of less than 1 seed, μm⁻².hour⁻¹ and a radial growth speed of 760 nm.min⁻¹.μM⁻¹.

Since Snf7 filaments can curl into circles or spirals (Hanson et al., 2008; Henne et al., 2012; Shen et al., 2014), we reasoned that one patch could be made of a single spiral filament growing from its tips at constant rate. In this case, however, the radial growth speed of the patch should slow down as the square root of time, and a dimmer fluorescence at the periphery of the patches would not be expected. To resolve this apparent contradiction, we studied the molecular structure of the Snf7 patches with atomic force microscopy (AFM) and electron microscopy (EM).

Snf7 Patches Are Made of Spiraling Filaments with Lateral Interactions

We first acquired images of Snf7 patches by AFM. GUVs composed of 60% DOPC and 40% DOPS were burst on a mica support (Figure S2A). After a four hours incubation of Snf7 at 1 μM, AFM images revealed that the micron-sized Snf7 patches consisted of packed arrays of Snf7 circular assemblies (Figure 2A; Figure S2B). Each assembly was formed by concentric circle-like structures. However, in these packed conditions, rather than being perfectly circular, each assembly was deformed into polygons with six neighbors on average (Figure 2B). The average external radius was 123 ± 35 nm (n = 120) in agreement with previous data (Pires et al., 2009; Shen et al., 2014). The thicker ones were double-stranded, with approximately twice the thickness (10.7 ± 0.7 nm; n = 7). The larger circular assemblies had an average radius of 110 ± 40 nm (n = 46). Following the path of the innermost filament in these assemblies revealed that they were made of a single filament, self-organized into a spiral (Figure 2H; Figure S2D). Within the spiral, we observed that the filament could associate laterally with itself, forming double-stranded filaments. These images showed that the spiraling nature of Snf7 filaments previously observed in solution (Shen et al., 2014) is conserved at the

(C) Patch nucleation rate as a function of [Snf7].
(D) Successive (from bright to dark green, every 10 min) Snf7 patch fluorescence profiles (circularly averaged) at [Snf7] = 200 nM.
(E) Snf7 patch edge fluorescence profile (average of 3 patches) as a function of [Snf7] (data for [Snf7] < 200 nM were obtained by first nucleating the patches at 350 nM for 5 min, and then [Snf7] was reduced to the desired value).
(F) Exchange of bulk Snf7-Alexa488 (green) with Snf7-Atto647N (red) at 200 nM. Inset: kymograph of the region selected (yellow box). The green line is the switch between green and red Snf7.
(G) Equatorial kymograph of the patch shown in B.
(H) Patch radial growth speed as a function of [Snf7]. The slope of the linear fit (gray line) is 760 nm.min⁻¹.μM⁻¹.

See also Figure S1.
Figure 2. Patches Are Made of Packed Snf7 Spirals
(A) AFM topographic image of the center of a Snf7 patch.
(B) Histogram of the number of neighbors per assembly.
(C) Histogram of the average outer radii of Snf7 assemblies. The average radius is 123 nm ± 33 nm.
(D) Histogram of the innermost circle radii. The average radius is 18 nm ± 3 nm.
(E) Histogram of the inter-circle distance. The average distance is b = 17 nm ± 3 nm.
(F) TEM image of a negatively stained, Snf7-coated LUV.

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membrane. It also suggests that previously reported circular as-
sembles could also be spirals: in vitro, a Snf7 mutant protein that
polymerizes spontaneously onto membranes (Henne et al.,
2012) or, in fibroblasts, the circular structures found upon over-
expression of CHMP4A and CHMP4B (Hanson et al., 2008).
The formation of rings suggests that Snf7 filaments grow with a
preferred curvature and the formation of spiral suggests that
Snf7 filaments can depart from this preferred curvature with
some flexibility.

The AFM images also revealed that filaments could split within
the disk assemblies (Figure 2I): thick filaments that appeared as
concentric circles are actually interconnected by thinner fila-
ments (Figure 2J, arrows). This is consistent with the spiral struc-
ture seen by EM: double-stranded parts are combined with sin-
gle-stranded connections (Figure 2H). The thinner filaments had
an average thickness of 4.9 nm ± 1.5 nm (n = 25), probably cor-
responding to single strands, whereas thicker ones had a thick-
ness of 10.6 ± 1.2 nm (n = 25), consistent with double-strands.
The AFM and EM analysis suggested a structure of the Snf7 as-
semblies, where a single spiral filament interacted laterally with
itself to form double-stranded filaments. Occasionally, spirals
were directly observed by AFM (Figure 2I, subpanel 6).

These observations prompted the question of how Snf7
patches were formed. Patch nucleation could, for instance, start
from a single closed ring, like those seen by EM. It is conceivable
that such rings could be prone to break open, thus freeing fila-
ment tips that could further grow into a spiral. How would then
this initial spiral transform into a patch? A possible scenario
consists of a two-step growth mechanism (Figure 3A). First,
new spirals are nucleated in the vicinity of existing spirals
(termed below spiral nucleation). Rupture of filaments would
separate the newly formed spirals from the initial spiral. Second,
these spirals would grow independently through the addition of
monomers at their filament tips. This scenario accounts for the
observed growth dynamics of Snf7 patches: the constant speed
of the radial growth of the patches implies that the density of
growing filament tips at their rim stays constant. The formation
of new spirals generates new tips, maintaining a constant density
of growing tips.

To explore quantitatively the implications of such scenario, we
developed a mathematical description of the dynamics of sur-
face coverage by growing Snf7 spirals (Figure 3B and Figure S3).
In the model, Snf7 spirals are represented by hard disks depos-
ited on a surface representing a small (micron-sized) piece of
membrane. As initial conditions, a few disks with radius
were present, corresponding to events of initial patch nucleation.
Patch nucleation is then neglected (set to zero) during the rest
of the dynamics. New disks are thus generated only by spiral
nucleation.

Disks growth corresponds to an area gain w per unit time
(in nm².s⁻¹), as expected if the Snf7 filaments are elongating
from their tips at a constant speed w/b (in nm.s⁻¹, where b =
17 nm is the distance between Snf7 filaments in a single spiral).
As the Snf7 spiral grows, its perimeter increases, offering an
increasing number of potential spiral nucleation sites. We model
this by stochastic nucleation of new disks with radius
at a rate
, where
 denotes the total perimeter of the existing disks,
and λ a constant spiral nucleation rate (expressed in number of
nucleation events per second per micrometer). In the model,
nucleation is prevented if the new disk location is already occu-
pied by an existing disk. In addition, both nucleation and
polymerization stop when the surface is completely covered
with disks (Figure 3C). Solving the model in a mean-field approx-
imation using the value obtained experimentally for r₀ gives a
final distribution of disk sizes with one unknown parameter
(w/λ) (Supplemental Information, Supplemental Mathematical
Modeling part 1). Fitting (w/λ) = 9.8 ± 1.5x10⁻³ µm³, we find
the distribution in excellent agreement with the experimental
size distribution (Figure 3D).

We then tested experimentally whether three key features of
our theoretical model are indeed fulfilled during the generation
of Snf7 assemblies. First, both the existence of initial single rings
and the growth into spirals imply that Snf7 filaments have a
preferred high curvature. Since the average radius of rings is in
the range 25–30 nm, this might correspond to the preferred cur-
vature. Second, the proposal of secondary nucleation of spirals
implies that new spirals can form from existing ones. Third, poly-
merization arrest should be correlated with contacts between
neighboring disks. Finally, we also sought to determine indepen-
dently the parameters w and λ, whose ratio (w/λ) had been esti-
mated from the fit of the size distribution of spirals (Figure 3D).
To address all this, we studied the molecular dynamics of spiral
growth by total internal reflection fluorescence microscopy
(TIRFM) and by dynamic high-speed AFM (HS-AFM) imaging
(Casuso et al., 2012).

Snf7 Filament Dynamics Uncovers the Intrinsic Filament
Curvature and the Mode of Nucleation

To characterize the initial events leading to patch formation, we
first studied the early steps of Snf7 patch nucleation by TIRFM,
which allowed us to quantify the approximate number of Snf7
molecules in diffraction limited spots from their fluorescence in-
tensity (see Experimental Procedures for quantification). Patch
nucleation started with the appearance of a fluorescent diffra-
tion limited spot (nucleus, Figure 4A) containing 50 ± 20 mono-
mers (n = 9). At a Snf7 concentration of 300 nM, the intensity
of the nuclei remained constant for several minutes until these
nuclei started to grow (Figure 4A). Under these conditions,
the number of nuclei is very low (Figure 1C). To increase the
number of nuclei and to obtain more robust statistics, we nucleated Snf7
assemblies by adding 1 µM ESCRT-II and 1 µM Vps20 to a 75 nM
Snf7 solution. Under these conditions, many Snf7 nuclei

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(Q) TEM images of Snf7 rings, single (upper row) and double (lower row) stranded.
(H) Top: TEM image of a single Snf7 spiral. The Snf7 filament is underlined in green (resp. red) when double stranded (resp. single stranded). Bottom: color code of the filament path from the most inner turn (red) to the most outer turn (purple). See also Figure S2D.
(I) AFM images of connections between filaments: 1 to 3, split filaments connecting two spirals – 4 and 5, filament split within a spiral – 6, a spiral filament.
(J) High resolution AFM topographic image of Snf7 filament splitting and branching within a single Snf7 spiral.
See also Figure S2.
appeared on the membrane surface (Figure 4B and Figure S4A) and remained stable for several tens of minutes, consistent with our observations with Snf7 alone (see Figure 4A).

The Snf7 nuclei have an average number of molecules of 60 ± 46 (mean ± SD; n = 1856, Figure 4C). If arranged in a closed ring, it would generate a circle of about 30 nm radius, considering a distance of 3.2 nm between Snf7 monomers in the filaments (Shen et al., 2014) (radius = perimeter / (2π) = (60 × 3.2)/2 × 3.14) ~32 nm. This radius corresponds to those observed in rings seen by EM (see Figure 2G), implying that the arrested nuclei observed by TIRF could be closed rings. This supports that Snf7 patch nucleation starts by the appearance of a single, highly curved Snf7 ring that would break to form a spiral. To further confirm this hypothesis, we further showed through photobleaching experiments that breakage of the nuclei induces patch formation (Figures S4B and S4C).

The formation of highly curved nucleation rings suggests that Snf7 filaments have a preferred radius of curvature in the 25–30 nm range. Indeed, we measured radii of a 27 nm radius by EM and estimated a 32 nm radius by TIRF, in line with other studies (21 nm (Shen et al., 2014) and 32 nm (Henne et al., 2012)). In spirals, this preference should not be satisfied, as outer turns are under-curved (123 nm radius, Figure 2C) and inner turns are over-curved (18 nm radius, Figure 2D). These forced suboptimal radii of curvature may induce significant mechanical stresses in the Snf7 filaments, which might in turn underlie Snf7 biological function in membrane deformation.

To explore the existence of such internal stresses, we used the HS-AFM tip as a nanodissector (Scheuring et al., 2003), briefly applying strong forces to partially break a densely packed array of Snf7 spirals. After breakage, 8 large disks were transformed into 29 small circles (Figure 4D), with a radius of 17 ± 5 nm (see size distribution Figure S4D), This is consistent with a scenario where Snf7 filaments were excised by rupture from the outer circles and curled back to a radius closer to their preferred radius. The nanodissector experiment indicated that even if polymerized at low radius of curvature, Snf7 filaments kept their ability to curl into smaller rings.

The radius of these broken filaments (17 nm) is smaller than the one of the initial ring (~27 nm), but very close to the size of the inner turns in large spirals (18 nm; see Figure 2D). This could be consistent with a preferred radius of curvature of 17 nm. Alternatively, the nanodissector-induced rings experienced the lateral pressure of the neighbors, forcing them to a smaller radius. To discriminate these two hypotheses, we studied the nucleation of new spirals from pre-existing Snf7 assemblies at molecular resolution by HS-AFM. We focused on areas where Snf7 spirals were already packed, but free membrane was still available (Figures 4E and F; Movies S5 and S6). Our image sequences showed that newly formed spirals were mainly initiated from filaments protruding from pre-existing Snf7 spirals. After breakage, 8 large disks were transformed into 29 small circles (Figure 4D), with a radius of 17 ± 5 nm (see size distribution Figure S4D). This is consistent with a scenario where Snf7 filaments were excised by rupture from the outer circles and curled back to a radius closer to their preferred radius. The nanodissector experiment indicated that even if polymerized at low radius of curvature, Snf7 filaments kept their ability to curl into smaller rings.

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Strikingly, when increasing from 2 to 3 concentric circles, the radius of the innermost circle was reduced from 22 nm to 14 nm (Figure 4G). These observations strongly support our hypothesis that the preferred radius of curvature of Snf7 oligomers is about 25 nm, but that lateral pressure can induce higher curvatures.

The radial growth of free spirals was initially rapid, but then slowed down (Figure 4I), roughly following a dependence on...
the square root of time as expected from our model. From these data, we measured an area growth rate \( w \) of 80 ± 36 \( \text{nm}^2 \cdot \text{s}^{-1} \) (\( n = 5 \); see Experimental Procedures for quantification) at 1 \( \mu \text{M} \). We estimated from \( w \) a growth rate of approximately 3 subunits.s\(^{-1}\).\(\mu\text{M}^{-1}\) (Supplemental Information, Supplemental Mathematical Modeling). The estimated filament growth rate is in the range of other filament rates: actin and tubulin are in the range of 5–10 subunits.s\(^{-1}\).\(\mu\text{M}^{-1}\). Using our previous estimate \( (w/\lambda) = 9.8 \times 10^{-3} \text{ \mu m}^2 / \text{s} \) and this experimental value for \( w \), we obtained the spiral nucleation rate as \( \lambda = 8.2 \times 10^{-3} \text{ spirals.\mu m}^{-1}.\text{s}^{-1} \) at [Snf7] = 1 \( \mu \text{M} \). This secondary spiral nucleation rate is 500 times larger than the initial patch nucleation rate (Supplemental Information, Supplemental Mathematical Modeling). It validated our assumption that the patch nucleation was negligible in our theoretical model.

In summary, these observations indicated (1) a preferred high curvature of the Snf7 filaments and (2) the mechanism of spiral nucleation from existing spirals. We then set up to study the third feature of our model, whether lateral contacts between spirals can inhibit their growth.

**Polymerization of Snf7 Filaments Induces Compression of the Spirals**

AFM images of packed arrays of spirals showed that filaments at the contact zone between spirals were flattened, resulting in spirals acquiring a polygonal shape that was more pronounced for longer incubation times (Figure 5A). Moreover, the central area of some of these polygons was pushed toward the substrate, as seen in the height profile of AFM images (Figure 5B). Also, the centers of these spirals were often found to be stiffer as seen in AFM mechanical maps (Figure 5B). We reasoned that this deformation reflected lateral compression of the spirals as the membrane became covered with Snf7. To study the correlation between polymerization rate and lateral compression of the Snf7 assemblies, we ought to measure them simultaneously.

We started by reconstituting Snf7 polymerization on GUVs adhered to a glass surface, to avoid any displacement during time-lapse imaging (Figure S5A) and followed the increase of Snf7 fluorescence on the vesicle with time. In supported bilayers, which have smaller lipid mobility than free bilayers, diffusion of Snf7 assemblies is very limited. In contrast, Snf7 assemblies had a higher diffusion on GUVs explaining that fluorescence showed a homogeneous distribution (Figure 5C).

The dynamics of saturation of Snf7 polymerization over the entire GUV (Figure 5C, bottom) was similar to the dynamics of coverage at a single point in the supported bilayer experiments (Figure S5B): after an approximately exponential increase of the fluorescence, the dynamics of coverage saturated through a progressive slow down phase. If polymerization rate is independent of lateral compression, as it is in our theoretical model, an abrupt arrest of growth is expected (Figure 5C). Conversely, a polymerization rate dependent on lateral compression would cause a progressive slow down until reaching saturation.

As an indication of lateral compression, we noticed that coated GUVs underwent dramatic morphological changes upon several hours of Snf7 polymerization: GUVs were not spherical anymore and instead showed extreme irregular shapes, similar to rigid punched table tennis balls (Figure 5D).

Upon aspiration into a micropipette, they deformed plastically (Figure 5E and Figure S5D), showing that the Snf7 solidifies the membrane. These observations are consistent with a scenario where a rigid Snf7 coat generates pressure on the GUVs, stretching their membrane.

We reasoned that the accumulation of lateral compression in the Snf7 coat would stretch the underlying membrane, increasing its tension (Figure 5F). In order to follow the dynamics of accumulation of lateral compression within the Snf7 layer, we directly measured membrane tension generation during Snf7 polymerization. Using optical tweezers, we pulled a thin tether from a GUV held in an aspiration pipette, that allows to monitor membrane tension through the measurement of the force \( F \) exerted on the optical tweezers’ bead (Cuvelier et al., 2005) (Figure 5G). We then flowed a 500 nM solution of Snf7 using an injection pipette, which triggered protein assembly onto the membrane (Figure 5H). Monitoring Snf7 fluorescence, we found that its membrane binding dynamics was identical to that measured previously, reaching the saturation value after 10–20 min (Figure 5I, top). Concomitant with this saturation, we observed an increase in the force exerted by the tube, indicating an increase of membrane tension (Figure 5I, bottom). These data show that Snf7 polymerization occurs at a slower rate as compression increases within the Snf7 layer, suggesting a coupling between polymerization and compression. The dependence of the polymerization energy \( \mu \) (polymerization energy per unit surface) on the changes in tube force is captured by the expression (Supplemental Information, Supplemental Mathematical Modeling part 4.3):

\[
\mu = \frac{F_f^2 - F_i^2}{8 \pi^2 k}
\]

where \( F_f = 8 \text{ pN} \) and \( F_i = 35 \text{ pN} \) are the tube forces before and after Snf7 polymerization and \( k = 4.8 \times 10^{-20} \text{ J} \), is the membrane bending rigidity (Figure S5E). These values yield \( \mu = 3.1 \times 10^{-4} \text{ J.m}^{-2} \), which is a fairly high polymerization energy, twice that of clathrin (Saleem et al., 2015), which by itself is able to cause membrane deformation during endocytosis. This indicates that the polymerization force of Snf7 can plausibly cause membrane deformation.

These data are therefore compatible with a scenario where Snf7 spirals act as 2D springs that load themselves through polymerization. Constrained by their neighbors, Snf7 spirals would deform significantly during polymerization, from a disk-like to a polygonal shape, until deformation and polymerization force are balanced. To test quantitatively whether the polymerization force is sufficient to deform Snf7 spirals, we established a theoretical elastic model of spiral compression.

In the model, the Snf7 layer is approximated to a hexagonal lattice of individual spirals. Each spiral is composed of a collection of concentric filaments spaced by a distance \( b = 17 \text{ nm} \) up to a typical radius \( R = 130 \text{ nm} \) similar to the experimental data shown in Figure 2C. The filaments are circular in the absence of lateral compression, but may deform into hexagons with rounded vertices, to accommodate tighter packing, as illustrated in Figures 5A and 5F. The amount of energy required to deform circular filaments into more polygonal shapes depend on the
Figure 4. Nucleation and Growth of Snf7 Spirals on Supported Membranes

(A) Left: TIRF microscopy kymographs of the nucleation of single Snf7 patches (green) at [Snf7] = 300 nM. Arrows indicate single ring to multiple spirals transition as postulated from the interpretation of these observations (right).

(B) TIRF microscopy image of Snf7-Alexa488 dots (green) nucleated by ESCRT-II, [Snf7] = 75 nM, [Vps20] = 1 μM, [ESCRT-II] = 1 μM. Inset: zoom on 4 diffraction-limited spots (scale bar, 2 μm).

(C) Histogram of the estimated number of Snf7 molecules within the dots nucleated by ESCRT-II (n = 1856).

(D) HS-AFM nanodissection experiment (see text) of Snf7 spirals. 2 cycles of high AFM force were applied, between 0 s and 10 s, and between 10 s and 20 s.

(E) HS-AFM time-lapse sequence showing the apparition of a new Snf7 spiral from pre-existing ones. Arrowheads show: filament protruding from a spiral (t = 8.5 s), filament curling from its tip (t = 17.0 s), and forming a small spiral (t = 37.4 s), growth of a second turn in the spiral (t = 152.2 s) and filament rearrangements (t = 164.9 s).

(legend continued on next page)
stiffness of the Snf7 filaments, which is characterized by its persistence length \( \ell_p \). We estimated this persistence length from the amplitude of the thermal fluctuations of isolated Snf7 filaments on a supported bilayer observed by HS-AFM. We obtained \( \ell_p = 260 \text{ nm} \) (Supplemental Information, Supplemental Mathematical Modeling part 2). This is in order-of-magnitude agreement with an estimate from a numerical model of Snf7 flexion (\( \ell_p = 800 \text{ nm} \)) (Shen et al., 2014). This value is higher than for DNA (\( \ell_p = 50 \text{ nm} \)), but smaller than for cytoskeletal filaments (\( \ell_p = 15 \mu\text{m} \) for actin and \( \ell_p = 6 \text{ nm} \) for microtubules) (Howard, 2001), implying that they are intrinsically soft enough to be deformed by moderate forces.

In our theoretical 2D spring model, spirals become significantly deformed when \( \mu \) exceeds a threshold surface energy \( \mu^* = 1/(2 - \pi/\sqrt{3})(k_B T \ell_p^2/R^2) \log(R/b) \) (where \( k_B \) is the Boltzmann constant and \( T \) the temperature, see Supplemental Information, Supplemental Mathematical Modeling part 3). Our experimental estimates for \( R \approx 130 \text{ nm} \), \( b = 17 \text{ nm} \) and \( \ell_p = 260 \text{ nm} \) implied \( \mu^* = 4 \times 10^{-5} \text{ J}\cdot\text{m}^{-2} \), 8 times smaller than the measured experimental \( \mu \). It indicates that the Snf7 polymerization energy, even if underestimated, is sufficient to induce strong deformations. Our data show that the Snf7 spirals can deform as spiral springs, and can self-load through a mechanism where deformation is mostly generated by growth of a filament.

**Snf7 Spirals Expansion Leads to Membrane Deformation**

Having established that the Snf7 spirals can deform to store significant elastic energy, we wondered if they could release this energy to deform the membrane. After several hours of Snf7 incubation, holes (called “pores” in the following) spontaneously appeared in a few GUVs, releasing membrane tension. Surprisingly, instead of bursting, the GUV membrane shrank from the rim of the pore toward the opposite side of the GUV (Movie S7). Occasionally, this process stopped before the vesicle had fully collapsed, and stable vesicles with open pores were observed (Figure 6A; Movie S8). In this case, a stronger fluorescence signal is seen at the rim.

To understand the stronger signal of the membrane marker at the rim, we imaged these opened GUVs by thin-section EM: the membrane at the rim of the pore was rolled toward the interior of the vesicle (Figure 6B). Occasionally, this process stopped before the vesicle had fully collapsed, and stable vesicles with open pores were observed (Figure 6A; Movie S8). In this case, a stronger fluorescence signal is seen at the rim.

We hypothesized that curling could be driven by the expansion of the previously compressed Snf7 layer following pore formation. To quantitatively study the plausibility of such scenario, we used our theoretical model to compute the ratio of the surface occupied by a compressed spiral (\( A_{\text{compressed}} \)) to that of a relaxed state (\( A_{\text{relaxed}} \)) (see Supplemental Information, Supplemental Mathematical Modeling part 3) as

\[
\frac{A_{\text{compressed}}}{A_{\text{relaxed}}} = \frac{\pi}{2\sqrt{3}} + \left(1 - \frac{\pi}{2\sqrt{3}}\right) \left[\frac{\mu^*}{\mu}\right]^2 = 94\%
\]

The value of this ratio implies that, during stress release, the Snf7 layer would expand by 6% relative to the underlying, almost inextensible lipid bilayer (Figure 6C). As a result, the membrane of the GUV would curl inward (Figure 6D), consistently with our observations in fluorescence and electron microscopy. The preferred curvature of the curl can be estimated as \( \rho_c = (d/2)(A_{\text{relaxed}} + A_{\text{compressed}})/(A_{\text{relaxed}} - A_{\text{compressed}}) \), where \( 2d \) is the total thickness of the lipid-Snf7 sandwich. Using AFM to measure the thickness of the Snf7 coated membranes, we estimated \( 2d = 9 \text{ nm} \) (5 nm for the membrane plus 4 nm for the Snf7 coat). With this value, we can estimate \( \rho_c = 37 \text{ nm} \). Considering previous studies (Callan-Jones et al., 2012), \( \rho_c \) corresponds to the curvature of the innermost roll observed (Figure 6D). Experimentally, we find a mean radius of \( \rho_c = 39 \pm 6 \text{ nm} \) \((n = 9)\). Therefore curling in the opened vesicles can be explained by the expansion of the Snf7 spiral springs. During this expansion, the spirals release their compression energy accumulated during polymerization.

We wondered how much energy was stored in a single spiral, as compared to the energy required for budding of a vesicle. Our elastic model implies that the lateral compression of a single Snf7 spiral corresponds to the accumulation of an elastic energy \( \Delta E = 170 k_B T = 7.0 \times 10^{-18} \text{ J} \), which is bigger than the bending energy \( 4 \pi k = 160 k_B T \) required to form a spherical membrane bud (see Supplemental Information, Supplemental Mathematical Modeling). Thus, a single spiral can accumulate enough elastic energy to form a spherical bud when released.

In summary, we show here that Snf7 filaments display the ability to act as spiral springs that load through polymerization. The release of the compression stress accumulated during the deformation of the Snf7 spiral is sufficient to drive membrane deformation.

**DISCUSSION**

In this study, we first showed that lipid membranes trigger the formation of wild-type Snf7 assemblies at their surface through a process of nucleation-growth. The patch nucleation rate is low (less than 1 seed,\( \mu\text{m}^{-2}\cdot\text{hour}^{-1} \)) which explains the necessity of a polymerization-activated mutant to observe the same assemblies by EM in previous studies (Henne et al., 2012). We found that the circular arrays formed by Snf7 on these membranes are spirals made of a single filament looping and interacting onto itself. This confirms that the interaction with membrane in vitro retains the...
spiral structure recently observed in solution (Shen et al., 2014) or in vivo (Cashikar et al., 2014; Hanson et al., 2008). We also find that Snf7 filaments can bundle into double-stranded filaments, probably through parallel lateral interactions.

In our assay, spirals become tightly packed into polygonal lattice at the surface of the supported bilayers. The packing of these spirals is correlated with the increase of lateral compression within the Snf7 coat. These data implies that ESCRT-III spirals, because of the relatively high flexibility of the Snf7 filament can be deformed by lateral compression. Moreover, we show that the expansion of compressed spirals can lead to membrane deformation if confinement is released (Figure 6). These observations imply that Snf7 can work as a two-dimensional spring, being able to compress and expand. In the following, we discuss how this spring-like activity highlighted by our study is relevant for the in vivo situation.

In vivo, it is unlikely that a densely packed array of ESCRT-III spirals is present at the surface of membranes, which may question the physiological relevance of the spiral compression observed in large patches of ESCRT-III. But the confinement required for such lateral compression might come from other membrane proteins, which may provide walls into which single spirals could be confined. In the membrane of MVBs, Lamp1 and 2 are particularly enriched (Bissig and Gruenberg, 2014) and may provide a scaffold onto which ESCRT-III spirals could be compressed. Of course, compression being isotropic...
in this in vivo case, compressed spirals would stay circular, instead of polygonal.

But another source of lateral compression is intrinsic to the spiral structure, and is present in single Snf7 spiral even in the absence of any external confining structures. We show that the filaments curl spontaneously at 20–30 nm, implying that if they grow at a different radius, they are under mechanical stress. Indeed, when spirals are broken, all the pieces of filaments further curl to a smaller radius. Thus, in the spiral structure, filaments with a radius larger than 25 nm are stretched, compressing the inner filaments. Accordingly, we find that even for non-laterally constrained spirals, the inner turn of the spirals tightens when the number of turns in the spirals goes above three.

Whatever the source of compression is, our observations show the ability of Snf7 spirals to deform elastically and accumulate potential energy that can be used for membrane deformation. But how would such energy drive membrane budding? It was previously proposed that the polymerization of ESCRT-III could enclose a patch of membrane and then, by reduction of the length of the Snf7 rim, the membrane would be folded into a bud in the middle of the Snf7 polymer. The ESCRT-III rim reduction has been proposed to be mediated by depolymerization of the Snf7 spiral (lasso model, (Saksena et al., 2009)) or by further inward polymerization of the Snf7 spiral (Cashikar et al., 2014). Our data suggests that Snf7 spiral spring could mediate the rim reduction by its elastic compression down to a 14 nm radius. However in this scenario, because the membrane is fluid, it is difficult to picture how the force of the spiral spring would be transmitted to the membrane. We propose that cargoes play an essential role for the force transmission (Figure 7, left): the rim reduction would lead to compaction of enclosed membrane cargoes. Theoretical (Derganc et al., 2013) and experimental studies (Stachowiak et al., 2012) indicate that highly dense cargoes could mediate budding by asymmetric crowding. This is consistent with a recent in vivo study (Mageswaran et al., 2015) where ILVs budding was critically dependent on accumulation of cargoes within ESCRT-III assemblies.

Another possibility is that the out-of-plane buckling of the spring itself might drive the invagination of the membrane (Figure 7, right) (Lenz et al., 2009), which could explain the formation...
of membrane tubules by overexpression of the human homolog of Snf7 (Hanson et al., 2008). This model implies that the flat membrane could be a metastable state of this elastic system: the stored elastic energy could be suddenly released upon external activation. Our observation that the spirals adopt a curved inverted dome shape (Figure 5B) is consistent with this model, and an estimate of the energy stored in one spiral further confirms that this energy is sufficient to bud the membrane into a sphere.

The spring-like properties of Snf7 filaments also nourish our understanding of ESCRT-III role in membrane fission. Because of their high flexibility, Snf7 filaments can grow at radii different from their preferred radius of curvature if steric or mechanical constraints force them to do so. This feature explains how Snf7 filaments could adapt to the wide range of radii observed in the various ESCRT-III mediated fission reactions: from microns in abscission and hundreds of nanometers in virus budding, down to tens of nanometers in ILV formation and membrane repair (Jimenez et al., 2014). However, the smallest size of the inner turn is on average 18 nm radius, which is far from the 1.4 nm observed with dynamin to finalize fission (Sundborger et al., 2014). This raises questions regarding the mechanism necessary to provide fission and pore closure and supports the role of other ESCRT-III proteins and lipids in these reactions.

**EXPERIMENTAL PROCEDURES**

**Protein Purification and Labeling**

Snf7 (Addgene plasmid no. 21492), Escrt-II (Addgene plasmid no. 17633) and Vps20 (Addgene plasmid no. 21490) were purified as previously described (Herrero et al., 2004; Wollert et al., 2009). Snf7 stock solution was 2.5 mM in 20 mM HEPES, 100 mM NaCl (pH 8). Snf7 was labeled either with TFP-Alexa-488 (Life technologie product no. A37563) or with NHS-Atto 647N (Atto-tec product no. AD 647N-3). Escrt-II (20 mM stock solution in 50 mM Tris, 150 mM NaCl, 5 mM i-mercaptoethanol [pH 7.5]) and Vps20 (10 μM stock solution in 20 mM HEPES 100 mM NaCl [pH 7.5]) were kept unlabeled.

**Giant Unilamellar Vesicles and Large Unilamellar Vesicles Preparation**

GUVs were prepared by electroformation using DOPC and DOPE mixtures, purchased from Avanti Polar Lipids (Alabaster, USA). When necessary, 0.1% fluorescent lipids (1,2-dioleoyl-sn-glycero-3-phosphoethanolamine-N-(lissamine rhodamine B sulfonyl)) (Rhodamine-PE) were added.

LVs were prepared by evaporating in a round-bottom glass tube, a volume of lipid mix (DOPC: DOPE, 6:4, mol:mol) containing 1 mg of total lipids. After addition of 200 μl of buffer, the tube was vortexed and freeze-thaw 3 times. This solution is kept at −20°C until use. Unless otherwise noted, the buffer used for all experiments is composed of 20 mM Tris HCl (pH 6.8), 200 mM NaCl, 1 mM MgCl2.

**Optical Microscopy of Membrane Assays**

For confocal and TIRF imaging, a coverslip is cleaned with water and ethanol, and then plasma-cleaned for 2 min (PDC-32G, Harrick Plasma, NY, USA). The coverslip is assembled to a flow chamber (sticky-Slide VI 0.4, Ibidi, Munich, Germany), with one entry connected to a syringe pump (Aladdin, World Precision Instruments, Sarasota, FL, USA), and the other left open for sequential introduction of other solutions. The flow chamber is initially filled by 200 μl of buffer. 5 μl of GUVs are flushed in the flow chamber (see Extended Experimental Procedures for methods to get supported membranes or partially adhered vesicles).

Imaging is performed using an inverted microscope assembled by 3i (Intelligent Imaging Innovation, Denver, USA) and Nikon (Eclipse C1, Nikon, Tokyo, Japan). For SDC imaging, a 2-μm-thick volume stack (1 μm above and below the supported membrane) is acquired then rendered to 2D by maximum intensity projection. TIRF Imaging is performed using a motorized Nikon TIRF system. The number of molecules within Snf7 oligomers is estimated by calibrating the microscope with commercially available fluorescent DNA origamis (GATTA-Brightness 9R and 18R, GATTAquant, Braunschweig, Germany) (Figure S6).

**Optical Tweezers Tube Pulling Experiment**

A modified version of a published setup (Morlot et al., 2012) allows simultaneous brightfield imaging, SDC microscopy, and optical tweezing on an inverted Nikon eclipse Ti microscope. A GUV is aspirated within a micropipette connected to a motorized micromanipulator (MP-285, Sutter Instrument, Novato, CA, USA) and a pressure control system (MFCS-VAC – 69 mbar, Fluigent, Villejuif, France) that sets the aspiration pressure ΔP. A membrane nanotube is then pulled out from the GUV through a streptavidin-coated bead (3.05 μm diameter, Spherotec, Lake Forest, IL, USA) held in a fixed optical trap. The optical trap was custom-made with a continuous 5 W 1064 nm fiber laser (ML5-CW-P-TKS-OTS, Manlight, Lannion, France) focused through a 100X 1.3 NA oil immersion objective. The force F exerted on the bead was calculated from the Hooke’s law: F = kΔx, where k is the stiffness of the trap (k = 60 pN·μm−1) and Δx the displacement of the bead from its...
equilibrium position. Snf7 was injected close to the nanotube with a second microinjection system, using a micropipette 

equipped with a Teflon-coated silicon tip.

Electron Microscopy

For negative stain EM observations, LUVs were incubated with Snf7 in suspension, spun down (4 min at 4,000 g), washed and then adsorbed onto glow-discharged Formvar coated EM grids. The samples were negatively stained for 30 s with 2% uranyl acetate before visualization.

Ultra-thin sectioning of Snf7-bound LUVs fixed in epon was performed using a microtome (Leica Ultracut) at a cutting angle of 6°. Sections were put on glow-discharged carbon-coated Formvar grids and imaged with a Tecnai G2 Sphera (FEI) electron microscope.

REFERENCES

Adell, M.A.Y., Vogel, G.F., Pakdel, M., Müller, M., Lindner, H., Hess, M.W., and Teis, D. (2014). Coordinated binding of Vps4 to ESCRT-III drives membrane neck constriction during MVB vesicle formation. J. Cell Biol. 205, 33–49.

Babst, M., Katzmann, D.J., Estepa-Sabal, E.J., Meerloo, T., and Emr, S.D. (2002). Escrt-III: an endosome-associated hetero-oligomeric protein complex required for mvb sorting. Dev. Cell 3, 271–282.

Bissig, C., and Gruenberg, J. (2014). ALIX and the multivesicular endosome: ALIX in Wonderland. Trends Cell Biol. 24, 19–25.

Buchkovich, N.J., Henne, W.M., Tang, S., and Emr, S.D. (2013). Essential N-terminal insertion motif anchors the ESCRT-III filament during MVb vesicle formation. Dev. Cell 27, 201–214.

Callan-Jones, A., Albarran Arriagada, O.E., Massiera, G., Lorman, V., and Abkarian, M. (2012). Red blood cell membrane dynamics during malaria parasite egress. Biophys. J. 103, 2475–2483.

Ashwell, J., Kao, J., Charn, M., Paul-Gilloteaux, P., Husain, M., Deneau, J.-P., Stahlberg, H., Sturgis, J.N., and Scheuring, S. (2012). Characterization of the motion of membrane proteins using high-speed atomic force microscopy. Nat. Nanotechnol. 7, 525–529.

Cookridge, E.M., and Stevens, T.H. (2010). The yeast vps class E mutants: the beginning of the molecular genetic analysis of multivesicular body biogenesis. Mol. Biol. Cell 21, 4057–4060.

Curie, C., Derényi, I., Bassereau, P., and Nassy, P. (2005). Coalescence of membrane tethers: experiments, theory, and applications. Biophys. J. 88, 2714–2726.

Derganc, J., Antonny, B., and Copic, A. (2013). Membrane bending: the power of protein imbalance. Trends Biochem. Sci. 38, 576–584.

Elia, N., Sougrat, R., Spurlin, T.A., Hurley, J.H., and Lippincott-Schwartz, J. (2011). Dynamics of endosomal sorting complex required for transport (ESCRT) machinery during cytokinesis and its role in abscission. Proc. Natl. Acad. Sci. USA 108, 4846–4851.

Fabrikant, G., Lata, S., Riches, J.D., Briggs, J.A., Weisshorn, W., and Koziol, M.M. (2009). Computational model of membrane fission catalyzed by ESCRT-III. PLoS Comput. Biol. 5, e1000575.

Guizetti, J., Schermelleh, L., Mäntler, J., Maar, S., Poser, I., Leonhardt, H., Müller-Reichert, T., and Gerlich, D.W. (2014). Cortical constriction during abscission involves helices of ESCRT-III-deficient filaments. Science 344, 1616–1620.

Hanson, P.I., Roth, R., Lin, Y., and Heuser, J.E. (2008). Plasma membrane deformation by circular arrays of ESCRT-protein filaments. J. Cell Biol. 180, 389–402.

Henne, W.M., Buchkovich, N.J., Zhao, Y., and Emr, S.D. (2012). The endosomal sorting complex ESCRT-II mediates the assembly and architecture of ESCRT-II helices. Cell 151, 356–371.

Hierro, A., Sun, J., Rusnak, A.S., Kim, J., Prag, G., Emr, S.D., and Hurley, J.H. (2004). Structure of the ESCRT-II endosomal trafficking complex. Nature 431, 221–225.

Howard, J. (2001). Mechanics of Motor Proteins and the Cytoskeleton (Sunderland, Massachusetts: Sinauer Associates).

Jimenez, A.J., Maiuri, P., Lafaurie-Janvore, J., Divoux, S., Piel, M., and Perez, F. (2014). ESCRT machinery is required for plasma membrane repair. Science 343, 1247136.

Lata, S., Schoehn, G., Jain, A., Pires, R., Piehler, J., Gottlinger, H.G., and Weisshorn, W. (2008). Helical structures of ESCRT-III are disassembled by VPS4. Science 327, 1354–1357.
Lenz, M., Crow, D.J., and Joanny, J.-F. (2009). Membrane buckling induced by curved filaments. Phys. Rev. Lett. 103, 038101.

Mabrouk, E., Cuvelier, D., Brochard-Wyart, F., Nassoy, P., and Li, M.-H. (2009). Bursting of sensitive polymesomes induced by curling. Proc. Natl. Acad. Sci. USA 106, 7294–7298.

Mageswaran, S.K., Johnson, N.K., Odorizzi, G., and Babst, M. (2015). Constitutively active ESCRT-II suppresses the MVB-sorting phenotype of ESCRT-0 and ESCRT-I mutants. Mol. Biol. Cell 26, 554–568.

McCullough, J., Colf, L.A., and Sundquist, W.I. (2013). Membrane fission reactions of the mammalian ESCRT pathway. Annu. Rev. Biochem. 82, 663–692.

Morlot, S., Galli, V., Klein, M., Chiaruttini, N., Manzi, J., Humbert, F., Dinis, L., Lenz, M., Cappello, G., and Roux, A. (2012). Membrane shape at the edge of the dynamin helix sets location and duration of the fission reaction. Cell 151, 619–629.

Pires, R., Hartlieb, B., Signor, L., Schoehn, G., Lata, S., Roessle, M., Moriscot, C., Popov, S., Hinz, A., Jamin, M., et al. (2009). A crescent-shaped ALIX dimer targets ESCRT-III CHMP4 filaments. Structure 17, 843–856.

Saksena, S., Wahlman, J., Teis, D., Johnson, A.E., and Emr, S.D. (2009). Functional reconstitution of ESCRT-III assembly and disassembly. Cell 136, 97–109.

Saleem, M., Morlot, S., Hohendahl, A., Manzi, J., Lenz, M., and Roux, A. (2015). A balance between membrane elasticity and polymerization energy sets the shape of spherical clathrin coats. Nat. Commun. 6, 6249.

Scheuring, S., Seguin, J., Marco, S., Lévy, D., Robert, B., and Rigaud, J.-L. (2003). Nanodissection and high-resolution imaging of the Rhodopseudomonas viridis photosynthetic core complex in native membranes by AFM. Atomic force microscopy. Proc. Natl. Acad. Sci. USA 100, 1690–1693.

Shen, Q.-T., Schuh, A.L., Zheng, Y., Quinney, K., Wang, L., Hanna, M., Mitchell, J.C., Otegui, M.S., Ahloquist, P., Cui, Q., and Audhya, A. (2014). Structural analysis and modeling reveals new mechanisms governing ESCRT-III spiral filament assembly. J. Cell Biol. 206, 763–777.

Stachowiak, J.C., Schmid, E.M., Ryan, C.J., Ann, H.S., Sasaki, D.Y., Sherman, M.B., Geissler, P.L., Fletcher, D.A., and Hayden, C.C. (2012). Membrane bending by protein-protein crowding. Nat. Cell Biol. 14, 944–949.

Sundborger, A.C., Fang, S., Heymann, J.A., Ray, P., Chappie, J.S., and Hinthaw, J.E. (2014). A dynamin mutant defines a superconstricted prefission state. Cell Rep. 8, 734–742.

Wollert, T., and Hurley, J.H. (2010). Molecular mechanism of multivesicular body biogenesis by ESCRT complexes. Nature 464, 864–869.

Wollert, T., Wunder, C., Lippincott-Schwartz, J., and Hurley, J.H. (2009). Membrane scission by the ESCRT-III complex. Nature 458, 172–177.