

## on releasing tension

Vivienne Baillie Gerritsen

Like life, cells are subject to continuous change. Nothing in the vicinity of a cell remains still – unless death has interrupted its course. And the same goes for the inside of each cell. All sorts of molecules are being shuttled from one part to another, after having been created or on their way to being degraded. The cell membrane is also a very dynamic and supple structure, with molecules wandering through it constantly. One means of transport for shifting molecules around are known as endosomes. Endosomes are formed by the invagination of one part of the cell membrane – like a bubble budding towards the inside of the cell – which is then cleaved and able to float free in the cell’s cytoplasm. Invagination then occurs on the surface of the endosomes themselves to form even smaller bubbles – or vesicles – that are in turn also set free. Like endosomes, these inner vesicles are a means of molecular transport too. It’s a sort of Russian doll experience... The art of invagination *per se* may sound simple but it involves a lot of imagination on behalf of biology. Recently, scientists discovered a protein coined Snf7 that is providing hints as to how endosome invagination may occur, and the subsequent creation of vesicles.



by Zaq Guimarães, 2016

Courtesy of the artist

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All kinds of proteins and protein complexes are involved in the making of endosomes and vesicles. Vesicle formation is driven by what are known as “endosomal sorting complexes required for transport”, or ESCRTs. There are three ESCRTs (I, II and III) which, between them, relay cargo to where invagination will take place, promote invagination and then set the nascent vesicle free by severing it from the endosomal membrane. The ESCRT machinery and its role in budding were first observed in yeast for transporting molecules destined for degradation. However, it is now known that such budding events are similar to those used for HIV release and daughter cell abscission. Snf7 is the most abundant protein in the ESCRT-III complex and seems to be

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Snf7 has a highly structured core domain of four alpha-helices, with a C-terminal alpha-helix that folds back onto the core domain. When the C-terminal end unfolds, Snf7 adopts an elongated “open” form that promotes protein-protein interactions; this is believed to be at the heart of membrane budding. What happens is Snf7 monomers can spontaneously fold into a single closed ring – or nucleation ring – which will, at one point, open. The “open form” induces Snf7b polymerisation; concomitantly, Snf7 is bent into a curve which is not “natural”. When a second Snf7 monomer is added, it is also bent into an unnatural curve and so on. As the filament grows, the geometrical shape that eventually emerges is a two-dimensional spiral filament of Snf7 monomers on the cytoplasmic side of the endosome. Moreover, since the curvature of each open Snf7 monomer is unnatural, there is a mechanical stress which is heightened as the spiral filament elongates. And mechanical stress spells locked-up energy – in this case elastic energy.

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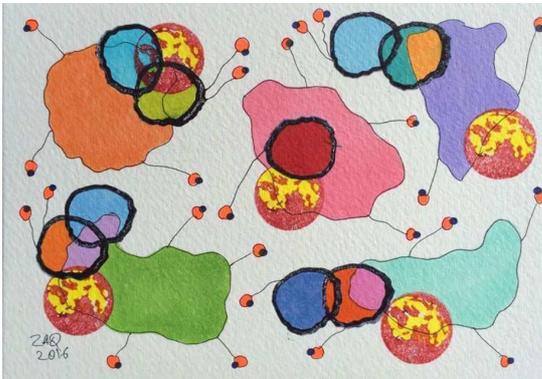
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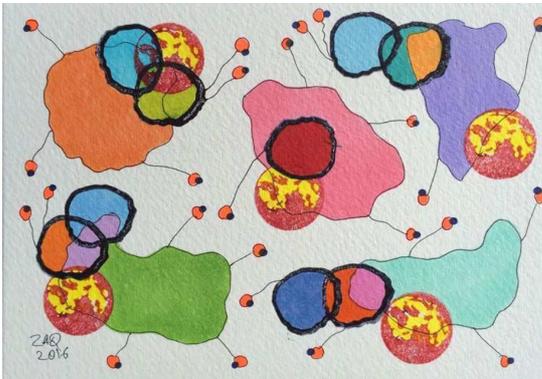
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