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Recent data pointing to a novel function of Cyclin A2 add another component to the complex regulatory network that involves cell cycle regulators and cytoskeletal structures participating in the control of cell movement. As already known, Cyclin A2 is a key regulator of cell division, since it controls both S phase and G2/M transition in association with CDK2 and CDK1, respectively. During S phase, Cyclin A2 regulates the initiation and progression of DNA synthesis, and at the G2/M transition, it plays a critical role in triggering Cyclin B1-CDK1 activation. In mice, it is essential in embryonic cells and in the hematopoietic lineage, yet dispensable in fibroblasts.

Surprisingly, depletion of Cyclin A2 is sufficient to increase cell motility of fibroblasts in 2D assays and cooperates with oncogenic transformation to increase their invasiveness in 3D collagen matrices. Cyclin A2-deficient cells contain a perturbed cytoskeleton, where Actin filaments are cortical and the distribution of focal adhesions is altered. Interestingly, these defects are corrected by a Cyclin A2 mutant unable to activate its cognate kinases, CDK1 and CDK2. This is associated with a downregulation of the RhoA-ROCK pathway and decreased phosphorylation of Cofilin, which is involved in the reorganization of Actin filaments, consecutively leading to an increased cell migration and invasion. Importantly, pharmacological inhibition of ROCK in control fibroblasts leads to an increase in migration velocity similar to that of Cyclin A2-depleted cells.

Cyclin D1 and the CDK inhibitors p21, p27 and p57 had also previously been shown to impinge upon the RhoA/ROCK pathway. Cyclin D1 binds directly to p27 and thereby blocks RhoA activation by inhibiting interaction with its GEFs. Similarly, cytoplasmic p21 has been shown to bind and inhibit ROCK1, which promotes neurite extension by neuroblastoma cells and hippocampal neurons, while p57 was shown to sequester LIMK (Fig. 1). Within this scheme, Cyclin A2 binds directly to RhoA and facilitate its GTP loading by GEFs. This is consistent
with the involvement of this GTPase in early mitosis, since its increased activity at that time leads to cortical retraction and cell rounding via its downstream effector ROCK.14 Moreover, formation of the contractile ring during cytokinesis has also been shown to depend upon RhoA activation in a precise zone at the cell equator.15

Knockdown of Cyclin A2 and inhibition of CDK2 prevent cells from forming stable attachments of their mitotic spindle to the cell cortex.16 This resulted in the spindles failing to locate to the central position in the cells and undergo dramatic rotation. Moreover, Cyclin A2-CDK2 specifically associated with APC in late G2 phase and phosphorylated it on Ser1360. Mutation of this serine to alanine results in identical off-centered mitotic spindles.17

Thus, this Cyclin A2-CDK2-dependent phosphorylation within the mutation cluster region of APC affects astral microtubule attachment to the cortical surface in mitosis.

Another potential player in this complex ballet appears to be the Golgi apparatus. Recent studies suggest the existence of functional interactions between this organelle and the centrosome, the structure responsible for the nucleation of the mitotic spindle. The Golgi and the centrosome are in close proximity at interphase and this intimacy is transiently lost during mitosis, when the Golgi is fragmented, and the two newly duplicated centrosomes migrate. Golgi-centrosomes interactions may play a central role in cell polarization, since both structures undergo reorientation toward the leading edge of a migrating cell (reviewed in ref. 17).

Proteins of the Golgi apparatus are likely to be instrumental in centrosome organization and positioning, and microtubules nucleated at the centrosome seem to play the same role for the Golgi. CyclinA2 has also been shown to localize to the centrosome in a CDK-independent manner, and through binding of MCM5 and Orc1, prevents the formation of supernumerary centrosomes.18 Since Cdc42 plays a central role in cell polarization, this raises the question of its role in the potential coupling of this process to centrosome duplication and spindle orientation. A partial answer to this question was provided recently. Bray et al. showed that Cdc42 controls spindle orientation to position the apical surface during epithelial morphogenesis, at least in vitro.19

These data highlight thus another aspect of Cyclin A2 function in the control of cytoskeleton dynamics, which places this cyclin at the crossroads of intracellular processes, such as mitosis, and extracellular cues emanating from neighboring cells or from the extracellular matrix (Fig. 2).

If we place these observations within the context of the epithelium, the orientation of the mitotic spindle with respect to the lamina is a major issue, since this has been proposed to be instrumental in the determination of the fate of the two daughter cells. Moreover, in tumors, basal extrusion of a daughter cell could initiate metastasis in some situations.20 Therefore, mutations that couple spindle misorientation with oncogene activation could facilitate tumor progression and metastasis spreading.

Until recently, Cyclin A2 was solely considered as a proliferation marker since, consistent with its known functions during the cell cycle, it is frequently overexpressed in highly proliferative cancers. However, our observations indicate that Cyclin A2 protein levels are significantly lower in an invasive colon carcinoma cell line derived from lymph node metastasis relative to a less-invasive counterpart issued from the primary site. More importantly, the same is observed upon metastasis of colon adenocarcinoma.1 Studies of renal, colorectal carcinoma and prostate cancer, found that proliferative tumors with low levels of Cyclin A2 were more aggressive than those with high Cyclin A2 expression.21-23 Moreover, Wang et al. established that Cyclin A2 levels were inversely correlated with invasiveness of oral squamous cell carcinoma (OSCC) both in vitro and in vivo.24

Altogether, these data indicate that Cyclin A2 downregulation could be an important step in the acquisition of an invasive property by epithelial cells, and thereby call for more studies on the involvement of this cyclin in the morphological changes that occur during metastasis. Along these lines, the epithelial to mesenchymal transition (EMT) appears to be instrumental in pathological situations such as fibrosis, tumor development and metastasis spreading.25-27 In the context of epithelial cancer, EMT provides one mechanism for tumor cells to invade the local tissue and blood vessels, setting the stage for metastatic spread. Therefore, EMT is hypothesized to contribute to tumor progression and clinical evidence suggests that upregulation of EMT regulators in cancer cells correlates with poor patient outcome and tumor aggressiveness.28-30
likely to be triggered by complex networks of signals emanating from the tumor stroma, along with a variety of cytokines and growth factors, such as transforming growth factor β (TGFβ), epidermal growth factor (EGF), fibroblast growth factor (FGF) and hepatocyte growth factor (HGF). Consistent with this, Ras and TGFβ have been shown to cooperate in promoting EMT. During this process, epithelial cells undergo morphological remodeling toward a more elongated and fibroblastic morphology. Among these changes, cells lose cell-to-cell junctions, undergo Actin cytoskeleton rearrangement and acquire enhanced invasive properties. At the molecular level, EMT is characterized by a loss of E-cadherin, and the acquisition of mesenchymal markers, such as Vimentin, Fibronectin or N-cadherin. Transcription factors, such as Snail, Slug, Twist or Zeb-1 and 2, have been implicated in this process. For some cell types, the acquisition of a more motile phenotype during EMT has been attributed to the increased expression and activation of Rho GTPases, such as Rac1, Cdc42 and RhoC.

Accordingly, overexpression of RhoC enhances the ability of melanoma cells to exit the circulatory system and colonize the lungs. Consistent with this, loss of RhoC does not affect tumor development while it leads to a drastic inhibition of metastasis. Nevertheless, RhoC-deficient mice are viable, indicating that this GTPase is dispensable for embryonic and post-natal development. RhoC and RhoA share more than 95% sequence similarity, thus it is not surprising that Cyclin A2 interacts as well with both GTPases. Indeed, when Cyclin A2 is knocked down in epithelial cells, such as normal mouse mammary epithelial cells (NMuMG), they exhibit a strong downregulation of RhoA activity and an increase in RhoC activity (our unpublished observation).

Several reports support a possible role for RhoC in the EMT-related invasion and in metastasis spreading. As mentioned earlier, the breeding of RhoC-deficient mice to the tumor and metastasis-prone MMTV-PyMT strain established the requirement for RhoC in metastasis to the lungs. Consistent with this, increased RhoC expression in adenocarcinoma of pancreas, along with hepatocellular, breast, ovarian, bladder and esophageal cancers, has been observed and correlated with progression and poor prognosis. Moreover, invasion by several cell types, such as breast, colon carcinoma, or prostate cancer cell lines, was shown to be dependent on RhoC expression and activation. With regards to EMT, recent studies have established that increased expression and/or RhoC activation promotes invasion, whereas Rac1 and Cdc42 were generally considered as major players in this process. In these studies, increased RhoC activity is correlated with an increased transcription of the gene. This transcription is induced by upregulation of Twist and miR-10b, or is dependent on the transcription factor Ets-1 in breast cancer and colon carcinoma cell lines, respectively. In conclusion, these data highlight the intricate relationship between Cyclin A2 expression and Rho GTPases activity within the context of cell adhesion and motility. While Cdc42 had previously been shown to be instrumental for anchorage-independent expression of the Cyclin A2 gene in primary mouse embryonic fibroblasts, this cyclin appears now to feed back on the activity of other GTPases, such as RhoA and RhoC. Altogether, these observations indicate that CyclinA2 is a potential novel player in the complex regulation of EMT, moreover, they reinforce the idea that cell cycle regulators are much more than just cell cycle regulators.

**Figure 2.** Cyclin A2 is at the interface between cytoplasmic and extracellular cues. When Cyclin A2 level is high, the two daughter cells orientate their bipolar mitotic spindles according to both their mother footprint and cues emanating from the lamina. If Cyclin A2 is abnormally downregulated, cell division is non-localized, cell spreading is isotropic and cells show increased motility.

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