

5th International ACC Symposium: An Outlook to Current and Future Research on the Biology of Adrenocortical Carcinoma: Diagnostic and Therapeutic Applications

Enzo Lalli^{1,2,3} · Hironobu Sasano⁴

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Abstract Groundbreaking progress has been recently made in elucidating the signaling pathways that are altered in adrenocortical carcinoma (ACC), an endocrine malignancy that still has an unfavorable prognosis, and in understanding its genomic structure. These advances need now to be translated to create cellular and animal models more relevant to human disease in order to develop new and more effective diagnostic procedures and targeted therapies against this deadly malignancy.

Adrenocortical carcinoma (ACC) is a rare endocrine malignancy with a poor prognosis. Surgery could be curative in the great majority of cases with localized disease, but ACC often relapses or is already metastatic at the time of diagnosis. In those cases, therapeutic options are limited to the use of mitotane, an adrenolytic drug whose introduction in ACC treatment dates back to 1960s, that can be associated to systemic chemotherapy [1, 2]. However, therapeutic results have

generally remained poor and a consensus exists about the urgent need for novel drugs that are more efficient to fight against this deadly malignancy. A steadily increasing number of laboratories worldwide is therefore actively engaged in basic and translational research to develop better therapies for ACC. A critical element for those developments consists in a better understanding of the biology of ACC, and in particular of the molecular mechanisms that regulate its growth and spread, through approaches that include the identification of new biomarkers for disease diagnosis and relapse, the generation of new cell and animal models and the preclinical testing of new therapies. The last session (*Closing the gap at the bench: molecules to organisms*) at the recent 5th International Adrenal Cancer Symposium (Ann Arbor MI, October 14–15, 2015) provided a unique opportunity for both basic scientists and clinicians to discuss the latest developments in the field. Here, we will make a brief discussion of the presentations in Ann Arbor trying to put them in the context of current research in the domain of ACC biology.

5th International ACC Symposium Session: Closing the Gap at the Bench: Molecules to Organisms

✉ Enzo Lalli
ninino@ipmc.cnrs.fr

✉ Hironobu Sasano
hsasano@patholo2.med.tohoku.ac.jp

¹ Institut de Pharmacologie Moléculaire et Cellulaire CNRS, 660 route des Lucioles, 06560 Valbonne, France

² NEOGENEX CNRS International Associated Laboratory, Valbonne, France

³ Université de Nice-Sophia Antipolis, Nice, France

⁴ Department of Pathology, Tohoku University School of Medicine, 2-1 Seiryomachi, Aoba-ku Sendai 980-8575, Japan

New Biomarkers for ACC

MicroRNA (miRNA) is a class of evolutionarily conserved, small non-coding eukaryotic RNA molecules which regulate gene expression at the level of translation. They are implicated in virtually every biological process and also have a relevant role in tumorigenesis. Several studies have reported that a distinct microRNA expression signature can differentiate ACC from adrenocortical benign tumors. In addition, the expression levels of specific miRNAs in the tumor are prognostic markers reviewed in [3]. Recent studies have revealed that circulating miRNAs may represent potential biomarkers of malignancy in ACC patients [4–6]. However, further studies on larger cohorts are needed to precisely assess their

diagnostic and prognostic role, in particular to monitor tumor recurrence. A new twist in the field of miRNA is their potential use as therapeutic tools. At the Ann Arbor meeting, Dr. Stan Sidhu (University of Sydney, Australia) presented results of a recently published study from his group showing that miR-7, which has a tumor suppressor role in ACC, can reduce growth of H295R and primary ACC cells xenografts when administered in the form of targeted, clinically safe delivery vesicles (nanocells) [7].

Another promising area of research is the study of circulating tumor cells (CTCs) in ACC patients. CTCs are neoplastic cells originating from either the primary tumor or metastases, which can be detected and isolated in the peripheral blood of cancer patients. CTCs are a reliable tool for prognosis and follow-up in several solid human malignancies [8]. A preliminary study has revealed that CTCs are present in the circulation of ACC patients but not in patients with benign adrenocortical tumors. In addition, a significant decrease in the number of CTCs has been reported following surgery compared to presurgical samples [9]. CTCs isolation may allow to obtain a “liquid biopsy” of ACC, which could be especially useful to monitor the occurrence of relapses and to study the evolution of ACC genomes during disease progression or chemotherapeutic treatments. Other approaches that may provide in the near future minimally invasive access to ACC genetic information for diagnostic and prognostic purposes include the study of cell-free DNA and exosomes in the bloodstream [10], while differential analysis of the urinary steroid metabolome may help to distinguish malignant from benign adrenal tumors [11, 12].

Cellular Models for ACC

Compared to most other malignancies, one serious shortcoming of ACC is the availability of virtually only one differentiated human adrenocortical cell line, the NCI-H295 and its derivatives reviewed in [13]. Another human cell line (SW-13), which has been used in several studies as a model for ACC, lacks expression of adrenocortical markers and does not produce steroids [13, 14]. Therefore, its usefulness as an ACC model cell line is highly questionable. As an additional tool, the SJ-ACC3 human adrenocortical cell line has been established from a pediatric tumor. This cell line only grows as subcutaneous xenografts in immunodeficient mice [15]. More recently, another pediatric cell line growing as xenografts has been established in the same laboratory from a different patient (G. Zambetti, personal communication). At the Ann Arbor meeting, Dr. Constanze Hantel (University of Munich, Germany) described a new human adrenocortical xenograft model (MUC-1), also growing as a cell line, established in her laboratory from a neck metastasis of an adult ACC. These new ACC cell lines will be very important in the near

future to expand and diversify the studies on adrenocortical cells beyond the NCI-H295 model.

Animal Models for ACC

A few animal models have been reported for ACC. Some of them exploit injection of adrenocortical cell lines (most commonly NCI-H295) in immunodeficient mice, while others are derived from genetically modified animals.

- Xenograft models: the first report of a subcutaneous xenograft animal model for ACC dates back to 2000 [16]. Since that time, numerous studies have used this model to monitor its sensitivity to different drug treatments reviewed in [17]. A major drawback of this model is the absence of the capacity to metastasize, which is in sharp contrast to human ACC. This renders the subcutaneous xenograft model not convenient for studies aimed at investigation of the cellular and molecular mechanisms of metastasis formation in ACC and to test therapies for metastatic disease. An orthotopic model of ACC consisting in normal bovine adrenocortical cells implanted under the renal capsule of immunodeficient mice has also been described. Genetic modification of those cells with an oncogenic version of Ras and a dominant-negative p53 induced malignant tumor formation with local dissemination [18]. Possibilities to boost the metastatic capacities of adrenocortical cells in future studies may include the use of new administration routes (intravenous, etc.) and the modulation of the expression of endogenous factors (adhesion molecules, cytokines, etc.) favoring metastasis formation.
- Genetic models: genetic models of adrenocortical neoplasia have been described, but in general, they exhibit only limited malignant character.
 - a) Gonadectomy can induce neoplasia in the adrenal cortex of the domestic ferret and in some strains of mice, as well as in some lines of genetically modified mice in a non-neoplasia-prone background reviewed in [19]. Transgenic mice harboring multiple copies of the *Sf-1* (*Nr5a1*) gene also develop adrenocortical neoplasia in the absence of gonadectomy [20]. The common features of those models are proliferation of neoplastic cells in a subcapsular location in the adrenal expressing gonadal markers, most probably derived from undifferentiated adrenogonadal progenitors [21] and little tendency to spread beyond the locoregional location.
 - b) Other genetic models of adrenocortical neoplasia are based on genes and signaling pathways that are altered in ACC.

Wnt/ β -Catenin

A seminal study by Tissier et al. first showed that activation of the Wnt/ β -catenin signaling pathway is a hallmark of both benign and malignant adrenocortical tumors [22]. Further genomic studies have confirmed that gene mutations that have the consequence to activate this pathway are encountered at high frequency both in adult [23] and children [24] adrenocortical tumors. Remarkably, the gene most frequently mutated or deleted in adult ACC is *ZNRF3*, encoding a E3 ubiquitin-protein ligase that acts as a negative regulator of the Wnt signaling pathway by mediating the ubiquitination and degradation of the Wnt receptor complex components Frizzled and LRP6. Studies in various mouse models have shown that β -catenin activation in the adrenal has relatively weak oncogenic effects, even when combined with overexpression of the IGF2 growth factor [25–27], suggesting the requirement of further genetic hits for tumor development. Based on the results in human ACC, it will be interesting to study the effect of mouse *Znrf3* inactivation on adrenocortical tumor development. However, β -catenin antagonism either by drugs [28] or by RNA interference [29, 30] is able to significantly decrease ACC cell proliferation in vitro and their growth as xenografts. Those data suggest that counteracting β -catenin activity in ACC could be a promising therapeutic strategy to be assessed by future studies. However, it will very important to tailor the choice of Wnt pathway inhibitors, targeting with drugs either upstream components (porcupine and tankyrase inhibitors) or downstream effectors (CBP, p300) of the pathway [31], according to the specific genetic alterations present in the individual patient with ACC.

TP53

Germline mutations in the tumor suppressor *TP53* gene are associated with Li-Fraumeni syndrome, a multiple tumor syndrome one of whose hallmarks is the high frequency of adrenocortical tumors [32]. On the other side, adrenocortical tumors in children are often associated with germline *TP53* mutations in the absence of other neoplasms [33, 34]. A peculiar epidemiological situation is present in southern Brazil, where the R337H *TP53* mutation is present at a very high frequency (0.3 %) in the population and is associated with adrenocortical tumors in children with low penetrance [35, 36]. Conversely, *TP53* germline mutations are uncommonly found in patients diagnosed with ACC as adults [37, 38] but somatic alterations in the p53/Rb pathway were identified in about 30 % of adult ACC [23]. This pathway then appears to represent a major driver of oncogenic transformation in ACC, similarly to many other malignancies. A new transgenic mouse model was presented at the Ann Arbor meeting by Dr. Pierre Val (University of Clermont, Clermont-Ferrand, France) where expression the SV40 T antigen, which binds to and inactivates both the p53

and the Rb tumor suppressor proteins, was targeted to the adrenal cortex under the control of the *Akr1b7* promoter ([39] and A.M. Lefrançois-Martinez, personal communication). Adrenal T transgenic mice develop adrenocortical carcinomas that metastasize with high frequency and will provide a very useful model to test novel therapies against metastatic ACC. Studies are also in progress in Dr. Gerard Zambetti's laboratory (St. Jude Children's Research Hospital, Memphis TN, USA) to characterize the phenotype of mice bearing the targeted R334H *Tp53* mutation, homologous to human R337H (G. Zambetti, personal communication).

Novel Drugs and Strategies for ACC Treatment

The recent conspicuous advances in our understanding of ACC biology have led to the proposal of a series of new therapies targeting some among the major actors in ACC pathogenesis; see [40–42] for review. However, for the moment, novel drugs have not gone beyond the preclinical stage of investigation or have produced unsatisfactory results when introduced in the clinic. This has been the case for IGF-1R inhibitors, whose trial in ACC patients was based on promising preclinical data [43, 44]. However, a clinical trial with the orally available IGF-1R inhibitor linsitinib (OSI-906) did not show positive effects of the drug on overall or progression-free survival in ACC patients [45]. Several elements can be invoked to account for those disappointing results:

- The trial involved patients with advanced disease who already received intense chemotherapeutic treatment.
- Since a partial response was observed in a small group of patients in that study, the identification of predictive biomarkers may help to identify patients that will benefit from IGF-1R inhibitors therapy.
- Other drugs may need to be associated to IGF-1R inhibitors to completely block downstream signaling pathways regulating proliferation and apoptosis of ACC cells. Another study in fact observed disease stabilization in ACC patients treated with a combination of the IGF-1R antibody cixutumumab and the mTOR inhibitor temsirolimus [46].

Recent studies suggest that the SOAT1 (Sterol-O-Acetyl Transferase 1) enzyme is a novel emerging therapeutic target in ACC. In Ann Arbor, Dr. Matthias Kroiss (University of Würzburg, Germany) presented the results of a groundbreaking study from his group that for the first time identified a molecular target for mitotane. Mitotane was found to inhibit SOAT1, causing accumulation of free cholesterol, oxysterols and fatty acids inside adrenocortical cells, induction of ER stress and, ultimately, cell death. The study by the Würzburg team has also shown that SOAT1 expression levels are

variable in ACC, being low or absent in about one third of cases [47]. This provides a rationale to screen patients with ACC for SOAT1 expression in their tumor to predict their response to mitotane. In another session of the symposium, Dr. Gary Hammer presented data showing the efficacy of ATR-101, a new orally available SOAT1 inhibitor, in preclinical models of ACC. A phase I clinical trial for safety and tolerability of ATR-101 is currently ongoing (<https://clinicaltrials.gov/ct2/show/NCT01898715>).

Conclusion

The recent great advances in our understanding of ACC biology need now to be translated to create cellular and animal models that are closer to the human disease in order to develop new, more effective diagnostic procedures and targeted therapies against this deadly neoplasm. Close collaboration between basic scientists and clinicians as well as maintenance and further expansion of international ACC patients registries is considered essential for the success of this endeavor.

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Compliance with Ethical Standards

Conflict of Interest The authors declare that they have no competing interests.

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