

(injurious and beneficial) and possible management strategies. Modifiable factors such as ventilator settings (3), respiratory drive (4), level of consciousness, and prescribed sedatives and opiates may predispose differentially to reverse triggering depending on the mechanism. Regarding clinical significance, some degree of permissive dyssynchrony is probably benign or even beneficial for patients at low risk of lung or diaphragm injury. For patients at highest risk, including those with moderate and severe ARDS, clinical consequences of reverse triggering likely depend on the subtype, duration of exposure, and tradeoffs of interventions such as neuromuscular blockade.

Despite the uncertainties surrounding reverse triggering, the following is clear: a rapidly expanding literature indicates reverse triggering occurs often in mechanically ventilated patients at risk of injury and might be underrecognized at the bedside. What, if anything, should be done clinically remains to be determined. ■

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Ⓔ Molecular Imaging of Pulmonary Fibrosis: Another Step Forward

Molecular imaging enables *in vivo* visualization of molecular processes within a tissue or organ of interest using targeted molecular probes. Because of advancements in molecular imaging

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in fields ranging from neurology to oncology, there are now multiple U.S. Food and Drug Administration–approved molecular probes for use in clinical care. Within the field of fibrotic diseases, the application of molecular imaging in the preclinical arena and early-phase clinical trials is growing (1, 2). Over the past year, the results of the first in-human studies using the $\alpha\text{v}\beta 6$ integrin–targeted positron emission technology (PET) probes [¹⁸F]FB-A20FMDV2 and [¹⁸F]FP-R₀1-MG-F2 and the type 1 collagen–targeted PET probe ⁶⁸Ga-CBP8 demonstrated increased PET signal in patients with pulmonary fibrosis consistent with increased $\alpha\text{v}\beta 6$ integrin expression and type 1 collagen, respectively (3–5). In addition, ¹⁸F-fluorodeoxyglucose

PET/computed tomography (CT) was used in a phase I study of the PI3K/mTOR inhibitor omipalisib in idiopathic pulmonary fibrosis (IPF) to confirm target engagement (6). With ongoing development, molecular imaging is poised to advance the field of pulmonary fibrosis in several notable and important ways.

First, molecular imaging can provide information as to fibrosis pathobiology that is not otherwise available (7). Most of the knowledge as to aberrant molecular pathways driving lung fibrosis has come from studies using the bleomycin mouse model. Validating these findings in patients with pulmonary fibrosis is difficult. Although the ability to obtain tissue for molecular phenotyping has revolutionized the field of oncology, obtaining lung tissue carries considerable risks for patients with pulmonary fibrosis, and this precludes its use for investigational purposes. Thus, molecular imaging emerges as a noninvasive alternative to invasive tissue sampling. It can enable the degree of activation of a molecular process of interest to be quantified and, at the same time, reveal the anatomic distribution where the process of interest is occurring. Second, and perhaps most notably, molecular imaging could be incorporated into clinical trials for the assessment of target engagement and expression, cohort enrichment, or the detection of treatment response (1), and in doing so, it could address several unmet needs that have hampered the development of IPF therapies. Molecular imaging could be used to stratify patients based on the degree of abnormality in the pathway of interest and enable clinical trials to be enriched with subjects with active disease. A molecular probe that targets a molecule or process along the pathway of fibrogenesis could serve as a much-needed early marker of treatment response overall, improving trial feasibility and accelerating the identification of effective therapies.

In this issue of the *Journal*, Brody and colleagues (pp. 78–89) add to the expanding field of molecular imaging by applying the CCR2 (chemokine receptor 2)-targeted PET probe ^{64}Cu -DOTA-ECL1i to detect C-C motif CCR2-positive macrophages and monocytes in pulmonary fibrosis (8). Studies have demonstrated an important role for CCR2 signaling in pulmonary fibrosis pathogenesis (9). This peptide-based probe binds the extracellular loop 1 of the CCR2 and has been used to assess inflammation in mouse models and explanted human tissue of other pulmonary diseases (10, 11). Multiple macrophage/monocyte-targeting probes have been developed; however, this probe is unique in that it recognizes CCR2⁺ immune cells. Other macrophage-targeting probes applied to pulmonary fibrosis have targeted the folate receptor- β or cysteine cathepsin as markers of activated macrophages (12, 13). It should be noted that these probes detect inflammation in the context of fibrosis and are not specific for fibrosis.

This study has several important findings. First, in mice injured with bleomycin, ^{64}Cu -DOTA-ECL1i uptake in the lung was increased at Day 14 and 28 in a pattern similar to the degree of elevation of CCR2⁺ cells. Increased probe uptake was also seen in explanted human fibrotic lungs with good correlation between probe accumulation determined by autoradiography and CCR2 intensity. Second, ^{64}Cu -DOTA-ECL1i uptake decreased in response to both an IL-1 β antibody and pirfenidone in the bleomycin model. It is important to note that other probes have detected response to fibrosis-targeting treatment (14–16), and although ^{18}F -fluorodeoxyglucose detected a response to

pirfenidone in mice, it was unable to do so in patients with IPF (16). Third, dosimetry studies demonstrated that ^{64}Cu -DOTA-ECL1i is primarily renally cleared, with minimal background uptake in healthy lungs, an important feature for detecting pulmonary pathology. The persistent high signal in the liver adds complexity to quantifying signal uptake in the surrounding lower lung. Lastly, performing ^{64}Cu -DOTA-ECL1i PET-CT in subjects with IPF was feasible. Areas of increased lung PET signal were seen visually in the four subjects with IPF compared with one healthy volunteer. Within the patients with IPF, areas of increased signal appeared to correspond with areas of fibrosis on CT. Although promising, these findings should be considered preliminary, as the number of subjects with IPF imaged ($n = 4$) was small, the results were presented in comparison with only one healthy volunteer, and no statistical analyses were performed.

Several important issues remain to be addressed regarding the potential for the clinical translation of this probe. Larger studies are needed to validate the ability of this probe to detect increased PET signal in patients with IPF even when adjusting for potential confounders given the presence of monocytes and macrophages in other pulmonary diseases. Although increased PET signal was detected within areas of fibrosis on corresponding CT, whether or not the degree of signal will provide additive information to CT and pulmonary function tests and inform as to the risk of disease progression, thus serving as a viable disease activity measure, remains to be determined. Performing test-retest studies over a small time interval will be essential to determine the ability of this probe to detect small changes that may occur with IPF therapy. Lastly, should this probe detect a decrease in PET signal in the setting of IPF treatment(s), ongoing research would be needed to determine whether a reduction in CCR2⁺ cells corresponds with therapeutic efficacy.

The application of CCR2-targeted PET imaging to pulmonary fibrosis represents another step forward in the emerging field of molecular imaging of fibrosis. Given the overlapping molecular processes across both subtypes of pulmonary fibrosis and fibrotic diseases in general, advancements in molecular imaging of IPF may prove beneficial to patients with other forms of pulmonary fibrosis and other fibrotic diseases. ■

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Therapeutic Interception of Early Lung Adenocarcinoma Progression: Not Just How, but When?

In this issue of the *Journal*, Treekitkarnmongkol and colleagues (pp. 90–101) present data on a pathway of early adenocarcinogenesis of the lung, demonstrating the effect of lipocalin-2 on the development of KRAS-mutated lung adenocarcinoma (KM-LUAD) and suggesting that airways damaged by chronic obstructive pulmonary disease (COPD) show the effect of a high-lipocalin-2 microenvironment (1). Their experiments were based on an observed elevation of *Lcn2* in the nonneoplastic airways of mice with knockout of *Gprc5a* (2) followed by studies examining lipocalin-2 in these mice exposed to tobacco carcinogen, increased tumor development in *Lcn2*^{-/-} mice, and translation into human tissues with COPD and LUAD.

The review of the literature of the relationship of LCN2 to human cancer would identify it as having a protumoral function (3–6), by mechanisms that include survival advantage through iron scavenging and enhancement of migration. In some tumor types, it is proposed as a target for therapy or, at a minimum, a biomarker for diagnosis or of aggressive behavior (7).

Although LUAD showed an increase in lipocalin-2, the authors show through their mouse model that loss of lipocalin-2 reduced antitumoral immune responses and enhanced a protumoral immune environment, such that the loss of *Lcn2* enhanced tumor

formation. The induction of LCN2 was not seen in squamous carcinoma, occurring in KM-LUAD preferentially and by immunohistochemistry in tumor cells. An inverse relationship between NKX2-1 (TTF1) reactivity and LCN2 reactivity suggested differentiation away from club cell and type 2 pneumocyte, toward a gastric-type cell differentiation. In addition, the upregulation of LCN2 was seen in COPD airways when compared with smokers without COPD but importantly included elevation in KM-LUAD from patients with COPD.

At first glance, there is a paradox—LCN2 is protumoral in several tumor types in prior reports yet here antitumoral. Also, LCN2 levels are high in LUAD, especially KM-LUAD. There are potential explanations for this type of paradox, which can include cell type or differentiation, differences in tumor microenvironment, and/or temporal switches in the stepwise progression toward neoplasia. Even among T1 adenocarcinomas, the antitumoral effect seems already lost; it is possible that the role of LCN2 in tumor initiation is different than a later role in cell migration. Additionally, it remains possible that non-tumor cell LCN2 becomes a factor in different phases of the disease. When we observe a tumor mass lesion, we are seeing the outcome of numerous potentially antagonistic tumor–host events that have occurred over time, which eventually fail to control tumor growth and spread. Such an explanation allows for an antitumoral effect for LCN2 that creates a balance between preneoplasia and cell death, a balance that accounts for a period of latency to adenocarcinoma development in COPD not seen for squamous or small cell carcinoma.

This could allow for risk stratification for the development of KM-LUAD that are smoking-associated LUAD among patients

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