Redrawing the Map to Novel DILI Biomarkers in Circulation: Where Are We, Where Should We Go, and How Can We Get There?

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Abstract: Circulating biomarkers of drug-induced liver injury (DILI) have been a focus of research in hepatology over the last decade, and several novel DILI biomarkers that hold promise for certain applications have been identified. For example, glutamate dehydrogenase holds promise as a specific biomarker of liver injury in patients with concomitant muscle damage. It may also be a specific indicator of mitochondrial damage. In addition, microRNA-122 is sensitive for early detection of liver injury in acetaminophen overdose patients. However, recent events in the field of DILI biomarker research have provided us with an opportunity to step back, consider how biomarker discovery has been done thus far, and determine how to move forward in a way that will optimize the discovery process. This is important because major challenges remain in the DILI field and related areas that could be overcome in part by new biomarkers. In this short review, we briefly describe recent progress in DILI biomarker discovery and development, identify current needs, and suggest a general approach to move forward.

Keywords: hepatotoxicity; drug-induced liver injury; acute liver failure; diagnosis; prognosis; regulation

1. Introduction

Liver injury is a life-threatening adverse effect of some drugs. In fact, drug-induced liver injury (DILI) is one of the most common reasons for pre-approval termination of new drugs and for post-approval withdrawal. There are two basic forms of DILI: intrinsic and idiosyncratic. Intrinsic DILI is caused solely by physico-chemical properties (e.g., reactivity) that are intrinsic to the drug itself (or to a metabolite) and usually displays a clear dose-response as a result. Idiosyncratic DILI (usually abbreviated IDILI), on the other hand, depends in part upon “host” factors (e.g., poor immunotolerance) and has much weaker dose-response, and therefore poor predictability as a result. Clinically, intrinsic DILI and IDILI together make up the primary cause of acute liver failure (ALF) in the US [1]. From a regulatory point of view, IDILI is also notoriously difficult to predict or diagnose. As a result, regulatory agencies and clinicians err on the side of caution, adopting an approach based on monitoring of circulating alanine aminotransferase (ALT) and other conventional liver markers within the context of the greater clinical picture. However, weaknesses in that approach are increasingly clear. To address that, recent years have seen considerable effort invested in development of novel circulating biomarkers of liver injury.

In 2018–2019, DILI biomarker researchers were shaken by the revelation that a prominent investigator committed scientific misconduct [2]. Although investigation revealed the impropriety was limited to one individual and one analyte, the consequences are far-reaching. For example, the European Medicines Agency retracted a letter of support for development of several DILI biomarkers for regulatory use. We view this as an opportunity to evaluate where we are in the field, how we got here, and how we can move forward in a...
way that optimizes the biomarker discovery process and reduces potential bias. Here, we discuss current DILI biomarkers and then look to the future to identify areas of continued need and general approaches to meet those needs.

2. Points of Departure
2.1. Where Are We?

Many interesting liver injury biomarkers have been identified, including keratin 18 (K18), caspase-cleaved K18 (ccK18), high-mobility group box 1 protein (HMGB1), microRNA-122 (miR-122), glutamate dehydrogenase (GLDH), osteopontin (OPN), macrophage colony-stimulating factor receptor (MCSFR), and numerous others [3–11]. Some clearly show promise for specific applications. For example, some provide mechanistic insight: ccK18 indicates mode of cell death (apoptosis vs. necrosis) [3]; MCSFR may indicate inflammation [9]; GLDH, mitochondrial DNA, carbamoyl phosphate synthetase 1 and acylcarnitines may indicate mitochondrial damage [5,12–15]; OPN, alpha-fetoprotein (AFP), and others reflect regeneration [9,16]. Others have clinical benefits: miR-122 seems to predict later liver injury in early-presenting acetaminophen (APAP) overdose patients [8], while GLDH is helpful to distinguish liver and muscle damage in patients at risk of both [9,17]. Indeed, the Critical Path Institute and associated researchers are making great strides to promote the widespread adoption of GLDH as a liver injury biomarker in the context of clinical trials to test treatments for musculoskeletal disorders. However, these successes have largely been achieved by adopting biomarkers developed for other diseases and therefore often lacking specificity. To some degree, there have also been developments of convenience—being tested only after introduction of simple, commercially-available test kits. It is tempting to imagine the great strides that could be made with more focused, systematic efforts beginning with untargeted ‘omics or modern artificial intelligence methods.

2.2. How Did We Get Here?

So far, despite the achievements listed above, the process of discovery of novel liver injury biomarkers has proceeded more-or-less laissez-faire. Biomarkers intended for use in other diseases have been co-opted for liver injury. For example, ccK18 was originally described as an immunocytochemical marker of apoptosis in cancer cells [18]. ELISAs were then developed and used to test it as a circulating biomarker of cancer [19,20]. Years passed between the identification of ccK18 as an apoptosis marker in cells and the first application in liver injury patients, and then the earliest studies focused on chronic liver diseases [21,22]. It is immediately clear from the history that ccK18 and K18 are not specific for the liver and certainly not for DILI. Similarly, HMGB1 was first measured in serum from sepsis patients [23] and later characterized as a damage-associated molecular pattern released from various cell types after necrosis [24], before being applied to liver injury. Those biomarkers became popular and were applied to liver injury only after introduction of convenient methods (e.g., commercial ELISAs) to measure them. For other biomarkers, such as malate dehydrogenase, convenient measurement methods have already existed for decades, but those markers have been tested and found to be elevated in numerous contexts, not just liver injury. Even ALT, the clinical and regulatory gold-standard for detection of liver injury, was intended as a biomarker of myocardial damage [25,26]. In fact, among the current crop of DILI biomarkers, only miR-122 was discovered systematically with a focus on liver damage [27], and even miR-122 has somewhat limited specificity, for example, being elevated in metabolic diseases and renal cell carcinoma as well [28,29]. Limiting biomarker exploration to biomarkers identified in other contexts and/or that have convenient test methods immediately limits the specificity for DILI and fails to optimize sensitivity. In addition, there may be more pressure to report positive results when only a small number of potential biomarkers and test methods are available.
2.3. How Should We Move Forward?

We contend that more systematic approaches are needed to fill the gaps in DILI biomarker development. Experiments should be designed to meet specific needs in the DILI field, moving forward from the need to the biomarker and not vice versa. This could be coupled with untargeted measurements for advanced, unbiased, ‘omics- or artificial intelligence-based approaches for the identification of many novel biomarkers, as recently discussed in detail elsewhere [30]. Those results should then be validated in additional models and in humans using appropriate samples.

3. DILI Destinations

So, what are the current needs? It has been argued that biomarkers could be useful for (1) diagnosis of DILI due to any drugs; (2) prediction of IDILI, specifically, prior to drug use; (3) prognosis in intrinsic DILI, IDILI, and acute liver injury more generally; and (4) prediction of hepatotoxic liability during pre-clinical drug development and early clinical trials. Here, we evaluate each and identify specific approaches that may work. We also argue that two of these goals are especially important and realistically achievable.

3.1. Biomarkers for Diagnosis

Clinically, recent reports have noted that diagnosis of DILI is challenging due to low prevalence [10,31–33]. It is unlikely that a biomarker can be developed that will reliably rule-in IDILI because the low prevalence (typically ≤1% and often as low as ≤0.1%) of IDILI among users of any given IDILI-causing drug means that no biomarker can achieve sufficient positive post-test probability (or positive predictive value) to prove a diagnosis (Figure 1) [31,33]. For the typical range of IDILI incidence, positive post-test probability maxes out around 10%, even for a test with high clinical sensitivity and specificity (Figure 1).

Although high negative post-test probability (and negative predictive value) can certainly be achieved (solid line in Figure 1B), the pre-test probability of IDILI is already low due to low prevalence, so a negative test result would have questionable benefit. However, it may be possible to diagnose causes of intrinsic DILI. Indeed, serum APAP-protein adducts appear to be useful to diagnose APAP overdose [33,34], which is the most common cause of DILI overall. Unfortunately, because other drugs are less common causes of liver injury individually, it is not clear how well this approach would actually work beyond APAP, especially in the case of IDILI-causing drugs.

![Figure 1. Relationship of test performance and pre-test probability to post-test probability. (A) Prevalence (pre-test probability) is shown on the x-axis. Post-test probability is shown on the y-axis. Each line displays the post-test probability](image-url)

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*Figure 1.* Relationship of test performance and pre-test probability to post-test probability. (A) Prevalence (pre-test probability) is shown on the x-axis. Post-test probability is shown on the y-axis. Each line displays the post-test probability for different values of test performance (sensitivity and specificity).

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on the y-axis as a function of prevalence (pre-test probability) at ascending test sensitivity (50%, 60%, 70%, 80%, 90% and 99%) and at fixed values for test specificity of 80% (red series), 90% (purple series), and 95% (blue series). The vertical line shows a typical upper limit of IDILI prevalence. (B) Fagan nomogram displaying prevalence (pre-test probability) on the left, likelihood ratio (LR) in the middle, and post-test probability on the right. The dashed blue line shows the relationship with positive post-test probability for a biomarker that has >90% sensitivity and >90% specificity for DILI (LR+ = 9–10). The solid yellow line shows the relationship with negative post-test probability for a biomarker that has 90% sensitivity and 90% specificity for DILI (LR− = 0.11).

3.2. Biomarkers for Prediction

Many researchers have proposed that genetic and circulating biomarkers for prediction could be used to manage risk of IDILI, specifically, by determining the likelihood that a drug will cause injury before a patient takes it. Numerous gene variants are associated with IDILI [35], but currently only one is used clinically for this purpose. The link between HLA B*5701 and abacavir reactions is useful because ~5% of abacavir users experience an adverse event, which can include IDILI [33,36]. It is unlikely that many other biomarkers will gain purchase for prediction of IDILI for the same reason it is difficult to diagnose IDILI. The prevalence of liver injury among users of most IDILI-associated drugs is low, usually ≤1% and often even ≤0.1%. Thus, even for a very good theoretical biomarker with 95% clinical specificity for IDILI, the rate of false positives will always be much greater than the rate of true positives, resulting in very low positive predictive value or positive post-test probability (Figure 1). Although it is theoretically possible to predict IDILI in the case of drugs with relatively high incidence of injury (e.g., ximelagatran [32,33]) or for drugs with high prevalence of adverse events in general—not limited to liver damage (like abacavir)—those examples are relatively rare. In addition, most genetic associations with IDILI are somewhat weak, reducing their clinical specificity in the first place.

3.3. Biomarkers for Prognosis

Biomarkers for prognosis in DILI overall (whether intrinsic or idiosyncratic) are sorely needed; first, to predict development of injury in patients presenting with normal-to-low ALT values, and second, to predict death in patients with drug-induced ALF. Recent studies have demonstrated that several biomarkers, but especially miR-122, can predict later injury in early-presenting APAP overdose patients despite its considerable biological variation [8]. In fact, miR-122 is probably approaching the limit of what is possible in that respect, based on the prevalence of injury among those who present early with suspected APAP overdose [33]. However, we lack biomarkers to predict death after acute liver injury, and particularly ALF. ALF has high mortality (>25%) [37–39] and availability of transplantable livers is limited. biomarkers to predict death could greatly improve clinical decision-making by helping physicians determine which patients need a new liver to survive. Theoretically, it should be easy to identify biomarkers for death in ALF that have high predictive values due to the high pre-test probability of death in ALF. However, less effort has been directed this way compared to other areas of DILI biomarker research. Still, some interesting results are available. For example, circulating AFP predicts survival in ALF, though it changes late in the course of ALF when it may be too late for a liver transplant [16]. Recent data indicate that OPN and K18 could also be useful to predict mortality [9], and a recent systematic review indicated that the combination of K18 with other measures may be especially helpful [40], though additional studies need to validate these early results. In addition, biomarkers that may reflect the severity or extent of underlying injury may be promising for prognostic use, as worse injury may be expected to lead to worse outcomes. Some recent examples of such biomarkers include advanced oxidation protein products and ischemia-modified albumin, which are elevated in DILI patients [41]. Overall, considerably more work should be done in this area.
3.4. Biomarkers for Hepatotoxic Liability

Better safety biomarkers for regulation are also still needed. The current practice of screening for hepatotoxicity with ALT and other conventional markers during clinical trials comes from the basic principles Hy’s Law and Temple’s Corollary. In essence, Hy’s Law states that an elevation in ALT accompanied by jaundice is a signal of serious liver injury, while Temple’s Corollary states that a drug that causes modest, transient ALT elevations in many patients is likely to cause Hy’s Law cases in at least a few. Thus, moderate ALT elevations in more than one subject or mild elevation of both ALT and bilirubin in even a single subject are considered signs of potentially serious hepatocellular damage that warrant termination of new drugs.

A problem with using ALT in this way is that it lacks specificity for DILI, but in different contexts. Pre-clinical models have poor sensitivity to predict IDILI in humans [41], while the modest ALT elevations frequently observed in clinical trials are sensitive but lack specificity for severe injury. The US FDA acknowledges that some drugs meet the criteria for hepatotoxicity concern in clinical testing, but have never caused a case of severe DILI in the general population [42]. Indeed, many drugs such as APAP, heparins, and cholestyramine can cause non-progressive ALT elevations in many users with no other evidence of liver damage [43–50]. Furthermore, there is evidence that ALT lacks specificity for any tissue damage, be it in the liver or elsewhere, in certain diseases [51,52]. We recently tried to address this issue by combining models of benign serum ALT elevations and models of liver injury with untargeted proteomics to identify serum biomarkers that can differentiate between them [52]. We propose that such a biomarker could be used in an algorithmic “screen-and-confirm” approach to determine if minor ALT elevations detected during screening for injury in clinical trials reflect actual liver injury. As for improving the sensitivity of pre-clinical models, one interesting strategy attempted was to compare serum biomarkers between animals treated with pairs of drugs with similar pharmacology, one of which was known to cause DILI in humans and one of which was not [53,54]. Those studies succeeded in identifying analytes that may be specific for the DILI-causing drugs, demonstrating that such an approach has promise.

Overall, the areas of greatest need and potential for further biomarker development now appear to be biomarkers of prognosis to predict death in DILI-related ALF (and ALF more generally) and biomarkers for hepatotoxic liability during drug development. There is potential for additional developments in the areas of diagnosis and prediction as well, but the issue of low prevalence will be a major hurdle to overcome.

4. Conclusions

We have an opportunity to re-examine the DILI biomarker field and consider how to move forward. Continued progress is important because significant needs remain. Biomarkers of diagnosis and prediction are theoretically possible and could have major clinical value, but will be very challenging to develop. On the other hand, biomarkers are urgently needed to predict patient outcomes in DILI, and especially DILI-induced ALF, and should be relatively simple to develop. Furthermore, biomarkers that are specific for liver injury are needed to determine if modest ALT elevations observed in clinical trials or clinical practice reflect actual liver injury, and we suggest using them in a screen-and-confirm approach. Finally, regardless of the focus, more systematic approaches such as applying ‘omics methods to specific models of liver injury or to samples from patients could help us meet these needs in a less biased way.

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