Full Title: **Dabigatran prevents lipopolysaccharide mediated apoptosis in zebrafish through**

a thrombin independent mechanism

Short Title: **Dabigatran prevents lipopolysaccharide induced toxicity**

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Abstract: Endotoxemia is a feature of sepsis pathogenesis and has also been found to mediate

the pathophysiology of multiple chronic inflammatory conditions. In this work, we use a

lipopolysaccharide (LPS) induced endotoxemia model in zebrafish to identify novel mediators of

LPS toxicity. We performed transcriptomic studies on LPS-treated larvae, followed by *in silico*

analysis, which revealed associations between the signatures of LPS-treated embryos and those

of drugs involving diverse pathways. In parallel, we performed an *in vivo* screen using >1,500

FDA-approved compounds and identified multiple novel small molecules that reduced

inflammation and prevented LPS toxicity. We focused on the direct thrombin inhibitor

dabigatran, which was identified through both the *in vivo* and *in silico* analyses. We found that

dabigatran co-administration significantly reduced the expression of inflammatory cytokines and

completely protected zebrafish from endotoxemic death due to LPS. Surprisingly, we found that

this protection occurs in prothrombin mutant fish, proving that protection from endotoxemia

occurs independently of the anticoagulant function of dabigatran. We additionally found that

dabigatran administration significantly decreased nitric oxide production and apoptosis

compared to LPS treatment alone, suggesting possible mechanisms by which protection from

endotoxemia is achieved. In summary, we identify several novel small molecules that prevent

LPS-induced endotoxemia and show that one such small molecule, dabigatran, exerts a

thrombin-independent effect on nitric oxide production and apoptosis. This and the other

identified small molecules warrant further exploration in inflammatory conditions including

sepsis.

Author Summary: Sepsis is a common and deadly illness caused when there is an

uncoordinated and overwhelming response to an infection. Endotoxemia is a feature of sepsis

and occurs when part of the outer cell wall of bacteria called lipopolysaccharide (LPS) is

released into the bloodstream, causing an excessive inflammatory response. To identify

candidate drugs that could be repurposed to combat human disease, we screened through over

1500 FDA approved small molecules for protection against LPS induced toxicity using zebrafish

as a model organism. In parallel, we undertook an *in silico* approach and compared the gene

signature of LPS-treated embryos to the gene signature of small molecules published in the

literature. We identified several novel drugs that could protect against LPS using these

approaches. We further found that one such drug, the anticoagulant dabigatran, blocks LPS

mediated cell death through a novel mechanism, independent of its primary target. Our work sets

the stage for dabigatran and the other candidate drugs to be explored as therapeutics in human

inflammatory conditions including sepsis.

Introduction:

Lipopolysaccharide (LPS), a cell wall component of gram-negative bacteria, induces an immune response upon recognition by Toll-Like Receptors (TLRs) (1, 2). LPS accounts for the relative impermeability of the outer membrane of gram-negative bacteria to hydrophobic molecules, making them resistant to many antimicrobials (3, 4). LPS-mediated endotoxemia mimics many features of sepsis pathogenesis and might also play a causative role in the pathophysiology of multiple chronic inflammatory disorders, including Parkinsons, Alzheimers, obesity, and atherosclerosis (5). Previous work in mammals suggests that the translocation of gut-derived LPS into the circulation might mediate some of these disease states (6, 7). Additionally, LPS contributes to disease pathogenesis through its action as a prothrombotic molecule; LPS-induced inflammatory mediators activate procoagulant pathways, induce platelet activation, downregulate anticoagulant pathways and promote disseminated intravascular coagulation (8, 9). Moreover, macrophages undergoing pyroptosis release tissue factor, further activating the coagulation cascade (5, 9).

Given its role in multiple diseases, efforts have been made to block LPS biogenesis, neutralize endotoxin, and hinder LPS-mediated immune activation (10-13), with many of these attempts made in the context of sepsis. However, there is the possibility that therapeutics targeting LPS signaling can also mitigate the development of chronic inflammatory conditions such as those listed above.

In this work, we undertook large-scale *in silico* and *in vivo* analyses to identify novel small molecule mediators of LPS toxicity. Murine and other mammalian models are cost-prohibitive for the same types of large-scale *in vivo* screening studies. Zebrafish, on the other

hand, allow for a speed of discovery not possible with mammals as they combine the scalability

of *in vitro* systems with the complexity of a whole vertebrate organism.

In zebrafish, LPS-induced inflammation has been provoked either by injection into the

embryonic yolk sac or immersion of embryos and larvae into LPS-containing media (14-16).

Incubation with LPS-containing media allows uniform, rapid treatment of large numbers of fish.

A response remarkably similar to that in mammals is observed, including inflammation, platelet

activation, vascular leakage, and organ damage (16). Thus, many of the cardinal features of

sepsis pathogenesis are recapitulated by this model.

We performed novel high throughput studies in a zebrafish model of LPS-provoked

endotoxemia with the goal of identifying drugs that could be repurposed to combat acute and

chronic inflammatory conditions. We conducted an *in silico* analysis, where we compared RNA-

seq data from LPS-treated fish to publicly available RNA-seq datasets, revealing associations

between LPS and drugs involving disparate pathways. In parallel, an *in vivo* pathway analysis

was performed using >1,500 FDA-approved compounds to find those that mitigated LPS-

induced endotoxemia, uncovering multiple novel small molecules. We focused on the sole

anticoagulant (dabigatran) identified in both the *in silico* and *in vivo* analyses. Interestingly, the

administration of dabigatran reduced the upregulation of inflammatory cytokines, significantly

reduced nitric oxide production and apoptosis and abolished larval mortality from endotoxemia

due to LPS. Surprisingly, this protection was found to occur independent of its inhibitory effect

on thrombin. These data implicate dabigatran and other novel small molecules as anti-

inflammatory agents that can be explored as potential therapeutics in acute and chronic

conditions.

Results:

LPS incubation results in inflammatory and coagulopathic changes in zebrafish embryos

S. typhi LPS was administered to zebrafish by incubation starting at 3 days post fertilization (dpf) in accordance with published data (16). A dose-dependent effect on lethality was noted over 24 hours. Hereafter, the lethal dose of LPS refers to the smallest amount that caused uniform lethality by 4 dpf, which was slightly different (ranging from 40-50 µg/mL) depending on the batch. A new dose-response curve was made for each batch of LPS tested, and an example is shown in Fig. 1A. In addition to increased mortality, heart rate was significantly decreased (Fig. 1B), and *grk5,* a gene involved in heart contractility, was significantly downregulated 3 hours after LPS administration (Fig. 1C).

Figure 1: LPS incubation results in dose-dependent lethality and signs of end-organ damage in zebrafish larvae. (A) Percent survival of 4 days post fertilization (dpf) zebrafish larvae incubated with *S. typhi* LPS at 3 dpf. 3 clutches of 15 fish (total of 45) were used per concentration. (B) Heart rate (beats per minute) was counted 3 hours after LPS (40 µg/mL) administration and compared to controls. Each point represents a different fish, and the results include 3 separate experiments. (C) *grk5*, a gene involved in heart rate contractility, was measured by qPCR in LPS-treated fish compared to controls. Larvae from two or more clutches were pooled, where each point represents a separate clutch, and the average fold change in LPS-treated fish vs. controls was graphed. 30-45 total fish were used per time point. Bars indicate mean and standard deviation. Student's t-test (two-tailed) on $\Delta\Delta C_t$ values was used to determine significance. *$P<0.05$.

To determine the significant biological pathways altered upon LPS administration, RNA-

seq was performed on treated fish compared to controls as previously described (Guirgis et al.,

2023). Kyoto Encyclopedia of Genes and Genomes (KEGG) pathway analysis was performed

and revealed that "Pathogenic *E. coli* Infection" was the most significantly altered pathway in

LPS-treated fish (Fig. 2A, complete analysis is shown in Table S1), confirming that

administration induces a response akin to that provoked by bacteria. As expected, immune and

interferon pathways such as TNF signaling were significantly changed (Fig. 2A). A subset of

genes from these pathways were confirmed by qPCR and trended toward upregulation, with the

genes *il1b*, *il6* and *irg1l* significantly upregulated (Fig. 2B). In addition to provoking a

proinflammatory response, LPS is also known to be a prothrombotic molecule (7, 9). RNA-seq

data also showed that the complement, coagulation, and platelet activation pathways were

significantly altered in LPS-treated zebrafish compared to controls (Fig. 2A). A subset of

procoagulant genes from these pathways was upregulated, which was confirmed by qPCR (Fig.

2B). Interestingly, tissue factor b (gene *f3b*), but not tissue factor a (gene *f3a*) was significantly

upregulated in LPS-treated fish compared to controls, suggesting possible arterial injury as *f3b* is

thought to play a primary role in arterial, rather than venous, circulation (17). Time to occlusion

(TTO) after venous and arterial laser-mediated endothelial injury was measured as previously

described (17, 18) and is illustrated in Fig. 2D. The TTO was found to be significantly increased

in fish administered LPS for both systems (Fig. 2E, F). This change suggests a consumptive

coagulopathy phenotype similar to what we have observed with the loss of natural anticoagulants

(18, 19).

Figure 2: The immune and interferon signaling pathways and procoagulant pathways are

significantly upregulated in LPS-treated fish compared to controls. (A) KEGG pathway

analysis of RNA-seq data from whole LPS treated fish compared to controls. Significantly

changed genes (adjusted p-value <0.05) from RNA-seq data on pooled 3 dpf LPS-treated vs.

control fish (consisting of 30-45 embryos per category) performed in triplicate were used for

further analyses. These genes were converted to human versions and then submitted for KEGG

Pathway Analysis using the Enrichr platform, a publicly available tool developed by the

Ma'ayan lab at Icahn School of Mount Sinai, NY (Chen et al., 2013, Kuleshov et al., 2016).

Overlap score is the number of significantly changed genes over the total number of genes in the

pathway. Combined score considers the z-score (a value's relationship to the mean) in addition

to the adjusted p-value. The combined score is used to assess the significance of gene set

enrichment findings, with higher values being more significant. Adjusted p-value <0.05 is

considered significant. Immune and interferon-related and complement and coagulation

pathways are shown. For complete analysis, see Table S1. (B) A subset of genes in the immune

and interferon pathways were tested by qPCR, using 3 dpf larvae treated with 40 µg/mL LPS vs.

control conditions. Larvae from two or more clutches were pooled, where each point represents a

separate clutch, and the average fold change in LPS-treated fish vs. controls is graphed. Bars

represent mean and standard deviation. Student's t-test (two-tailed) on $\Delta\Delta C_t$ values was used to

determine significance. *P<0.05. (C) A subset of procoagulant genes was tested by qPCR, using

3 dpf larvae treated with 40 µg/mL LPS vs. control conditions. Larvae from two or more clutches

were pooled, where each point represents a separate clutch, and the average fold change in LPS-

treated fish vs. controls is graphed. Bars indicate mean and standard deviation. Student's t-test

(two-tailed) on $\Delta\Delta C_t$ values was used to determine significance. *P<0.05. A schematic of the

laser injury assay used to measure time to occlusion is shown in (D), where the time to formation

of an occlusive thrombus after laser pulses were delivered to the posterior cardinal vein or dorsal

aorta was measured. 120 seconds was the maximum time each larva was observed. Time to

occlusion (TTO) after laser injury in the venous (E) and arterial (F) circulation was measured in

3 dpf zebrafish larvae 3 hours after LPS administration and compared to controls. Each point

represents a separate fish, and the observer was blinded to the treatment condition. $P < 0.05$ is

considered significant by Mann-Whitney *U* testing.

In silico and in vivo analysis reveals small molecule mediators of LPS-mediated endotoxemia

To identify possible mediators of LPS toxicity, we initially undertook an *in silico* analysis

comparing the RNA-seq signature of LPS-treated embryos to those of genes and compounds

from publicly available databases (Table S2). After obtaining the list of small molecules related

to LPS based on the gene expression profiles (Table S2), we further looked into mechanisms of

action for each of these unique molecules as provided by the Connectivity Map (CMap)

database. Even though some compounds did not have an annotated MoA, the data showed an

intriguing pattern among the ones with detailed annotations. Notably, neuroactive compounds

such as dopamine and serotonin receptor antagonists, along with ion channel and kinase

inhibitors, emerged as the most commonly shared classes amongst the LPS-associated

compounds (Table 1).

Mechanism of Action (MoA)	How often MoA occurs among candidate drugs from Table S2
Dopamine receptor antagonist	10
EGFR inhibitor	9
Tubulin inhibitor	8
Adrenergic receptor antagonist	6
MEK inhibitor	6
PI3K inhibitor	6
Serotonin receptor antagonist	6
Calcium channel blocker	5
KIT inhibitor	5
Na/K-ATPase inhibitor	5
RAF inhibitor	5
JAK inhibitor	4
PDGFR inhibitor	4
PKC inhibitor	4
Topoisomerase inhibitor	4

Table 1: MoA of small molecules from Table S2 sorted by the number of times the MoA appears among the list of candidate drugs (only occurrences >= 4 are shown)

In parallel, we performed an *in vivo* analysis in zebrafish embryos. A lethal dose of LPS was co-administered at 3 dpf with an FDA-approved drug repurposing set consisting of >1500 compounds. Embryos were examined for survival 24 hours post-treatment, and each drug that was a 'hit' (i.e., enabled survival of at least one embryo) was validated in a larger cohort to confirm rescue. Approximately 20 small molecules were identified in this manner (Table 2). Most hits completely protected against LPS toxicity during the embryonic/early larval period, and others provided partial protection (Table 2, Fig. S1 A-T). These small molecule mediators of

LPS toxicity spanned multiple categories, most heavily represented by serotonin and dopamine

receptor modulators, chemotherapeutics, and anti-infectious agents. Similar to the *in silico*

analysis, small molecules modulating the dopamine and serotonin pathways and those affecting

tyrosine kinases were heavily represented *in vivo*. Four drugs were hits in both the *in silico* and

in vivo analyses; these were amsacrine, conivaptan, vilazodone and dabigatran etexilate (the

prodrug formulation of dabigatran, hereafter referred to simply as dabigatran) (Table S2, Table

2). Approximately half of the drugs identified *in vivo* (berberine, ketanserin, sunitinib, ceritinib,

fedratinib, nelfinavir, saquinavir, indacaterol, dabigatran) have been shown to ameliorate LPS

effects, and to function as mammalian anti-sepsis agents (20-40), suggesting conservation

between zebrafish and mammals in terms of the effects of these small molecules. The remaining

hits from the *in vivo* screen – ziprasidone, pergolide, vilazodone, bedaquiline, rucaparib,

pralsetinib, amsacrine, conivaptan, deserpidine, and lomitapide – are, to our knowledge, novel

mediators of LPS toxicity (Table 2). Despite some variability, there was an overall trend toward

decreased inflammatory cytokine production when a subset of these drugs were co-administered

with LPS and compared to LPS treatment alone (Fig. S2 A-L). Thus, these novel mediators of

LPS toxicity may function as anti-inflammatory agents.

Drug Category	Drug identified from current zebrafish LPS screen	Main Mechanism of Action	Drug previously tested with LPS in mammals?/ Novel	Drug previously tested in sepsis?/ Novel
Dopamine and Serotonin Modulators	Ziprasidone	Dopamine and serotonin antagonism	Novel	Novel
	Pergolide	Dopamine receptor agonist	Novel	Novel
	Vilazodone	Inhibits serotonin transporter, partial agonist at serotonin receptors	Novel	Novel
	Brexpiprazole	Dopamine and serotonin receptor modulation	Novel	Novel
	Ketanserin	Serotonin receptor antagonist	Mouse (Liu et al., 2013, Sugino et al., 2009)	Clinical trial for septic shock (Vellinga et al., 2015)
Chemotherapeutics	Rucaparib	Inhibits poly(ADP-ribose) polymerase	Novel	Novel
	Sunitinib	Inhibits receptor tyrosine kinases	Mouse (Zhao et al., 2019)	Novel
	Fedratinib	Inhibits janus receptor kinase 2	Human macrophages (Akcora et al., 2019)	Novel
	Pralsetinib	Rearranged during transfection kinase inhibitor	Novel	Novel
	Ceritinib	Inhibits autophosphorylation of anaplastic lymphoma kinase	Human endothelial cells (Dayang et al., 2021, Hu et al., 2019)	CLP induced sepsis in mice (Hu et al., 2019)
	Amsacrine	DNA intercalator	Novel	Novel
Antiviral/ Antibacterial Agents	Nelfinavir	Protease inhibitor	Human macrophages (Wallet et al., 2012)	CLP induced sepsis in mice (Weaver et al., 2004)
	Saquinavir	Protease inhibitor	Mouse (Yao et al., 2023, Tong et al., 2021), Human lung cells (Zhang et al., 2018)	CLP induced sepsis in mice (Tong et al., 2021)
	Ledipasivir	Inhibits viral phosphorylation	Novel	Novel
	Bedaquiline	Blocks proton pump for ATP synthase	Novel	Novel

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234   **Table 2: Small molecules protecting zebrafish embryos from LPS toxicity**

Drug Category	Drug identified from current zebrafish LPS screen	Main Mechanism of Action	Drug previously tested with LPS in mammals?/ Novel	Drug previously tested in sepsis?/ Novel
Miscellaneous	Indacaterol	β-adrenoreceptor agonist	Guinea pigs (Aritake et al., 2021), Human lung macrophages (Maarsingh et al., 2021)	Novel
	Conivaptan	Vasopressin receptor antagonist	Novel	Novel
	Deserpidine	Vesicular monoamine transporter blocker	Novel	Novel
	Lomitapide	Microsomal triglyceride transfer protein inhibitor	Novel	Novel
	Dabigatran	Thrombin inhibitor	Mouse (Kimura et al., 2020)	Mouse sepsis models, including CLP (Lei et al., 2021), *S. aureus* bacteremia in humans (Butt et al., 2021)
	Berberine	Activates AMPK (increases glycolysis)	Mouse (Chen et al., 2022), rats (Chen et al., 2021, Zhao et al., 2023), mouse macrophage cells (Zhao et al., 2023), bovine endometrial epithelial cells (Fu et al., 2023)	CLP-induced sepsis in mice (Chen et al., 2023), cecal slurry-induced sepsis (Li et al., 2021), *S. aureus*-induced septic arthritis in mice (Asila et al., 2022)

236   **Table 2: Small molecules protecting zebrafish embryos from LPS toxicity**
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Dabigatran reduces LPS mediated inflammation and mortality

Given the availability of tools to study coagulation in zebrafish (41, 42) we decided to

characterize further the one anticoagulant, dabigatran, that was identified through both *in silico*

and *in vivo* analyses. Although a prior study has shown only a modest effect of dabigatran on

lethality in sepsis models (31), we found that it completely averted LPS-induced lethality (Fig.

3A). Expression of inflammatory cytokines and procoagulant genes were downregulated in fish

co-treated with LPS and dabigatran compared to LPS alone (Fig. 3B). As previously shown (43),

time to venous occlusion was increased in dabigatran treated fish, indicating successful

anticoagulation (Fig. 3C). There was no additive effect on occlusion with co-treatment.

Figure 3: The anticoagulant dabigatran mitigates LPS-induced lethality and reduces the

expression of inflammatory genes. (A) Zebrafish were administered a lethal dose of LPS with

25 μM dabigatran along with appropriate controls and survival was tracked daily from 3 through

6 dpf. (B) Expression of inflammatory genes was evaluated by qPCR 3 hours after drug

administration in 3 dpf larvae treated with LPS and dabigatran. Pooled larvae (15-45) from a

clutch were used, where each point represents a separate clutch, and the average fold change in

LPS-treated fish vs. controls is graphed. Bars indicate mean and standard deviation. Student's t-

test (two-tailed) on $\Delta\Delta C_t$ values was used to determine significance. *$P<0.05$. (C) Time to

occlusion (TTO) after laser injury in the venous circulation was measured in 3 dpf zebrafish

larvae 3 hours after LPS and dabigatran administration. Each point represents a single larva, and

the observer was blinded to the condition. $P<0.05$ was considered significant by Mann-Whitney

U testing.

We next determined whether dabigatran administration could produce a more general

anti-inflammatory effect. Tail transection, an established method to induce inflammation, was

performed on 3 dpf larvae. We found that dabigatran did not reduce neutrophil or macrophage

recruitment to the tail transection site, suggesting that its protection might be more specific to

LPS rather than representing a general anti-inflammatory response (Fig. S3).

Dabigatran protects zebrafish from LPS toxicity independent of its anticoagulant function

An anti-sepsis effect of dabigatran mediated through fibrinogen-like protein 2 (Fgl2) has been

described (31). To determine whether the protection from LPS toxicity was dependent on the

anticoagulant effect of dabigatran, we evaluated three other commonly used anticoagulants – two

affecting factor Xa, rivaroxaban and apixaban, as well as another thrombin inhibitor, argatroban.

Rivaroxaban and apixaban were previously shown to be active in zebrafish and were used here at

the same concentration (43). However, we required shorter incubation times due to the rapid

lethality induced by LPS. As expected, all anticoagulants increased the TTO (Fig. 4A, Fig. S4 A,

B). Interestingly, some of the anticoagulants appeared to have a synergistic effect with LPS,

significantly increasing the time to occlusion when compared to the drug alone (Fig. 4A, Fig. S4

A, B). However, none of these conferred protection from LPS toxicity (Fig. 4B, Fig. S4 C, D),

suggesting that anticoagulation might not be the mechanism through which dabigatran mitigates

LPS-induced endotoxemia.

Figure 4: Dabigatran protection from LPS toxicity occurs independently of prothrombin.

(A) Time to occlusion (TTO) after laser injury in the venous circulation was measured in 3 dpf

zebrafish larvae 3 hours after LPS and argatroban administration. Each point represents a single

larva, and the observer was blinded to the condition. P<0.05 was considered significant by

Mann-Whitney *U* testing. (B) Zebrafish were treated with argatroban and/or LPS at 3 dpf and

observed daily until 6 dpf for survival. (C) Three dpf larvae from *f2* heterozygous incrosses were

divided into four groups: untreated controls, 40 μg/mL LPS, 25 μM dabigatran, and

LPS/dabigatran. All groups were followed for 3 days for survival. (D) Genotype distribution of

surviving larvae.

To further test this hypothesis, we employed a prothrombin knockout background (null

mutation in the *f2* gene (43). Larvae from *f2* heterozygous incrosses were divided into four

groups: untreated controls, LPS, dabigatran, and LPS + dabigatran. Each group was followed for

survival over 3 days. LPS alone was lethal in all genotypes (Fig. 4C), revealing that thrombin

inhibition is not necessary for protection from LPS. The presence of homozygous mutants was

confirmed by genotyping all clutches, except for LPS alone due to their rapid degradation (Fig.

4D). Thus, dabigatran administration protected *f2* homozygous mutants from lethal LPS,

proving that its mechanism of protection occurs independently of thrombin inhibition.

Dabigatran administration reduces LPS mediated nitric oxide production and apoptosis

Nitric oxide (NO) production can contribute to the hypotension seen in septic shock (44), and

LPS has previously been showed to increase NO (44). We confirmed that LPS administration

increased NO production in our system (Fig. 5A, B). Additionally, dabigatran has previously

been found to suppress nitric oxide synthase (45). We therefore tested whether dabigatran

treatment could reduce NO production and found this to be the case (Fig. 5A, B).

Figure 5: Dabigatran reduces LPS mediated nitric oxide production.

(A) Representative images of 3 dpf fish stained with DAF-FM DA to visualize nitric oxide

production after the indicated treatment. (B) ImageJ analysis showing mean fluorescence

intensity of images of 3 dpf DAF-FM DA stained fish with the indicated treatment; observers

were blinded to condition. Each point represents a different fish, and the results include 3

separate experiments. Student's t-test (two-tailed) on $\Delta\Delta C_t$ values was used to determine

significance. *$P<0.05$.

RNA sequencing data suggested that apoptosis related pathways were increased in LPS-

treated fish (Fig. 6A). As NO production has been shown to lead to apoptosis after LPS exposure

(46), we next tested whether this represented a mechanism by which protection from LPS is

conferred. We found that cell death (Fig. 6B, C) and apoptosis (Fig. 6D, E) were significantly

decreased in dabigatran and LPS co-treated fish. These data suggest that decreased NO

production and apoptosis might be mechanisms by which dabigatran protects against LPS

toxicity.

Figure 6: Dabigatran reduces LPS mediated cell death and apoptosis.

(A) KEGG pathway analysis of RNA-seq data from whole LPS treated fish compared to

controls. Significantly changed genes (adjusted p-value <0.05) from RNA-seq data on pooled 3

dpf LPS-treated vs. control fish (consisting of 30-45 embryos per category) performed in

triplicate were used for further analyses. These genes were converted to human versions and then

submitted for KEGG Pathway Analysis using the Enrichr platform, a publicly available tool

developed by the Ma'ayan lab at Icahn School of Mount Sinai, NY (Chen et al., 2013, Kuleshov

et al., 2016). Overlap score is the number of significantly changed genes over the total number

of genes in the pathway. Combined score considers the z-score (a value's relationship to the

mean) in addition to the adjusted p-value. The combined score is used to assess the significance

of gene set enrichment findings, with higher values being more significant. Adjusted p-value

<0.05 is considered significant. Apoptosis related pathways are shown. For complete analysis,

see Table S1. (B) Representative images of 3 dpf fish stained with acridine orange after the

indicated treatment. (C) ImageJ analysis showing mean fluorescence intensity of images of 3 dpf

acridine orange stained fish with the indicated treatment; observers were blinded to condition.

Each point represents a different fish, and the results include 3 separate experiments. Student's t-

test (two-tailed) on $\Delta\Delta C_t$ values was used to determine significance. *P<0.05. (D) Representative

images of 3 dpf fish stained with TUNEL after the indicated treatment. (E) Apoptosis was scored

in 3 dpf TUNEL stained fish with the indicated treatement; observers were blinded to condition.

Each point represents a different fish, and the results include at least 2 separate experiments.

Student's t-test (two-tailed) on $\Delta\Delta C_t$ values was used to determine significance. *P<0.05.

Discussion:

We have identified several small molecule mediators of LPS toxicity that can act as anti-

inflammatory agents. Approximately half had been previously explored in mammalian LPS

models, and all have been shown to reduce inflammation and ameliorate toxicity (20-23, 25-29,

32, 34, 35, 39, 40, 47-49), suggesting strong concordance between zebrafish and mammals.

These mediators were found to act through multiple mechanisms, including inhibition of TLR

and NFκB activation (25, 48) as well as inhibition of inflammation through NFκB-independent

mechanisms (26, 27). Other potential mechanisms include effects on apoptosis (22) and NOS

(nitric oxide synthase) activation (23). A smaller number of the identified drugs have also been

investigated in mammalian sepsis models (24, 31, 33, 35-37, 39, 50), all with beneficial effects,

indicating further concordance with our zebrafish data. The novel mediators of LPS toxicity that

have not previously been tested in other studies – amsacrine, bedaquiline, conivaptan,

deserpidine, lomitapide, pergolide, prasetinib, rucaparib, vilazodone, ziprasidone – have as yet

undetermined mechanisms of action and represent avenues for further research in both

mammalian inflammation and sepsis.

Clinical trials using anticoagulants in the context of sepsis have proven equivocal at best

(51-60), although it is still possible that specific subtypes of sepsis patients may benefit from

anticoagulant therapy. Previous studies have shown that dabigatran reduces inflammation ((31,

32) and can enhance survival in mouse models of sepsis (31), congruent with a cohort study

revealing that patients on dabigatran had a lower incidence of *Staphylococcus aureus* bacteremia

(33). Dabigatran has also been found to mitigate inflammation in multiple contexts, including

diet (61), oxysterols (62), and acute myocardial infarction (45). However, it is unclear whether

these beneficial effects were due to the anticoagulant function of dabigatran. Our data suggest

that dabigatran reduces inflammation and promotes survival from LPS toxicity through a novel

mechanism independent of its direct anticoagulant effect on thrombin.

We found that dabigatran administration reduces LPS mediated NO production and

apoptosis, suggesting that this is one possible mechanism by which dabigatran confers protection

from LPS. Dabigatran has previously been shown to suppress nitric oxide synthase in acute

myocardial infarction (45) and future experiments will clarify the mechanism by which

dabigatran reduces LPS mediated nitric oxide production.

There are several limitations of this study. Due to the stringency of the screen (mortality),

it is likely that other small molecules conferring beneficial effects but not affecting mortality

(e.g., those that reduce inflammation) were not identified. This study also does not necessarily

identify the mechanism by which each drug acts to reduce LPS toxicity. It is possible that the

effect is mediated through a non-primary or previously unknown mechanism of action (as is the

case with dabigatran).

Our data suggest that the novel small molecule mediators of LPS toxicity uncovered

through our screen can be explored as anti-inflammatory and anti-sepsis agents in mammals.

Given that the hits previously examined in mammalian systems were also found to have

beneficial effects in this zebrafish model, we expect that at least some of the novel drugs will

translate to mammalian models.

Previous work and our data suggest that dabigatran can act as an anti-inflammatory and

potential anti-sepsis agent (31-33). We show here definitive proof that dabigatran acts

independently of thrombin to exert its rescue of lethal endotoxemia. Venous thromboembolism

prophylaxis is the standard of care for most adult hospitalized inpatients, especially those

admitted for sepsis (63, 64). The most common agents used are unfractionated and low

molecular weight heparins, but the data from our study suggest the possibility that dabigatran

could be an alternative with a dual benefit –both anticoagulant and completely distinct anti-LPS

functions. Randomized controlled trials in human inflammatory conditions or inpatients would

be relatively easy to support given the successful use of direct oral anticoagulants and the

indication for anticoagulation in hospitalized patients (63, 64).

Materials and Methods:

Animal care and husbandry:

Zebrafish (*Danio rerio*) were raised in accordance with animal care guidelines as approved by

the University of Michigan Animal Care and Use Committee. Wild-type fish were a hybrid line

generated by crossing AB and TL strain zebrafish acquired from the Zebrafish International

Resource Center. Genetic knockouts of prothrombin (*f2*) have been previously described (43).

Tricaine (Western Chemical) was used for anesthesia and rapid chilling in an ice water bath for

euthanasia.

Small molecule treatments:

Small molecule screens (MedChemExpress FDA drug repurposing set) were started at 3 dpf,

where 3 larvae were plated per well in 96-well plates containing embryo media. All drugs were

initially tested at a concentration of 25 μM for resistance to LPS toxicity, and confirmatory

studies were performed with a larger group of fish (9-45) representing at least three different

clutches (3 separate experiments), where the drugs were tested at various concentrations.

Heart rate measurements:

Heart rate was measured 3 hours after lethal LPS (40 μg/mL) administration and compared to

untreated controls. Larvae were mounted in 0.8% low melting point agarose and visualized using

an Olympus IX73 microscope. Heart rate was manually counted over 60 seconds for each larva.

Each experiment consisted of at least 15 larvae. At least three different clutches (3 different

experiments) were utilized.

Real-time quantitative polymerase chain reaction (RT-qPCR):

RT-qPCR was performed as described previously (65). Total RNA was extracted from 3 dpf

whole larvae 3 hours post treatment using the RNeasy Mini Plus kit (Qiagen) and transcribed

into cDNA using Superscript III cDNA synthesis kit (Invitrogen). Transcript levels of genes of

interest were quantified by qPCR (Mastercycler realplex^2, Eppendorf) using Fast SYBR Green

Mix (Applied Biosystems,   Waltham, MA). Three equal pools of whole larvae (15-45 embryos

per pool) were tested. Primer sequences are provided in Table S3. All expression data were

normalized to the *actb2* gene, and significance was analyzed using the double delta C_t method as

described (66).

RNA-seq preparation and analysis:

RNA-seq was performed as described previously (65). Three biological replicate pools of 30-45

3 dpf whole embryos were used per condition. Total RNA was extracted, and cDNA was

prepared as above. Samples were sent to Novogene (Sacramento, CA) for further processing.

Paired-end 150 base pair (bp) sequencing was obtained with the Illumina NovaSeq 6000

Sequencing System. Reads were mapped to exonic regions based on the zebrafish reference

genome (GRCz11) using STAR36 (v.2.7.9a) and featureCounts37 (v.2.0.3). Differential

expression analysis was conducted using the R package DESeq2 ((67). All significant

differentially expressed genes (FDR adjusted p-values) were converted to human genes using the

R package babelgene. KEGG Pathway Analysis was then performed using the Enrichr platform,

a publicly available tool (68, 69). Overlap score is the number of significantly changed genes

over the total number of genes in the pathway. The combined score considers the z-score (a

value's relationship to the mean) in addition to the adjusted p-value; this score has been found to

outperform other ranking metrics (69). The combined score is used to assess the significance of

gene set enrichment findings, with higher values being more significant; values > 30 have been

considered significant in literature (18, 70).

Laser-mediated vascular endothelial injury:

Larvae were treated with LPS and drug starting at 3 dpf and collected after 3 hours for

assessment of time to clot formation, performed as previously described (17, 18, 41, 43). Briefly,

3 dpf larvae were mounted in 0.8% low melting point agarose and visualized using an Olympus

IX73 microscope with an attached Micropoint focusing system and pulsed-dye laser (Andor).

For venous laser injury, 99 laser pulses were administered to the ventral surface of the posterior

cardinal vein, 5 somites caudal to the anal pore. For arterial laser injury, 99 laser pulses were

administered 5 somites caudal to the anal pore on the dorsal surface of the dorsal aorta. Time to

occlusion (formation of an occlusive thrombus) was measured, where 120 seconds was the

maximum time each larva was observed. Three biological replicates were performed for each

experiment.

In silico analysis:

Differential gene expression data were retrieved from the RNAseq results as previously

described (65). The human orthologues of the respective genes were obtained from the Zebrafish

Information Network database (https://zfin.org/downloads/archive/2023.04.26) and the CLUE

software platform was utilized in performing Connectivity Map-based analyses

(https://www.sciencedirect.com/science/article/pii/S0092867417313090 (access date April 26,

2023)). Genes were ranked based on the adjusted p-values for the gene expression level changes,

and the top 150 upregulated or down-regulated genes were tested for the gene connectivity

evaluations across 64 human cell lines and disparate perturbation profiles. The connectivity data

were processed by using cmapR library in R (v4.2.2) environment (Natoli T (2023). cmapR:

Cmap Tools in R. R package version 1.12.0, https://github.com/cmap/cmapR). The data were

further filtered down to 27 cell lines of which most were hematopoietic/hepatic/pulmonary/non-

tumorigenic origins, or endothelial phenotypes (A375, A549, ASC, BJAB, CD34, HA1E,

HCC515, HEK293, HELA, HEPG2, HPTEC, HUH7, HUVEC, JURKAT, K562, MCF10A,

NCIH508, NCIH596, NOMO1, OCILY19, OCILY3, PHH, THP1, TMD8, U2OS, U937,

YAPC).

Acridine Orange Staining:

Larvae were incubated in 10 µg/ml Acridine Orange stain (Sigma) for 15 minutes in embryo

media   with LPS treatment and then rinsed three times before mounting in 0.8% agarose

and imaged with an Olympus IX73 microscope. Quantification of intensity was performed in

ImageJ.

TUNEL Staining:

The TUNEL assay was performed according to the instructions provided in the In Situ Cell

Death Detection Kit, TMR red (Roche) for cryopreserved samples, with embryos cryopreserved

in Optimal Cutting Temperature Compound (OCT) and cryosectioned prior to labeling.

Images were quantified by an observer blinded to condition using a visual scoring system. The

visual scoring system consisted of grades 0 through 3 (0 = 0-9% fluorescence in visual field, 1 =

10-20% fluorescence, 2 = 30-70% fluorescence, 3 = 80-100% fluorescence).

NO Staining:

Larvae were split into four treatment groups: control, LPS, dabigatran, and LPS with dabigatran.

Each treatment group was simultaneously incubated with 5 uM of DAF-FM-DA solution in the

dark at 25°C as previously described (Lepiller et al., 2007). After 2 hours larvae were washed

with embryo media and mounted in 0.8% agarose. Imaging was done using an Olympus IX73

microscope. Fluorescence quantification was obtained utilizing ImageJ.

Tail transection:

Tg(lyzC:GFP)[nz117] and TgBAC(mpeg1.1:EGFP)[vcc7] transgenic larvae were anesthetized at 5 days post fertilization and tail transection was performed with a sterile scalpel blade. Wounded larvae were transferred into fresh embryo media with 25 µM dabigatran and incubated at 28.5 °C for 5 hours prior to imaging using a Nikon SMZ25 stereoscope. Leukocyte recruitment was quantified by measuring the fluorescent pixel area within 100 µm of the wound site in FIJI/ImageJ.

Statistics:

Statistics were performed using the GraphPad PRISM package. $P<0.05$ was considered significant. For qPCR and RNA-seq data, Student's t-test on $\Delta\Delta C_t$ values was used to determine significance. For time to occlusion (TTO) data, Mann-Whitney U testing was used to determine significance.

Study Approval:

Approval for zebrafish studies was granted by the University of Michigan Institutional Animal Care and Use Committee, identification number PRO00010679 and the A*STAR Institutional Animal Care and Use Committee, identification numbers 211667 and 221694.

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Competing Interests:

J.A.S. has been a consultant for Sanofi, Takeda, Genentech, CSL Behring, Pfizer, Medexus and NovoNordisk. The other authors declare that they have no competing interests to disclose.

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Data Availability:

RNAseq data have been submitted to NCBI Gene Expression Omnibus and are available under accession number GSE245299. Any other original data can be obtained by contacting the corresponding author at vijacob@umich.edu.

541 **Author Contributions:**

542 V.J., J.A.S., S.O. and A.F. conceived and planned experiments. V.J., A.F., R.M., W.W. and X.S.

543 performed and analyzed all experiments and M.Y. performed *in silico* analysis. V.J. wrote the

544 manuscript, with input from all authors. J.A.S. revised the manuscript.

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Figure S1: Multiple FDA-approved drugs protect zebrafish larvae from a lethal dose of LPS. (A)-(U) Zebrafish were administered a lethal dose of LPS along with compounds from an FDA-approved repurposing library, and survival was tracked daily from 3 through 6 dpf. Any experimental controls comprised the same solvent volume of each drug, either DMSO or water.

Figure S2: Co-administration of identified drugs with LPS decreases inflammatory cytokine expression. (A)–(L) Expression of inflammatory genes was evaluated by qPCR 3 hours after drug administration in 3 dpf larvae treated with LPS, and each represented small molecule. Pooled larvae (15-45) from a single clutch were used, where each point represents a separate clutch, and the average fold change in LPS-treated fish vs. controls is graphed. At least two biological replicates were performed. Bars indicate mean and +/- 1 standard deviation. Student's t-test on $\Delta\Delta C_t$ values was used to determine significance. *$P<0.05$.

Figure S3: Tail transection does not result in increased neutrophil or macrophage migration to the site of injury. 5-day-post-fertilisation (dpf) Tg(lyzC:GFP)[nz117] and TgBAC (mpeg1.1: EGFP)[vcc7] transgenic embryos were anesthetized and tail transaction was performed using a sterile surgical blade at the distal end of the notochord. After transection, embryos were transferred into fresh E3 medium containing 25 μM dabigatran and kept at 28.5 °C. The recruitment of neutrophils (A) and macrophages (B) to the wound site was imaged at 5 hours post transection and quanitified by measuring fluorescence at the wound site.

Figure S4: Other direct anticoagulants do not protect zebrafish larvae from LPS toxicity.

(A),(B). Time to occlusion (TTO) after laser injury in the venous circulation was measured in 3

dpf zebrafish larvae 3 hours after LPS and anticoagulant administration. Each point represents a

single larva, and the observer was blinded to the condition. $P<0.05$ was considered significant by

Mann-Whitney U testing. (C), (D) Zebrafish were treated with apixaban or rivaroxaban and/or

LPS at 3 dpf and observed daily until 6 dpf for survival.

Figure 1

A

B

C

Figure 2

A

Name of Pathway	Overall Ranking (out of 98 significantly changed pathways)	Overlap Score	Adjusted P-value	Combined Score
Pathogenic *E.coli* Infection	1	38/197	2.59E-10	127.749924
Leukocyte Transendothelial Migration	9	21/114	8.13E-06	65.1873171
Complement and Coagulation Cascades	8	18/85	7.42E-06	78.4936729
TNF Signaling	12	20/112	1.84E-05	57.8681584
Platelet Activation	70	14/124	0.017	13.16661224

B

C

D

E

F

Figure 3

Figure 4

Figure 5

A

DMSO

100 µm

LPS

100 µm

Dabigatran

100 µm

LPS + Dabigatran

100 µm

B

****

Figure 6

A

Name of Pathway	Overall Ranking (out of 98 significantly changed pathways)	Overlap Score	Adjusted P-value	Combined Score
Apoptosis	11	23/142	1.84 E-05	51.783249

B

DMSO

LPS Control

Dabigatran

Dabigatran + LPS

C

***

Mean Fluorescence Intensity

DMSO   LPS   Dabigatran   Dab.+LPS

D

DMSO   100 µm

LPS   100 µm

Dabigatran 100 µm

Dabigatran + LPS   100 µm

E

**

Flourescence Assesment Score

DMSO   LPS   Dabigatran   Dab. + LPS