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ILLUSTRATED REVIEW



Fishing for answers to hemostatic and thrombotic disease: Genome editing in zebrafish

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Abstract

Over the past two decades, the teleost vertebrate Danio rerio (zebrafish) has emerged as a model for hemostasis and thrombosis. At genomic and functional levels, there is a high degree of conservation of the hemostatic system with that of mammals. Numerous features of the fish model offer unique advantages for investigating hemostasis and thrombosis. These include high fecundity, rapid and external development, optical transparency, and extensive functional homology with mammalian hemostasis and thrombosis. Zebrafish are particularly suited to genome-wide mutagenesis experiments for the study of modifier genes. They are also amenable to whole-organism small-molecule screens, a feature that is exceptionally relevant to hemostasis and thrombosis. Zebrafish coagulation factor knockouts that are in utero or neonatal lethal in mammals survive into adulthood before succumbing to hemorrhage or thrombosis, enabling studies not possible in mammals. In this illustrated review, we outline how zebrafish have been employed for the study of hemostasis and thrombosis using modern genome editing techniques, coagulation assays in larvae, and in vivo evaluation of patient-specific variants to infer causality and demonstrate pathogenicity. Zebrafish hemostasis and thrombosis models will continue to serve as a clinically directed basic research tool and powerful alternative to mammals for the development of new diagnostic markers and novel therapeutics for coagulation disorders through high-throughput genetic and small-molecule studies.

KEYWORDS

coagulation, genetics, genome editing, hemostasis, thrombosis, zebrafish

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Graphical Abstract

This graphical abstract outlines our review of the use of zebrafish to study disorders of hemostasis and thrombosis. We initially introduce the zebrafish and its associated technologies. This is followed by a review of the models of hemostatic and thrombotic disease that have been produced using genome editing. Finally, we review application of zebrafish to studies of patient coagulation factor variation.

Essentials

- The zebrafish hemostatic system is highly conserved with mammals genomically and functionally.
- Hemostatic disease models produced using genome editing show conservation with human disorders.
- Zebrafish coagulation factor mutants that are early lethal in mammals survive into adulthood.
- Potential disease-causing variants can be rapidly assessed in vivo using zebrafish knockouts.



The teleost-specific genome duplication with the presence of duplicated paralogs gives a unique opportunity to identify previously undiscovered gene functions.³⁶

Comparison of platelets and thrombocytes Platelets Thr ♦ Nucleated with dense chromatin, Anucleate with granular cytoplasm⁴¹ cytoplasmic projections44 ♦ Produced in large numbers vastly out of Relative numbers similar to white cells⁴⁵ proportion to nucleated blood cells ♦ Function similar to platelets in adherence ♦ Plasma membrane invades the platelet and aggregation⁴⁶ interior to make a surface-connected ♦ Stimulated by multiple platelet agonists: canalicular system⁴² eg, ADP, arachidonic acid, collagen⁴⁷ ♦ Receptors mediate hemostasis and other ♦ Receptors for adhesion and aggregation functions43 are conserved48 Unique to mammals \diamond Present in fish, birds, reptiles, and TROMBOC amphibians Platelet Megakaryocyte Thrombocyte Zebrafish thrombocyte spreading on fibrinogen is similar to human platelets.⁴⁹ During activation, human platelets form filopodia and lamellipodia extensions,⁵⁰ zebrafish thrombocytes also develop pseudopod-like protrusions after activation. Platelet Spreading in Humans Late activation Inactive Early activation Active

cDNA rescue: Both human and zebrafish at3 cDNA injected into at3^{-/-} single-cell

embryos rescue the delayed occlusion after laser-mediated venous endothelial injury.59

cDNA transgene

The role of fibrinogen in zebrafish hemostasis Fibrin clot Fibrin(ogen) incorporates into Loss of fibrinogen results in hemorrhage in a developing clot in zebrafish multiple tissues larvae fga mutants exhibit spontaneous hemorrhage in Fluorescently tagged fibrinogen the jaw, abdomen, muscle, and fin, as well as the demonstrates that fibrin is the forebrain, midbrain, and hindbrain (inset).35,66 primary component of thrombi Hemorrhage occurs later than common pathway produced through endothelial mutations, with partial lethality by one year of age. injury in the venous circulation.¹¹ fga^{-/-};nfe2^{-/-} Loss of both fga and nuclear fga-/-;nfe2-/factor erythroid 2 (nfe2) exhibit synthetic lethality fga-/-:nfe2-/- double mutant zebrafish die by one year of age due to Incubation in afibrinogenemia and fga cDNA/human fibrinogen infusion ε-aminocaproic acid thrombocytopenia.35 posterior cardinal vein occlusion fga^{-/-} no occlusion fga-/fga mutant larval fish exhibit a hemostatic defect As in mammals, targeted mutation of fga results in loss of fibrinogen, and absence of

thrombus formation following endothelial injury.^{35,67} This can be rescued by the expression of zebrafish *fga* cDNA or human fibrinogen infusion. Surprisingly, incubation with ε -aminocaproic acid also partially reverses this hemostatic defect.³⁵

Vwf in zebrafish: zebrafish Vwf forms high molecular multimers and is packaged into Weibel-Palade bodies.⁷⁰ Loss of Vwf results in a higher time to occlusion in *vwf^{-/-}* mutant fish than the wild type larvae in both the arterial and venous circulation.⁷¹

. zebrafish include:

- absence of birth trauma
- L differential hemostatic challenges in an aquatic environment
- lower blood pressures in zebrafish
- □ species-specific genetic differences resulting in a differential baseline hemostatic balance^{60,72}

(black and red lines indicate average age of death for each mutant line)

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Evaluation of human genetic variants in zebrafish hemostasis and thrombosis knockout models

AT3 thrombosis variants	Affected domain(s)	Ability to rescue <i>at3</i> mutant DIC phenotype ⁵⁹
Leu99Phe ^{73,74}	heparin binding domain	complete
Lys114Glu ^{75,76}	heparin binding domain	complete
Arg393His ^{77,78}	reactive center loop (RCL) : P1 Arg mutation	none
Ala404Asp ⁷⁹	pleiotropic effect on RCL and heparin binding domain	partial

F5 deficiency variants	Affected domain(s)	Ability to rescue <i>f5</i> mutant phenotype ³¹
Ser83Arg	A1	complete
Gly97Asp	A1	none
Tyr1702Cys ⁸⁰	A3	none
Arg2074Cys ⁸¹	C2	complete
Arg2187Cys	C2	complete

F10 deficiency variants	Affected domain(s)	Ability to rescue <i>f10</i> mutant phenotype ³⁴
Arg68Cys	EGF1	none
Gly173Typ	activation peptide	none
ΔT176_Q186 deletion	activation peptide	none
Gly262Asp	protease	none
lle323Met	protease	partial
Cys390Phe	protease	none
GIn416Leu	protease	partial

FGA variants	Affected domain(s)	Ability to rescue <i>fga</i> mutant phenotype ³⁵
Met1Val ⁸²	pre-peptide	none
Cys55Gly ⁸³	N-terminus	complete
Cys64Tyr ⁸⁴	N-terminus	complete
Tyr809Cys ⁸⁵	α _E C domain	none

Since hemostasis must occur within a closed circulatory system, consisting of flowing blood, proteins, cells, and endothelium, the zebrafish is superior to cell culture systems for assessing variants.

AUTHOR CONTRIBUTIONS

AR and JS developed the concepts and wrote the manuscript. AR and AF produced the illustrations. AR, AF, and JS approved the final content.

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RELATIONSHIP DISCLOSURE

JAS has been a consultant for Sanofi, Takeda, CSL Behring, HEMA Biologics, and Bayer. AR and ACF report no conflicts of interest.

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