

Subretinal fluid in rhegmatogenous retinal detachment: potential biomarkers and therapeutic targets for proliferative vitreoretinopathy

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Abstract

• Rhegmatogenous retinal detachment (RRD) is a serious ocular condition marked by the separation of the neuroretina from the retinal pigment epithelium (RPE). The pathogenesis of RRD involves intricate molecular and cellular mechanisms, including inflammation, cell migration, and the activation of proliferative signaling pathways. One of the most challenging complications of RRD is proliferative vitreoretinopathy (PVR), which refers to the proliferation and contraction of fibrocellular membranes on the retinal surface and in the vitreous cavity. PVR is a major cause of surgical failure in RRD, as it can lead to recurrent retinal detachment and severe vision loss. However, the pathogenesis of PVR is not yet fully understood, and the treatment options are quite limited. Recent advances in analytical techniques have offered valuable insights into the molecular alterations present in the subretinal fluid (SRF) of patients with RRD. This review seeks to consolidate the current knowledge regarding the SRF profile in RRD and PVR, emphasizing potential biomarkers and therapeutic targets.

• **KEYWORDS:** proliferative vitreoretinopathy; rhegmatogenous retinal detachment; subretinal fluid

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INTRODUCTION

Rhegmatogenous retinal detachment (RRD) is a major cause of visual impairment worldwide^[1], with incidence rates varying across different populations and age groups. The annual incidence of RRD is estimated to range from 6.3 to 17.9 cases per 100 000 individuals^[2]. In recent years, studies have shown that the incidence of RRD has been on the rise^[3]. Intraocular fluid, including subretinal fluid (SRF), plays a pivotal role in the pathophysiology of the disease^[4]. Under normal conditions, the vitreous humor helps maintain the structural integrity of the eye and facilitates nutrient transport^[5]. However, in RRD, changes in the vitreous and the accumulation of SRF are critical drivers of disease onset and progression^[6]. The biochemical characteristics of SRF changes with the duration of the disease^[7]. This fluid exacerbates retinal detachment by exerting hydrostatic pressure, while its biochemical components—including inflammatory mediators, growth factors, and proteins—play a significant role in cellular responses. These factors have a direct impact on the severity and prognosis of RRD^[8-10]. For instance, elevated levels of certain cytokines within SRF can promote inflammation and tissue remodeling, contributing to the development of proliferative vitreoretinopathy (PVR)^[11-12], a common and challenging complication of RRD. Additionally, analyzing the composition of SRF provides valuable insights into the cellular and molecular mechanisms underlying the disease. It is instrumental in identifying potential therapeutic targets and advancing treatment strategies^[13]. This review systematically summarizes the molecular pathological features, biomarkers, and clinical significance of SRF in the context of RRD, offering a comprehensive exploration of its role in the disease process.

MOLECULAR COMPONENT ANALYSIS OF SRF IN RRD

Cytokines and Chemokines Interleukin-6 (IL-6) is a pleiotropic cytokine that plays a critical role in both

physiological and pathological processes. Research has demonstrated that elevated IL-6 production by vitreous cells can stimulate the proliferation of vascular endothelial cells, thereby contributing to the pathophysiology of PVR^[14]. Study reported significantly higher levels of IL-6 in both SRF and the vitreous humor^[15]. Furthermore, evidence indicates a strong correlation between IL-6 levels and the matrix metalloproteinases (MMP)/tissue inhibitors of metalloproteinases (TIMP) ratio in SRF, suggesting that IL-6 may drive the increased MMP/TIMP ratio observed during PVR progression^[16].

Chemokines are small proteins that play a critical role in regulating the migration of various leukocyte types to sites of inflammation. Research indicates that several chemokines C-C motif ligand (CCL), such as CCL2, CCL3, CCL8, CCL9, CCL10, CCL11, CCL17, CCL18, CCL19, CCL22, and macrophage migration inhibitory factor (MIF), exhibit significantly elevated levels in SRF^[17].

It has been observed that patients who developed PVR exhibited significantly higher total levels of vascular endothelial growth factor (VEGF) in the SRF compared to RRD patients who did not^[18]. It was found that VEGF is alternatively spliced to form the angiogenic VEGF_{xxx} and antiangiogenic VEGF_{xxx}b family of isoforms. Total VEGF levels are elevated in the PVR group, with the anti-angiogenic VEGF_{xxx}b isoform being predominant in SRF^[19]. However, another study reported no significant differences in VEGF levels among patients with PVR, while noting that pigment epithelium-derived factor (PEDF) levels were significantly elevated compared to the control group^[20].

Transforming growth factor-beta (TGF- β) serves a wide range of functions, including promoting the production of the extracellular matrix. Furthermore, TGF- β accelerates fibrous tissue formation and contributes to the progression of various disease processes^[21]. Notably, TGF- β_2 is produced by retinal pigment epithelial (RPE) cells and Müller cells. Elevated levels of TGF- β_2 have been detected in the SRF and are closely associated with PVR^[22].

Adipose tissue generates a diverse array of cytokines, collectively referred to as adipokines, which participate in both physiological and pathological processes^[23]. These adipokines have gained recognition as a distinct group of mediators that play pivotal roles in regulating inflammation, modulating immune responses, and facilitating wound healing. Studies revealed that levels of adiponectin, cathepsin S, and leptin are significantly elevated in the PVR group^[24].

Cytoskeleton, Adhesion and Migration PVR is recognized as a pathological wound-healing process, defined by the formation of fibrous membranes on both the inner and outer surfaces of the detached neuroretina. The MMP family exhibits

significant heterogeneity and consists of 24 distinct proteolytic enzymes. MMPs and their natural inhibitors, TIMPs, are believed to play crucial roles in the development of PVR. Research has demonstrated that the levels of MMPs and TIMPs in the vitreous humor and SRF of patients with RRD are elevated^[25], showing a strong correlation with the severity of PVR.

Fibulins (Fblns) are extracellular matrix proteins that play a crucial role as key components of elastic fibers and basement membranes. A study found that the upregulation of Fbln2, coupled with the downregulation of Fbln5 during RRD, may contribute to an imbalance in the adhesive interactions between the RPE and the neuroretina^[26].

Syndecan-1 (also known as CD138) is a crucial transmembrane glycoprotein belonging to the heparan sulfate proteoglycan (HSPG) family. It is predominantly found in epithelial cells and is characterized by heparan sulfate chains that facilitate interactions with various peptides, including extracellular matrix components and is potent mediators of proliferation, adhesion, and migration^[27]. Syndecans play vital roles in regulating wound repair processes, as well as in the formation and signaling of chemokines and growth factors. RRD is linked to a marked increase in the levels of soluble syndecan-1 within the vitreous humor and SRF. Moreover, the elevated concentration of soluble syndecan-1 has been shown to positively correlate with the duration of retinal detachment^[28]. Intercellular adhesion molecule-1 (ICAM-1) and vascular cell adhesion molecule-1 (VCAM-1) are critical in mediating leukocyte adhesion and migration to endothelial cells. These molecules also play a pivotal role in inflammatory responses and the activation of immune cells^[29]. Study has revealed a significant increase in the levels of soluble ICAM-1 and soluble VCAM-1 in patients with PVR^[30].

Lipid Metabolites Studies on lipid metabolites have revealed that the glycerophosphocholine lipid profile in SRF closely mirrors that of outer segments of the retina cone cells, despite the fact that human retina containing a higher proportion of rod cells. This observation suggests that lipids derived from cones may play a pivotal role in the pathogenesis of RRD^[31].

Lipid peroxidation is a well-established mechanism of cell membrane damage linked to various ocular diseases. One of its byproducts, malondialdehyde (MDA), is a secondary product of lipid peroxidation with a relatively long half-life, making it a reliable marker of peroxidative damage to cell membranes^[32]. Research has shown that MDA concentrations in SRF are correlated with the severity of retinal detachment, highlighting its potential as an indicator of oxidative stress in RRD patients^[33].

Amino Acids Amino acids serve as the essential building blocks of proteins and peptides, playing a crucial role in

cellular signaling and metabolic pathways that facilitate tissue repair. A deficiency in certain amino acids has been shown to trigger endoplasmic reticulum stress and promote the expression of specific growth factors^[34]. Notably, one study revealed significant alterations in amino acid concentrations in patients with RRD. Researchers found that compared to the vitreous humor of the control group, the levels of aspartate, arginine, glutamate, and glycine in the SRF of RRD patients were markedly elevated, while the concentrations of alanine, isoleucine, leucine, methionine, phenylalanine, threonine, tyrosine, and valine were reduced^[18]. Another study further highlighted that glutamate levels in both the SRF and vitreous humor of RRD patients were significantly higher than those in the control group^[35]. Excessive glutamate is believed to contribute to toxicity through mechanisms involving osmotic and calcium damage. Elevated glutamate levels can overstimulate sodium influx into cells, which, combined with the entry of water and chloride, may result in osmotic cell lysis and neuronal swelling^[36].

MicroRNA Expression The microRNA (miRNA) are key regulators of gene expression, controlling cellular functions under both physiological and pathophysiological conditions^[37]. miR-210, plays an important role in controlling lipid metabolism and neuronal function^[38]. The expression levels of miR21 and miR34 exhibit a positive correlation with the duration of symptoms in retinal detachment cases. Conversely, miR146a expression is notably reduced in patients diagnosed with PVR. Additionally, miR21 serves as a predictor for both the time elapsed from symptom onset to treatment and the number of retinal breaks. Furthermore, miR26a demonstrates a correlation with the best-corrected visual acuity (BCVA) following surgery. The expression of specific miRNAs is closely linked to risk factors for PVR, highlighting their potential as biomarkers and therapeutic targets in the management of this condition^[39].

Coagulation Cascade The activation of the coagulation cascade has been identified as a potential contributor to the development of PVR^[40]. Intravitreal thrombin stimulates retinal pigment epithelial cells to secrete pro-inflammatory and pro-fibrotic mediators, indicating its role in PVR pathogenesis. Studies have shown that levels of coagulation factor XII and the thrombin-antithrombin complex are elevated in SRF, reflecting significant thrombin generation in SRF of patients with RRD^[41]. Tissue factor (TF), a transmembrane glycoprotein and the primary initiator of blood coagulation in normal hemostasis, also plays a critical role. Research revealed that thrombin generation is induced in SRF, and the addition of TF-targeting antibodies almost completely neutralized this thrombin-activating effect^[42].

Other Biomarkers Soluble Fas (sFas) and soluble Fas ligand

(sFasL) play a crucial role in the development of PVR by regulating the apoptosis of RPE cells. Studies have revealed that sFas and sFasL levels are significantly elevated in individuals with PVR^[30].

The occurrence of PVR was also found to be correlated with several early apoptosis factors, including caspase-8, caspase-9, and B-cell lymphoma 2 (Bcl-2) associated death promoter. Additionally, early apoptosis factors and signaling proteins, such as ERK1/2, were shown to impact OCT features three months postoperatively^[43].

ATP levels showed an increase in SRF, followed by a gradual decline over time. This reduction aligned with the upregulation of ectonucleoside triphosphate diphosphohydrolase-1. These observed kinetics shed light on the pathological mechanisms driving the excessive accumulation of extracellular ATP after retinal detachment^[44].

Study has revealed that the concentration of transthyretin (TTR) is notably higher in SRF compared to serum and cerebrospinal fluid. TTR, which is abundantly expressed in the RPE, plays an essential role in transporting retinol, lipids, and other low-solubility molecules. This function may contribute to lipofuscin accumulation by influencing retinol-binding capacity. Additionally, modifications in TTR, such as glutathionylation, has been detected at significantly elevated levels in SRF compared to serum, suggesting a potential involvement in the pathogenesis of RRD^[4].

Researchers have identified that proteins involved in proteolysis play a significant role within the SRF in RRD patients. Notably, chitinase 3-like protein 1 (CHI3L1) and galectin-3-binding protein appear to be distinctive components of SRF^[45]. CHI3L1 belongs to the glycosyl hydrolase 18 (GH18) gene family, which has remained conserved across species and throughout evolution. This protein is known to be dysregulated in various conditions, including inflammatory, infectious, remodeling, and neoplastic disorders^[46].

S100B is a calcium-binding protein expressed in astrocytes, playing a key role in regulating various intracellular processes, including proliferation and differentiation. Research has shown that S100B concentrations in SRF increase following rhegmatogenous RRD, establishing it as a reliable marker of retinal stress. Furthermore, study has demonstrated a positive correlation between S100B levels and the duration of RRD^[47].

Evidence also indicates that the concentration of neuron-specific enolase (NSE), a marker of neuronal injury, is significantly higher in the SRF compared to its levels in the serum and vitreous of the control group^[48].

CLINICAL THERAPEUTICS FOR PVR

PVR remains a critical complication following RRD, contributing to a considerable proportion of recurrent detachments, even after seemingly successful surgical

repair^[49]. The identification of specific proteins and metabolites in SRF holds substantial therapeutic potential. Targeting inflammatory pathways could pave the way for innovative treatment strategies for managing RRD^[12,50]. Furthermore, the development of targeted therapies rooted in specific biomarkers offers the promise of more personalized and effective treatment options for RRD patients^[51].

Corticosteroids Corticosteroids are utilized to address vitreoretinal scarring due to their anti-inflammatory and antiproliferative properties, as well as their capacity to prevent the breakdown of the blood-ocular barrier. Research has demonstrated that a slow-release 0.7 mg dexamethasone intravitreal implant (Ozurdex) significantly reduces cystoid macular edema following PPV^[52]. However, the effectiveness of intravitreal triamcinolone acetonide injections or oral corticosteroids in preventing PVR remains uncertain^[53].

Methotrexate Methotrexate (MTX) has shown inhibitory effects on cell proliferation and cytokine-producing immune cells, highlighting its potential role in managing PVR^[54]. This potential is believed to stem from its ability to inhibit RPE metaplasia, as well as the proliferation of myocontractile cells and glial elements within the retina. Research has demonstrated a low incidence of recurrent PVR and favorable single-surgery success rates associated with MTX use. Furthermore, most studies have not reported severe complications directly attributable to MTX^[55-56].

Other Pharmacologic Agents Intravitreal agents such as anti-VEGF^[57] and 5-fluorouracil^[58], among others, have not demonstrated proven efficacy in the treatment or prevention of PVR.

CONCLUSION

The analysis of SRF in RRD has yielded significant insights into the molecular mechanisms driving this condition. Various proteins, lipids, and metabolites have been identified, reflecting the pathological state of the retina and highlighting their potential as biomarkers for disease diagnosis and monitoring. Moving forward, research should prioritize a deeper understanding of these molecules' roles in the pathogenesis of RRD while investigating their viability as therapeutic targets. The development of targeted therapies derived from specific biomarkers holds promise for enabling more effective and personalized treatments for RRD patients, ultimately enhancing visual outcomes and overall quality of life.

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