Safety and Feasibility of Quantitative Multiplexed Cytokine Analysis From Office-Based Vitreous Aspiration

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PURPOSE. The goals of this study were to evaluate the safety of office-based vitreous sampling, and determine the utility of these samples with multiplex cytokine analysis.

METHODS. Vitreous samples were collected from office-based needle aspiration and the rate of adverse events during follow-up was reviewed. The vitreous cytokine concentrations in a subset of patients with diabetic macular edema (DME) were analyzed using a 42plex cytokine bead array. These results were compared with vitreous cytokine concentrations in proliferative diabetic retinopathy (PDR) and controls (macular hole, epiretinal membrane, symptomatic vitreous floaters) from pars plana vitrectomy.

RESULTS. An adequate volume of vitreous fluid (100–200 μL) was obtained in 52 (88%) of 59 office-based sampling attempts. The average length of follow-up was 300 days (range, 42–926 days). There were no complications, including cataract, retinal tear or detachment, and endophthalmitis. Two patients (3%) had posterior vitreous detachments within 3 months. Vitreous cytokine concentrations were measured in 44 patients: 14 controls, 13 with DME, and 17 with PDR. The concentration of ADAM11, CXCL-10, IL-8, and PDGF-A were higher in PDR compared with controls and DME. The concentration of IL-6 was higher in PDR compared with controls, but not compared with DME.

CONCLUSIONS. Office-based vitreous aspiration is safe and yields high-quality samples for multiplex vitreous cytokine analysis. Significant elevations of vitreous cytokines were found in PDR compared with DME and controls, including the novel finding of elevated ADAM11. As such, office-based aspiration is a safe and effective means to identify vitreous factors associated with vitreoretinal disease.

Keywords: vitreous humor, cytokine, diabetic retinopathy, growth factors

Diabetic retinopathy is a common complication occurring in approximately 35% of patients with diabetes.1 Vision loss from diabetic retinopathy primarily occurs from either diabetic macular edema (DME) or proliferative diabetic retinopathy (PDR). The role of VEGF has been established in the pathogenesis of both DME and PDR.2 The intravitreal use of anti-VEGF drugs, such as bevacizumab, ranibizumab, and aflibercept, is standard practice in the management of patients with diabetic retinopathy, but there are limitations. Many patients are refractory to anti-VEGF pharmacotherapy or demonstrate only a partial response despite multiple, frequent injections.3–5 The reasons for inadequate response are currently poorly understood. It has been suggested that poor responders to anti-VEGF therapy may have additional inflammatory mediators that need to be targeted in addition to VEGF.6

The study of inflammatory mediators in the human vitreous has been gaining increasing attention. Several reports have examined vitreous proteomic changes in diseases such as diabetic retinopathy, AMD, retinal vein occlusion, and retinal detachment.2,7–15 Although human vitreous could serve as an important source of biomarkers for the diagnosis and management of vitreoretinal disease, further clinical application of vitreous cytokine analysis has been limited by two factors: (1) practical approaches to sample intraocular fluid, and (2) methods to analyze very small volume samples. Previous studies have primarily used operating room–based pars plana vitrectomy to obtain vitreous specimen.2,7–15 Attempts to examine cytokine and growth factors levels from more easily accessible anterior chamber samples do not reliably reflect those levels found in the vitreous.16 For vitreous cytokine analysis to be established as a part of the diagnosis and management of patients, a safe and practical approach to obtaining a vitreous sample must be established. One proposed method to more easily sample vitreous fluid is in-office aspiration of vitreous fluid. Giansanti et al.17 reported that microsampling of small volumes (10–15 μL) in rabbit eyes is a useful technique in vitreous pharmacokinetic studies. Pfahler et al.18 found very few complications with a large number of in-office vitreous sampling attempts, but did not report on the ability to analyze samples acquired in this manner.

Although avoiding the need for the operating room, in-office vitreous aspiration is limited by the small volume of a safely
obtainable sample. The primary methods of analyzing cytokines in biofluid have been ELISA and Western blot, which are limited to single factor analysis. The recent development of multiplexed cytometric bead assay has allowed for broader vitreous cytokine analysis and can be performed on sample volumes as small as 25 μL. There have been a few reports using cytometric bead assay on vitreous samples from patients undergoing vitrectomy. The practicality of in-office vitreous aspiration paired with multiplexed cytometric bead assay, however, may unlock the true power of vitreous analysis in our understanding of vitreoretinal disease. The clinical potential of proteomic analyses is within reach, but like any clinical tool, must first be validated. The present report examines the safety of office-based vitreous sampling techniques and the feasibility of using this fluid for multiplexed cytokine analysis.

**Methods**

This study was a prospective case series at the University of Michigan Kellogg Eye Center conducted between November 2011 and May 2015. This study was approved by the University of Michigan Institutional Review Board and adhered to the tenets of the Declaration of Helsinki. Every patient in the study had a vitreoretinal condition (Table 1) that required intraocular injection or surgical intervention. Informed consent was obtained from all subjects undergoing in-office vitreous aspiration after explanation of the nature and possible consequences of the study. Vitreous samples also were collected in the operating room before clinically indicated vitrectomy as part of a larger protocol establishing a vitreous biorepository at the University of Michigan. After vitreous sampling by either method, the patients were observed for follow-up as clinically indicated by their underlying condition. The charts for follow-up office visits were reviewed and all safety and adverse events were recorded.

**Office-Based Vitreous Sampling**

Topical anesthesia was initiated with a drop of proparacaine 1% followed by placement of cotton-tip applicators soaked in lidocaine 4% solution on the superotemporal conjunctiva. Subconjunctival lidocaine 1% was infiltrated in the area and the eyelashes and upper and lower eyelids were gently scrubbed with two 10% povidone-iodine sticks. A sterile speculum was used to place an inferotemporal infusion cannula and a 25-gauge, 5/8-inch needle was introduced through the marked site into the midvitreous cavity. A maximum of 0.2 mL vitreous fluid was gently aspirated into a 1-mL syringe (Supplementary Video S1). If clinically warranted, the patient immediately underwent injection of 0.05 to 0.1 mL therapeutic agent near the same area as vitreous sampling (e.g., an anti-VEGF agent or steroid injection). Optic nerve perfusion was confirmed by counting fingers vision and the patient was discharged without topical antibiotics. Each vitreous sample was stored in a −80°C freezer until cytokine analysis.

**Operating Room-Based Vitreous Sampling**

The patient was taken to the operating room at the Kellogg Eye Center and either general anesthesia was induced or posterior sub-Tenon's local anesthetic (2% lidocaine mixed 50/50 with 0.5% Marcaine) was injected. Trocars of 23, 25, or 27 gauge were used to place an inferotemporal infusion cannula and another superotemporal cannula 4.0 mm and 3.5 mm posterior to the corneoscleral limbus for phakic and pseudophakic patients, respectively. With the infusion off, a microvitrector was used to cut in the midvitreous cavity while an assistant applied gentle aspiration to an attached 3-mL syringe. Approximately 0.2 mL vitreous fluid was collected, after which the position of the infusion line was checked and the infusion initiated. Any immediate complications from vitreous sampling were recorded. The vitreous fluid was stored in a −80°C freezer until cytokine analysis and the remainder of the vitreoretinal surgical procedure was then performed as planned.

**Cytokine Analysis**

To examine the feasibility of cytometric bead assay analysis of office-based vitreous aspirates, three groups of patients were identified for comparison. A subset of vitreous samples from both the office-based and operating room-based groups was selected for cytokine analysis. Thirteen patients with DME had vitreous aspirated in the office. Seventeen patients with PDR and 14 age-matched controls without DME or PDR had vitreous sampling in the operating room by vitrectomy. Control patients were defined as those undergoing vitrectomy for a diagnosis of epiretinal membrane, symptomatic vitreous floater, or macular hole. Cytokines and growth factors were assessed from each vitreous sample using 42-plex cytometric bead immunoassay (EMD Millipore, Darmstadt, Germany) according to the manufacturer's instruction manual. Each sample was assessed using 25 μL vitreous in triplicate. The mean concentration of each cytokine was compared among groups with Kruskal-Wallis tests, and P values were adjusted by Holms method for multiple comparisons. Post hoc pairwise group comparisons were performed using 2-sample Wilcoxon tests.

**Results**

A total of 264 vitreous samplings were attempted: 63 from office-based sampling and 201 from operating room-based sampling. Of the office-based group, four patients with choroidal melanoma were excluded due to not having follow-up. Three patients had the eye enucleated and one patient did not return for follow-up. Fifty-nine office-based vitreous aspirations from 48 patients were included for safety and feasibility analysis. Of these 48 patients, 38 patients had one office-based vitreous sample, 9 patients had two samples at different visits, and 1 patient had three samples at different visits. Twenty-eight (58%) were male and the average age was 61 (range, 24–90). Thirty-three eyes (69%) were phakic and 15 (31%) were pseudophakic. Patient demographics and underly-
There were no intraoperative complications from vitreous sampling during operating room–based procedures. There were no additional retinal breaks or signs of hypotony, such as corneal striae, choroidal folds, or choroidal detachment.

Vitreous cytokines were measured from 44 patients in three groups: DME ($n = 13$), PDR ($n = 17$), and controls ($n = 14$). All 13 DME samples were obtained by office-based vitreous aspiration, and the 17 PDR samples and 14 controls (epiretinal membrane, symptomatic vitreous floaters, and macular holes) were obtained by office-based vitrectomy. Subjects were selected chronologically by date of vitreous sampling from each group for inclusion in the subset of patients undergoing cytokine analysis. The cytokines included in the multiplex bead assay were the following: chemokine ligand 2 (CCL2/monocyte chemotactic protein 1), CCL3, CCL4, CCL5, CCL7 (monocyte chemottractant protein 3), CX-C motif chemokine 1 (CXCL1/growth-regulated alpha protein), CX-C motif chemokine 10 (CXCL10/IP-10), disintegrin and metalloproteinase domain-containing protein 11 (ADAM11), fractalkine, eotaxin, epidermal growth factor (EGF), FGF-fms-related tyrosine kinase 2 ligand, granulocyte-colony stimulating factor (G-CSF), granulocyte-macrophage colony stimulating factor (GM-CSF), IFN-α, IFN-γ, IL-1 receptor antagonist protein, IL-1α, IL-1β, IL-2, IL-3, IL-4, IL-5, IL-6, IL-7, IL-8, IL-9, IL-10, IL-12, IL-13, IL-15, IL-17A, platelet-derived growth factor (PDGF)-A, PDGF-B, soluble CD 40 ligand, TGF-α, TNF-α, TNF-β (lymphotoxin-2), and VEGF. The mean concentrations of each cytokine stratified by group (DME, PDR, and control) are shown in Table 3.

After Holm’s adjustment for multiple comparisons, there were statistically significant differences between groups ($P < 0.05$) observed for five vitreous cytokines: ADAM11, CXCL2, IL-6, IL-8, and PDGF-A. The pairwise comparisons between groups for significant cytokines are shown in Table 4. The vitreous concentrations of each of the five cytokines were significantly higher in patients with PDR compared with controls. In addition, the average concentration of each cytokine except for IL-6 was significantly higher in the PDR group compared with the DME group. The Figure shows a boxplot comparing the distribution of vitreous concentrations of the novel cytokine ADAM11 for each group.

**TABLE 2. Patient Characteristics of Operating Room–Based Vitreous Sampling**

| Number of samples | 201 |
|-------------------|-----|
| Number of subjects | 199 |
| Mean age (range) | 62 (8–89) |
| Sex: male (%) | 109 (55) |
| Diagnoses (%) | |
| PDR | 57 (28) |
| Rhegmatogenous retinal detachment | 40 (20) |
| Epiretinal membrane | 37 (18) |
| Macular hole | 17 (8) |
| Choroidal melanoma | 14 (7) |
| IOI, dislocations and lens fragments | 11 (5) |
| Ocular lymphoma | 5 (2) |
| Diagnostic vitrectomy | 5 (2) |
| Nondiabetic nonclearing vitreous hemorrhage | 5 (2) |
| PPV for pars plana glaucoma tube | 3 (1) |
| Symptomatic PVD (floaters) | 3 (1) |
| IOFB removal | 2 (1) |
| Other | 2 (1) |

**Notes:**
- IOFB, intraocular foreign body; PPV, pars plana vitrectomy; PVD, posterior vitreous detachment.
- The mean age was 62 (range, 8–89).
- The underlying vitreoretinal diagnoses of subjects in the office-based group are listed in Table 1.

A total of 201 operating room–based vitreous samples were performed in 199 patients. Two patients with PDR had both eyes included. Of this group, 109 (55%) were male and the average age was 62 (range, 8–89). The underlying vitreoretinal diagnoses for which study subjects were being treated are presented in Table 2.

Of the 59 office-based vitreous samples included with follow-up, 52 (88%) had adequate volume of aspirated vitreous. There were four “dry” samples (7%) in which vitreous fluid could not be aspirated, and three samples (5%) in which only a very small volume could be obtained. Of the 52 samples, 9 (17%) had a prior posterior vitreous detachment documented. Of the seven inadequate samples, four (57%) had a prior posterior vitreous detachment documented. This was a statistically significant association by Fisher’s exact test ($P = 0.0360$). For all office-based vitreous samples, the average length of follow-up was 300 days (range, 42–926 days; SD, 176 days).

There were no documented instances of cataract, retinal tear, retinal detachment, or endophthalmitis in patients undergoing vitreous aspiration. No patients with “dry” or inadequate vitreous sampling attempt had any complications. There were two (3%) patients with posterior vitreous detachments documented within 3 months after vitreous aspiration. One patient was found to have asymptomatic posterior vitreous detachment 2 months after vitreous aspiration. The other patient presented 14 days after vitreous aspiration with complaints of flashing photopsias and a new floater in the study eye. Both patients had no retinal tears or detachments on complete dilated fundus examination with scleral depression, and continued to do well without complications after 3 and 13 months of follow-up, respectively.

Four patients (7%) had new posterior vitreous detachment documented between 3 and 6 months. Five patients (8%) had new posterior vitreous detachment documented between 6 and 12 months. All patients with new posterior vitreous detachment after 3 months had additional intraventricular injections after vitreous aspiration. No retinal tears or detachments were noted in any patients with new documented posterior vitreous detachment. Of the 11 patients with new posterior vitreous detachment within 12 months of vitreous aspiration, 9 (82%) were asymptomatic.

**Discussion**

The identification of protein biomarkers in the vitreous could elucidate pathophysiologic mechanisms of vitreoretinal disease and potentially serve as targets for future therapies. The proximity of vitreous body to the retina and its interaction in various disease states provides a unique means of studying the retina itself. Several reports have characterized vitreous cytokines and have shown the concentrations of various inflammatory and angiogenic factors to be altered in a number of disease states.10,11,16,20–23

**Vitreous Cytokines**
setting and is associated with side effects typical of vitrectomy, such as cataract progression. A few reports have examined office-based vitrectomy systems (Intrector [Insight Instruments, Stuart, FL, USA] and VersVIT [Synergetics, O’Fallon, MO, USA]) for various retinal procedures, but these systems have not been widely adopted. Current reports examining vitreous proteins have mostly used typical techniques of diagnostic pars plana vitrectomy in the operating room to acquire vitreous sample. Office-based aspiration of vitreous sample for proteomic analysis has less commonly been used. Despite theoretical risks of vitreoretinal traction, a large review of in-office vitreous sampling by Pfahler et al. showed that aspiration of small volumes was a safe and reproducible procedure. Lobo and Lightman reported a similar rate, 92%, of obtaining adequate sample from vitreous aspiration and few complications in diagnosing patients with infectious, malignant, and inflammatory causes of uveitis. Feasibility of proteomic analysis from their vitreous aspiration samples, however, was not discussed.

### Table 3. Mean Concentrations (pg/mL) and SDs of Vitreous Cytokines Compared Among Control (Macular Holes, Symptomatic Vitreous Floaters, and Epiretinal Membranes), DME, and PDR Patients

| Cytokine | Controls, n = 14 | DME, n = 13 | PDR, n = 17 | P* | P† |
|----------|-----------------|-------------|-------------|----|----|
|          | Mean | SD | Mean | SD | Mean | SD | Mean | SD | Mean | SD |
| EGF      | 0.55 | 0.07 | 0.57 | 0.10 | 1.18 | 1.76 | 0.5700 | 1.0000 |
| Eotaxin  | 2.35 | 1.73 | 4.02 | 2.69 | 13.10 | 19.58 | 0.0018 | 0.0066 |
| FGF2     | 21.27 | 37.73 | 11.04 | 2.99 | 30.02 | 65.06 | 0.2743 | 1.0000 |
| Flt-3L   | 13.08 | 7.23 | 16.54 | 7.97 | 27.59 | 24.85 | 0.0127 | 0.4191 |
| Fractalkine | 18.77 | 19.53 | 16.22 | 14.10 | 75.34 | 158.92 | 0.1383 | 1.0000 |
| G-CSF    | 1.65 | 1.89 | 1.46 | 1.99 | 19.05 | 70.46 | 0.8851 | 1.0000 |
| GM-CSF   | 0.32 | 0.21 | 0.72 | 0.68 | 0.65 | 0.74 | 0.3128 | 1.0000 |
| GRO      | 19.23 | 15.19 | 19.20 | 14.81 | 55.41 | 52.99 | 0.0266 | 0.7770 |
| IFN-α    | 0.34 | 0.10 | 0.41 | 0.18 | 0.71 | 1.34 | 0.5855 | 1.0000 |
| IFN-γ    | 0.42 | 0.13 | 0.40 | 0.16 | 1.34 | 1.61 | 0.0259 | 0.7770 |
| IL-1α    | 0.42 | 0.13 | 0.40 | 0.16 | 1.34 | 1.61 | 0.0259 | 0.7770 |
| IL-1β    | 0.43 | 0.11 | 0.43 | 0.09 | 0.49 | 0.16 | 0.4158 | 1.0000 |
| IL-1RA   | 1.42 | 0.66 | 6.00 | 15.69 | 19.97 | 52.15 | 0.2802 | 1.0000 |
| IL-2     | 0.45 | 0.45 | 0.53 | 0.43 | 0.86 | 1.54 | 0.7180 | 1.0000 |
| IL-3     | 0.35 | 0.03 | 0.35 | 0.03 | 0.47 | 0.31 | 0.3233 | 1.0000 |
| IL-4     | 0.81 | 0.44 | 0.82 | 0.42 | 1.06 | 1.02 | 0.9483 | 1.0000 |
| IL-5     | 0.28 | 0.02 | 0.27 | 0.06 | 0.40 | 0.19 | 0.0042 | 0.1470 |
| IL-6     | 4.72 | 3.47 | 17.23 | 16.15 | 59.37 | 87.22 | 0.0003 | 0.0114 |
| IL-7     | 4.46 | 3.66 | 5.05 | 3.78 | 15.54 | 49.76 | 0.7530 | 1.0000 |
| IL-8     | 7.45 | 3.95 | 20.69 | 24.90 | 87.89 | 88.29 | 0.0001 | 0.0042 |
| CXCL10   | 300.49 | 295.77 | 217.45 | 138.08 | 2241.95 | 2772.64 | <0.0001 | <0.0001 |
| CCL2     | 1609.94 | 1158.55 | 1532.56 | 100.05 | 5449.81 | 2095.08 | 0.0088 | 0.2992 |
| CCL7     | 2.96 | 1.42 | 4.20 | 5.05 | 6.25 | 4.70 | 0.0806 | 1.0000 |
| ADAM11   | 15.17 | 13.33 | 34.88 | 34.63 | 91.84 | 68.60 | 0.0002 | 0.0078 |
| CCL3     | 2.52 | 2.15 | 3.00 | 2.60 | 31.43 | 102.13 | 0.1326 | 1.0000 |
| CCL4     | 1.88 | 1.45 | 2.54 | 2.52 | 3.77 | 3.84 | 0.5148 | 1.0000 |
| PDGF-AA  | 29.42 | 5.18 | 44.77 | 29.49 | 131.75 | 119.69 | <0.0001 | 0.0042 |
| PDGF-BB  | 0.84 | 0.58 | 0.66 | 0.13 | 15.64 | 47.92 | 0.0234 | 0.7254 |
| CCL5     | 1.29 | 0.51 | 1.84 | 1.64 | 3.81 | 9.78 | 0.5626 | 1.0000 |
| sCD40L   | 0.91 | 0.56 | 0.65 | 0.29 | 3.79 | 12.62 | 0.7994 | 1.0000 |
| TGF-β    | 0.48 | 0.11 | 0.48 | 0.11 | 1.90 | 5.25 | 0.1041 | 1.0000 |
| TNF-α    | 0.50 | 0.19 | 0.60 | 0.27 | 0.79 | 0.65 | 0.6116 | 1.0000 |
| VEGF     | 7.36 | 7.56 | 259.82 | 297.47 | 240.18 | 459.51 | 0.0031 | 0.1116 |

Values in bold are statistically significant (P < 0.05 Holm’s adjustment).
* Kruskal-Wallis test.
† P values Holm’s adjustment for multiple comparisons.

### Table 4. P Values for Pairwise Comparisons Using Wilcoxon 2-Sample Test of Control, DME, and PDR Patients

| Cytokine | Controls vs. DME | Controls vs. PDR | DME vs. PDR |
|----------|------------------|-----------------|-------------|
| ADAM11   | 0.0765           | 0.0008          | 0.0109      |
| CXCL10   | 0.5816           | 0.0016          | 0.0008      |
| IL-6     | 0.0663           | 0.0005          | 0.0588      |
| IL-8     | 0.1213           | 0.0003          | 0.0059      |
| PDGF-AA  | 0.1642           | 0.0004          | 0.0048      |

MO, USA) for various retinal procedures, but these systems have not been widely adopted. Current reports examining vitreous proteins have mostly used typical techniques of diagnostic pars plana vitrectomy in the operating room to acquire vitreous sample. Office-based aspiration of vitreous sample for proteomic analysis has less commonly been used. Despite theoretical risks of vitreoretinal traction, a large review of in-office vitreous sampling by Pfahler et al. showed that aspiration of small volumes was a safe and reproducible procedure. Lobo and Lightman reported a similar rate, 92%, of obtaining adequate sample from vitreous aspiration and few complications in diagnosing patients with infectious, malignant, and inflammatory causes of uveitis. Feasibility of proteomic analysis from their vitreous aspiration samples, however, was not discussed.
and colleagues described learning and motor coordination. It is mainly expressed in the nervous system but has been played a role in cell-cell or cell–extracellular matrix interactions.

Significantly higher levels of IL-6, IL-8, CXCL10, and PDGF-AA were found compared with DME and controls. Similar to previous reports, the present study found that patients with PDR have statistically significant differences in certain cytokine concentrations among PDR, DME, and controls. ADAM11 was originally identified as a candidate tumor suppressor gene for breast cancer. It is thought to be an adhesion molecule that is involved in the neuropathic changes frequently associated with diabetes.

There are several limitations to this study. This is a small pilot study to assess safety and efficacy of vitreous cytokine analysis through office- versus operating room-based vitreous sampling. Larger randomized studies could help transition office-based vitreous aspiration from a research tool into a standard procedure in the diagnosis and management of vitreoretinal disease, following validation of specific vitreous biomarkers. Second, we did not implement a standardized follow-up protocol in determining complications associated with office-based vitreous sampling. Follow-up intervals were determined by the patient’s underlying clinical diagnosis. Despite this, there was an adequate overall length of follow-up with the average patient followed 300 days after office-based vitreous sampling. Third, it is possible that office-based vitreous aspiration may capture only soluble or nonbound proteins, whereas vitreous cutter-based biopsies may gather additional insoluble or bound factors. However, Skeie and colleagues found that proteins collected from sequential vitreous needle and vitreous cutter biopsies were nearly equivalent.

In conclusion, this study supports the safety and feasibility of obtaining vitreous by office-based aspiration for analysis with cytometric bead array. Our results show a very low rate of complications from in-office vitreous sampling of a small volume of fluid with the method described. A single patient experienced an uncomplicated acute posterior vitreous detachment. The procedure was well tolerated and there was sufficient length of follow-up to rule out complications, such as endophthalmitis, retina tear or detachment, and cataract. At present, we believe that office-based vitreous aspiration can be a safe, effective way to acquire vitreous fluid for research with increasing insight into the role of vitreous biomarkers, office-based vitreous sampling paired with multiplexed cytokine analysis could one day be clinically useful in diagnosing, monitoring, and treating vitreoretinal disease, such as diabetic retinopathy.

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