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ORIGINAL ARTICLE

Hypoxia Increases the Potential for Neutrophil-mediated Endothelial Damage in Chronic Obstructive Pulmonary Disease

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Abstract

Rationale: Patients with chronic obstructive pulmonary disease (COPD) experience excess cardiovascular morbidity and mortality, and exacerbations further increase the risk of such events. COPD is associated with persistent blood and airway neutrophilia and systemic and tissue hypoxia. Hypoxia augments neutrophil elastase release, enhancing capacity for tissue injury.

Objective: To determine whether hypoxia-driven neutrophil protein secretion contributes to endothelial damage in COPD.

Methods: The healthy human neutrophil secretome generated under normoxic or hypoxic conditions was characterized by quantitative mass spectrometry, and the capacity for neutrophil-mediated endothelial damage was assessed. Histotoxic protein concentrations were measured in normoxic versus hypoxic neutrophil supernatants and plasma from patients experiencing COPD exacerbation and healthy control subjects.

Measurements and Main Results: Hypoxia promoted PI3Kγ-dependent neutrophil elastase secretion, with greater release seen in neutrophils from patients with COPD. Supernatants from neutrophils incubated under hypoxia caused pulmonary endothelial cell damage, and identical supernatants from COPD neutrophils increased neutrophil adherence to endothelial cells. Proteomics revealed differential neutrophil protein secretion under hypoxia and normoxia, and hypoxia augmented secretion of a subset of histotoxic granule and cytosolic proteins, with significantly greater release seen in COPD neutrophils. The plasma of patients with COPD had higher content of hypoxia-upregulated neutrophil-derived proteins and protease activity, and vascular injury markers.

Conclusions: Hypoxia drives a destructive "hypersecretory" neutrophil phenotype conferring enhanced capacity for endothelial injury, with a corresponding signature of neutrophil degranulation and vascular injury identified in plasma of patients with COPD. Thus, hypoxic enhancement of neutrophil degranulation may contribute to increased cardiovascular risk in COPD. These insights may identify new therapeutic opportunities for endothelial damage in COPD.

Keywords: cell degranulation; neutrophil elastase; vascular endothelium; cardiovascular disease

At a Glance Commentary

Scientific Knowledge on the Subject: Chronic obstructive pulmonary disease (COPD) is characterized by persistent neutrophilia in the setting of local and systemic hypoxia and is associated with excess cardiovascular disease, even allowing for known risk factors. Neutrophils accumulating in areas of inflammation and microcirculatory impairment experience profound hypoxia, which prolongs their survival and increases their secretory responses. Thus, hypoxic neutrophils have increased potential to cause endothelial injury, but their role in mediating the increased cardiovascular risk in COPD is poorly understood.

What This Study Adds to the Field: Herein we show that hypoxia augments the ability of neutrophils to selectively secrete a subset of histotoxic proteins capable of damaging endothelial cells. Hypoxia further enhanced release of these proteins from neutrophils from patients experiencing COPD exacerbation, with elevated concentrations also detected in patient plasma. This study suggests that hypoxic enhancement of neutrophil degranulation may contribute to increased cardiovascular risk in COPD and that the hypoxic neutrophil secretome proteins may represent new therapeutic targets to alleviate endothelial dysfunction in COPD.

Chronic obstructive pulmonary disease (COPD) is characterized by neutrophilic inflammation in the setting of tissue (and often systemic) hypoxia and by increased risk of cardiovascular disease and pulmonary

hypertension. Neutrophil elastase (NE) has been implicated in COPD pathogenesis (1), and we have previously shown that hypoxia markedly augments NE release from neutrophils to promote respiratory epithelial cell damage (2). However, the impact of hypoxia on the extended neutrophil secretome and the potential for "hypoxic neutrophils" to injure other disease-relevant cell types are currently unknown. Identifying novel targets implicated in driving COPD morbidity may provide new therapeutic opportunities.

Even in health, certain tissues (e.g., muscle) are hypoxic (3). This "physiological hypoxia" may be compounded during exercise, inducing neutrophil phenotypic changes (4). In disease, profound "pathological hypoxia" exists in inflamed or infected tissues and areas of microcirculatory impairment. Although patients with severe COPD are systemically hypoxic, significant tissue hypoxia (<1.3% oxygen) can occur even in mild disease, demonstrated in inflamed airways (5–7) and atherosclerotic vasculature (8), where neutrophils accumulate. Upregulation of hypoxia-inducible factors in neutrophils from patients with acute lung injury, including coronavirus disease (COVID-19) infection, indicates neutrophil exposure to hypoxia *in vivo* (9).

Neutrophil antimicrobial function depends on the fusion of preformed granules, containing cytotoxic proteins and proteases, with the pathogen-containing phagosome. However, highly activated neutrophils can release granule contents extracellularly (degranulation), with potential for collateral tissue damage (2). Platelet-activating factor (PAF), a physiological priming agent capable of substantially enhancing neutrophil degranulation in response to subsequent stimulation (2, 10), has been implicated in COPD pathogenesis and in endothelial damage and remote organ damage in the setting of hypoxia (11, 12). Bacteria release formylated peptides (*N*-formyl-methionyl-leucyl-phenylalanine [fMLP]), which potently activate neutrophils; these peptides are present in cigarette smoke and have been implicated in emphysema progression (13). Patients with COPD suffer recurrent infection-driven exacerbations, but

neutrophilic inflammation persists even in the absence of detectable infection, correlating with disease severity and progression (14). Despite evidence of NE-induced lung injury, translation of NE inhibitors has not led to significant benefit (15), perhaps reflecting the complex array of additional neutrophil-secreted proteins with damaging potential.

Patients with COPD have increased risk of cardiovascular morbidity and mortality even after adjusting for shared risk factors, including smoking (16), particularly after exacerbation (17). Pulmonary endothelial dysfunction in patients with COPD can induce pulmonary hypertension, which correlates with hypoxia (18). Accumulating evidence indicates inflammation, oxidative stress, and vascular tissue damage as key mechanisms linking COPD and cardiovascular disease (19), with neutrophil degranulation identified as an important pathway (20). Circulating neutrophils primed for enhanced degranulation have been identified in patients experiencing COPD exacerbation (21), with potential to contribute to endothelial injury.

Herein we show that hypoxia promotes neutrophil degranulation and neutrophil-induced endothelial damage. Proteomic analysis reveals hypoxia-driven secretion of highly histotoxic proteins from healthy neutrophils, and a subset of these are further increased from COPD neutrophils. Supernatants from hypoxic COPD neutrophils enhance neutrophil–endothelial adhesion. Finally, we identify increased concentrations of corresponding hypoxic neutrophil histotoxic granule proteins in COPD plasma, together with endothelial injury biomarkers. Some results have been reported previously in abstract form (22–24).

Methods

Ethics Statement
Written informed consent was obtained from all healthy volunteers and patients with COPD (06/Q0108/281, 08/H0308/281, 18/WM/0097, and 20/SS/0085). All studies

complied with the Declaration of Helsinki. All animal experiments were approved in accordance with the Animals (Scientific Procedures) Act 1986.

Human Neutrophils

Venous blood neutrophils were isolated by centrifugation over plasma-Percoll gradients (25). Neutrophils were resuspended in normoxic (atmospheric 21% O_2) or hypoxic (0.8% O_2 equating to media P_{O_2} of 3 kPa [26], 5% CO_2, Baker Ruskinn or Whitley hypoxia workstation) Iscove's modified Dulbecco's medium (IMDM). At 4 hours, neutrophils (11.1×10^6/ml) were treated with PAF (1 μM, 5 min) and then fMLP (100 nM, 10 min) (2). PI3K inhibitors were added before incubation: PI3Kγ-selective (AS605240, 3 μM) and PI3Kδ-selective (CAL-101, 100 nM).

Murine Neutrophils

Femoral bone marrow neutrophils were isolated by negative immunomagnetic selection from PI3Kδ-hyperactived E1020K heterozygote (PPL 80/2248 and P4802B8AC) (27), PI3Kδ-kinase-dead D910A homozygote (PPL 70/7661) (28), or PI3Kγ$^{-/-}$ (PPL 70/8100) mice, alongside age- and strain-matched wild-type control mice (E1020K/D910A: C57BL/6J, and PI3Kγ$^{-/-}$: C57BL/E129). Neutrophils were resuspended in normoxic or hypoxic IMDM before treatment at 4 hours with cytochalasin B (5 μg/ml, 5 min) and then fMLP (10 μM, 10 min).

Protein Secretion

Neutrophil supernatant and plasma protein content were measured by ELISA, chemiluminescence immunoassay, or activity assay. Plasma NE- and PR3-specific fibrinogen cleavage products were measured (1, 29).

Neutrophil Secretome Preparation for Tandem Mass Tag–Mass Spectrometry

Neutrophils were resuspended in normoxic or hypoxic IMDM (4 h) containing ethylenediaminetetraacetic acid (1 mM) and sivelestat (10 μM) and treated with PAF and fMLP as above. Concentrated protein supernatants underwent tandem mass tag–mass spectrometry (TMT-MS).

Endothelial Cell Survival

Confluent human pulmonary artery endothelial cells (hPAECs) (Lonza) or human pulmonary microvascular endothelial cells (hPMECs) (Promocell) were treated with neutrophil supernatants, with or without alpha-1-antitrypsin (α1AT, 46 μg/ml, 10 min). Cell detachment of rhodamine phalloidin- and DAPI-stained fixed hPAECs was assessed by immunofluorescence. Viability and apoptosis of unfixed hPAECs or hPMECs was assessed by MTT assay or annexin V positivity.

Endothelial–Neutrophil Interaction

Confluent hPMECs were treated with neutrophil supernatants. At 4 hours, hPMECs were perfused with neutrophils (1×10^6 cells/ml, 4 min, 0.1 Pa shear stress) and neutrophil adhesion or transmigration assessed.

Statistical Analysis

Data were analyzed using GraphPad Prism version 9 software, reported as mean ± SEM from (n) independent experiments. Gaussian data were analyzed by t test or two-way ANOVA with Sidak's correction. Non-Gaussian data were analyzed by Mann-Whitney test. TMT-MS–generated P values were adjusted by Benjamini-Hochberg procedure. A P value of less than 0.05 was considered statistically significant.

For further information, see EXPANDED MATERIALS AND METHODS in the online supplement.

Results

Patients Experiencing COPD Exacerbation Have Increased Circulating Evidence of Neutrophil Protease Activity

Although neutrophil proteases are implicated in COPD lung parenchymal destruction, the extent of systemic release of neutrophil granule proteins and their potential role in endothelial injury are unclear. We show that patients experiencing COPD exacerbation (see Table E1 in the online supplement) have higher plasma concentrations of fibrinogen cleavage products AαVal[360] and AαVal[541] than age- and sex-matched healthy control subjects (Figures 1A and 1B). These footprints specifically indicate increased activity of neutrophil azurophil granule proteases NE and proteinase 3, respectively, secreted on neutrophil activation (which may occur in the circulation, during adherence to vascular endothelium, or after migration into
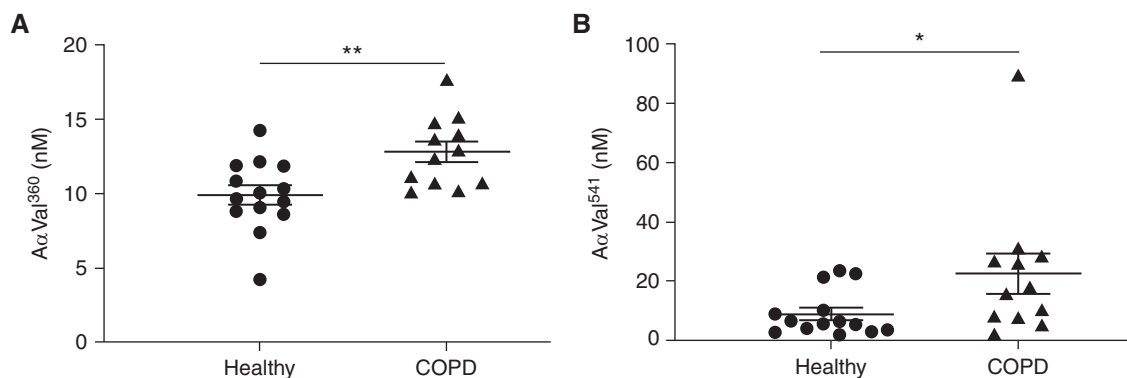
Figure 1. Patients experiencing chronic obstructive pulmonary disease (COPD) exacerbation have a circulating signature indicating increased protease activity. Plasma from patients experiencing COPD exacerbation or age- and sex-matched healthy control subjects was assessed for content of (A) neutrophil elastase-specific fibrinogen cleavage product AαVal[360] or (B) PR3-specific fibrinogen cleavage product AαVal[541] by immunoassay (n = 12 COPD, n = 14 healthy; cohort 1). Results represent mean ± SEM, (A) unpaired t test and (B) Mann-Whitney test. *P < 0.05 and **P < 0.01.
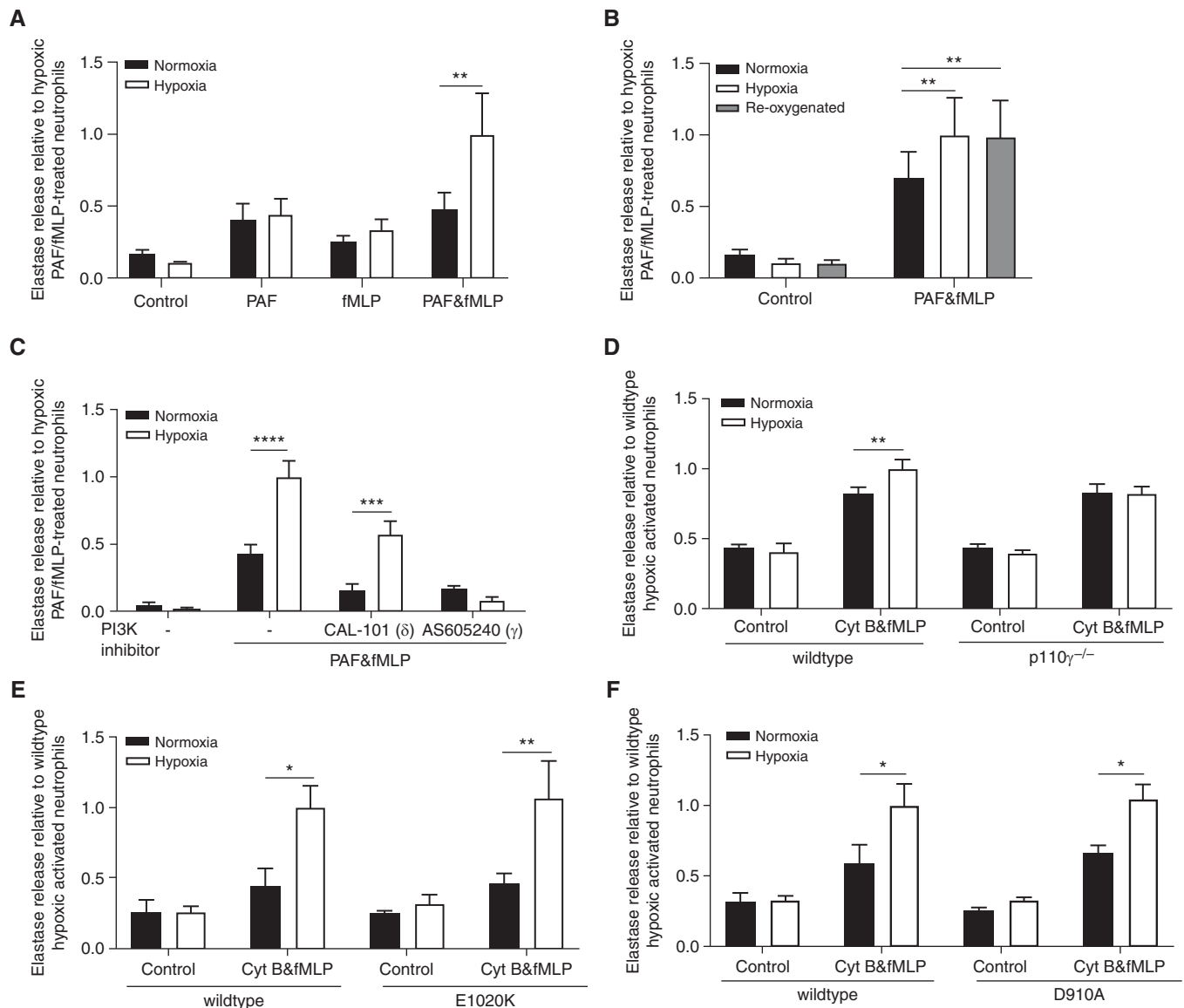
Figure 2. Hypoxia increases elastase release from platelet-activating factor (PAF)-primed neutrophils in a PI3Kγ-dependent manner. (*A–C*) Neutrophils from healthy human donors were incubated under normoxia or hypoxia in the presence or absence of PI3Kγ-selective inhibitor (AS605240, 3 μM) or PI3Kδ-selective inhibitor (CAL-101, 100 nM) as indicated. After 4 hours, cells were treated with PAF (1 μM, 5 min) and/or *N*-formyl-methionyl-leucyl-phenylalanine (fMLP) (100 nM, 10 min) or vehicle control as indicated. For reoxygenation, unstimulated hypoxic cells were moved to normoxia with the addition of twice-volume normoxic media for 30 minutes before treatment with PAF and fMLP. Supernatant neutrophil elastase (NE) activity was measured and is expressed as fold change relative to hypoxic activated neutrophils (*A*: $n = 5$, *B*: $n = 4$, *C*: $n = 4$–6). (*D–F*) Femoral bone marrow neutrophils were isolated from PI3Kγ-null (PI3Kγ$^{-/-}$), PI3Kδ-hyperactive (E1020K), PI3Kδ-kinase dead (D910A), or wild-type mice from the relevant genetic background. After 4 hours, cells were treated with cytochalasin B (Cyt B; 5 μg/ml, 5 min) and fMLP (10 μM, 10 min) or vehicle control. Supernatant NE activity was measured and is expressed as fold change relative to wild-type hypoxic activated neutrophils (*D*: 3–4 mice per genotype per experiment, $n = 5$ independent experiments; *E* and *F*: 3–4 mice per genotype per experiment, $n = 3$ independent experiments). Results represent mean ± SEM, two-way ANOVA. *$P < 0.05$, **$P < 0.01$, ***$P < 0.001$, and ****$P < 0.0001$.

tissues) before inactivation by circulating antiproteases, such as α1AT. Given this systemic signature of enhanced neutrophil protease activity during COPD exacerbation, established evidence of pathological hypoxia during inflammation, and our previous results demonstrating

hypoxia-augmented NE release from GM-CSF (granulocyte–macrophage colony–stimulating factor)–primed neutrophils (2), we next examined the ability of inflammatory mediators relevant to COPD and hypoxia to influence neutrophil degranulation.

Hypoxia Augments NE Release from PAF-primed Neutrophils in a PI3Kγ-Dependent Manner
Neutrophils treated with PAF and fMLP in combination (but not alone) released up to threefold more active NE when incubated under hypoxia
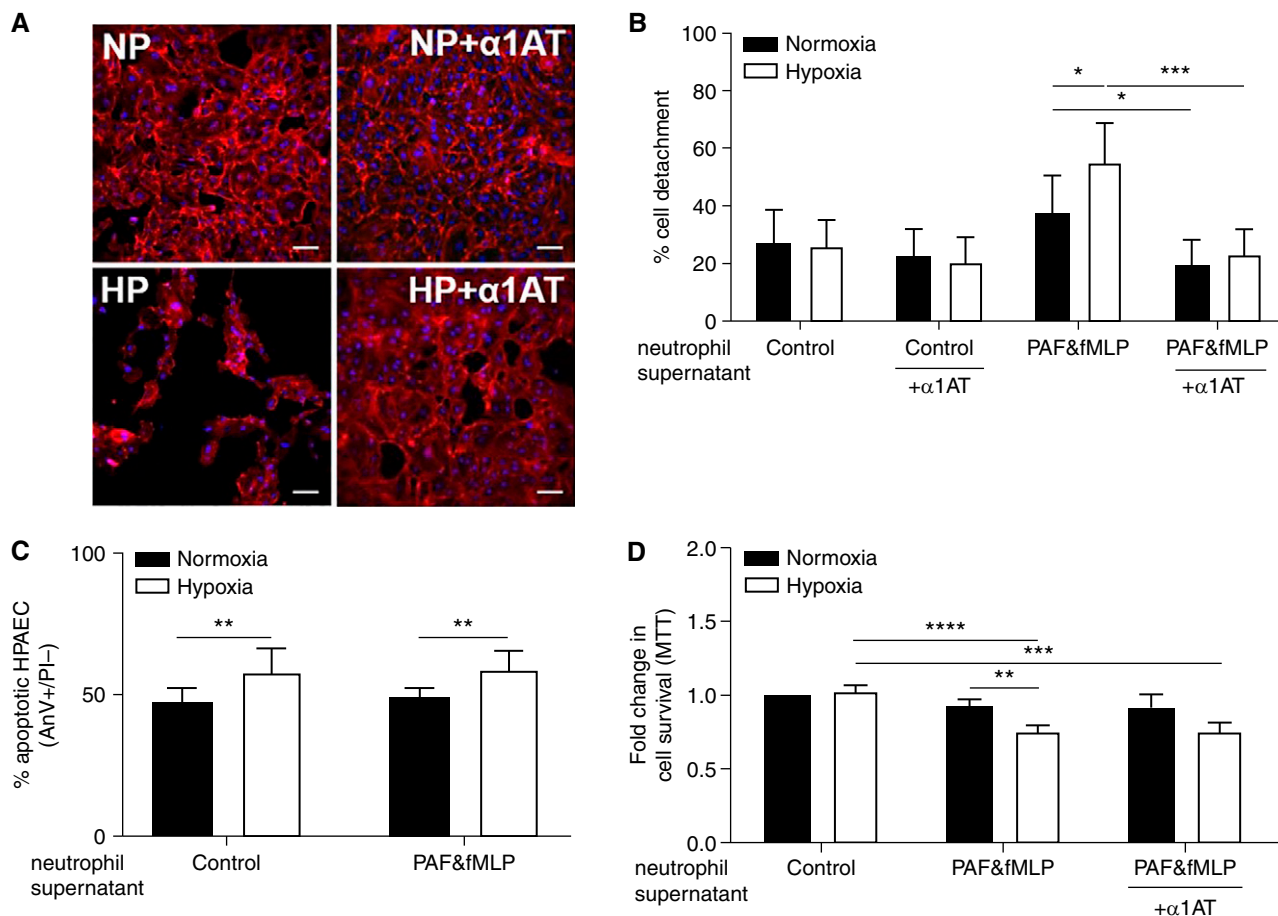
Figure 3. Supernatants from hypoxic activated neutrophils cause increased endothelial cell damage in a partially protease-dependent manner. Neutrophils from healthy donors were incubated under normoxia or hypoxia for 4 hours and then treated with platelet-activating factor (PAF) (1 μM, 5 min) and N-formyl-methionyl-leucyl-phenylalanine (fMLP) (100 nM, 10 min) or vehicle control. Supernatants from normoxic versus hypoxic, PAF/fMLP versus vehicle control-treated neutrophils were incubated with confluent human pulmonary artery endothelial cells (hPAECs) for (A and B) 24 hours, (C) 6 hours, or (D) 48 hours in the presence or absence of alpha-1-antitrypsin (α1AT, 46 μg/ml) as indicated. (A and B) hPAECs were fixed and stained with rhodamine-phalloidin and DAPI. Supernatants were from normoxic (NP) or hypoxic (HP) PAF/fMLP-treated neutrophils. Cell detachment was quantified using ImageJ, expressed as percentage detachment of whole field of view. (A) Representative confocal images from (B) five independent experiments; scale bars, approximately 20 μm. (C) hPAECs were stained with FITC-AnV and propidium iodide (PI) for flow cytometric assessment of apoptosis with apoptotic (AnV$^+$PI$^-$) cells expressed as percentage of total population (n = 4). (D) Survival of hPAECs was measured by MTT assay (n = 6–12). Results represent mean ± SEM, two-way ANOVA. *$P < 0.05$, **$P < 0.01$, ***$P < 0.001$, and ****$P < 0.0001$. AnV = annexin V; FITC = fluorescein isothiocyanate.

(0.8% O_2, 3 kPa) compared with normoxia (21 kPa) (Figure 2A). This enhanced secretion was not reversed by reoxygenation (Figure 2B). Given the known role of the PI3K pathway in the hypoxic upregulation of degranulation from GM-CSF–primed neutrophils, and the aberrant chemotaxis of neutrophils from patients with COPD, which could be corrected by PI3K inhibition (30), we explored whether inhibition of PI3K signaling pathways modulated the hypoxic response of PAF-primed neutrophils using PI3K isoform–selective small molecule inhibitors. PI3Kγ-selective inhibition abrogated the

hypoxic uplift of NE release from PAF-primed neutrophils; this effect was not seen with the PI3Kδ-selective inhibitor (Figure 2C). These results were replicated in a cohort of patients with COPD (Table E2 and Figure E1). As PI3Kγ inhibition also markedly inhibited NE release from stimulated normoxic cells, we further explored whether PI3K signaling was simply essential for overall degranulation or had a specific role in hypoxia-mediated degranulation, using transgenic mice with abolished or enhanced activity of PI3Kγ/δ isoforms. As PAF did not elicit a detectable priming response in murine neutrophils

(data not shown), these cells were treated with cytochalasin B and fMLP to liberate NE. The hypoxic increase in NE release from murine neutrophils deficient in PI3Kγ was abolished, with preserved ability to degranulate under normoxia (Figure 2D). In contrast, the hypoxic enhancement of NE release from murine neutrophils with either activating or kinase-dead PI3Kδ mutations was unaffected (Figures 2E and 2F). Together, these data indicate that the augmented degranulation observed under hypoxia from human or murine neutrophils requires PI3Kγ but not PI3Kδ activity.
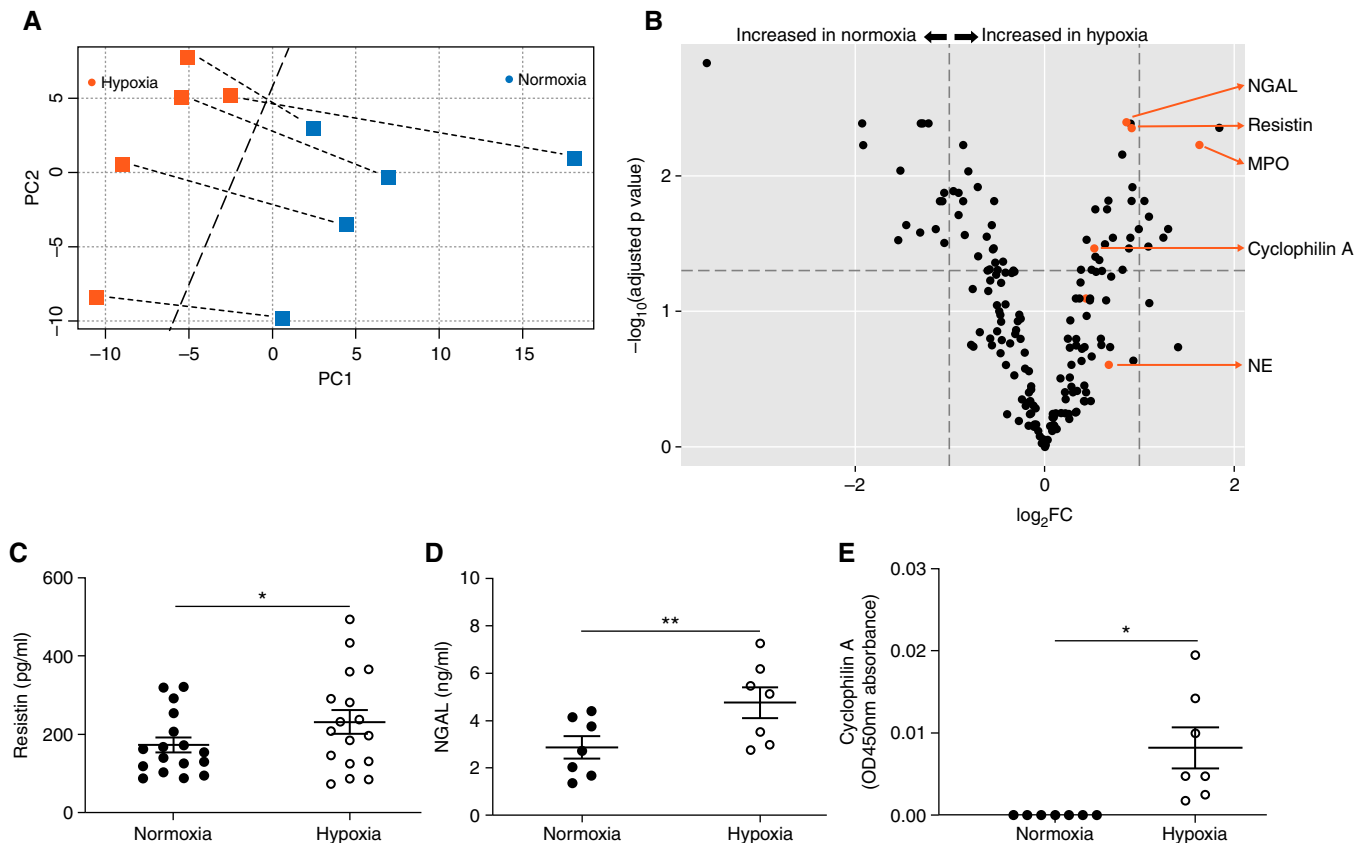
Figure 4. Hypoxia selectively increases granule and cytoplasmic histotoxic protein secretion from activated neutrophils. Neutrophils from healthy donors were incubated under normoxia or hypoxia in the (A and B) presence or (C–E) absence of ethylenediaminetetraacetic acid (1 mM) and sivelestat (10 μM) for 4 hours and then treated with platelet-activating factor (PAF) (1 μM, 5 min) and N-formyl-methionyl-leucyl-phenylalanine (fMLP) (100 nM, 10 min). (A and B) Trypsin-digested supernatants were individually labeled with isobaric tags and subjected to tandem mass spectrometry. (A) Principal component analysis showed separation of normoxic versus hypoxic supernatant samples by PC1 (dashed line) with samples from individual donors indicated by dashed lines connecting hypoxic and normoxic samples ($n = 5$). (B) Volcano plot representation of differential protein expression between paired normoxic and hypoxic supernatants where the vertical dashed lines represent log$_2$ fold change (FC) of protein abundance $= \pm 1$, and the horizontal dashed line represents adjusted P value $= 0.05$ ($n = 5$). (C–E) Neutrophil supernatant content of resistin (C: $n = 17$), NGAL (neutrophil gelatinase-associated lipocalin) (D: $n = 7$), and cyclophilin A (E: $n = 7$) was measured from independent samples by ELISA. Results represent mean ± SEM, (B) paired t test with P value adjusted by Benjamini-Hochberg procedure and (C–E) paired t test. *$P < 0.05$ and **$P < 0.01$. MPO = myeloperoxidase; NE = neutrophil elastase; PC = principal component.

Hypoxia Increases the Capacity for Neutrophil Supernatants to Damage Endothelial Cells

As patients with COPD suffer increased cardiovascular morbidity compared with healthy control subjects and display a footprint of increased circulating protease activity (Figure 1), we investigated whether hypoxia increases the potential for neutrophil-mediated endothelial damage. We incubated supernatants from normoxic versus hypoxic activated (PAF/fMLP) neutrophils with hPAEC and hPMEC monolayers in the presence or absence of α1AT and assessed cell integrity and survival. Supernatants from activated hypoxic neutrophils induced more endothelial detachment (Figures 3A and 3B) and death

(Figures 3C, 3D, and E2) than their normoxic counterparts, which was not completely rescued by coincubation with α1AT (Figure 3D).

Hypoxia Differentially Regulates Protein Release from Activated Neutrophils

Because the antiprotease strategy did not completely mitigate neutrophil-induced endothelial damage (Figure 3D), and multiple neutrophil-derived granule products have potentially damaging actions, we investigated the effect of hypoxia on the total detectable proteome released by activated neutrophils. Mass spectrometry characterization of the normoxic versus hypoxic neutrophil secretome revealed clear

separation by principal component analysis (Figure 4A). TMT-MS identified 1,245 proteins, 717 of which were present in all samples. Of these 717 proteins, 199 had a false discovery rate < 0.01, and 63 were differentially regulated (adjusted P value < 0.05) between normoxia and hypoxia (Figure 4B). Of these 63 proteins, 35 were more abundant in normoxic (Table 1) and 28 were increased in hypoxic (Table 2) neutrophil supernatants. The majority of proteins upregulated in hypoxic supernatants were granule proteins, whereas those more abundant in normoxic supernatants were predominantly cytoplasmic. However, some granule-associated proteins were increased in normoxic samples (e.g., leukocyte-specific

Table 1. Proteins Significantly Increased in Normoxic Neutrophil Supernatants

Accession	Description	Adjusted P Value	FC	Location
Q5TCU8	Tropomyosin β chain	0.001	11.855	CYT
P10599	Thioredoxin	0.017	3.909	CYT
E7EX29	14-3-3 Protein zeta/delta	0.015	3.844	S/G
P52566	Rho GDP-dissociation inhibitor 2	0.004	3.802	CYT
P08670	Vimentin	0.006	3.766	CYT
O00299	Chloride intracellular channel protein 1	0.030	2.918	?
P11021	78-kDa glucose-regulated protein	0.009	2.873	CYT
E9PK25	Cofilin-1	0.023	2.753	CYT
E7EMB3	Calmodulin	0.026	2.488	?
P62993	Growth factor receptor-bound protein 2	0.004	2.478	CYT
P20700	Lamin-B1	0.004	2.435	NUC
Q32MZ4	Leucine-rich repeat flightless-interacting protein 1	0.004	2.337	NUC/CYT
P06737	Glycogen phosphorylase, liver form	0.025	2.217	?
P32942	Intercellular adhesion molecule 3	0.015	2.149	S/G
Q9Y490	Talin-1	0.015	2.116	S/G
O15144	Actin-related protein 2/3 complex subunit 2	0.031	2.086	CYT
P06702	Protein S100-A9	0.013	2.084	CYT
P52209	6-Phosphogluconate dehydrogenase, decarboxylating	0.013	1.950	CYT
P26038	Moesin	0.019	1.878	S
P33241	Leukocyte-specific protein 1	0.013	1.878	S/G
P18206	Vinculin	0.015	1.816	CYT
P30740	Leukocyte elastase inhibitor	0.006	1.813	A
Q96C19	EF-hand domain-containing protein D2	0.027	1.794	?
A6NIZ1	Ras-related protein Rap-1b-like protein	0.009	1.749	SV
P35579	Myosin-9	0.012	1.631	CYT
Q29963	HLA class I histocompatibility antigen, Cw-6 α chain	0.039	1.627	PM
P61247	40S ribosomal protein S3a	0.028	1.528	?
P02042	Hemoglobin subunit delta	0.049	1.492	?
E7EQR4	Ezrin	0.023	1.473	?
P08133	Annexin A6	0.035	1.458	?
P31146	Coronin-1A	0.034	1.451	S/G
Q15907	Ras-related protein Rab-11B	0.015	1.447	G
P46781	40S ribosomal protein S9	0.044	1.438	CYT
P46940	Ras GTPase-activating-like protein IQGAP1	0.049	1.411	G
P39687	Acidic leucine-rich nuclear phosphoprotein 32 family member A	0.043	1.358	?

Definition of abbreviations: A = azurophil granules; CYT = cytoplasm; FC = fold change; G = gelatinase granules; NUC = nucleus; PM = plasma membrane; S = specific granules; SV = secretory vesicles.
Proteins significantly increased in supernatants from normoxic neutrophils (adjusted P value < 0.05), which were present in all 10 samples with a false-discovery rate < 0.01, are listed in order of the magnitude of the FC. Location data were compiled from Reference 50 and the Uniprot database (www.uniprot.org). For some proteins, the location within neutrophils is currently uncertain or unknown (?).

protein 1), and certain cytoplasmic (e.g., cyclophilin A) and nuclear (e.g., histone H4) proteins were increased by hypoxia. Selected hypoxia-upregulated protein targets with the potential to play a role in endothelial damage were biochemically validated using supernatants from independent healthy donors. Concentrations of the azurophil granule protein resistin (Figure 4C), the specific granule protein NGAL (neutrophil gelatinase-associated lipocalin); (Figure 4D), and the cytoplasmic protein cyclophilin A (Figure 4E) were

significantly elevated in hypoxic versus normoxic neutrophil supernatants.

Investigation of Cytoplasmic and Nuclear Protein Secretion

As cyclophilin A is cytoplasmic rather than granule associated, we examined the release of neutrophil-derived microvesicles (NMVs), which contain components derived from parent cells, as a potential source of protein release in addition to degranulation. However, we found no difference in NMV numbers in hypoxic

versus normoxic supernatants (Figure E3A) or in plasma from healthy patients versus patients with COPD (Figure E3B). Furthermore, there was no difference in the content of cyclophilin A between NMVs generated from normoxic versus hypoxic cells, and cyclophilin A was also detected in microvesicle-depleted neutrophil supernatants (Figures E3C and E3D).

Because the nuclear protein, histone H4, was increased in the hypoxic supernatant proteome (although other histones were not likewise increased), we also investigated the release of neutrophil extracellular traps (NETs), which release both nuclear and granule proteins into the extracellular space. However, there was no difference in NETosis from normoxic versus hypoxic neutrophils (Figure E4). Overall, our data do not support a contribution of NMVs or NETs to the differential spectrum of cytoplasmic or nuclear proteins released from neutrophils under hypoxia.

Hypoxic Neutrophils from Patients Experiencing COPD Exacerbation Release More Histotoxic Proteins

During COPD exacerbations (which are associated with excess cardiovascular morbidity), neutrophils are subject to intensified local and systemic hypoxia in addition to a markedly proinflammatory microenvironment. Because hypoxia enables even "healthy" neutrophils to release multiple proteins capable of causing endothelial damage, we studied the effect of hypoxia on neutrophils isolated from patients experiencing COPD exacerbation (Table E1). COPD neutrophils were not basally shape-changed (indicating no priming or activation) and responded identically to healthy control neutrophils after fMLP stimulation (Figure 5A). Furthermore, there was no difference in the release of NE from unstimulated neutrophils obtained from patients experiencing COPD exacerbation versus age- and sex-matched healthy control subjects under normoxic or hypoxic conditions (Figure 5B). Together, these data indicate that, in our cohort 1, circulating neutrophils from patients with COPD were not primed during exacerbations. Despite this, when incubated under hypoxia, stimulated COPD neutrophils released up to threefold more active NE than equivalent healthy control cells (Figure 5B). Likewise, secretion of selected granule and cytoplasmic protein targets identified by proteomics NGAL

Table 2. Proteins Significantly Increased in Hypoxic Neutrophil Supernatants

Accession	Description	Adjusted P Value	FC	Location
P62805	Histone H4	0.004	3.587	NUC
P05164	Myeloperoxidase	0.006	3.107	A
A6NC48	ADP-ribosyl cyclase/cyclic ADP-ribose hydrolase 2	0.025	2.468	SV
O75083	WD repeat-containing protein 1	0.029	2.382	CYT
Q92820	γ-Glutamyl hydrolase	0.020	2.143	S
P02788	Lactotransferrin	0.033	2.136	S
A5A3E0	POTE ankyrin domain family member F	0.015	2.074	CYT
Q0VD83	Apolipoprotein B receptor	0.025	1.993	S/G
P10124	Serglycin	0.012	1.904	CYT
Q9HD89	Resistin	0.004	1.898	A/S
P07737	Profilin-1	0.015	1.890	CYT
A0A087WXL1	Folate receptor γ	0.029	1.877	S
P11215	Integrin α-M	0.004	1.872	S
P16035	Metalloproteinase inhibitor 2	0.034	1.853	G
X6R8F3	Neutrophil gelatinase-associated lipocalin	0.004	1.842	S
V9GYM3	Apolipoprotein A-II	0.049	1.769	?
G3V3D1	Epididymal secretory protein E1	0.007	1.765	A
P10153	Nonsecretory RNase	0.029	1.650	?
P04217	α-1B-glycoprotein	0.015	1.594	?
P05107	Integrin β-2	0.018	1.580	G
P01024	Complement C3	0.032	1.556	?
P20061	Transcobalamin-1	0.042	1.492	S
P78324	Tyrosine-protein phosphatase nonreceptor type substrate 1	0.018	1.449	SV
J3KNB4	Cathelicidin antimicrobial peptide	0.039	1.446	S/G
P62937	Peptidyl-prolyl cis-trans isomerase A/Cyclophilin A	0.035	1.438	CYT
P30086	Phosphatidylethanolamine-binding protein 1	0.049	1.406	?
A0A075B6H6	Ig kappa chain C region	0.030	1.358	?
A0A075B6K9	Ig lambda-2 chain C regions	0.049	1.305	?

Definition of abbreviations: A = azurophil granules; CYT = cytoplasm; FC = fold change; G = gelatinase granules; NUC = nucleus; PM = plasma membrane; S = specific granules; SV = secretory vesicles.
Proteins significantly increased in supernatants from hypoxic neutrophils (adjusted *P* value < 0.05), which were present in all 10 samples with a false discovery rate < 0.01, are listed in order of the magnitude of the FC. Location data were compiled from Reference 50 and the Uniprot database (www.uniprot.org). For some proteins, the location within neutrophils is currently uncertain or unknown (?).

(Figure 5C) and cyclophilin A (Figure 5D) was increased 1.5- and 5-fold, respectively, from stimulated hypoxic neutrophils from patients with COPD versus healthy control subjects, with a similar pattern demonstrated for resistin release (Figure 5E). In contrast, although secretion of the azurophil granule protein MPO (myeloperoxidase) was consistently increased in hypoxia versus normoxia, it was not further enhanced when comparing COPD and healthy control neutrophils (Figure 5F).

Hypoxia Promotes Endothelial–Neutrophil Interaction Induced by COPD Patient Neutrophil Supernatants

To investigate whether neutrophils from patients with COPD have increased capacity for endothelial cell injury and/or activation, we applied supernatants from normoxic or hypoxic activated COPD versus healthy neutrophils to hPMEC monolayers and assessed neutrophil recruitment (rolling, adhesion, and transmigration) in a biologically relevant *in vitro* flow system (Figures 6A and 6B). Treatment with hypoxic COPD neutrophil supernatants resulted in a marked increase in neutrophil rolling and adhesion compared with both normoxic COPD supernatants and hypoxic healthy supernatants (Figure 6C).

Hypoxia May Synergize with Inflammatory Mediators to Promote Upregulation of Circulating Histotoxic Neutrophil Granule Proteins in Patients with COPD

Because hypoxia enhanced histotoxic protein release from COPD neutrophils (Figure 5),

and supernatants from these cells promoted neutrophil–endothelial interaction (Figure 6), we examined whether there was a circulating signature of hypoxia-induced neutrophil protein secretion. We detected significantly increased concentrations of the neutrophil granule proteins NE (Figure 7A), MPO (Figure 7B), and NGAL (Figure 7C) in patients experiencing COPD exacerbation versus healthy control plasma (derived from an independent cohort 3) (Table E3), although there was no difference in the plasma content of resistin (Figure 7D). Despite the observation of increased release of the cytoplasmic protein cyclophilin A from isolated COPD neutrophils (Figure 5E), unexpectedly, the plasma content of cyclophilin A was higher for healthy control subjects than patients with COPD (Figure 7E). Biomarkers of vascular injury or activation ICAM-1 (intercellular adhesion molecule-1) (Figure 7F) and VCAM-1 (vascular cell adhesion molecule-1) (Figure 7G) and inflammation (Figures E5A and E5B) were increased in plasma from patients experiencing COPD exacerbation. In contrast, the COPD plasma content of angiogenesis biomarkers was predominantly unchanged compared with that of healthy control subjects (Figures E5C–E5J), suggesting a damage phenotype that may enhance endothelial–neutrophil interaction but without a corresponding increase in vascular regeneration ability.

Discussion

Our work demonstrates that hypoxic neutrophils display a hypersecretory phenotype with enhanced capacity to both activate and injure cultured endothelial cells. Hypoxia-driven release of histotoxic proteins was observed from healthy donor neutrophils and translated to a pathophysiologically relevant cohort of patients experiencing COPD exacerbation, confirming even further augmented histotoxic protein secretion under hypoxia, together with higher circulating concentrations of selected neutrophil-derived proteins and a plasma signature of increased neutrophil protease activity and vascular injury.

Our data from human neutrophils treated with PI3K isoform-selective inhibitors and from transgenic murine cells support a nonredundant role for PI3Kγ in the hypoxic augmentation of NE release.
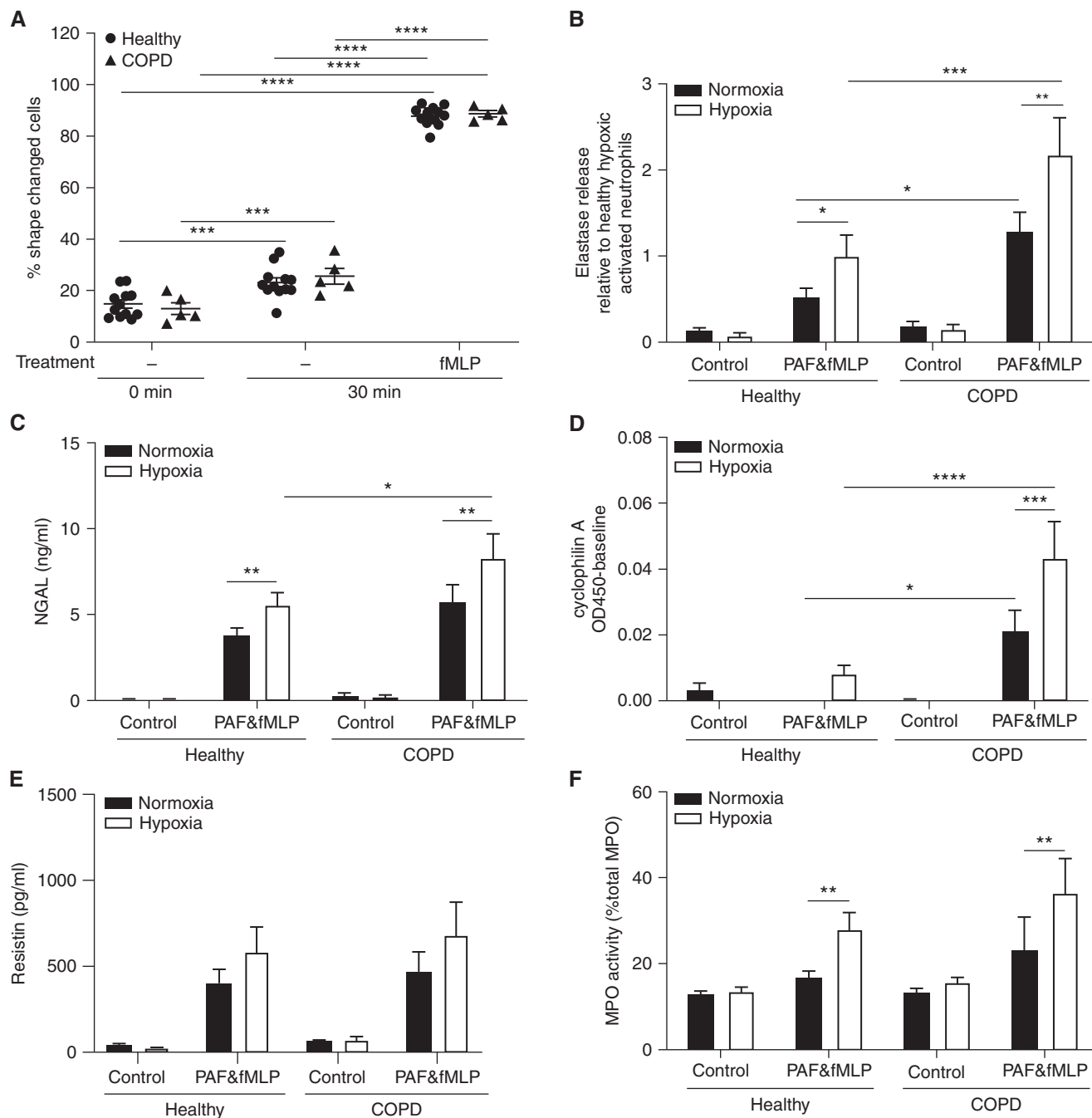
Figure 5. Hypoxia further augments histotoxic protein release from chronic obstructive pulmonary disease (COPD) versus healthy neutrophils. (*A*) Neutrophils from healthy donors or patients experiencing COPD exacerbation were incubated under normoxia and treated with *N*-formyl-methionyl-leucyl-phenylalanine (fMLP) (100 nM, 30 min) or vehicle control. Shape change was assessed by flow cytometric analysis of forward scatter, expressed as percentage shape-changed cells of total population ($n = 5$–12). (*B*–*F*) Neutrophils from healthy donors or patients experiencing COPD exacerbation were incubated under normoxia or hypoxia for 4 hours and then treated with platelet-activating factor (PAF) (1 μM, 5 min) and fMLP (100 nM, 10 min) or vehicle control. Supernatant content of elastase (*B*: $n = 7$–14), NGAL (neutrophil gelatinase-associated lipocalin) (*C*: $n = 7$–12), cyclophilin A (*D*: $n = 3$–7), resistin (*E*: $n = 7$–12), and MPO (myeloperoxidase) (*F*: $n = 3$–6) was measured by ELISA or activity assay. Supernatant neutrophil elastase activity is expressed as fold change relative to healthy hypoxic activated neutrophils. All samples were obtained from cohort 1. Results represent mean ± SEM, two-way ANOVA. *$P < 0.05$, **$P < 0.01$, ***$P < 0.001$, and ****$P < 0.0001$.
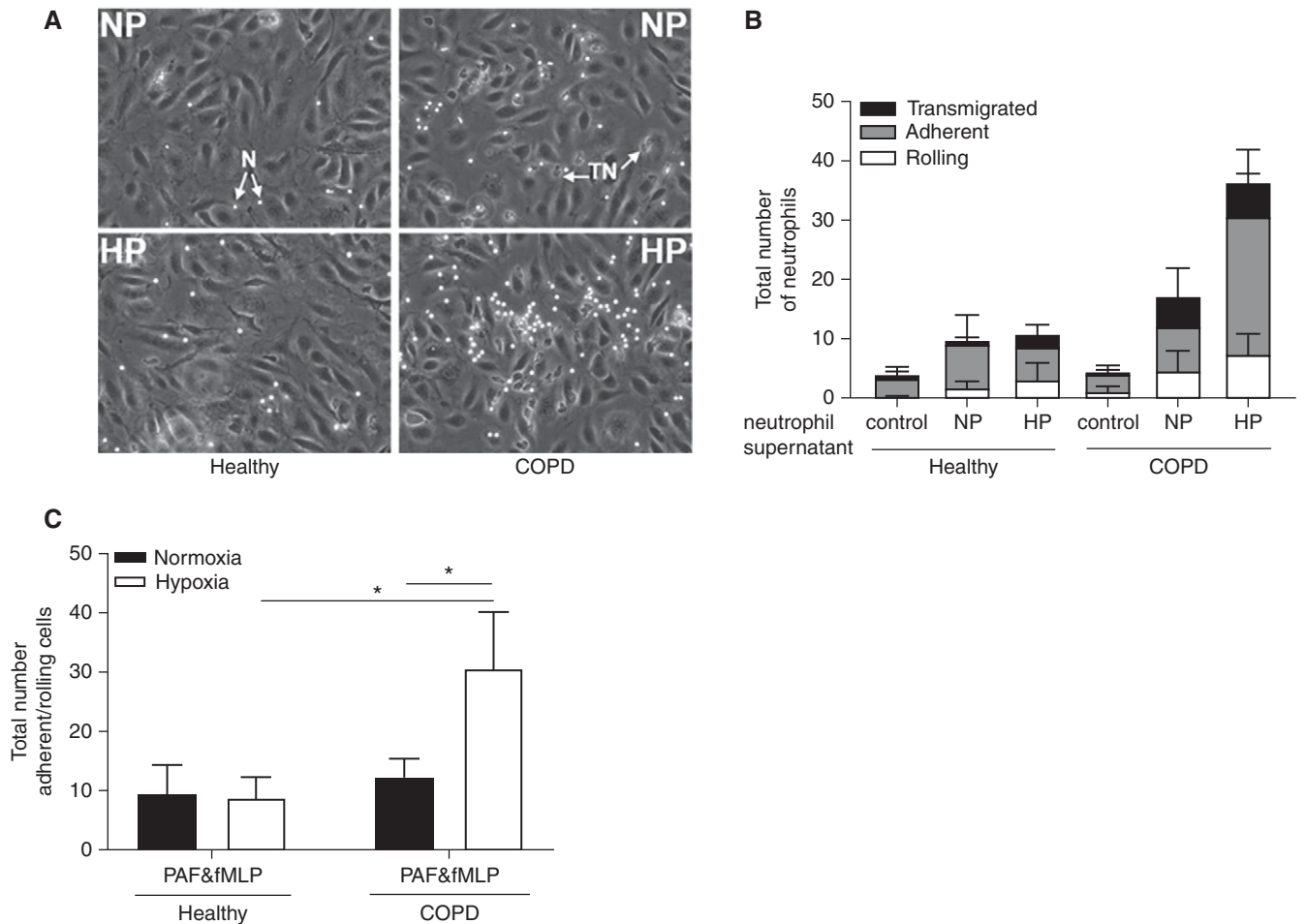
Figure 6. Hypoxia accentuates endothelial–neutrophil interaction induced by chronic obstructive pulmonary disease (COPD) versus healthy neutrophil supernatants. Neutrophils from healthy donors or patients experiencing COPD exacerbation were incubated under normoxia or hypoxia for 4 hours and then treated with platelet-activating factor (PAF) (1 μM, 5 min) and *N*-formyl-methionyl-leucyl-phenylalanine (fMLP) (100 nM, 10 min). Supernatants from normoxic (NP) or hypoxic (HP) PAF- or fMLP-treated neutrophils were incubated with confluent human pulmonary microvascular endothelial cells (hPMECs) for 4 hours in the presence of serum (2%). Washed hPMECs were perfused with healthy neutrophils. (*A*) Representative images (original magnification, ×100) showing arrested/rolling neutrophils (N) as bright phase and transmigrated neutrophils (TN) as dark phase. (*B*) Endothelial–neutrophil interactions (total number of rolling, adhered, and transmigrated neutrophils after bolus neutrophil injection) were captured with time-lapse imaging (*n* = 3). (*C*) Quantification of neutrophil rolling and adherence was performed using ImagePro software (*n* = 3). All neutrophil supernatant samples were obtained from cohort 1. Results represent mean ± SEM, two-way ANOVA. *$P < 0.05$.

Dysregulated PI3K signaling has previously been associated with COPD, with the impaired neutrophil migratory accuracy improved by PI3Kγ/δ inhibition (30), although we found no role for the δ isoform in hypoxic degranulation. Our results suggest that PI3Kγ is required for hypoxia-potentiated neutrophil degranulation, whether in response to tyrosine kinase– (2) or G-protein–coupled agonists, such as PAF. Consistent with a role for PI3Kγ-dependent PAF/hypoxia-mediated neutrophil degranulation in vascular insults, PI3Kγ/δ inhibition limited both PAF-induced hindlimb inflammation and ischemic cardiac infarct size (31), and PI3Kγ-null mice had improved cardiac recovery after ischemia

(32). Hence, this signaling pathway could conceivably be targeted to mitigate neutrophil-mediated endothelial damage in COPD and other diseases underpinned by hypoxia, inflammation, and vascular damage.

Hypoxia enhances neutrophil degranulation, but, surprisingly, hypoxia-upregulated proteins identified by proteomic analysis did not segregate precisely with discrete granule populations. We have previously shown that hypoxia promotes differential secretion from eosinophil granules (26), with similar results reported using mast cells (33). Our results imply a comparable "differential degranulation" may be occurring from neutrophils in the

setting of hypoxia. A limited number of studies analyzing the neutrophil secretome generated under normoxic conditions have shown variation in protein content according to the inciting stimulus (34), suggesting that the (potentially hypoxic) inflammatory environment can influence the precise composition of secreted granule proteins. This may also explain our observation that release of the azurophilic granule proteins NE, resistin, and MPO from hypoxic COPD neutrophils did not fully mirror each other.

In addition to enhanced degranulation, our proteomic data further suggest active secretion of selected cytoplasmic and nuclear proteins under
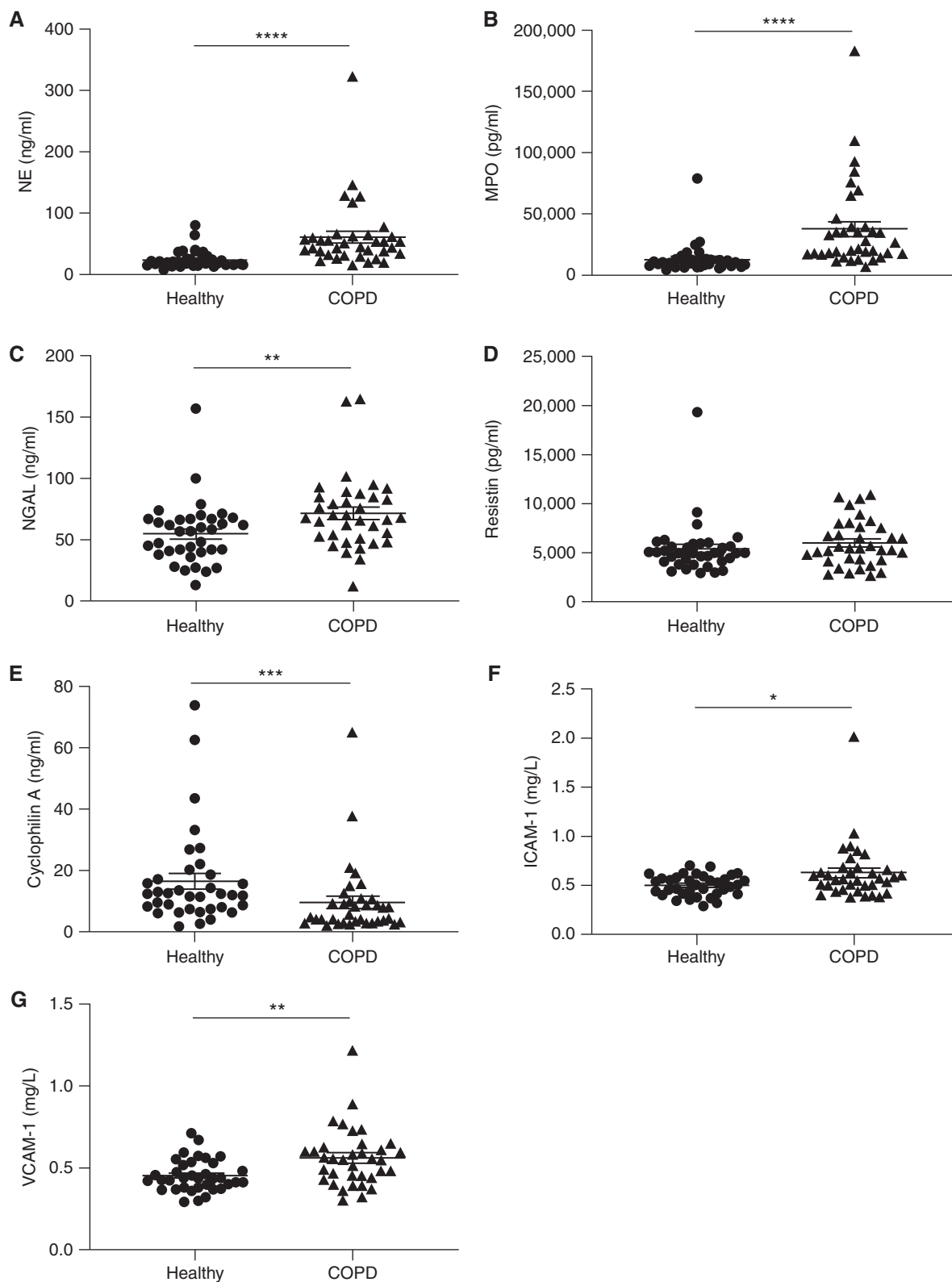
Figure 7. Plasma from patients with chronic obstructive pulmonary disease (COPD) has increased content of hypoxia-upregulated histotoxic granule proteins and vascular injury biomarkers. Plasma from healthy donors or patients experiencing COPD exacerbation was assessed for content of (*A*) neutrophil elastase (NE), (*B*) MPO (myeloperoxidase), (*C*) NGAL (neutrophil gelatinase-associated lipocalin), (*D*) resistin, (*E*) cyclophilin A, (*F*) ICAM-1 (intercellular adhesion molecule-1), and (*G*) VCAM-1 (vascular cell adhesion molecule-1) by ELISA (NE, NGAL, and cyclophilin A) or chemiluminescence immunoassay (MPO, resistin, ICAM-1, and VCAM-1) (*n* = 36 healthy, *n* = 36 COPD; 4 samples from cohort 1 and 32 samples from cohort 3). Results represent mean ± SEM, Mann-Whitney test. *$P < 0.05$, **$P < 0.01$, ***$P < 0.001$, and ****$P < 0.0001$.

hypoxia. We provide the first description of neutrophilic secretion of the proinflammatory cytoplasmic protein cyclophilin A, aligning to a previous study demonstrating its hypoxia-driven secretion from cardiac myocytes (35). Our data also demonstrated increased release of the nuclear protein histone H4 (although no other histones) in hypoxic supernatants. The source of this protein remains unclear, as we did not observe any difference in NETosis between normoxic and hypoxic neutrophils.

We have established that peripheral blood neutrophils from patients experiencing COPD exacerbation display markedly greater hypoxic release of NE, NGAL, and cyclophilin A relative to matched healthy control subjects. These proteins have been previously implicated in endothelial dysfunction or atherosclerosis (36, 37), with raised circulating levels of NGAL associated with cardiovascular events and with hypoxemia in COPD (38, 39). Although we found no relationship between admission or venesection oxygen levels and protein release (data not shown), this may have been confounded by prior exposure to supplemental oxygen. In keeping with a potential *in vivo* role for hypoxic augmentation of neutrophil degranulation, we found higher levels of the granule proteins NE, MPO, and NGAL and a footprint of increased NE and PR3 activity in COPD versus healthy plasma, aligning with recent severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) studies, where NGAL in particular was associated with mortality (40). Our results suggest that enhanced neutrophil degranulation in the setting of hypoxia may contribute to systemic endothelial injury in patients experiencing COPD exacerbation and potentially other conditions, such as SARS-CoV-2 infection.

Primed circulating neutrophils with heightened potential for cellular damage have been identified in patients with COPD (21). However, we did not observe significant shape change or enhanced degranulation (both features of priming) from unstimulated COPD neutrophils, suggesting that the enhanced hypoxic release of granule proteins from these cells is not simply a consequence of priming and may "substitute" for this process in promoting damage. A potential explanation for enhanced protein release

is a change in protein abundance in COPD neutrophils, with one study showing increased NE activity in COPD versus healthy neutrophil lysates (41). However, this result may represent a protease/antiprotease imbalance, as leukocyte elastase inhibitor was reduced, a finding consistent with our proteomic data and that may explain the discrepancy between these results and the NE activity assay.

Our study has some limitations. Although our patients and control subjects were age and sex matched, the patients had a significant smoking history and more comorbidities than the healthy volunteers. The patients with COPD were studied during exacerbations, a high-risk period for acute cardiovascular events (17), but with variation in terms of exacerbation etiology and severity; however, the use of two separate cohorts in different institutions mitigates this limitation. It is possible that acute or long-term medications could affect neutrophil function in our patient cohorts. For example, all patients with COPD in cohort 1 were taking inhaled combination corticosteroid and long-acting β-agonists and were treated with oral prednisolone during exacerbation with varying duration before venesection. However, although glucocorticoids are known to delay neutrophil apoptosis, this effect is lost in hypoxia (42), and inhaled corticosteroids have been shown not to affect neutrophil protein secretion (43). Given the variation in comorbidities (Table E4), and thus treatments, within our cohorts, there is unlikely to be any consistent impact on neutrophil function.

Although our results are predominantly consistent with previous studies, we note that despite reports of enhanced circulating levels of resistin and cyclophilin A in patients with COPD versus healthy control subjects (44, 45), we did not detect such an increase in our COPD cohort. These discrepant results may reflect differences in the timing of sampling, medications, patient heterogeneity, or obscuration of a neutrophil-specific signal, because both proteins have multiple cellular sources (46, 47). Both cyclophilin A and its receptor have been detected at high levels in atherosclerotic plaques (48). Hence, in COPD, cyclophilin A may already be membrane bound, preventing its detection in plasma. As patients with COPD have

increased atherosclerotic burden, hypoxia within plaques may promote enhanced local release of cyclophilin A from neutrophils in this and other microenvironments. This would be consistent with our *in vitro* data and with a previous study demonstrating increased cyclophilin A expression in the lung tissue of patients with COPD compared with either smoking or nonsmoking control subjects (49).

Overall, our data demonstrate that patients experiencing COPD exacerbation have an enhanced footprint of circulating neutrophil protease activity and that neutrophils from these patients exhibit a hypoxia-driven hypersecretory phenotype with enhanced capacity for endothelial damage. We provide the first description of the ability of hypoxia to augment the secretion of histotoxic proteins from COPD neutrophils *in vitro* and have identified a corresponding increase in the plasma concentrations of selected granule proteins and markers of vascular injury in patients with COPD. Our findings may illuminate novel therapeutic targets in treatment-recalcitrant neutrophil-mediated inflammatory diseases such as COPD. ∎

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