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### Journal

Environmental Science and Technology, 58(14)

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### Publication Date

2024-04-09

### DOI

10.1021/acs.est.3c06289

Peer reviewed

# Oxidative Transformation of Nafion-Related Fluorinated Ether Sulfonates: Comparison with Legacy PFAS Structures and Opportunities of Acidic Persulfate Digestion for PFAS Precursor Analysis

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Cite This: *Environ. Sci. Technol.* 2024, 58, 6415–6424



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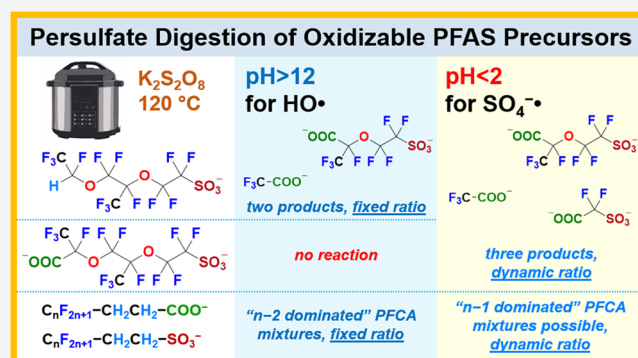
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**ABSTRACT:** The total oxidizable precursor (TOP) assay has been extensively used for detecting PFAS pollutants that do not have analytical standards. It uses hydroxyl radicals ( $\text{HO}^\bullet$ ) from the heat activation of persulfate under alkaline pH to convert H-containing precursors to perfluoroalkyl carboxylates (PFCAs) for target analysis. However, the current TOP assay oxidation method does not apply to emerging PFAS because (i) many structures do not contain C–H bonds for  $\text{HO}^\bullet$  attack and (ii) the transformation products are not necessarily PFCAs. In this study, we explored the use of classic acidic persulfate digestion, which generates sulfate radicals ( $\text{SO}_4^{\bullet-}$ ), to extend the capability of the TOP assay. We examined the oxidation of Nafion-related ether sulfonates that contain C–H or  $-\text{COO}^-$ , characterized the oxidation products, and quantified the F atom balance. The  $\text{SO}_4^{\bullet-}$  oxidation greatly expanded the scope of oxidizable precursors. The transformation was initiated by decarboxylation, followed by various spontaneous steps, such as HF elimination and ester hydrolysis. We further compared the oxidation of legacy fluorotelomers using  $\text{SO}_4^{\bullet-}$  versus  $\text{HO}^\bullet$ . The results suggest novel product distribution patterns, depending on the functional group and oxidant dose. The general trends and strategies were also validated by analyzing a mixture of 100000- or 10000-fold diluted aqueous film-forming foam (containing various fluorotelomer surfactants and organics) and a spiked Nafion precursor. Therefore, (1) the combined use of  $\text{SO}_4^{\bullet-}$  and  $\text{HO}^\bullet$  oxidation, (2) the expanded list of standard chemicals, and (3) further elucidation of  $\text{SO}_4^{\bullet-}$  oxidation mechanisms will provide more critical information to probe emerging PFAS pollutants.

**KEYWORDS:** total oxidizable precursor (TOP) assay, Nafion byproducts (NBPs), acid persulfate digestion, fluorotelomer, sulfate radical, hydroxyl radical, aqueous film-forming foam (AFFF), oxidative defluorination



## INTRODUCTION

The study of per- and polyfluoroalkyl substance (PFAS) pollutants has been extended from the early focus on all-carbon perfluoroalkyl carboxylates (PFCAs,  $\text{C}_n\text{F}_{2n+1}-\text{COO}^-$ ) and perfluoroalkanesulfonates (PFSAs,  $\text{C}_n\text{F}_{2n+1}-\text{SO}_3^-$ ) to a large variety of "novel" structures with various heteroatoms, branching patterns, and chain lengths.<sup>1–3</sup> For example, many fluorinated ether structures have been detected in water and soil worldwide<sup>4–7</sup> and shown various toxicities.<sup>8–13</sup> The belated attention to these structures is primarily attributed to (1) the information gap between fluorine chemistry and environmental chemistry and (2) the lack of analytical standards for targeted analysis.<sup>14</sup>

For the detection of telomer ( $\text{C}_n\text{F}_{2n+1}-(\text{CH}_2)_m-\text{X}$ , X = highly diverse organic functional groups) and sulfonamide ( $\text{C}_n\text{F}_{2n+1}-\text{SO}_2\text{NH}-\text{X}$ ) surfactants, a total oxidizable precursor (TOP)

assay has been widely applied.<sup>14</sup> The H atoms in the "precursor" molecules allow oxidation by hydroxyl radicals ( $\text{HO}^\bullet$ , generated from persulfate at pH > 12 at 85 °C), yielding one or multiple PFCAs for targeted analysis.<sup>15,16</sup> Our lab further modified the reaction condition (e.g.,  $[\text{NaOH}]:[\text{K}_2\text{S}_2\text{O}_8] = 5:1$  and 120 °C), extended the substrate scope (e.g.,  $\text{C}_n\text{F}_{2n+1}-(\text{CH}_2)_2-\text{COO}^-$ ,  $\text{C}_n\text{F}_{2n+1}-(\text{CH}_2)_2-\text{SO}_3^-$ , and  $\text{HC}_n\text{F}_{2n}-\text{COO}^-$  with variable n) and quantified short-chain products (e.g.,  $\text{CF}_3-\text{COO}^-$ ,  $\text{C}_2\text{F}_5-$

Received: August 4, 2023

Revised: March 9, 2024

Accepted: March 11, 2024

Published: March 26, 2024



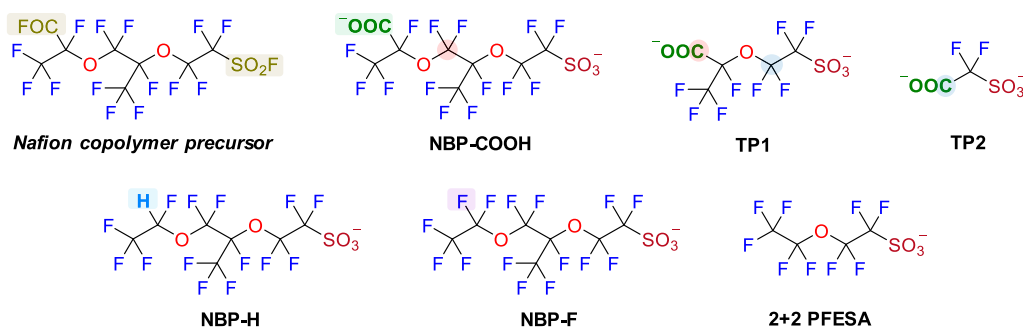


Figure 1. Nafion-related PFAS structures studied in this work.

COO<sup>-</sup>, and <sup>-</sup>OOC-CF<sub>2</sub>-COO<sup>-</sup>).<sup>17,18</sup> Notably, structures with only one C-H bond in HCF<sub>2</sub>- and multiple C-H bonds in -(CH<sub>2</sub>)<sub>2</sub>- showed distinct transformation patterns.<sup>18</sup> A previous TOP assay of various fluoro ether structures by Zhang et al.<sup>19</sup> found that only H-containing structures could be converted by HO<sup>•</sup>, but the product information remained largely unknown.<sup>20</sup>

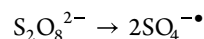
The findings presented above reflect several limitations (but not necessarily disadvantages) of the current TOP assay. First, HO<sup>•</sup> oxidation requires H atoms in the precursor and does not transform either PFCAs or PFSA. In contrast, oxidation using sulfate radicals (SO<sub>4</sub><sup>-•</sup>, generated from persulfate and preserved at pH < 2) is a well-adopted pretreatment for water sample analysis. For example, the measurement of total phosphorus uses acidic persulfate digestion to convert various organic phosphate esters into orthophosphate.<sup>21,22</sup> SO<sub>4</sub><sup>-•</sup> can also oxidize PFCAs via decarboxylation.<sup>23–26</sup> Second, “novel” PFAS structures do not necessarily produce linear PFCAs,<sup>14,19</sup> which the current TOP assay relies on. However, while the reaction kinetics under oxidative conditions have been extensively quantified, the structural transformation of emerging PFAS has not been adequately understood.<sup>20,26</sup> Hence, it is important to elucidate the transformation of novel PFAS “precursors” by both SO<sub>4</sub><sup>-•</sup> and HO<sup>•</sup> for environmental detection and source tracking.

In this work, we examined the oxidative transformation of perfluorinated and polyfluorinated ether sulfonates (PFESAs) related to Nafion polymer production. Nafion membranes are extensively used in the chloralkali electrochemical process<sup>27</sup> and fuel cells.<sup>28</sup> Various “Nafion byproduct” PFESAs have been detected in surface water,<sup>4,5</sup> aquatic animals,<sup>8,29,30</sup> and humans.<sup>31</sup> However, these PFESAs with ether bonds and -CF<sub>3</sub> branches (Figure 1) cannot be detected by the current TOP assay.<sup>14</sup> The findings from PFESA oxidation are further compared with those of the extensively studied legacy PFAS to (1) reveal novel mechanistic insights and (2) add new tools for the TOP assay of emerging PFAS pollutants.

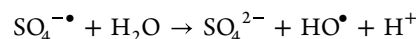
## MATERIALS AND METHODS

**Chemicals.** Information on PFAS chemicals, the preparation of stock solutions, and the conversion of structures containing acyl fluoride (-COF) and/or sulfonyl fluoride (-SO<sub>2</sub>F) into the corresponding carboxylate (-COO<sup>-</sup>) and/or sulfonate (-SO<sub>3</sub><sup>-</sup>) are described in the Supporting Information (SI). Potassium persulfate (K<sub>2</sub>S<sub>2</sub>O<sub>8</sub>), sulfuric acid (H<sub>2</sub>SO<sub>4</sub>), and sodium hydroxide (NaOH) were purchased from Fisher Chemical.

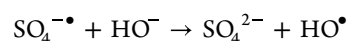
**Oxidative Reactions.** At the elevated temperature, each S<sub>2</sub>O<sub>8</sub><sup>2-</sup> decomposes into two sulfate radicals (SO<sub>4</sub><sup>-•</sup>), which are relatively abundant at pH < 2:



For the oxidation using SO<sub>4</sub><sup>-•</sup>, a 35 mL heavy-walled glass pressure vessel (Synthware Glass, #P160002D) was loaded with 30 mL of the aqueous solution containing 0.5 mM of individual PFAS and variable concentrations of K<sub>2</sub>S<sub>2</sub>O<sub>8</sub> (5–100 mM, 10–200 equiv to PFAS). The initial pH was adjusted by H<sub>2</sub>SO<sub>4</sub> to 2.0. The pressure vessels were covered by a threaded cap, which was not closely tightened to avoid pressure buildup. Multiple vessels were vertically immersed in water inside a 250 mL glass beaker. The beaker was placed in a pressure cooker (Farberware 6 Quart, loaded with 1 L of water), which was heated to 120 °C within 20 min and maintained at 120 °C for 40 min. The final pH of the reaction mixture was below 2.0. Without an adequate base, the aqueous solution can be further acidified:



For the oxidation using HO<sup>•</sup>, the reaction settings remained the same except that five molar equivalents of NaOH (relative to K<sub>2</sub>S<sub>2</sub>O<sub>8</sub>) were also added to ensure pH > 12.0 throughout the reaction. At alkaline pH, SO<sub>4</sub><sup>-•</sup> radicals are primarily converted to hydroxyl radicals (HO<sup>•</sup>):



**Water Sample Analysis.** PFAS parent compounds and transformation products (TPs) were measured by high-performance liquid chromatography equipped with a high-resolution quadrupole Orbitrap mass spectrometer (HPLC-HRMS/MS, Q Exactive, Thermo Fisher Scientific).<sup>17</sup> The instrument settings and quality assurance/control details are provided in the SI. The released fluoride ion (F<sup>-</sup>) was measured with an ion-selective electrode (ISE, Fisherbrand Accumet) with a Thermo Scientific Orion Versa Star Pro meter. Each sample was added with an equal volume of total ionic strength adjustment buffer (TISAB for fluoride electrode, Thermo Scientific). The measurement accuracy in the persulfate oxidation matrix has been validated in our previous report.<sup>17</sup> The F<sup>-</sup> released from -COF and -SO<sub>2</sub>F in the hydrolysis pretreatment was not counted for the oxidative defluorination percentage (deF%).

## RESULTS AND DISCUSSION

**Oxidation of PFESAs by HO<sup>•</sup>.** We tested six structurally relevant PFESAs (Figure 1). NBP-COOH is the hydrolysis product of the Nafion copolymer precursor.<sup>5</sup> NBP-H (commonly named “Nafion byproduct 2”) is the decarboxylation product of NBP-COOH. TP1 is a shorter analogue of NBP-COOH. TP2 can be considered as the further shortened “analogue” of TP1. NBP-F is a fully fluorinated structure without either a -COO<sup>-</sup>

group or C–H bond on the terminal. **2** + **2** PFESA can be considered a shorter and linear analogue of NBP-F. Instead of meticulously measuring second-order rate constants,<sup>20</sup> we evaluated the reactivity using F<sup>−</sup> release as an established rapid probe of oxidative degradation.<sup>17</sup>

As expected, the five structures without a C–H bond for HO• attack allowed <5% of defluorination (Table 1, entries 2, 6, 11,

**Table 1. Oxidative Defluorination of Various PFAS Chemicals**

| entry | PFAS chemical (name used in this paper)   | [S <sub>2</sub> O <sub>8</sub> <sup>2−</sup> ]: [PFAS] molar ratio | dominant radical              | overall defF (%) <sup>a</sup> |
|-------|---|--|-------------------------------|-------------------------------|
| 1     | <sup>−</sup> OOC–CF(CF <sub>3</sub> )–O–CF <sub>2</sub> CF(CF <sub>3</sub> )–O–CF <sub>2</sub> CF <sub>2</sub> –SO <sub>3</sub> <sup>−</sup> (NBP-COOH) | 100:1  | SO <sub>4</sub> <sup>•−</sup> | 46.4                          |
| 2     |   | 100:1  | HO•                           | 2.8 <sup>b</sup>              |
| 3     | H–CF(CF <sub>3</sub> )–O–CF <sub>2</sub> CF(CF <sub>3</sub> )–O–CF <sub>2</sub> CF <sub>2</sub> –SO <sub>3</sub> <sup>−</sup> (NBP-H)                   | 100:1  | SO <sub>4</sub> <sup>•−</sup> | 79.7                          |
| 4     |   | 100:1  | HO•                           | 25.0                          |
| 5     | <sup>−</sup> OOC–CF(CF <sub>3</sub> )–O–CF <sub>2</sub> CF <sub>2</sub> –SO <sub>3</sub> <sup>−</sup> (TP1)   | 100:1  | SO <sub>4</sub> <sup>•−</sup> | 76.5                          |
| 6     |   | 100:1  | HO•                           | 3.0 <sup>b</sup>              |
| 7     | <sup>−</sup> OOC–CF <sub>2</sub> –SO <sub>3</sub> <sup>−</sup> (TP2)  | 10:1   | SO <sub>4</sub> <sup>•−</sup> | 77.9                          |
| 8     |   | 20:1   | SO <sub>4</sub> <sup>•−</sup> | 85.8                          |
| 9     |   | 50:1   | SO <sub>4</sub> <sup>•−</sup> | 88.4                          |
| 10    |   | 100:1  | SO <sub>4</sub> <sup>•−</sup> | 89.5                          |
| 11    |   | 100:1  | HO•                           | 1.1 <sup>b</sup>              |
| 12    | CF <sub>3</sub> –COO <sup>−</sup> (TFA)   | 10:1   | SO <sub>4</sub> <sup>•−</sup> | 53.3                          |
| 13    |   | 20:1   | SO <sub>4</sub> <sup>•−</sup> | 61.8                          |
| 14    |   | 50:1   | SO <sub>4</sub> <sup>•−</sup> | 79.6                          |
| 15    |   | 100:1  | SO <sub>4</sub> <sup>•−</sup> | 90.9                          |
| 16    |   | 100:1  | HO•                           | 0.9 <sup>b</sup>              |
| 17    | C <sub>6</sub> F <sub>13</sub> –CH <sub>2</sub> CH <sub>2</sub> –SO <sub>3</sub> <sup>−</sup> (6:2 FTSA)  | 100:1  | SO <sub>4</sub> <sup>•−</sup> | 8.3                           |
| 18    |   | 100:1  | HO•                           | 54.2                          |
| 19    | C <sub>6</sub> F <sub>13</sub> –CH <sub>2</sub> CH <sub>2</sub> –COO <sup>−</sup> (6:3 FTCA)  | 100:1  | SO <sub>4</sub> <sup>•−</sup> | 41.5                          |
| 20    |   | 100:1  | HO•                           | 49.5                          |
| 21    | CF <sub>3</sub> CF <sub>2</sub> –O–CF <sub>2</sub> CF(CF <sub>3</sub> )–O–CF <sub>2</sub> CF <sub>2</sub> –SO <sub>3</sub> <sup>−</sup> (NBP-F)         | 100:1  | SO <sub>4</sub> <sup>•−</sup> | 4.8 <sup>c</sup>              |
| 22    |   | 100:1  | HO•                           | 4.7 <sup>c</sup>              |
| 23    | CF <sub>3</sub> CF <sub>2</sub> –O–CF <sub>2</sub> CF <sub>2</sub> –SO <sub>3</sub> <sup>−</sup> (2 + 2 PFESA)  | 100:1  | SO <sub>4</sub> <sup>•−</sup> | 1.0 <sup>c</sup>              |
| 24    |   | 100:1  | HO•                           | 0.4 <sup>c</sup>              |

<sup>a</sup>The calculation considered the total F in the initial PFAS (0.5 mM) to reflect the general efficacy of oxidation. In comparison, the “molecular oxidative defF” in Tables 2 and 3 indicate the average defluorination from each degraded molecule. <sup>b</sup>The limited but significant defluorination at pH > 12 could be attributed to (1) the reaction with a small fraction of SO<sub>4</sub><sup>•−</sup> radicals before they were quenched by HO<sup>−</sup> to yield HO• or (2) impurities containing C–H or other functional groups that are reactive with HO•. <sup>c</sup>The limited but significant defluorination could be attributed to impurities containing C–H or other functional groups that are reactive with HO• and SO<sub>4</sub><sup>•−</sup>.

22, and 24), whereas NBP-H allowed up to 25% of defluorination at [S<sub>2</sub>O<sub>8</sub><sup>2−</sup>]:[NBP-H] = 100:1 (Table 1, entry 4). The previous study did not identify the oxidation product from NBP-H.<sup>20</sup> In this work, following the established “H-abstraction” mechanism by HO•, we predicted that the branched terminal •CF(CF<sub>3</sub>)–O–R<sub>F</sub> radical (Scheme 1a) would bind with another HO• to yield HO–CF(CF<sub>3</sub>)–O–R<sub>F</sub>. This structure with one –F and one –OH on the same carbon would undergo HF elimination to yield the ester CF<sub>3</sub>CO–O–R<sub>F</sub>, which would further hydrolyze under the alkaline condition<sup>32</sup> to afford CF<sub>3</sub>COO<sup>−</sup> (TFA, inert to HO• as shown in Table 1, entry 16) and HO–R<sub>F</sub>. The HO–R<sub>F</sub>, in this case, would further undergo HF elimination and subsequent hydrolysis to yield TP1. With TP1 as the analytical standard, we verified our prediction and quantified the formation of both TP1 and TFA (Table 2, zone A). At various [S<sub>2</sub>O<sub>8</sub><sup>2−</sup>]:[NBP-H] ratios from 10:1 to 100:1, the F atom balance calculated from the residual NBP-H, the released F<sup>−</sup>, and the TP1 and TFA products reached 92–99%, indicating the dominance of the simple oxidative pathway under HO• oxidation. As the [S<sub>2</sub>O<sub>8</sub><sup>2−</sup>]:[NBP-H] ratio went over 20:1, the formation of TP1 (0.43–0.47 mM) and TFA (0.42–0.45 mM) were nearly stoichiometric from the initial NBP-H (0.5 mM).

**Oxidation of PFESAs by SO<sub>4</sub><sup>•−</sup>.** When the oxidant was switched to SO<sub>4</sub><sup>•−</sup>, all structures containing –COO<sup>−</sup> allowed significant defluorination (Table 1, entries 1, 3, 5, 10, and 15). As expected, fully fluorinated NBP-F and 2 + 2 PFESA were not reactive (Table 1, entries 21 and 23). Following the established “decarboxylation” mechanism by SO<sub>4</sub><sup>•−</sup>, we predicted that NBP-COOH would yield the same •CF(CF<sub>3</sub>)–O–R<sub>F</sub> radical (Scheme 1b) as that from the reaction between NBP-H and HO•. The carbon radical would bind with another SO<sub>4</sub><sup>•−</sup> to yield <sup>−</sup>O<sub>3</sub>S–O–CF(CF<sub>3</sub>)–O–R<sub>F</sub>, which would further hydrolyze under the acidic condition<sup>33</sup> to HO–CF(CF<sub>3</sub>)–O–R<sub>F</sub>. The following steps would yield TP1 and TFA as described above. However, SO<sub>4</sub><sup>•−</sup> could further decarboxylate TP1 following the same set of mechanisms to yield TP2 and another equivalent of TFA. We verified our prediction and quantified the generation of TP1, TP2, and TFA from both NBP-COOH (Table 2, zone C) and NBP-H (Table 2, zone B). The SO<sub>4</sub><sup>•−</sup> oxidation of TP1 also generated expected TP2 and TFA (Table 2, zone D).

A closer data examination led to a series of mechanistic insights. First, when a C–H bond is present in NBP-H, the degradation by SO<sub>4</sub><sup>•−</sup> was much more efficient than by HO•. At the lowest [S<sub>2</sub>O<sub>8</sub><sup>2−</sup>]:[NBP-H] ratio of 10:1, SO<sub>4</sub><sup>•−</sup> led to 97.3% degradation of parent NBP-H, while HO• merely resulted in 66.2% degradation. The defluorination from the degraded portion of NBP-H by SO<sub>4</sub><sup>•−</sup> (48.0%) was much higher than that by HO• (17.7%) because all three transformation products were also reactive with SO<sub>4</sub><sup>•−</sup>.

Second, for the –COO<sup>−</sup>-containing PFESA structures without a C–H bond, the degradation became more difficult. Within the same time frame, the ratios of [S<sub>2</sub>O<sub>8</sub><sup>2−</sup>]:[NBP-COOH] and [S<sub>2</sub>O<sub>8</sub><sup>2−</sup>]:[TP1] needed to be 200:1 to achieve >95% degradation of the parent structures. In particular, the degradation of NBP-COOH and NBP-H by SO<sub>4</sub><sup>•−</sup> only differed in the first step (i.e., decarboxylation versus H-abstraction, Scheme 1b), but the defluorination from NBP-COOH with all S<sub>2</sub>O<sub>8</sub><sup>2−</sup> doses were lower than from NBP-H (Table 2, zone C versus B). TP1 from NBP-COOH or as the starting material (Table 2, zone D) also showed higher recalcitrance than that from NBP-H.

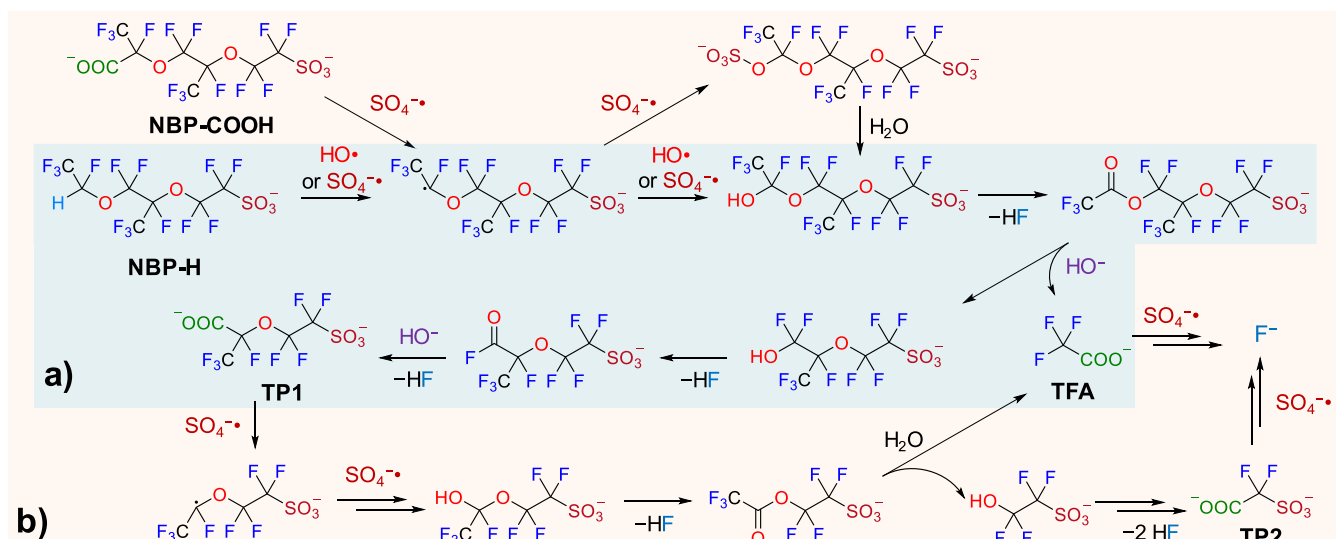
Scheme 1. Oxidative Pathways for PFESA Structures by (a) HO• and (b) SO<sub>4</sub><sup>-•</sup> Radicals

Table 2. Oxidative Transformation of the PFESA Chemicals

| [S <sub>2</sub> O <sub>8</sub> <sup>2-</sup> ]:[PFAS] molar ratio   | parent PFESA degraded (%) | F <sup>-</sup> ion released (mM) | molecular oxidative deF (%) <sup>a</sup> | TP1 (mM) | TP2 (mM) | TFA (mM) | C–F remaining in parent PFESA and targeted TPs (mM) | F atom balance (%) <sup>b</sup> |
|---|---------------------------|----------------------------------|--|----------|----------|----------|---|---------------------------------|
| H–CF(CF <sub>3</sub> )–O–CF <sub>2</sub> CF(CF <sub>3</sub> )–O–CF <sub>2</sub> CF <sub>2</sub> –SO <sub>3</sub> <sup>-</sup> (NBP–H) (0.5 mM, containing 7.0 mM C–F)       |                           |                                  |  |          |          |          |   |                                 |
| A. by HO• (pH > 12)   |                           |                                  |  |          |          |          |   |                                 |
| 10:1  | 66.2                      | 0.82                             | 17.7                                     | 0.31     | n.d.     | 0.32     | 5.80  | 95.0                            |
| 20:1  | 96.0                      | 1.41                             | 20.9                                     | 0.43     | n.d.     | 0.42     | 4.98  | 91.9                            |
| 50:1  | 98.6                      | 1.61                             | 23.3                                     | 0.45     | n.d.     | 0.44     | 5.02  | 95.0                            |
| 100:1   | 99.4                      | 1.74                             | 25.0                                     | 0.47     | n.d.     | 0.45     | 5.15  | 99.1                            |
| B. by SO <sub>4</sub> <sup>-•</sup> (pH < 2)  |                           |                                  |  |          |          |          |   |                                 |
| 10:1  | 97.3                      | 3.27                             | 48.0                                     | 0.17     | 0.13     | 0.30     | 2.71  | 84.9                            |
| 20:1  | 98.8                      | 3.83                             | 55.3                                     | 0.10     | 0.06     | 0.26     | 1.78  | 79.9                            |
| 50:1  | 99.3                      | 4.60                             | 66.2                                     | 0.08     | 0.06     | 0.16     | 1.29  | 84.0                            |
| 100:1   | 99.6                      | 5.56                             | 79.7                                     | 0.01     | 0.03     | 0.08     | 0.41  | 85.9                            |
| 200:1   | 99.3                      | 6.08                             | 87.5                                     | n.d.     | n.d.     | 0.01     | 0.08  | 88.1                            |
| -OOC–CF(CF <sub>3</sub> )–O–CF <sub>2</sub> CF(CF <sub>3</sub> )–O–CF <sub>2</sub> CF <sub>2</sub> –SO <sub>3</sub> <sup>-</sup> (NBP–COOH) (0.5 mM, containing 7.0 mM C–F) |                           |                                  |  |          |          |          |   |                                 |
| C. by SO <sub>4</sub> <sup>-•</sup> (pH < 2)  |                           |                                  |  |          |          |          |   |                                 |
| 10:1  | 60.0                      | 1.91 <sup>c</sup>                | 18.9                                     | 0.07     | 0.05     | 0.06     | 3.66  | 65.3                            |
| 20:1  | 67.0                      | 2.34 <sup>c</sup>                | 24.9                                     | 0.11     | 0.09     | 0.04     | 3.52  | 69.4                            |
| 50:1  | 78.8                      | 3.52 <sup>c</sup>                | 39.9                                     | 0.20     | 0.09     | 0.07     | 3.44  | 85.1                            |
| 100:1   | 88.5                      | 4.71 <sup>c</sup>                | 52.4                                     | 0.22     | 0.07     | 0.06     | 2.87  | 94.0                            |
| 200:1   | 96.7                      | 5.82 <sup>c</sup>                | 62.3                                     | 0.11     | 0.03     | 0.10     | 1.45  | 89.5                            |
| -OOC–CF(CF <sub>3</sub> )–O–CF <sub>2</sub> CF <sub>2</sub> –SO <sub>3</sub> <sup>-</sup> (TP1) (0.5 mM, containing 4.0 mM C–F)   |                           |                                  |  |          |          |          |   |                                 |
| D. by SO <sub>4</sub> <sup>-•</sup> (pH < 2)  |                           |                                  |  |          |          |          |   |                                 |
| 10:1  | 56.0                      | 4.28 <sup>c</sup>                | 59.8                                     | 0.22     | 0.06     | 0.16     | 2.36  | 100.7                           |
| 20:1  | 67.5                      | 4.75 <sup>c</sup>                | 58.2                                     | 0.16     | 0.10     | 0.12     | 1.86  | 95.7                            |
| 50:1  | 75.8                      | 5.40 <sup>c</sup>                | 62.7                                     | 0.12     | 0.09     | 0.12     | 1.49  | 96.7                            |
| 100:1   | 89.8                      | 6.50 <sup>c</sup>                | 68.2                                     | 0.05     | 0.03     | 0.09     | 0.74  | 94.9                            |
| 200:1   | 96.7                      | 6.87 <sup>c</sup>                | 68.1                                     | 0.02     | 0.002    | n.d.     | 0.14  | 85.8                            |

<sup>a</sup>The calculation considered the degraded portion rather than the total initial PFESA. For NBP–COOH and TP1, the F<sup>-</sup> released from hydrolysis was excluded (see footnote c). <sup>b</sup>Including both oxidation-released F<sup>-</sup> and all C–F bonds in the remaining parent PFAS and targeted TPs. A significant gap from 100% suggests the formation of unknown TPs. <sup>c</sup>Including F<sup>-</sup> released from the hydrolysis of sulfonyl fluoride (–SO<sub>2</sub>F) and acyl fluoride (–COF) in the precursor chemicals for NBP–COOH and TP1. The 2 equiv of F<sup>-</sup> (1.0 mM) was not considered in the calculation of oxidative defluorination and F atom balance.

Third, following the known mechanisms, oxidative defluorination is realized via a series of “spontaneous” pathways after the oxidants add a –OH to the fluorinated carbon. The pathways

include HF elimination from –C(OH)F–, hydrolysis of –C(O)F, and hydrolysis of ester CF<sub>3</sub>CO–O–R<sub>F</sub>. Thus, all transformation products are expected to include a –COO<sup>-</sup>,

**Table 3. Oxidative Transformation of C<sub>6</sub>F<sub>13</sub>–Fluorotelomer Carboxylate and Sulfonate**

| [S <sub>2</sub> O <sub>8</sub> <sup>2-</sup> ]:[PFAS]<br>molar ratio  | parent PFAS<br>degraded (%) | F <sup>-</sup> ion in the solution (mM) and<br>oxidative deF (%) <sup>a</sup> | C <sub>n</sub> F <sub>2n+1</sub> –COO <sup>-</sup> products (mM) |                  |                 |                  |                  |                  | F atom<br>balance (%) <sup>b</sup> |
|---|-----------------------------|---|--|------------------|-----------------|------------------|------------------|------------------|------------------------------------|
|   |                             |   | TFA<br>(n = 1)   | PFPrA<br>(n = 2) | PFBA<br>(n = 3) | PFPeA<br>(n = 4) | PFHxA<br>(n = 5) | PFHpA<br>(n = 6) |                                    |
| C <sub>6</sub> F <sub>13</sub> –CH <sub>2</sub> CH <sub>2</sub> –SO <sub>3</sub> <sup>-</sup> (6:2 FTSA) (0.5 mM, containing 6.5 mM C–F)  |                             |   |  |                  |                 |                  |                  |                  |                                    |
| E. by SO <sub>4</sub> <sup>-•</sup> (pH < 2)  |                             |   |  |                  |                 |                  |                  |                  |                                    |
| 10:1  | 31.0                        | 0.19 (9.6)  | n.d.   | n.d.             | n.d.            | 0.004            | 0.004            | 0.004            | 74.0                               |
| 20:1  | 44.5                        | 0.29 (9.9)  | n.d.   | n.d.             | 0.004           | 0.007            | 0.007            | 0.004            | 63.4                               |
| 50:1  | 58.8                        | 0.36 (9.5)  | n.d.   | n.d.             | 0.004           | 0.007            | 0.009            | 0.005            | 50.8                               |
| 100:1   | 83.3                        | 0.54 (10.0)   | n.d.   | n.d.             | 0.007           | 0.013            | 0.014            | 0.007            | 31.4                               |
| F. by HO <sup>•</sup> (pH > 12)   |                             |   |  |                  |                 |                  |                  |                  |                                    |
| 10:1  | 85.3                        | 2.44 (44.1)   | 0.025  | 0.042            | 0.055           | <b>0.101</b>     | 0.079            | 0.010            | 92.1                               |
| 20:1  | 99.4                        | 3.16 (48.9)   | 0.045  | 0.076            | 0.092           | <b>0.108</b>     | 0.042            | 0.004            | 90.0                               |
| 50:1  | >99.8                       | 3.50 (53.9)   | 0.051  | 0.057            | 0.064           | <b>0.123</b>     | 0.052            | 0.008            | 94.9                               |
| 100:1   | >99.8                       | 3.52 (54.2)   | 0.014  | 0.042            | 0.069           | <b>0.128</b>     | 0.064            | 0.019            | 97.7                               |
| C <sub>6</sub> F <sub>13</sub> –CH <sub>2</sub> CH <sub>2</sub> –COO <sup>-</sup> (6:3 FTCA) (0.5 mM, containing 6.5 mM C–F) <sup>c</sup> |                             |   |  |                  |                 |                  |                  |                  |                                    |
| G. by SO <sub>4</sub> <sup>-•</sup> (pH < 2)  |                             |   |  |                  |                 |                  |                  |                  |                                    |
| 10:1  | >99.8                       | 1.31 (20.1)   | 0.015  | 0.024            | 0.032           | 0.062            | 0.127            | 0.015            | 59.2                               |
| 20:1  | >99.8                       | 1.65 (25.3)   | 0.034  | 0.036            | 0.040           | 0.060            | 0.069            | 0.008            | 55.5                               |
| 50:1  | >99.8                       | 1.91 (29.4)   | 0.044  | 0.051            | 0.051           | 0.060            | 0.052            | 0.007            | 59.4                               |
| 100:1   | >99.8                       | 2.70 (41.5)   | 0.048  | 0.048            | 0.039           | 0.037            | 0.028            | 0.004            | 62.4                               |
| H. by HO <sup>•</sup> (pH > 12)   |                             |   |  |                  |                 |                  |                  |                  |                                    |
| 10:1  | >99.8                       | 3.36 (51.7)   | 0.049  | 0.048            | 0.062           | <b>0.128</b>     | 0.051            | 0.008            | 92.2                               |
| 20:1  | >99.8                       | 3.32 (51.0)   | 0.033  | 0.037            | 0.071           | <b>0.120</b>     | 0.052            | 0.013            | 91.0                               |
| 50:1  | >99.8                       | 3.33 (51.3)   | 0.024  | 0.035            | 0.069           | <b>0.115</b>     | 0.065            | 0.018            | 93.2                               |
| 100:1   | >99.8                       | 3.22 (49.5)   | 0.021  | 0.033            | 0.072           | <b>0.112</b>     | 0.066            | 0.023            | 92.0                               |

<sup>a</sup>The calculation considered the degraded portion rather than the total initial FTSA/FTCA. <sup>b</sup>Including both oxidation-released F<sup>-</sup> and all C–F bonds in the remaining parent PFAS and targeted TPs. A significant gap from 100% suggests the formation of unknown TPs. <sup>c</sup>Further processed data of previously reported experiments (Tables S3 and S5 of ref 17).

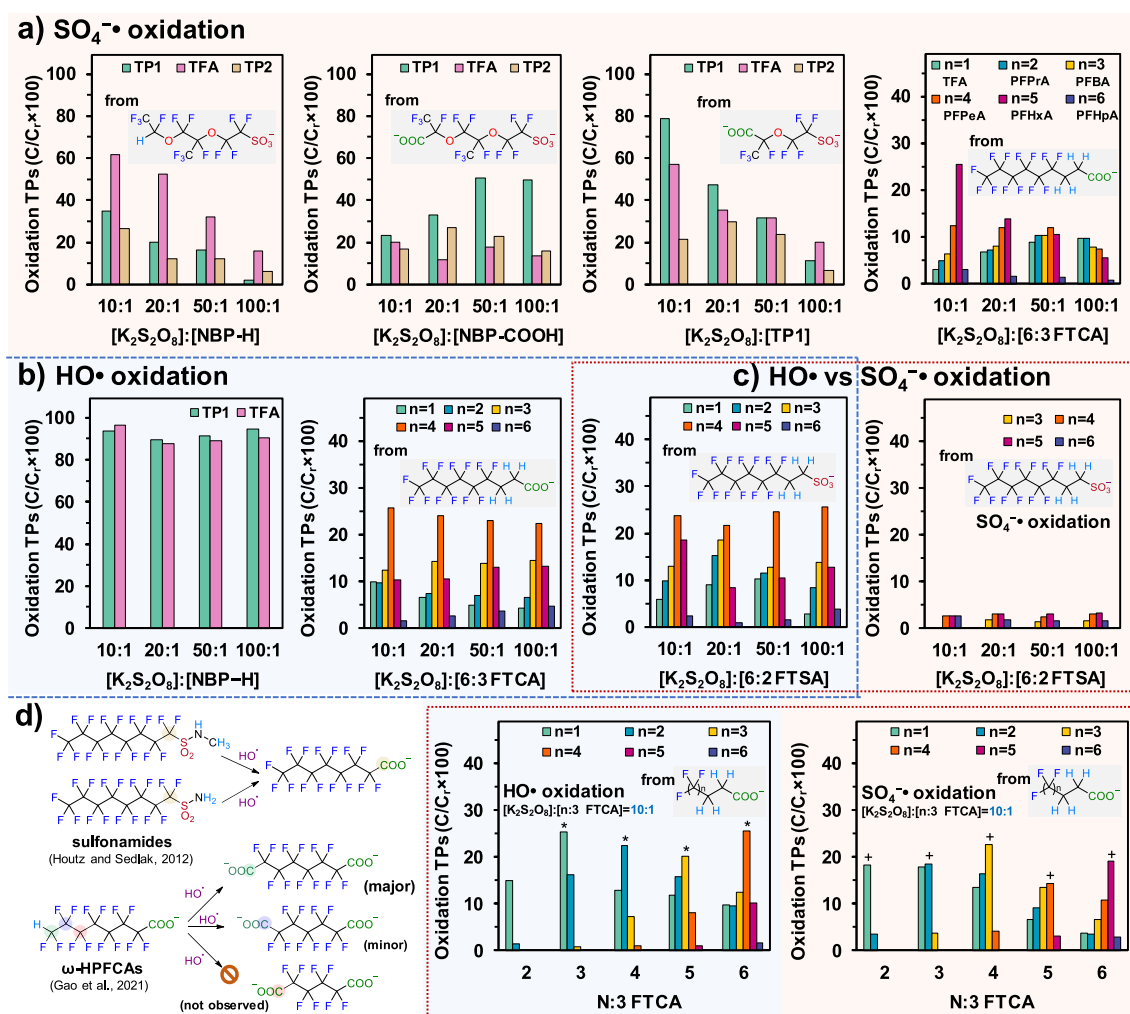
**Table 4. Oxidative Transformation of n:3 Fluorotelomer Carboxylates**

| C <sub>n</sub> F <sub>2n+1</sub> –CH <sub>2</sub> CH <sub>2</sub> –COO <sup>-</sup> (n:3 FTCA)<br>(0.5 mM, containing [n + 0.5] mM C–F) | parent FTCA<br>degraded (%) | F <sup>-</sup> ion in the solution<br>(mM) and oxidative deF<br>(%) <sup>a</sup> | C <sub>n</sub> F <sub>2n+1</sub> –COO <sup>-</sup> products (mM) |                  |                 |                  |                  |                  | F atom<br>balance (%) <sup>b</sup> |
|---|-----------------------------|--|--|------------------|-----------------|------------------|------------------|------------------|------------------------------------|
|   |                             |  | TFA<br>(n = 1)   | PFPrA<br>(n = 2) | PFBA<br>(n = 3) | PFPeA<br>(n = 4) | PFHxA<br>(n = 5) | PFHpA<br>(n = 6) |                                    |
| I. by SO <sub>4</sub> <sup>-•</sup> (pH < 2), [S <sub>2</sub> O <sub>8</sub> <sup>2-</sup> ]:[FTCA] = 10:1                              |                             |  |  |                  |                 |                  |                  |                  |                                    |
| n = 1   | >99.8                       | 1.17 (78.2)  | 0.053  |                  |                 |                  |                  |                  | 88.9                               |
| n = 2   | >99.8                       | 1.86 (74.3)  | <b>0.091</b>   | 0.017            |                 |                  |                  |                  | 88.6                               |
| n = 3   | >99.8                       | 2.09 (59.7)  | 0.089  | <b>0.092</b>     | 0.019           |                  |                  |                  | 84.3                               |
| n = 4   | >99.8                       | 2.18 (48.5)  | 0.067  | 0.082            | <b>0.113</b>    | 0.021            |                  |                  | 83.9                               |
| n = 5   | >99.8                       | 1.61 (29.2)  | 0.033  | 0.045            | 0.067           | <b>0.072</b>     | 0.015            |                  | 58.4                               |
| n = 6 <sup>c</sup>  | >99.8                       | 1.24 (19.0)  | 0.019  | 0.017            | 0.033           | 0.054            | <b>0.095</b>     | 0.014            | 51.2                               |
| J. by HO <sup>•</sup> (pH > 12), [S <sub>2</sub> O <sub>8</sub> <sup>2-</sup> ]:[FTCA] = 10:1 <sup>d</sup>                              |                             |  |  |                  |                 |                  |                  |                  |                                    |
| n = 1   | >99.8                       | 1.43 (95.0)  | n.d.   |                  |                 |                  |                  |                  | 95.0                               |
| n = 2   | >99.8                       | 1.90 (76.2)  | 0.075  | 0.007            |                 |                  |                  |                  | 86.5                               |
| n = 3   | >99.8                       | 2.46 (70.3)  | <b>0.127</b>   | 0.081            | 0.004           |                  |                  |                  | 93.6                               |
| n = 4   | >99.8                       | 2.91 (64.6)  | 0.064  | <b>0.112</b>     | 0.036           | 0.005            |                  |                  | 88.0                               |
| n = 5   | >99.8                       | 3.26 (59.2)  | 0.059  | 0.079            | <b>0.101</b>    | 0.040            | 0.005            |                  | 90.1                               |
| n = 6 <sup>c</sup>  | >99.8                       | 3.36 (51.7)  | 0.049  | 0.048            | 0.062           | <b>0.128</b>     | 0.051            | 0.008            | 92.2                               |

<sup>a</sup>The calculation considered the total initial FTCA because the parent structure degradation had reached >99.8%. <sup>b</sup>Including all C–F bonds in the targeted PFCA TPs because parent FTCA degradation had completed. A significant gap from 100% suggests the formation of unknown TPs. <sup>c</sup>This experiment had the same settings with the first line in Table 3 zone G. <sup>d</sup>Further processed data of previously reported experiments (Figure 2 of ref 17). <sup>e</sup>This experiment had the same settings with the first line in Table 3 zone H.

being subject to further degradation by SO<sub>4</sub><sup>-•</sup>. In theory, 100% oxidative defluorination can be expected, but experimental results found limited defluorination of NBP-COOH (62.3%) and TP1 (68.1%) at a high S<sub>2</sub>O<sub>8</sub><sup>2-</sup> dose of 200:1 (Table 2, zones

C and D). Even for the simple molecule TP2, the defluorination appeared to be limited at ~90% (Table 1, entries 7–10). Therefore, other unknown pathways and mechanisms exist for SO<sub>4</sub><sup>-•</sup> oxidation. This insight is further evidenced by the larger



**Figure 2.** Product distribution by (a)  $\text{SO}_4^{\bullet-}$  (orange background) and (b)  $\text{HO}^{\bullet}$  (blue background) oxidation at varying persulfate doses; (c) comparison of products by  $\text{HO}^{\bullet}$  versus  $\text{SO}_4^{\bullet-}$  from 6:2 FTSA at varying persulfate doses and from various FTCAs at  $[\text{S}_2\text{O}_8^{2-}]:[\text{PFAS}] = 10:1$ ; (d) two reported characteristic product formation from perfluoroalkane sulfonamides and omega-hydro PFCAs. Note: C, indicates the degraded portion of the parent PFAS.

gap of F atom balance for  $\text{SO}_4^{\bullet-}$  than for  $\text{HO}^{\bullet}$  (Table 2, zones B and C versus zone A). We did not detect other meaningful TPs via HRMS analyses.

**Comparison with the Oxidation of Fluorotelomers.** We compared the oxidative transformation patterns of fluorotelomers to those of the “legacy PFAS precursors”. For the extensively studied  $\text{C}_6\text{F}_{13}-\text{CH}_2\text{CH}_2-\text{SO}_3^-$  (6:2 FTSA), the degradation and defluorination by  $\text{SO}_4^{\bullet-}$  were much more difficult than by  $\text{HO}^{\bullet}$  (Table 3, zone E versus F). When  $\text{HO}^{\bullet}$  was used for the heated oxidation, all shorter-chain PFCAs were generated, with the  $n - 2$  PFCA as the most abundant product (e.g.,  $\text{C}_4\text{F}_9-\text{COO}^-$  from  $\text{C}_6\text{F}_{13}-\text{CH}_2\text{CH}_2-\text{SO}_3^-$ ).<sup>15,17</sup> The defluorination, PFCA product distribution, and F atom balance (90–98%) were rather consistent at various  $\text{S}_2\text{O}_8^{2-}$  doses. In contrast, when  $\text{SO}_4^{\bullet-}$  was used, short-chain PFCAs were negligible, and the defluorination appeared to have an  $\sim 10\%$  limit despite more 6:2 FTSA being degraded at higher  $\text{S}_2\text{O}_8^{2-}$  doses. An increasingly lower F atom balance (from 74 to 31%) was also observed, suggesting unknown oxidative pathways and mechanisms.

Notably, the carboxylate analogue  $\text{C}_6\text{F}_{13}-\text{CH}_2\text{CH}_2-\text{COO}^-$  (6:3 FTCA) showed different behaviors from 6:2 FTSA. A low oxidant dose at  $[\text{S}_2\text{O}_8^{2-}]:[6:3 \text{ FTCA}] = 10:1$  achieved complete

parent structure degradation for both  $\text{SO}_4^{\bullet-}$  and  $\text{HO}^{\bullet}$ . We attributed the much higher reactivity of FTCA than FTSA to the weaker C–H bond on the  $\alpha$  carbon adjacent to the  $-\text{COO}^-/-\text{SO}_3^-$  group.<sup>17</sup> However, we still cannot explain how this “local” structural difference caused vastly different levels of defluorination by  $\text{SO}_4^{\bullet-}$  (Table 3, zone G versus E). In the case of 6:3 FTCA, we observed an obvious shift of the PFCA product dominance from long chain to shorter chain when the  $\text{S}_2\text{O}_8^{2-}$  dose was raised. The rather consistent F atom balance (56–62%) and the increased deF% suggest the defluorination via decarboxylation and the subsequent chain-shortening of PFCA products by  $\text{SO}_4^{\bullet-}$ . Still, other unknown pathways and mechanisms are responsible for the  $\sim 40\%$  gap in the F atom balance. For comparison,  $\text{HO}^{\bullet}$  oxidation of FTCA and FTSA showed very similar results in deF%, PFCA product distribution, and high F atom balance (Table 3, zone H versus F).

The extended comparison using various chain lengths of the “most oxidizable”  $n:3$  FTCAs showed the dominance of  $n - 2$  PFCA in the products from  $\text{HO}^{\bullet}$  oxidation (Table 4, zone J). Elevating the  $\text{S}_2\text{O}_8^{2-}$  doses did not significantly alter the PFCA product distribution (Table 3, zones F and H) because  $\text{HO}^{\bullet}$  does not degrade any PFCA. In comparison,  $\text{SO}_4^{\bullet-}$  oxidation using the low dose of  $[\text{S}_2\text{O}_8^{2-}]:[n:3 \text{ FTCA}] = 10:1$  yielded the

Table 5. Oxidative Transformation of Diluted AFFF and Spiked NBP-COOH at Sub- $\mu\text{M}$  Levels

| K <sub>2</sub> S <sub>2</sub> O <sub>8</sub> (mM)  | NaOH (mM) | oxidation products ( $\mu\text{M}$ ) <sup>a</sup> |       |       |                      |                       |                       |                       |                      |
|--|-----------|---|-------|-------|----------------------|-----------------------|-----------------------|-----------------------|----------------------|
|  |           | NBP-COOH remained                                 | TP1   | TP2   | PFBA ( <i>n</i> = 3) | PFPeA ( <i>n</i> = 4) | PFHxA ( <i>n</i> = 5) | PFHpA ( <i>n</i> = 6) | PFOA ( <i>n</i> = 7) |
| K. by HO <sup>•</sup> (pH > 12), 100000× diluted AFFF + 0.35 $\mu\text{M}$ NBP-COOH, 9.5 mg L <sup>-1</sup> of organic carbon              |           |   |       |       |                      |                       |                       |                       |                      |
| 5  | 25        | 0.327   | 0.012 | n.d.  | 0.065                | <b>0.109</b>          | 0.050                 | 0.031                 | 0.020                |
| 20   | 100       | 0.351   | 0.029 | n.d.  | 0.064                | <b>0.106</b>          | 0.035                 | 0.025                 | 0.017                |
| 60   | 300       | 0.343   | n.d.  | n.d.  | 0.113                | <b>0.119</b>          | 0.050                 | 0.036                 | 0.016                |
| L. by SO <sub>4</sub> <sup>-•</sup> (pH < 2), 100000× diluted AFFF + 0.35 $\mu\text{M}$ NBP-COOH, 9.5 mg L <sup>-1</sup> of organic carbon |           |   |       |       |                      |                       |                       |                       |                      |
| 5  |           | 0.115   | 0.126 | 0.039 | 0.055                | 0.068                 | <b>0.095</b>          | 0.024                 | 0.017                |
| 20   |           | 0.031   | 0.050 | 0.008 | 0.025                | 0.023                 | 0.019                 | 0.008                 | 0.005                |
| 60   |           | 0.013   | 0.034 | n.d.  | 0.021                | 0.013                 | 0.008                 | 0.004                 | 0.002                |
| M. by SO <sub>4</sub> <sup>-•</sup> (pH < 2), 10000× diluted AFFF + 3.5 $\mu\text{M}$ NBP-COOH, 95 mg L <sup>-1</sup> of organic carbon    |           |   |       |       |                      |                       |                       |                       |                      |
| 5  |           | 3.283   | 1.792 | 0.533 | 0.392                | 0.475                 | 0.568                 | 0.168                 | 0.162                |
| 20   |           | 0.934   | 1.354 | 0.435 | 0.481                | 0.476                 | 0.450                 | 0.144                 | 0.126                |
| 60   |           | 0.242   | 0.492 | 0.048 | 0.494                | 0.376                 | 0.220                 | 0.093                 | 0.048                |
| N. by SO <sub>4</sub> <sup>-•</sup> (pH < 2), 10000× diluted AFFF, no NBP-COOH, 95 mg L <sup>-1</sup> of organic carbon                    |           |   |       |       |                      |                       |                       |                       |                      |
| 5  |           | n.d.  | n.d.  | n.d.  | 0.399                | 0.458                 | 0.664                 | 0.188                 | 0.188                |
| 20   |           | n.d.  | n.d.  | n.d.  | 0.459                | 0.450                 | 0.389                 | 0.154                 | 0.095                |
| 60   |           | n.d.  | n.d.  | n.d.  | 0.352                | 0.256                 | 0.166                 | 0.071                 | 0.048                |

<sup>a</sup>The LC-MS/MS instrument setting did not allow the detection of ultrashort PFASs, TFA (*n* = 1) and PFPrA (*n* = 2).

most dominant product as *n* - 1 PFCA. Elevating the S<sub>2</sub>O<sub>8</sub><sup>2-</sup> doses shifted the product distribution to short-chain PFASs (Table 3, zones G). However, due to the limited capability of SO<sub>4</sub><sup>-•</sup> oxidation, even with a large excess of S<sub>2</sub>O<sub>8</sub><sup>2-</sup> at 100:1, all PFCA products were still detected after oxidative digestion, leaving various evidence for “forensic analysis” of the parent structures.

**Implications to PFAS Precursor Analyses.** The results presented above suggest a series of similarities and differences between SO<sub>4</sub><sup>-•</sup> and HO<sup>•</sup> for the oxidation of legacy and emerging “PFAS precursors” and their potential value for PFAS reactive analyses. First of all, both radicals are destructive to the reactive PFAS structures. The defluorination from NBP-H, 6:2 FTSA, and 6:3 FTCA by HO<sup>•</sup> reached 25, 54, and 52%, respectively (parent compound degradation >99.8%). Because SO<sub>4</sub><sup>-•</sup> can also oxidize the C-H bond and further degrade PFASs, the molecular defluorination (i.e., calculated for the degraded portion) of NBP-H reached 48–88% with varying S<sub>2</sub>O<sub>8</sub><sup>2-</sup> doses. Therefore, the sum of target PFAS structures after oxidative sample treatment with either SO<sub>4</sub><sup>-•</sup> or HO<sup>•</sup> cannot indicate the total amount of C-F bonds in the original sample. It is also critical to note that different types of PFAS can allow very different levels of defluorination (e.g., NBP-COOH versus NBP-H by SO<sub>4</sub><sup>-•</sup>; C<sub>*n*</sub>F<sub>2*n*+1</sub>-CH<sub>2</sub>CH<sub>2</sub>-COO<sup>-</sup> versus HC<sub>2</sub>F<sub>2*n*</sub>-COO<sup>-</sup> by HO<sup>•</sup><sup>18</sup>). In addition, if the original unknown PFAS molecule does not contain a C-H bond or -COO<sup>-</sup>, it would escape from target analysis. For these reasons, the TOP assay is less accurate than combustion ion chromatography<sup>34</sup> for total F quantitation, but it can be very informative for qualitative analysis and may play a critical role in identifying original PFAS structures.

Second, experimental results have shown that a majority of PFAS structures cannot be fully mineralized by either SO<sub>4</sub><sup>-•</sup> or HO<sup>•</sup>. If a large excess of SO<sub>4</sub><sup>-•</sup> could deplete key -COO<sup>-</sup>-containing products, using different oxidant doses may alter the product presence or dominance (Figure 2a). This feature is unique to SO<sub>4</sub><sup>-•</sup> oxidation. For comparison, the oxidation of NBP-H, 6:3 FTCA, and 6:2 FTSA with HO<sup>•</sup> resulted in rather

consistent product spectra regardless of the oxidant dose (Figure 2b). It is also noteworthy that SO<sub>4</sub><sup>-•</sup> or HO<sup>•</sup> can have very different product dominance for the legacy fluorotelomers (Figure 2c). The complementary use of both radicals will lead to novel information toward deducing the structures of the original “PFAS precursors”. Other oxidation methods, such as cobalt-activated peroxydisulfate at 20 °C are also worth exploring. In a recent report, the oxidation of 6:2 FTSA (40  $\mu\text{M}$ ) using 5 mM of KHSO<sub>5</sub> and 50  $\mu\text{M}$  CoSO<sub>4</sub> yielded *n* = 5 PFHxA as the dominant TP.<sup>35</sup>

We highlight the importance of exploring the transformation mechanisms and pathways for oxidizable PFAS molecules. The detection of TFA as the only PFCA product from Nafion-related PFESAs is a novel example. The H-CF(CF<sub>3</sub>)-O- or -OOC-CF(CF<sub>3</sub>)-O- are oxidized and then transformed into CF<sub>3</sub>-C(O)-O- (Scheme 1), which further hydrolyzes to yield TFA. TP2 is another novel indicator for PFESAs with a -O-CF<sub>2</sub>CF<sub>2</sub>-SO<sub>3</sub><sup>-</sup> terminal, a common structure shared in Nafion-related (Figure 1), F-53B, and other novel structural analogues.<sup>5,6,36</sup> Previous studies also identified other PFAS transformation patterns under HO<sup>•</sup> oxidation, such as the exclusive formation of C<sub>7</sub>F<sub>15</sub>-COO<sup>-</sup> from C<sub>8</sub>F<sub>17</sub>-SO<sub>2</sub>NH<sub>2</sub><sup>15</sup> and the dominating formation of -OOC-C<sub>*n*-1</sub>F<sub>2*n*-2</sub>-COO<sup>-</sup> from H-CF<sub>2</sub>-C<sub>*n*-1</sub>F<sub>2*n*-2</sub>-COO<sup>-</sup> (Figure 2d).<sup>18</sup> We recommend (i) pretreating the samples with both HO<sup>•</sup> and SO<sub>4</sub><sup>-•</sup> (and sequential treatment, if needed) before target PFAS analysis, (ii) adding more known structures (e.g., -OOC-C<sub>*n*</sub>F<sub>2*n*</sub>-COO<sup>-</sup>, ether carboxylates/sulfonates, and other commercially available PFAS chemicals) to the target list of TOP assay, (iii) using advanced mass spectrometry methodologies,<sup>37</sup> data processing algorithms,<sup>38</sup> and additional spectroscopy methods (e.g., <sup>19</sup>F NMR) to assist structural determination, and (iv) modifying the existing methodologies with new structural transformation mechanisms. In order to satisfy the imminent need for the detection, monitoring, and treatment of “non-legacy” PFAS pollutants, it is imperative to expand our understanding of structure-transformation relationships for emerging PFAS chemicals.<sup>18–20,36</sup> The elucidation of additional



transformation pathways for  $\text{SO}_4^{\bullet-}$  oxidation is equally important and intriguing to environmental chemists. The currently unknown  $\text{SO}_4^{\bullet-}$  oxidation TPs might soon become the “signature” of certain PFAS pollutants.

Lastly, PFAS analytical works have not adopted the widely used acidic persulfate digestion for environmental samples. We expect this report to trigger interest in testing  $\text{SO}_4^{\bullet-}$  oxidation (simply adjust the  $\text{S}_2\text{O}_8^{2-}$  solution pH to 2.0 and run a few additional LC–MS samples) in specific PFAS analytical scenarios<sup>14</sup> to obtain more critical information on pollutant structures and profiles.

**Validation with an “Environmentally Relevant” Demonstration.** Here we validate the aforementioned trends of TP formation and the recommended strategies for probing PFAS precursors using acidic persulfate digestion. We set the scenario as analyzing a water sample contaminated by both legacy (from fire-fighting activities) and emerging PFAS (from fluoropolymer manufacturing) with a triple quadrupole mass spectrometer (LC–MS/MS) for quantifying target PFAS at low concentrations. An aqueous film-forming foam (AFFF, containing  $\sim 10 \text{ g L}^{-1}$  of organic fluorine<sup>34</sup> and  $\sim 950 \text{ g L}^{-1}$  of organic carbon) was diluted by 100000-fold and then spiked with  $0.35 \mu\text{M}$  of NBP-COOH.  $^{19}\text{F}$  NMR characterization (Figure S1) and degradation experiments of the AFFF have confirmed its dominant PFAS species as a  $\text{C}_6\text{F}_{13}$ -fluorotelomer structure.<sup>39</sup> Hence, the resulting solution contained  $93 \mu\text{g L}^{-1}$  of F from NBP-COOH,  $\sim 100 \mu\text{g L}^{-1}$  of F from fluorotelomer-based surfactants, and  $9.5 \text{ mg L}^{-1}$  of organic carbon. After oxidation by 5, 20, and 60 mM  $\text{K}_2\text{S}_2\text{O}_8$  at acidic and alkaline pH, the analyses by a third-party LC–MS/MS (Table 5) showed consistent results with our earlier findings from using 0.5 mM individual PFAS and 5–100 mM  $\text{K}_2\text{S}_2\text{O}_8$  without an organic matrix.

Because the current TOP assay protocol involved only alkaline persulfate digestion, the detection of a PFCA mixture (Table 5, zone K) was not sufficient to interpret the precursor structure. If one applies the “ $n - 2$  dominance” rule,<sup>15,17</sup> the dominance of PFPeA ( $\text{C}_4\text{F}_9\text{-COO}^-$ ) and relatively constant abundance of other PFCA products (regardless of the persulfate dose) could be attributed to a  $\text{C}_6\text{F}_{13}\text{-CH}_2\text{CH}_2\text{-X}$  fluorotelomer. However, it is also possible that each PFCA was individually derived from the corresponding sulfonamide (Figure 2d)<sup>15</sup> used in early AFFF products.<sup>40–42</sup> At this point, acidic persulfate digestion provided additional clues (Table 5, zone L). When the persulfate dose was low, the “ $n - 1$ ” PFHxA ( $\text{C}_5\text{F}_{11}\text{-COO}^-$ ) was the dominant product. As the persulfate dose increased, the product dominance shifted to shorter-chain PFCAs. This trend is consistent with our earlier results (Figure 2a for 6:3 FTCA at various persulfate doses; Figure 2c for various  $n$ :3 FTCA at a low persulfate dose). Therefore, one could be more confident in assigning the primary precursor as a  $\text{C}_6\text{F}_{13}\text{-CH}_2\text{CH}_2\text{-X}$  fluorotelomer. Notably, when the AFFF was diluted 10000 $\times$  (thus containing  $95 \text{ mg L}^{-1}$  of organic carbon and consuming more oxidants), the similar PFCA product profiles were observed (Table 5, zone M).

Regarding the detection of Nafion-related PFAS, TP2 ( $\text{-OOC-CF}_2\text{-SO}_3^-$ ) was not found in the samples after alkaline persulfate digestion (Table 5, zone K). The low concentration of TP1 could be attributed to a small chance of NBP-COOH oxidation by  $\text{SO}_4^{\bullet-}$  (before reacting with  $\text{HO}^-$ , also see Table 1, entry 2). In contrast, acid persulfate digestion generated significant yields of TP1 and TP2 (Table 5, zones L and M). Furthermore, TP2 was not produced from the acid persulfate digestion of AFFF (Table 5, zone N versus M).

Therefore, if the simple chemical TP2 is used as an analytical standard, its detection after acid persulfate digestion could indicate some novel PFAS beyond the legacy AFFF-relevant structures. For comparison, the current TOP assay protocol using alkaline persulfate digestion and conventional PFCA analytical standards would not catch Nafion-related PFAS.

Therefore, acidic persulfate digestion for the TOP assay allows the transformation of perfluorinated carboxylates into simpler PFAS structures for target analysis. Its combination with alkaline persulfate digestion will enhance flexibility and confidence in discovering and tracking emerging PFAS pollutants. Like the original TOP assay reported in 2012,<sup>15</sup> this “new” approach warrants further evaluation, development, and optimization for various scenarios and needs.

## ■ ASSOCIATED CONTENT

### Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acs.est.3c06289>.

Information on PFAS chemicals and alkaline hydrolysis of acyl fluoride and sulfonyl fluoride structures, details of instrument settings and QA/QC of HPLC–HRMS analyses, and additional data on defluorination of all PFAS structures by both  $\text{SO}_4^{\bullet-}$  and  $\text{HO}^{\bullet}$  (PDF)

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### Author Contributions

Z.L. conducted most experiments, analyzed the data, and drafted the manuscript; B.J. and Y.M. assisted in HPLC–HRMS

measurements; D.R. conducted tests of the diluted AFFF + NBP-COOH solution mixtures; M.J.B. characterized the AFFF in earlier studies; T.L. assisted in <sup>19</sup>F NMR characterization of NBP structures; J.G. contributed additional data on oxidative conversion; J.L. conceived the idea, supervised the research, and revised the manuscript.

## Notes

The authors declare no competing financial interest.

## ACKNOWLEDGMENTS

Financial support was provided by the Strategic Environmental Research and Development Program (WP21-1071 and ER22-3184 for Z.L. and J.L.; ER20-1541 for B.J. and Y.M.). Drs. Zijie Xia, John Brockgreitens, Terrance P. Smith, and Michelle Bellanca at the PFAS Analytical Lab of Claros Technologies Inc. supported the LC-MS/MS analyses in the section of method validation with diluted AFFF + NBP-COOH.

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