Supplemental Information

Molecular insights on aging and aqueous phase processing from ambient biomass burning emissions-influenced Po Valley fog and aerosol

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1. FT-ICR MS data processing and molecular formula assignment review

The individual transient scans of FT-ICR MS data for each sample were reviewed manually and the unacceptable scans with an abrupt change in the total ion current were removed; the remaining transient scans were co-added together to create the working file for each sample (this helped to increase signal to noise and enhance sensitivity). Molecular formula assignments were made as previously described (Mazzoleni et al., 2010; Putman et al., 2012; Zhao et al., 2013; Dzepina et al., 2015) using Sierra Analytics Composer software (version 1.0.5) within the limits of: C₂₋₂₀₀H₄₋₁₀₀₀O₁₋₂₀N₀₋₃S₀₋₁. The calculator uses a CH₂ Kendrick mass defect (KMD) analysis to sort homologous ion series and extend the molecular formula assignments to higher masses (Hughey et al., 2001; Kujawinski and Behn, 2006). A de novo cut-off at m/z 500 was applied and the minimum relative abundance required for molecular formula assignment was > 10 times the estimated signal-to-noise ratio, determined for each sample between m/z 900–1000. Only integer values up to 40 were allowed for the double bond equivalents (DBE). The data set was manually reviewed to remove: formulas with an absolute error > 3 ppm, elemental ratios that were not chemically sensible (such as O:C > 3 or H:C < 0.3), and formulas which violated the rule of 13 or violated the nitrogen rule. Molecular formulas that contained ¹³C or ³⁴S were also removed from the data set. Homologous series with large gaps in the DBE trend were removed, as well as homologous series with a length of one. The assigned formulas were also analyzed with consideration to the DBE and oxygen number trends, (Herzsprung et al., 2014) where unreliable formula assignments were also removed.

2. Ultrahigh resolution FT-ICR MS results

Reconstructed difference mass spectra of the assigned molecular formulas for both fog and aerosol samples are shown in Fig. S3. These difference mass spectra permit a direct comparison of the samples using normalized relative abundances. The individual relative abundances were normalized by the total abundance of the assigned molecular formulas identified in each of the samples. In Fig. S3, the individual masses with higher abundances in either the positive or negative direction were substantially greater in one of the two samples, whereas the masses of similar abundance tended to cancel each other. To enhance the interpretation of the compositional differences, the individual masses were color-coded to represent the number of oxygen atoms in the assigned formula. Overall, we observed higher numbers of oxygen in the masses of the two samples with aged biomass burning emissions influence compared to the two samples with fresh biomass burning emissions influence. The molecular formulas assigned to the fresh samples had approximately 0-5 oxygen atoms over the mass range of 50-250 Da, 5-10 oxygen atoms over 250-550 Da, and a few molecular formulas were assigned with 10-15 oxygen atoms over 500-600 Da. In contrast, the aged samples had a large number of molecular formulas with 10-15 oxygen atoms in the range of 400-550 Da. This clearly shows a greater amount of oxidation in the aged influenced samples compared to the fresh influenced samples.

KMD diagrams can be used as useful tools to visualize the relationships between the many molecular formulas of complex mixtures such as atmospheric samples. We used Kendrick mass defect to sort the molecular formulas into CH₂ homologous series of identical heteroatom content and DBE, where the formulas in the same series differ only by a number of CH₂ units (Stenson et al., 2003). It should be noted that the presence of multiple formulas

in the same homologous series does not necessarily imply a related chemical structure. The homologous series are visible as horizontal rows of formulas in Figs. S6 and S7. There were multiple homologous series per subclass, where the base formula for each series differ in DBE and increase in KMD to form an ensemble of "steps" within each subclass. In our samples individual CHO and CHNO subclasses had approximately 5-16 different homologous series, while CHOS and CHNOS subclasses had approximately 3-10 different homologous series. The number of homologous series in a subclass increased with oxygen number, and peaked near the median oxygen number, then decreased again towards the maximum number of oxygen; this led to fewer molecular formulas in subclasses with higher and lower oxygen numbers, and more formulas in subclasses near the median oxygen number. The subclasses with the highest numbers of molecular formulas per elemental group were: O₇, NO₈, O₇S and NO₉S. It was atypical for the unique formulas of a sample to be completely unrelated to other formulas across the data set; often the unique formulas were extensions of homologous series that appeared across samples.

3. FT-ICR MS data set

An abbreviated list of the complete FT-ICR MS dataset is provided and is available on Digital Commons: http://digitalcommons.mtu.edu/chemistry-fp/98/

4. Supplemental Figures and tables



Figure S1: Distributions of the molecular formulas within all 64 elemental group subclasses for CHO, CHNO, CHOS and CHNOS groups as indicated in the Figure. The total number of molecular formulas for each SPE-recovered WSOC sample were split into two groups of unique and non-unique formulas; the darker shade represents formulas unique to a sample, (denoted in the Figure legend with an asterisk after the sample name, e.g. "Fresh Fog*") while the lighter shade represents common formulas. The sample names Fresh Fog, Aged Fog, Fresh Aerosol, and Aged Aerosol correspond to SPC0106F, SPC0201F, BO0204N, and BO0213D, respectively.



Figure S2: van Krevelen diagrams for the SPE-recovered WSOC by elemental group (rows) and sample (columns) as indicated in the Figure. Dashed lines represent H:C = 1.2 (horizontal), O:C = 0.6 (vertical) and $OS_C = 0$ (diagonal) as described in Tu et al. (2016). Formulas are color scaled to the number of oxygen atoms in the assigned formula. The sample names Fresh Fog, Aged Fog, Fresh Aerosol, and Aged Aerosol correspond to SPC0106F, SPC0201F, BO0204N, and BO0213D, respectively.



Figure S3: Reconstructed difference mass spectra for theoretical masses of assigned molecular formulas in the Po Valley samples with normalized relative abundance. Fresh influenced samples (SPC0106F and BO0204N) are plotted with positive abundance and aged influenced samples (SPC0201F and BO0213D) are plotted with negative abundance. Molecular compositions in both samples with the same mass and similar normalized relative abundance are reduced toward zero. The peaks in the mass spectra are color scaled to the number of oxygen atoms in the assigned molecular formula, where it can be observed that the aged samples shift towards species with higher oxygen numbers at lower masses, compared to the fresh samples. The sample names Fresh Fog, Aged Fog, Fresh Aerosol, and Aged Aerosol correspond to SPC0106F, SPC0201F, BO0204N, and BO0213D, respectively.



Figure S4: Oxygen difference trends for aerosol (a) and fog (b) samples. Abundance trends were calculated as in Figure 6 of the main text, and then the respective aged sample normalized abundance was subtracted from the fresh sample normalized abundance for each oxygen number value. A positive difference of abundance indicates an enhanced abundance of formulas in the fresh sample compared to the aged sample. Similarly, a negative difference of abundance indicates an enhanced abundance of formulas in the aged sample compared to the fresh sample. The sample names Fresh Fog, Aged Fog, Fresh Aerosol, and Aged Aerosol correspond to SPC0106F, SPC0201F, BO0204N, and BO0213D, respectively.



Figure S5: Double bond equivalent difference trends for aerosol (a) and fog (b) samples. Abundance trends were calculated as in Figure 6 of the main text, and then the respective aged sample normalized abundance was subtracted from the fresh sample normalized abundance for each integer double bond equivalent value. A positive difference of abundance indicates an enhanced abundance of formulas in the fresh sample compared to the aged sample. Similarly, a negative difference of abundance indicates an enhanced abundance of formulas in the aged sample compared to the fresh sample. The sample names Fresh Fog, Aged Fog, Fresh Aerosol, and Aged Aerosol correspond to SPC0106F, SPC0201F, BO0204N, and BO0213D, respectively.



Figure S6: Kendrick mass defect diagrams for each of the Po Valley samples, partitioned by elemental group (rows) and sample (columns) as indicated in the Figure. The molecular formulas unique to each sample are color scaled to the number of oxygen atoms in the assigned formula; grey points represent formulas which are common. Homologous series of molecular formulas are visible as horizontal rows of points, where formulas which are unique to a sample may make up all or only part of an individual homologous series. The sample names Fresh Fog, Aged Fog, Fresh Aerosol, and Aged Aerosol correspond to SPC0106F, SPC0201F, BO0204N, and BO0213D, respectively.



Figure S7: Kendrick mass defect diagrams for each of the Po Valley samples, partitioned by elemental group (rows) and sample (columns) as indicated in the Figure. Molecular formulas are color scaled to the number of oxygen atoms in the assigned formula. The sample names Fresh Fog, Aged Fog, Fresh Aerosol, and Aged Aerosol correspond to SPC0106F, SPC0201F, BO0204N, and BO0213D, respectively.

Formula	SPC0106F	SPC0201F	BO0204N	BO0213D	Identity	Reference*
C ₄ H ₆ O ₅				Х	syringol aqSOA	a
C ₅ H ₆ O ₃				Х	Mt. Tai cloud water ambient OA	b
C ₅ H ₆ O ₄		Х		Х	syringol aqSOA	a, b
C ₅ H ₆ O ₅				Х	syringol aqSOA (2-ketoglutaric acid)	a, c-e
C ₅ H ₈ O ₄		Х		Х	Mt. Tai cloud water ambient OA (2-	b-f
					methylsuccinic acid and glutatric acid)	
C ₅ H ₈ O ₅				Х	Mt. Tai cloud water ambient OA (3-	b, g
					hydroxyglutaric acid)	
C ₆ H ₅ NO ₅			Х	Х	Mt. Tai cloud water ambient OA	b
C ₆ H ₈ O ₄	Х	Х		Х	Mt. Tai cloud water ambient OA	b
C ₆ H ₈ O ₆				Х	syringol aqSOA	a
C ₆ H ₁₀ O ₃		Х		Х	Mt. Tai cloud water ambient OA	b
C ₆ H ₁₀ O ₄		Х		Х	Mt. Tai cloud water ambient OA (3-	b-f
					methylglutaric acid - adipic acid)	
C ₆ H ₁₀ O ₆				Х	dimethyltartaric acid	h
C7H6N2O5	X	Х	Х	Х	Mt. Tai cloud water ambient OA	b
C7H6N2O6	X		Х	Х	Mt. Tai cloud water ambient OA	b
C7H6O4	Х	Х	Х	Х	phenol aqSOA	a
C7H6O5		Х		Х	syringol aqSOA	j
C ₇ H ₇ NO ₄	Х	Х	Х	Х	Mt. Tai cloud water ambient OA	b
C ₇ H ₇ NO ₅	Х	Х	Х	Х	Mt. Tai cloud water ambient OA	b
C7H10O4	Х	Х	Х	Х	methyl vinyl ketone SOA	i
C ₇ H ₁₀ O ₆	Х	Х	Х	Х	guaiacol aqSOA	a
$C_7H_{12}O_4$		Х		Х	methyl vinyl ketone SOA (pimelic	c-e, i
					acid)	
$C_7H_{12}O_6S$		Х		Х	<i>d</i> -limonene SOA	i
$C_7H_{12}O_7$	Х				syringol aqSOA	a
$C_7H_{14}O_5S$	Х	Х		Х	dodecane SOA	i
C ₇ H ₁₄ O ₆ S		Х		Х	d-limonene SOA	i
C ₈ H ₅ NO ₄	X	Х	Х	Х	Mt. Tai cloud water ambient OA	b
C ₈ H ₆ O ₃	X	Х	Х	Х	Mt. Tai cloud water ambient OA;	a, b
					guaiacol aqSOA	
C ₈ H ₆ O ₅	Х	Х		Х	phenol aqSOA	a

Table S1: Summary of the possible identified molecular formulas from the present study. Identical formulas from the literature are provided with their references.

Formula	SPC0106F	SPC0201F	BO0204N	BO0213D	Identity	Reference*
C ₈ H ₇ NO ₃	Х		Х	Х	Mt. Tai cloud water ambient OA	b
C ₈ H ₇ NO ₄	Х	Х	Х	Х	Mt. Tai cloud water ambient OA	b
C ₈ H ₇ NO ₅	Х	Х	Х	Х	Mt. Tai cloud water ambient OA	b
C ₈ H ₉ NO ₃	Х	Х	Х	Х	Mt. Tai cloud water ambient OA	b
C ₈ H ₉ NO ₄	Х	Х	Х	Х	Mt. Tai cloud water ambient OA	b
C ₈ H ₉ NO ₅	Х	Х	Х	Х	Mt. Tai cloud water ambient OA	b
C ₈ H ₁₂ O ₇ S		Х			d-limonene SOA	i
$C_8H_{12}O_8S$	Х	Х		Х	d-limonene SOA	i
C ₈ H ₁₄ O ₆ S	Х	Х		Х	d-limonene SOA	i
C ₈ H ₁₄ O ₇	Х				2-methylglyceric acid dimer	h
C ₈ H ₁₄ O ₇ S		Х			d-limonene SOA	i
C ₉ H ₇ NO ₄	Х	Х	Х	Х	Mt. Tai cloud water ambient OA	b
C ₉ H ₈ O ₂	Х	Х	Х		Mt. Tai cloud water ambient OA	b
C ₉ H ₉ NO ₃	Х		Х	Х	Mt. Tai cloud water ambient OA	b
C ₉ H ₉ NO ₄	Х	Х	Х	Х	Mt. Tai cloud water ambient OA	b
C ₉ H ₁₄ O ₇ S	Х	Х		Х	d-limonene SOA	i
C ₉ H ₁₄ O ₈ S	Х	Х		Х	d-limonene SOA	i
C ₉ H ₁₄ O ₉ S	Х	Х			d-limonene SOA	i
C ₉ H ₁₅ NO ₈ S	Х	Х	Х	Х	monoterpene SOA	i
C ₉ H ₁₆ O ₆ S	Х	Х		Х	d-limonene SOA	i
C9H16O7S	Х	Х		Х	d-limonene SOA	i
C ₉ H ₁₆ O ₈ S		Х			α-terpinene SOA	i
C ₉ H ₁₈ O ₆ S	Х	Х	Х	Х	marine SOA	k
C ₁₀ H ₈ O ₃	Х	Х	Х	Х	Syringol aqSOA	a
C ₁₀ H ₁₀ O ₄	Х	Х	Х		ferrulic acid	c-e
C ₁₀ H ₁₄ O ₅	Х	Х	Х	Х	α-pinene	i
$C_{10}H_{14}O_6$	Х	Х		Х	α-pinene	i
$C_{10}H_{14}O_7S$	Х				d-limonene SOA	i
$C_{10}H_{14}O_8S$	Х	Х			d-limonene SOA	i
C ₁₀ H ₁₆ O ₆ S	Х				d-limonene SOA	i
C ₁₀ H ₁₆ O ₇ S	Х	Х		Х	d-limonene SOA	i
C ₁₀ H ₁₆ O ₈ S		Х		Х	d-limonene SOA	i
C ₁₀ H ₁₆ O ₉ S	Х	Х			d-limonene SOA	i
C ₁₀ H ₁₇ NO ₁₀ S	Х	Х	Х	Х	monoterpene SOA	i
C ₁₀ H ₁₇ NO ₇ S	Х	Х	Х	Х	monoterpene SOA	i

Formula	SPC0106F	SPC0201F	BO0204N	BO0213D	Identity	Reference*
$C_{10}H_{18}O_5S$		Х			β-pinene SOA	i
$C_{10}H_{18}O_7S$	Х	Х		Х	d-limonene SOA	i
$C_{10}H_{18}O_8S$	Х	Х			d-limonene SOA	i
$C_{10}H_{20}O_6S$	Х	Х		Х	marine SOA	k
$C_{10}H_{20}O_7S$	Х	Х		Х	α-terpinene SOA	i
$C_{11}H_{10}O_8$		Х			phenol aqSOA	a
C ₁₁ H ₂₂ O ₆ S	Х	Х		Х	marine SOA	k
$C_{12}H_{10}N_2O_8$			Х		Mt. Tai cloud water ambient OA	b
$C_{12}H_{10}O_2$	Х		Х		phenol aqSOA	a, j
C ₁₂ H ₁₀ O ₃	Х		Х	Х	phenol aqSOA	a
$C_{12}H_{10}O_4$			Х	Х	phenol aqSOA	a
$C_{12}H_{10}O_7$	Х	Х	Х	Х	syringol aqSOA	j
$C_{12}H_{12}O_6$	Х	Х	Х	Х	syringol SOA	i, j
$C_{12}H_{12}O_7$	Х	Х	Х	Х	syringol aqSOA	a, i, j
C ₁₂ H ₁₄ O ₄	Х	Х	Х	Х	syringol aqSOA	a
$C_{12}H_{20}O_7S$	Х	Х		Х	dodecane SOA	i
C ₁₂ H ₂₄ O ₆ S	Х	Х		Х	marine SOA	k
C ₁₃ H ₁₀ O ₃				Х	guaiacol aqSOA	a
C ₁₃ H ₁₀ O ₄	Х	Х	Х		guaiacol aqSOA	a
C ₁₃ H ₁₀ O ₅	Х		Х	Х	guaiacol aqSOA	a
C ₁₃ H ₁₂ O ₄			Х	Х	guaiacol aqSOA	a
C ₁₃ H ₁₂ O ₆	Х	Х	Х	Х	guaiacol aqSOA	a, i
C ₁₃ H ₁₄ O ₅	Х	Х	Х	Х	syringol aqSOA	a
C ₁₃ H ₁₄ O ₇	Х	Х	Х	Х	syringol aqSOA	a
$C_{13}H_{16}O_8$	Х		Х		syringol aqSOA	a
$C_{13}H_{26}O_6S$	Х	Х	Х	Х	marine SOA	k
C ₁₄ H ₁₀ O ₅	Х	Х	Х	Х	phenol aqSOA	a
C ₁₄ H ₁₂ O ₆	Х	Х	Х	Х	guaiacol aqSOA	a, i
$C_{14}H_{12}O_7$	Х	Х		Х	syringol aqSOA	a
C ₁₄ H ₁₄ O ₄			Х	Х	guaiacol aqSOA	a, j, k
C ₁₄ H ₁₄ O ₅	X	Х	Х	Х	guaiacol aqSOA	a, i, j
C ₁₄ H ₁₄ O ₆	Х	Х	Х	Х	guaiacol aqSOA	a, i, j
C ₁₄ H ₁₄ O ₈	Х	Х		Х	syringol aqSOA	a
C ₁₄ H ₁₆ O ₁₀	Х	Х		Х	syringol aqSOA	a
$C_{14}H_{16}O_8$	Х	Х	Х	Х	syringol aqSOA	a

Formula	SPC0106F	SPC0201F	BO0204N	BO0213D	Identity	Reference*
C ₁₄ H ₁₆ O ₉	Х	Х		Х	syringol aqSOA	a
C ₁₄ H ₂₀ O ₉	Х	Х		Х	isoprene SOA	i
$C_{14}H_{24}O_8$		Х			isoprene SOA	i
C ₁₅ H ₁₄ O ₆	Х	Х	Х	Х	syringol aqSOA	i, j
C ₁₅ H ₁₄ O ₈	Х	Х	Х	Х	syringol aqSOA	j
C ₁₅ H ₁₆ O ₆	Х	Х	Х	Х	syringol aqSOA	a, i, j
C15H16O8	Х	Х		Х	syringol aqSOA	a
C ₁₅ H ₁₆ O ₉	Х	Х		Х	syringol aqSOA	a, i, j
C ₁₅ H ₁₈ O ₁₀	Х	Х		Х	syringol aqSOA	a
C15H18O7	Х	Х	Х	Х	syringol aqSOA	a, i, j
C ₁₅ H ₁₈ O ₉	Х	Х	Х	Х	syringol aqSOA	a
C15H24O9	Х	Х		Х	isoprene SOA	i
C ₁₆ H ₁₈ O ₆	Х	Х	Х		syringol aqSOA	a, i-k
C ₁₆ H ₁₈ O ₇	Х	Х	Х	Х	syringol aqSOA	a
C ₁₆ H ₁₈ O ₉	Х	Х		Х	syringol aqSOA	j
C ₁₆ H ₂₄ O ₁₁ S		Х			<i>d</i> -limonene SOA	i
C ₁₈ H ₁₂ O ₅	Х			Х	phenol aqSOA	a
C ₁₈ H ₁₄ O ₄	Х		Х	Х	phenol aqSOA	a
$C_{18}H_{26}O_{12}S$		Х			<i>d</i> -limonene SOA	i
$C_{18}H_{28}O_{11}S$	Х	Х			<i>d</i> -limonene SOA	i
$C_{19}H_{30}O_{12}S$	Х				<i>d</i> -limonene SOA	i
C ₂₀ H ₁₄ O ₆	Х		Х		phenol aqSOA	a
C ₂₀ H ₁₆ O ₇	Х		Х	Х	guaiacol aqSOA	a
C ₂₀ H ₁₈ O ₆	Х		Х	Х	guaiacol aqSOA	a
C ₂₀ H ₂₆ O ₃	Х		Х		7-oxodehydroabietic acid	f
C ₂₀ H ₂₈ O ₂			Х		dehydroabietic acid	f
C ₂₁ H ₁₈ O ₈	Х	Х	Х	Х	guaiacol aqSOA	a
C ₂₁ H ₂₀ O ₆	Х		Х	Х	guaiacol aqSOA	a, j
C ₂₁ H ₂₀ O ₈	Х		Х		guaiacol aqSOA	а
C ₂₈ H ₂₆ O ₈			Х	Х	guaiacol aqSOA	a

*References: (a) Yu et al. (2016); (b) Desyaterik et al. (2013); (c) Pietrogrande et al. (2014a); (d) Pietrogrande et al. (2014b); (e) Pietrogrande et al. (2015); (f) Mazzoleni et al. (2007); (g) He et al. (2014); (h) Herrmann et al. (2015); (i) Cook et al. (2017); (j) Yu et al. (2014); and (k) Dzepina et al. (2015).

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