

Supporting Information for

**A Helium Nanodroplet Setup for Mid and Far-Infrared Spectroscopy
using Pulsed-Free Electron Lasers: Vibrational Spectra of Propargyl
Alcohol.**

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Table S1. Coordinates of the optimized geometries of gauche and trans-propargyl alcohol at MP2/6-311+G(3df,2p) level.

Atom	gauche-PA			trans-PA		
	x (Å)	y(Å)	z(Å)	x(Å)	y(Å)	z(Å)
C	1.92120	-0.21663	-0.01240	-1.94127	-0.20695	0.00000
C	0.76867	0.16507	-0.00497	-0.77694	0.13287	0.00007
H	2.93150	-0.54653	-0.03079	-2.95810	-0.51574	-0.00020
C	-0.63161	0.59001	0.03550	0.61353	0.57254	0.00011
O	-1.54097	-0.49073	-0.09835	1.45463	-0.57321	0.00020
H	-0.83943	1.25870	-0.79788	0.79143	1.19006	0.88521
H	-0.81288	1.14468	0.96031	0.79151	1.19006	-0.88497
H	-1.30104	-1.16173	0.54640	2.36620	-0.26947	-0.00268

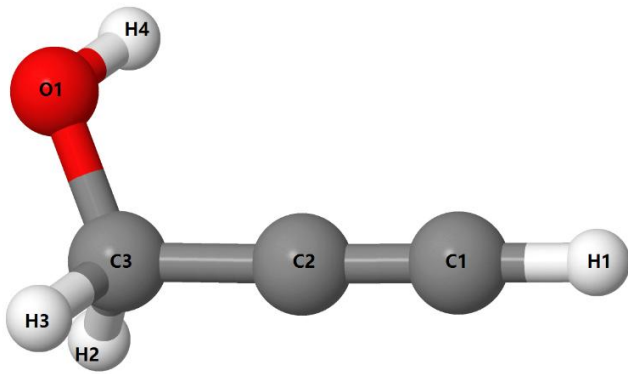


Figure S1. Optimized geometry of gauche-PA at MP2/6-311+G(3df,2p) level.

Table S2. Energies (E) in cm⁻¹ and Integrated intensities in km.mol⁻¹ for gauche and trans-PA for under harmonic and anharmonic approximations at MP2/6-311+G(3df,2p) level. An approximate description of corresponding normal modes is also given. Refer to Figure S1 for atom numberings.

Mode (Quanta)	gauche-PA				Trans-PA				Approximate mode description
	E (harm)	E (anharm)	I(harm)	I (anharm)	E (harm)	E (anharm)	I (harm)	I (anharm)	
1(1)	3869.4	3682.4	44.3	38.7	3871.4	3690.5	42.1	37.3	O-H stretch
2(1)	3484.4	3356.9	57.0	51.2	3487.4	3359.9	55.0	49.4	acetylenic C-H stretch
3(1)	3147.4	3009.1	5.2	5.8	3092.9	2953.3	14.5	17.3	antisymmetric CH ₂ stretch
4(1)	3066.5	2905.9	28.3	29.6	3049.6	2908.3	34.1	35.3	symmetric CH ₂ stretch
5(1)	2133.2	2094.1	1.8	1.6	2146.3	2104.0	2.6	2.5	C≡C stretch
6(1)	1518.5	1477.4	1.9	1.6	1526.0	1480.3	1.1	1.1	CH ₂ Scissoring
7(1)	1423.0	1382.2	51.0	44.0	1454.0	1410.0	25.1	19.4	CH ₂ wagging coupled with C-OH bending
8(1)	1361.8	1320.6	2.8	4.0	1276.8	1247.6	0.0	0.0	O1C3H3 bending coupled with C3O1H4bending
9(1)	1227.5	1198.9	17.1	13.9	1247.1	1236.0	65.2	58.0	CH ₂ twist coupled with C3O1H4 bend for gauche-PA And CH ₂ wagging coupled with C3O1H4 bend for trans-PA
10(1)	1080.6	1044.1	106.5	97.7	1081.5	1044.3	112.4	118.5	C3O1 stretch
11(1)	1006.0	989.1	18.4	17.5	1042.1	1014.9	4.1	3.3	CH ₂ rock coupled with C3O1H4 bend
12(1)	923.3	912.5	23.8	24.8	922.5	912.6	9.7	8.9	C2C3 stretch coupled with C3O1H4 bend
13(1)	673.2	662.8	37.5	36.4	675.7	664.7	36.1	35.4	C2≡C1-H1 bend in the C-C-O plane
14(1)	647.8	637.0	45.8	44.4	645.3	635.9	44.7	44.0	C2≡C1-H1 bend perpendicular to C-C-O plane
15(1)	557.9	549.9	12.5	9.2	549.8	551.1	3.1	1.7	carbon chain bend parallel to C3-O1 bond, coupled with large amplitude COH motion
16(1)	326.9	298.9	67.5	25.2	308.1	308.6	0.2	7.1	carbon chain bend perpendicular to C3-O1 bond, coupled with large amplitude O-H and CH ₂ motions
17(1)	263.2	242.7	61.7	104.4	225.5	181.2	128.1	110.7	carbon chain bend perpendicular to C3-O1 bond, coupled with large amplitude O-H and CH ₂ motions
18(1)	190.5	189.7	7.9	16.9	202.6	199.8	8.7	8.0	carbon chain bend parallel to C3-O1 bond, coupled with large amplitude COH motion