

# **ELECTRONIC SUPPLEMENTARY INFORMATION**

## **Impact of Speciation on the Electron Charge Transfer Properties of Nanodiamond Drug Carriers**

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# Thermodynamically Limited Boltzmann Distribution

Table 1: Boltzmann distribution, for samples that are thermodynamically limited: Shapes, with the probability distributed over all sizes.

	Expectation Values (eV)				Quality Factors (arb.)			
	IP	EA	$E_{gap}$	$E_{Fermi}$	IP	EA	$E_{gap}$	$E_{Fermi}$
Octahedron	5.063	4.153	0.910	-4.611	33.0	24.4	3.0	86.4
Rhombi-truncated Octahedron	5.380	4.569	0.810	-4.973	28.2	26.5	3.0	39.6
Truncated Octahedron	5.306	4.413	0.892	-4.859	47.6	23.0	3.9	44.9
Doubly-truncated Octahedron	5.145	4.337	0.808	-4.755	28.5	18.4	2.3	41.6
Rhombic Dodecahedron	5.614	4.823	0.791	-5.215	37.8	29.3	5.2	37.2
Truncated Dodecahedron	5.748	4.889	0.859	-5.320	33.4	21.7	6.1	28.1
Small Rhombicuboctahedron	5.458	4.497	0.961	-4.975	35.0	21.5	4.6	32.1
Doubly-truncated Dodecahedron	5.576	4.755	0.820	-5.164	37.7	17.0	2.9	30.4
Cube	6.052	5.060	0.991	-5.558	15.1	16.9	4.6	16.2
Truncated Cube	5.674	4.851	0.823	-5.262	36.2	31.3	3.2	58.7
Cuboctahedron	5.488	4.760	0.729	-5.096	24.2	16.0	4.7	31.8
Great Rhombicuboctahedron	5.570	4.630	0.940	-5.100	144.9	29.9	29.2	34.1
Mixture (All)	5.442	4.583	0.859	-5.012	18.9	14.6	3.3	18.9

Table 2: Boltzmann distribution, for samples that are thermodynamically limited: Facet-constrained samples, with the probability distributed over all sizes.

	Expectation Values (eV)				Quality Factors (arb.)			
	IP	EA	$E_{gap}$	$E_{Fermi}$	IP	EA	$E_{gap}$	$E_{Fermi}$
(111)-enriched	5.230	4.379	0.852	-4.809	25.9	18.1	2.9	29.2
(110)-enriched	5.582	4.710	0.873	-5.144	29.3	17.1	4.0	24.2
(100)-enriched	5.623	4.772	0.851	-5.188	22.1	17.8	3.7	22.1
Quasi-spherical	5.449	4.582	0.866	-5.018	21.0	16.4	3.2	21.7
Highly faceted	5.433	4.585	0.848	-5.004	16.8	12.8	3.5	16.3

Table 3: Boltzmann distribution, for samples that are thermodynamically limited: Speciation-constrained samples, with the probability distributed over all sizes.

	Expectation Values (eV)				Quality Factors (arb.)			
	IP	EA	$E_{gap}$	$E_{Fermi}$	IP	EA	$E_{gap}$	$E_{Fermi}$
sp <sup>3</sup> -enriched	5.563	4.727	0.837	-5.148	24.1	18.4	4.6	23.1
sp <sup>2</sup> -enriched	5.265	4.374	0.891	-4.822	24.7	19.8	3.0	30.4
sp <sup>2+x</sup> -enriched	5.515	4.663	0.852	-5.082	18.5	14.8	2.7	20.3
$N_{coord} < 3.5$	5.221	4.253	0.967	-4.737	22.0	19.7	3.3	27.8
$N_{coord} > 3.5$	5.509	4.683	0.826	-5.095	20.5	17.4	3.5	22.0

# Size-dependent Normal Distribution

Table 4: Normal distribution, for samples that are kinetically limited: Shapes, with the probability distributed over all sizes.

	Expectation Values (eV)				Quality Factors (arb.)			
	IP	EA	$E_{gap}$	$E_{Fermi}$	IP	EA	$E_{gap}$	$E_{Fermi}$
Octahedron	5.030	4.144	0.886	-4.589	32.1	27.4	3.0	101.9
Rhombi-truncated Octahedron	5.379	4.573	0.805	-4.977	26.1	23.3	2.8	34.8
Truncated Octahedron	5.311	4.350	0.961	-4.830	40.4	31.0	5.2	48.4
Doubly-truncated Octahedron	5.121	4.394	0.727	-4.761	25.3	20.7	2.0	45.7
Rhombic Dodecahedron	5.590	4.801	0.789	-5.190	34.5	25.5	4.8	32.9
Truncated Dodecahedron	5.755	4.870	0.884	-5.315	34.3	21.2	5.6	28.2
Small Rhombicuboctahedron	5.496	4.495	1.001	-4.994	35.1	20.4	5.2	29.9
Doubly-truncated Dodecahedron	5.583	4.773	0.811	-5.176	41.4	13.9	2.3	27.0
Cube	6.104	5.114	0.990	-5.611	15.7	16.4	4.7	16.5
Truncated Cube	5.681	4.849	0.832	-5.265	34.1	29.9	3.1	54.8
Cuboctahedron	5.430	4.716	0.714	-5.042	23.7	15.6	4.1	35.1
Great Rhombicuboctahedron	5.571	4.623	0.948	-5.096	178.6	30.8	33.2	34.6
Mixture (All)	5.464	4.601	0.863	-5.030	17.4	14.4	3.2	17.9

Table 5: Normal distribution, for samples that are kinetically limited: Facet-constrained, with the probability distributed over all sizes.

	Expectation Values (eV)				Quality Factors (arb.)			
	IP	EA	$E_{gap}$	$E_{Fermi}$	IP	EA	$E_{gap}$	$E_{Fermi}$
(111)-enriched	5.225	4.388	0.837	-4.808	23.4	19.3	2.8	28.5
(110)-enriched	5.595	4.708	0.887	-5.150	30.4	16.2	3.8	23.9
(100)-enriched	5.644	4.776	0.868	-5.203	20.1	17.2	3.6	20.1
Quasi-spherical	5.457	4.594	0.863	-5.026	20.1	17.0	3.0	21.8
Highly faceted	5.472	4.609	0.863	-5.035	15.1	12.3	3.6	14.9

Table 6: Normal distribution, for samples that are kinetically limited: Speciation-constrained, with the probability distributed over all sizes.

	Expectation Values (eV)				Quality Factors (arb.)			
	IP	EA	$E_{gap}$	$E_{Fermi}$	IP	EA	$E_{gap}$	$E_{Fermi}$
sp <sup>3</sup> -enriched	5.589	4.726	0.863	-5.159	25.9	18.7	5.0	23.8
sp <sup>2</sup> -enriched	5.251	4.387	0.863	-4.820	26.0	25.7	3.5	34.5
sp <sup>2+x</sup> -enriched	5.556	4.727	0.830	-5.140	20.3	19.0	3.0	23.5
$N_{coord} < 3.5$	5.205	4.272	0.934	-4.738	21.8	21.9	3.3	29.0
$N_{coord} > 3.5$	5.537	4.693	0.843	-5.113	18.8	16.5	3.2	20.3

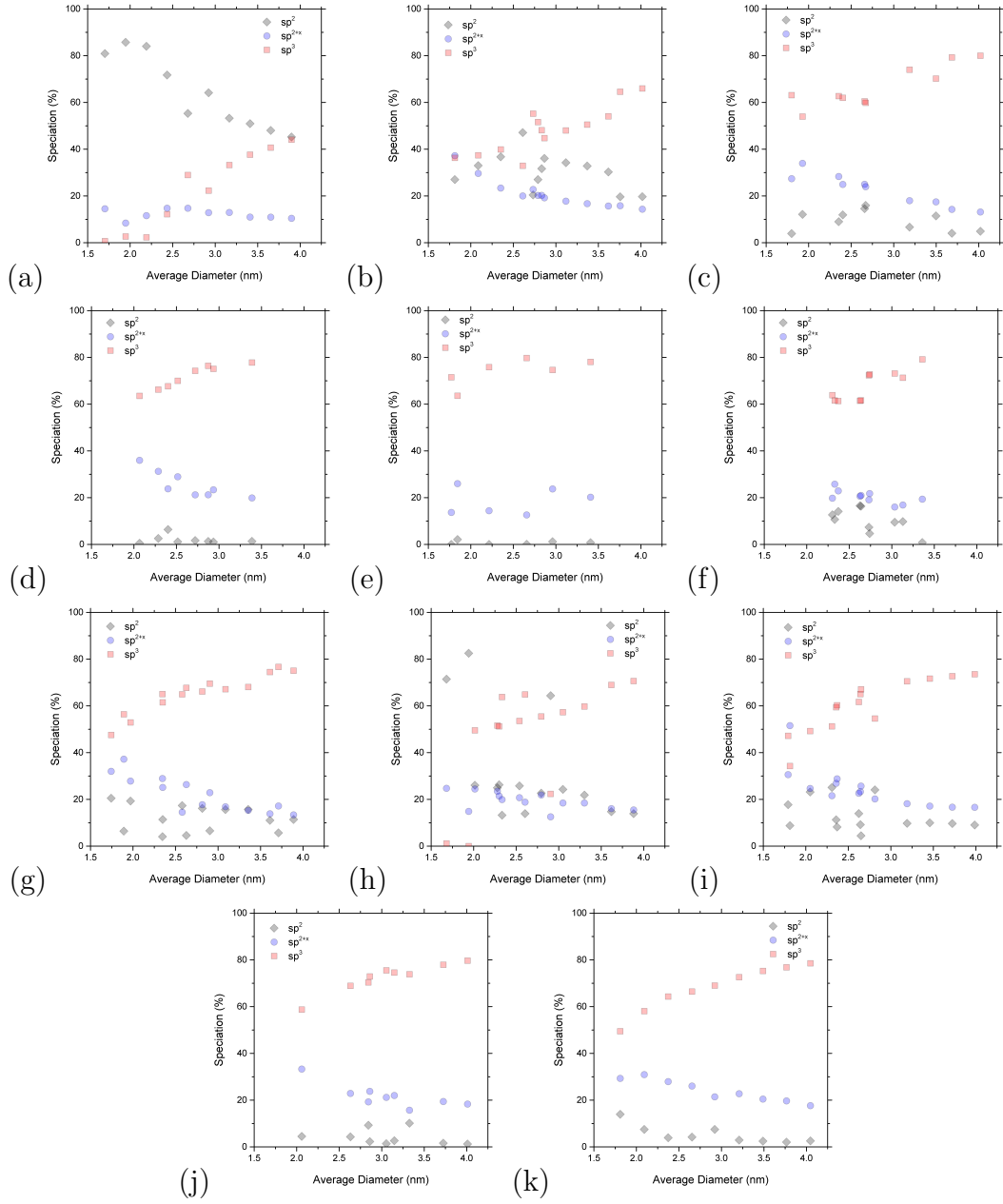


Figure 1: Size-dependent speciation for each of the shapes represented in the dataset used in this study: (a) the octahedron, (b) truncated octahedron, (c) cuboctahedron, (d) truncated cube, (e) cube, (f) great rhombicuboctahedron, (g) small rhombicuboctahedron, (h) doubly-truncated octahedron, (i) rhombi-truncated octahedron, (j) truncated dodecahedron and (k) the rhombic dodecahedron.

# Bond Length Distributions

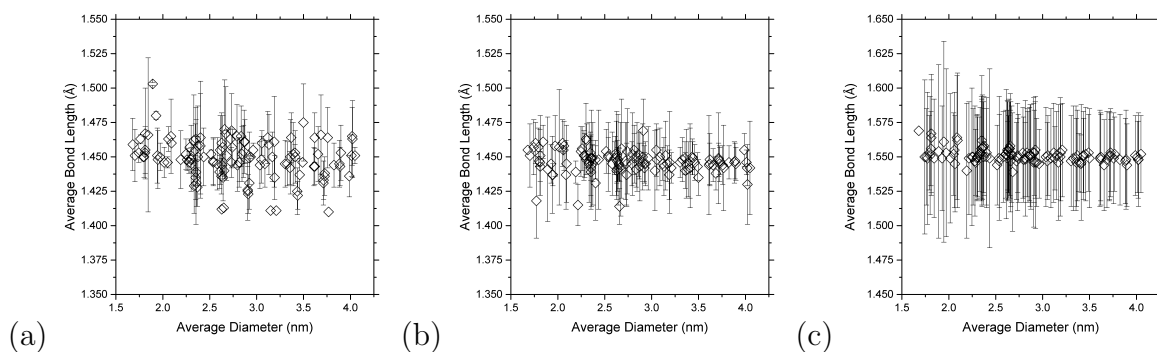


Figure 2: Bond lengths, with error bar representing the bond length distribution, for each bucky-diamond structure represented in this dataset, separated according to the speciation: (a)  $sp^2$  hybridized atoms, (b)  $sp^{2+x}$  hybridized atoms, and (c)  $sp^3$  hybridized atoms.

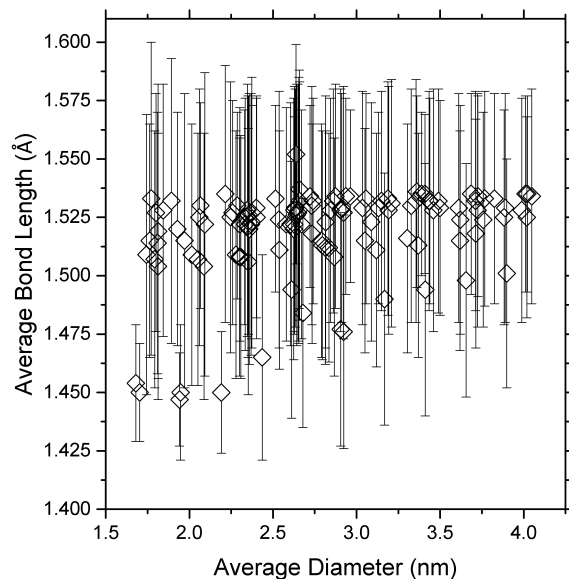


Figure 3: The average bond length, with error bar representing the bond length distribution, for each of the bucky-diamond structure represented in this dataset, averaging over all bond types.

# Bond Angle Distributions

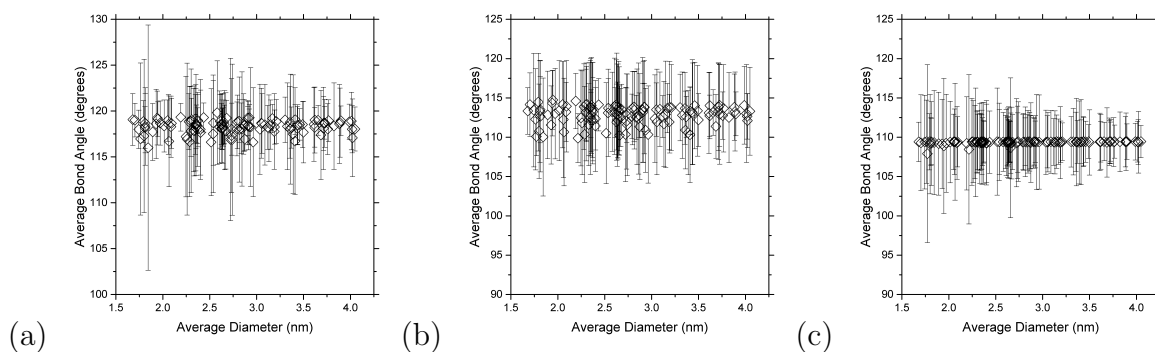


Figure 4: Bond angles, with error bar representing the bond length distribution, for each bucky-diamond structure represented in this dataset, separated according to the speciation: (a)  $sp^2$  hybridized atoms, (b)  $sp^{2+x}$  hybridized atoms, and (c)  $sp^3$  hybridized atoms.

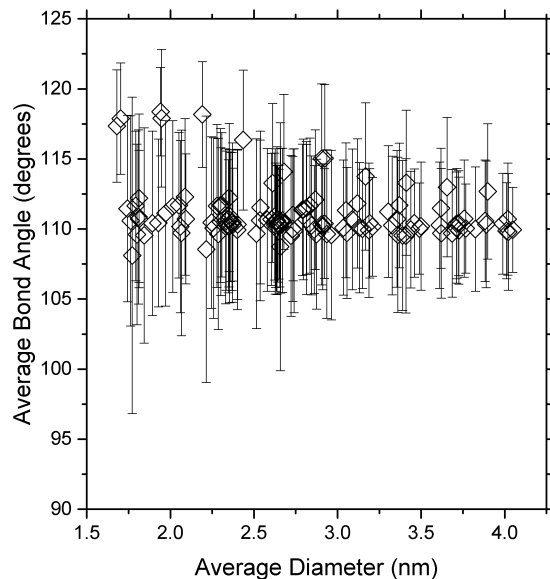


Figure 5: The average bond angle, with error bar representing the bond length distribution, for each of the bucky-diamond structure represented in this dataset, averaging over all bond types.

## Acknowledgement

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